Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

NAME	001-LA	002-LA	003-LB	004-LA	005-LB	006-LB	007-LA	008-LB	009-LB	010-LB	011-LA	012-LB	013-LBA	014-LB	015-LAR	016-LB	017-LB	018-LB
NA1	236.0	852.0	79.8	15.7	92.7	30.3	9.9	128.0	53.7	40.8	9.2	1.2	8.5	1.6	0.4	1.6	2.3	19.3
NA2	134.0	636.0	40.8	10.9	15.7	6.1	1.0	9.8	2.0	0.3	1.2	-0.1	0.5	0.3	-0.1	0.3	0.3	1.6
NAA1	203.0	657.0	45.6	11.2	42.3	12.8	3.4	43.8	18.2	11.7	1.7	0.7	3.2	0.5	0.3	0.5	1.5	7.0
NAA10	235.0	507.0	78.3	11.1	54.3	14.5	3.9	39.3	16.4	11.2	2.8	0.6	3.6	0.6	-1.1	0.2	0.9	6.8
NAA11	264.0	867.0	67.2	19.4	77.4	23.8	33.0	163.0	66.9	58.5	72.6	1.3	67.2	19.1	0.9	19.1	3.8	30.6
NAA2	465.0	2290.0	195.0	83.1	321.0	87.0	75.3	447.0	183.0	138.0	157.0	0.3	147.0	41.4	1.4	41.4	14.1	72.3
NAA3	414.0	1250.0	72.9	24.3	40.8	13.5	5.3	29.2	6.2	6.9	4.4	0.6	3.9	3.4	-0.1	3.4	0.9	6.3
NAA4	153.0	573.0	29.1	10.9	15.1	9.8	0.6	16.1	3.3	3.9	1.7	0.8	1.6	0.4	-0.1	0.4	0.6	3.1
NAA5	209.0	711.0	61.5	11.1	29.0	7.1	2.9	23.1	9.6	5.0	2.4	0.3	2.1	0.5	0.3	0.5	0.7	4.5
NAA6	248.0	699.0	48.6	12.0	27.8	9.4	2.7	36.3	3.4	9.9	2.9	0.6	2.5	0.9	-0.3	0.9	0.9	5.9
NAA7	115.0	552.0	36.3	9.4	19.7	12.9	2.4	18.8	3.9	4.0	1.5	0.3	1.4	0.4	-0.1	0.4	0.5	3.4
NAA8	123.0	504.0	36.9	8.5	22.0	14.3	2.4	19.5	3.9	4.3	1.4	0.2	1.7	0.3	-0.1	0.3	0.4	2.9
NAA9	147.0	630.0	93.3	9.1	64.8	16.1	4.4	54.6	22.7	14.1	3.6	0.5	3.4	0.6	0.3	0.6	1.1	8.0
NB1	167.0	759.0	50.7	14.1	34.2	12.0	3.2	45.0	4.6	13.2	5.6	0.7	4.4	0.7	-0.3	0.7	1.0	6.4
NB2	207.0	804.0	54.0	14.6	30.9	7.5	3.3	20.8	4.6	4.4	1.7	0.2	1.5	0.1	-0.1	0.1	0.5	3.1
NB3	212.0	870.0	46.5	13.4	26.4	7.7	3.2	19.8	4.1	4.9	2.3	0.2	1.6	0.4	-0.1	1.6	0.5	3.2
NB4	280.0	1160.0	54.0	20.1	36.0	12.3	4.9	28.4	12.0	10.2	3.2	0.5	1.1	0.4	-0.1	0.4	0.7	4.6
NBB1	185.0	579.0	85.8	9.1	60.0	17.3	2.8	49.2	20.5	14.1	2.6	0.6	2.3	4.5	-0.1	4.5	0.9	7.6
NBB10	154.0	645.0	55.5	8.7	30.0	9.0	4.4	28.0	11.7	6.8	4.5	0.6	2.2	0.5	-0.1	0.5	0.6	4.9
NBB11	218.0	699.0	21.3	8.6	23.3	8.5	6.9	68.7	28.2	19.3	5.9	1.1	5.6	10.1	0.3	10.1	2.9	14.7
NBB2	190.0	570.0	60.0	10.8	40.2	11.4	7.5	46.5	19.2	11.3	9.5	1.0	12.2	0.6	0.3	0.6	1.4	7.8
NBB2-R	210.0	609.0	59.7	11.5	38.1	9.9	7.9	45.3	18.8	11.0	10.4	0.8	12.4	4.8	-0.1	4.8	1.4	7.0
NBB3	278.0	804.0	111.0	20.1	84.0	21.9	12.8	90.6	6.3	27.1	18.6	1.2	13.5	1.5	0.3	1.5	2.7	13.3
NBB4	312.0	927.0	66.9	14.0	43.2	12.2	3.6	41.1	8.1	9.8	3.4	0.6	1.5	5.1	-0.1	5.1	3.0	6.7
NBB5	151.0	573.0	33.6	10.7	20.5	6.7	2.9	20.7	4.3	5.5	2.0	0.4	1.9	0.4	-0.1	0.4	0.6	3.6
NBB6	192.0	717.0	76.5	12.9	68.7	18.4	11.4	81.6	53.6	23.4	15.4	1.1	13.8	0.9	-0.3	0.9	2.1	13.2
NBB7	125.0	540.0	48.6	9.2	24.6	6.9	0.8	16.5	4.2	3.1	1.2	0.2	1.1	0.4	-0.1	0.4	0.3	2.2
NBB8	160.0	630.0	60.9	9.3	40.2	10.8	5.4	34.8	14.4	8.5	5.0	0.5	2.3	1.2	-0.3	0.7	0.7	5.4
NBB9	226.0	876.0	64.8	16.8	56.1	15.2	26.5	92.7	38.7	25.8	39.6	1.2	36.6	11.0	0.8	11.0	2.1	16.4
NC1	126.0	555.0	40.5	9.9	23.9	14.0	2.9	18.4	1.2	4.5	1.6	0.2	0.5	0.2	-0.1	0.2	0.3	2.6
NC2	121.0	564.0	36.6	10.3	23.0	6.8	2.3	20.1	5.2	4.7	1.9	0.2	1.5	0.3	-0.1	0.4	0.6	3.1
NC3	165.0	633.0	37.2	12.4	17.4	4.9	0.9	19.3	3.3	2.6	1.6	-0.1	2.2	0.4	-0.1	0.4	0.3	2.1
NC4	139.0	633.0	28.3	11.5	13.0	4.5	0.8	5.3	2.5	2.1	1.2	-0.1	1.6	0.4	-0.1	0.4	0.2	1.7
NC5	156.0	693.0	37.2	13.3	22.4	9.5	3.6	21.6	4.9	4.5	2.3	0.6	3.1	0.3	-0.1	0.3	0.7	3.4
NC6	143.0	699.0	50.1	1.7	41.1	12.2	5.3	46.8	11.2	11.4	3.5	0.4	3.2	0.8	0.3	0.8	1.3	7.3
NC7	106.0	567.0	34.2	10.7	18.4	11.8	0.8	16.8	3.7	3.6	1.7	0.2	0.7	0.5	-0.1	0.6	0.4	2.6
NC8	144.0	567.0	28.8	9.9	17.2	10.5	1.0	8.0	3.7	3.6	1.4	0.2	2.4	0.3	-0.1	0.3	0.4	2.7
NCC1	861.0	540.0	268.0	65.8	150.0	39.9	15.5	106.0	5.4	28.8	7.2	1.0	8.4	5.8	-0.2	5.8	1.5	10.8
NCC1-R	765.0	363.0	256.0	74.7	148.0	37.8	14.3	95.4	5.2	26.7	6.8	1.1	7.8	5.4	-0.1	5.4	1.4	9.9
NCC2	169.0	578.0	46.5	11.5	32.2	28.1	7.5	122.0	51.0	38.7	8.7	1.9	6.0	1.0	0.3	3.5	1.5	18.5
NCC3	181.0	621.0	46.8	11.5	51.3	16.6	6.2	70.8	29.4	19.4	4.5	0.6	4.6	6.4	0.3	6.4	1.5	9.7
NCC4	163.0	699.0	103.0	9.4	71.7	28.3	6.7	84.6	84.8	22.4	6.9	0.6	6.7	0.8	-0.2	0.8	1.9	14.6
NCC5	198.0	600.0	25.0	11.0	21.3	13.4	5.4	35.4	14.6	9.6	4.5	0.6	4.1	1.1	0.3	1.1	1.0	6.0
NCC6	245.0	753.0	54.3	11.5	22.1	6.2	0.7	22.3	2.2	5.0	1.8	0.4	1.3	0.3	-0.3	0.8	0.6	3.2
ND1	300.0	906.0	70.8	17.0	66.0	20.5	6.1	60.6	25.0	16.4	1.6	0.3	2.9	0.7	1.1	0.7	1.2	8.3
ND10	221.0	795.0	34.6	13.3	16.9	9.2	1.5	15.1	3.3	3.3	1.3	0.2	0.1	0.4	-0.1	0.3	0.4	2.5
ND2	276.0	810.0	63.0	15.3	45.9	16.0	5.6	48.3	20.0	10.6	4.6	0.3	5.4	0.8	1.2	0.8	0.9	6.8
ND3	168.0	663.0	42.9	11.6	19.7	12.1	0.7	11.5	3.1	2.5	1.4	-0.1	0.7	0.4	-0.1	0.4	0.2	1.9
ND4	357.0	1130.0	54.0	20.1	32.4	9.5	4.9	27.6	5.5	3.4	0.5	1.2	3.6	0.3	3.6	0.8	5.4
ND5	232.0	619.0	37.6	14.6	15.5	5.6	0.8	10.6	2.1	2.0	4.1	-0.1	0.6	0.5	-0.1	0.5	0.4	1.6
ND6	253.0	834.0	35.1	16.1	24.6	8.6	2.7	22.4	4.9	5.3	2.2	0.2	2.6	0.4	-0.1	0.4	0.6	3.8
ND7	272.0	963.0	35.1	15.1	32.2	22.4	4.4	40.5	3.3	17.5	2.2	0.4	1.5	0.6	-0.1	0.6	1.0	6.2
ND8	125.0	558.0	19.8	10.2	11.9	7.5	1.2	15.1	2.8	3.8	1.5	0.2	3.1	0.4	-0.1	0.4	0.5	3.2
ND8-R	155.0	648.0	23.0	13.1	14.9	8.4	3.4	17.7	7.4	4.6	2.2	0.4	1.9	0.4	-0.1	0.4	0.6	3.6
ND9	174.0	765.0	38.1	13.0	25.0	9.7	3.8	23.4	4.7	5.2	2.1	0.2	1.8	0.5	-0.1	0.5	0.7	4.1
NDD1	279.0	852.0	63.3	19.9	91.2	27.8	17.9	125.0	52.5	39.3	23.4	1.3	22.4	1.4	-0.5	1.4	2.6	20.6
NDD2	276.0	735.0	133.0	13.0	147.0	45.0	5.7	132.0	54.6	37.8	6.8	1.5	6.6	1.0	0.3	1.0	3.0	14.6
NDD3	210.0	691.0	39.6	9.9	41.4	12.1	4.4	50.4	20.9	13.4	4.2	0.3	4.4	0.6	-0.3	0.5	1.2	8.9
NDD4	183.0	576.0	27.2	9.7	23.7	8.3	2.7	31.5	13.0	9.0	3.0	0.5	2.6	0.5	-0.1	0.5	0.8	6.5
NDD5	176.0	564.0	20.1	9.1	18.3	11.4	0.8	14.9	2.7	3.5	1.6	0.2	0.7	0.6	-0.1	0.6	0.5	3.0
NE1	197.0	804.0	89.7	14.3	113.0	34.5	11.9	149.0	61.2	46.2	15.3	1.3	14.2	1.1	0.4	1.1	2.6	22.3
NE10	154.0	606.0	39.0	10.1	21.2	12.6	0.6	7.2	3.3	2.4	0.8	-0.1	0.7	0.4	-0.1	0.4	0.4	1.6
NE11	186.0	663.0	26.6	10.3	11.7	8.1	0.7	5.3	2.5	0.3	0.9	-0.1	0.3	0.3	-0.1	0.3	0.3	1.8
NE2	117.0	540.0	47.7	10.1	22.3	9.5	0.9	7.9	3.2	0.2	1.4	-0.1	0.5	0.4	-0.1	0.4	0.4	1.5
NE3	145.0	549.0	42.0	9.0	38.1	11.5	1.2	36.6	8.3	8.6	2.4	0.6	2.3	0.6	-0.1	0.6	0.7	4.7
NE4	137.0	435.0	39.3	9.2	34.2	10.9	1.0	34.2	7.5	9.2	2.0	0.6	1.7	0.4	-0.1	0.4	1.0	4.8
NE4-R	128.0	420.0	36.0	8.5	29.3	8.7	0.9	25.0	6.1	6.2	1.3	0.3	2.4	0.3	-0.1	0.3	0.4	3.4

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NE5	242.0	819.0	79.8	11.4	76.2	25.9	-4.0	89.1	34.2	24.9	4.1	1.2	3.6	1.1	1.2	1.1	2.5	12.5
NE6	137.0	570.0	22.5	10.7	10.6	6.0	0.6	10.4	2.3	2.2	1.3	-0.1	0.9	0.5	-0.1	0.5	0.5	2.2
NE7	169.0	741.0	103.0	13.4	91.8	26.4	1.8	73.2	3.9	19.5	2.0	0.2	1.6	0.8	-0.1	0.9	0.9	7.4
NE8	240.0	822.0	31.2	13.2	13.5	5.3	1.3	8.0	1.7	0.3	0.9	-0.1	0.8	0.4	-0.1	0.4	0.4	1.3
NE9	112.0	489.0	16.5	8.5	12.3	7.4	-0.4	19.3	8.0	5.4	1.2	-0.3	4.2	0.3	-0.1	0.5	-0.5	3.8
NEE1	192.0	546.0	25.0	8.9	15.8	11.0	1.0	17.7	3.8	3.3	1.5	0.3	0.3	0.3	-0.1	0.3	0.4	3.6
NEE2	216.0	684.0	74.4	12.7	63.0	15.7	7.9	57.3	13.2	12.7	9.3	0.7	10.6	4.7	-0.3	4.7	1.0	8.6
NEE3	205.0	645.0	49.2	11.5	36.6	9.4	4.3	35.7	7.6	8.2	3.4	0.5	3.0	0.3	0.3	0.3	0.8	5.7
NEE4	162.0	575.0	46.3	13.1	20.5	6.2	1.5	16.3	4.2	3.1	2.4	-0.3	2.0	0.1	-0.1	0.4	-0.5	3.3
NEE5	145.0	615.0	54.3	10.9	30.6	8.3	2.6	19.2	1.4	4.6	1.7	0.2	0.9	0.5	-0.1	0.5	0.3	2.7
NEE6	194.0	714.0	36.0	12.3	35.4	12.4	7.1	47.4	20.1	14.4	5.5	0.9	5.1	0.7	-0.3	0.7	1.3	10.5
NEE7	330.0	912.0	63.6	15.2	62.7	25.0	0.9	57.3	5.3	13.3	1.7	1.1	1.4	0.8	0.3	0.8	1.4	8.6
NF1	163.0	654.0	44.1	8.6	31.2	10.4	-3.9	36.0	15.1	9.8	2.2	-0.6	0.9	0.8	1.2	0.6	0.9	7.2
NF10	140.0	624.0	27.4	11.1	11.5	6.7	1.8	13.5	1.5	3.3	2.0	0.3	1.7	0.9	0.3	0.9	0.5	2.5
NF11	308.0	664.0	70.6	24.8	50.7	12.3	5.6	40.2	2.2	11.9	8.2	0.3	3.2	0.7	1.0	0.7	0.8	6.2
NF2	139.0	564.0	28.9	9.8	11.1	6.4	-0.1	4.6	2.3	0.6	1.1	-0.1	0.9	0.3	-0.1	0.3	0.4	1.3
NF3	130.0	555.0	34.5	10.3	20.1	12.5	-0.8	15.0	3.3	3.6	1.4	-0.1	0.2	0.3	-0.1	0.3	0.3	2.5
NF4	144.0	609.0	22.6	11.0	8.5	5.0	0.6	8.9	1.8	0.2	1.2	-0.1	1.6	0.3	-0.1	0.3	0.2	1.8
NF5	205.0	786.0	38.4	13.2	15.9	7.4	1.1	11.3	3.9	2.0	4.5	-0.1	1.2	0.4	-0.1	0.4	0.4	1.9
NF6	293.0	987.0	50.7	19.6	24.2	10.0	1.0	9.8	1.2	4.0	1.8	0.3	0.5	0.6	-0.1	0.6	0.4	2.4
NF7	195.0	545.0	26.5	10.3	34.5	13.4	7.7	80.4	34.2	22.7	1.6	1.3	5.6	11.5	1.3	11.6	3.9	17.9
NF8	119.0	561.0	48.3	9.2	26.0	8.2	0.9	25.5	1.7	6.8	1.3	0.3	2.2	0.4	-0.1	0.4	0.8	3.7
NF8-R	140.0	636.0	44.4	12.2	26.8	12.3	1.9	30.3	2.0	8.4	2.3	0.5	0.8	0.4	-0.3	0.4	1.1	5.0
NF9	130.0	555.0	26.8	11.5	15.4	5.6	0.8	16.2	3.3	4.0	1.9	0.4	1.7	1.6	-0.1	1.6	0.6	3.1
NFF1	126.0	555.0	44.1	9.5	26.6	17.5	-2.1	23.9	10.0	6.0	1.8	-0.2	1.9	0.4	-0.1	0.8	0.6	3.6
NFF2	345.0	918.0	60.3	17.2	51.6	19.2	10.4	60.0	24.9	18.6	7.6	0.9	6.7	0.7	0.2	0.7	1.5	10.8
NFF3	158.0	714.0	45.8	12.9	18.9	4.3	1.9	8.3	0.8	2.8	4.9	0.2	0.8	0.5	-0.1	0.5	0.3	2.3
NFF4	324.0	780.0	82.8	13.7	80.1	30.6	7.9	94.8	39.0	32.1	6.9	1.4	6.5	0.9	0.3	0.9	3.1	16.4
NFF5	367.0	579.0	74.4	13.0	57.9	21.9	4.6	62.7	26.0	17.6	3.0	0.5	3.4	1.1	1.1	1.4	1.3	9.2
NFF6	252.0	636.0	31.8	11.2	15.0	9.2	1.3	16.1	1.4	3.6	1.7	0.4	1.3	0.2	0.3	0.2	0.6	2.7
NFF6-R	242.0	633.0	30.0	11.5	14.0	8.8	0.7	15.8	1.4	3.8	4.7	0.3	1.2	1.1	-0.3	1.1	0.5	4.5
NFF7	312.0	561.0	72.3	14.1	49.2	12.4	3.4	33.9	7.4	7.3	2.9	0.6	2.6	0.4	-0.1	0.4	0.9	4.9
NFF8	192.0	599.0	46.3	9.5	25.7	7.8	3.3	24.8	4.6	5.3	2.6	0.5	2.0	0.3	0.3	0.8	0.9	4.4
NG1	163.0	654.0	40.8	8.3	14.2	8.6	0.6	4.2	1.9	0.9	0.9	-0.1	0.3	0.3	-0.1	0.3	0.3	1.0
NG10	202.0	681.0	44.1	10.3	20.0	6.1	1.1	15.7	3.5	3.2	2.4	0.2	2.0	0.4	-0.1	0.4	0.4	2.8
NG11	834.0	759.0	71.1	60.3	45.3	11.0	8.3	32.7	1.9	7.9	3.5	-0.1	3.8	0.3	-0.1	0.3	0.5	3.2
NG2	268.0	768.0	47.7	9.5	21.5	7.7	1.1	13.9	1.7	2.2	1.0	0.2	0.6	1.3	-0.1	1.8	1.2	2.3
NG2-R	281.0	804.0	48.3	10.8	19.9	12.0	0.9	12.1	2.2	2.2	0.9	0.2	0.6	0.9	-0.1	0.9	0.7	1.3
NG3	138.0	619.0	27.6	6.9	12.7	8.6	0.5	4.8	2.4	0.4	0.8	-0.1	0.5	0.3	-0.1	0.3	0.3	1.6
NG4	264.0	879.0	42.9	13.4	30.3	10.6	2.5	32.7	6.2	7.5	1.1	0.4	0.7	0.2	-0.1	0.2	0.5	4.7
NG5	207.0	684.0	66.0	9.2	52.8	20.4	6.0	76.8	32.7	21.4	5.5	1.1	4.7	0.9	0.3	0.9	2.9	13.0
NG6	154.0	597.0	34.8	7.5	16.6	11.0	2.3	18.4	3.4	4.3	1.1	0.3	0.4	1.8	0.2	1.8	0.4	3.5
NG7	167.0	621.0	50.1	7.3	29.0	7.4	0.1	8.4	0.9	3.4	0.7	-0.1	0.7	0.4	-0.1	0.4	0.4	1.6
NG8	201.0	756.0	41.7	11.0	19.4	5.9	1.1	13.2	2.7	3.0	0.9	-0.1	0.8	0.4	-0.1	0.4	0.4	2.3
NG9	183.0	636.0	60.0	8.2	40.8	14.1	2.1	44.4	9.4	10.5	2.7	0.6	2.7	0.7	-0.1	0.7	1.2	5.9
NGG1	205.0	660.0	29.8	10.6	15.7	10.9	1.9	14.6	2.8	3.5	1.9	0.3	2.4	0.3	-0.1	0.3	0.6	2.8
NGG10	175.0	372.0	86.4	11.9	56.4	16.9	4.7	46.2	2.5	11.1	6.5	0.7	6.4	0.8	-0.7	0.8	1.0	5.5
NGG2	163.0	558.0	24.1	9.4	11.6	8.0	0.7	4.7	2.3	2.5	1.2	-0.1	1.8	0.4	-0.1	0.4	0.4	2.0
NGG3	519.0	1280.0	124.0	43.9	90.9	35.4	5.7	88.2	36.9	27.1	6.1	1.2	5.7	10.2	0.2	10.2	1.9	13.3
NGG4	209.0	627.0	43.2	9.9	35.4	13.0	3.5	44.4	18.5	14.0	3.6	0.5	3.2	0.6	-0.1	0.6	1.2	7.6
NGG5	180.0	657.0	49.8	10.3	34.5	11.0	3.8	38.1	15.9	11.0	2.5	0.6	2.2	0.6	-0.1	0.6	1.0	6.3
NGG6	136.0	549.0	26.5	10.1	14.4	9.2	2.1	18.0	3.5	4.1	1.6	0.3	2.9	0.4	-0.1	0.4	0.6	3.3
NGG7	333.0	582.0	34.5	10.9	21.1	7.9	2.6	25.0	10.4	7.6	2.6	0.5	2.3	0.5	-0.3	0.5	1.0	5.5
NGG8	135.0	573.0	33.6	11.0	14.1	8.7	0.9	13.8	2.6	2.9	2.7	0.2	2.2	0.4	-0.1	0.4	0.6	3.1
NGG9	138.0	627.0	25.2	11.7	9.0	5.4	1.4	7.2	1.3	4.0	1.1	-0.1	1.2	0.4	-0.1	0.4	0.4	1.8
NH1	129.0	612.0	33.6	11.5	20.1	12.6	2.4	17.2	3.3	4.0	2.3	0.3	2.0	0.5	-0.1	0.5	0.5	3.3
NH10	278.0	582.0	126.0	12.8	130.0	42.6	4.8	91.8	19.2	26.3	3.4	1.1	4.2	5.6	-0.1	5.6	1.4	10.1
NH10-R	312.0	978.0	143.0	16.4	185.0	59.1	6.4	153.0	63.3	54.3	1.6	1.2	1.7	0.7	0.3	0.7	1.9	18.6
NH11	194.0	714.0	37.5	10.6	27.0	9.0	3.7	33.0	13.7	8.8	3.9	-0.6	5.0	0.7	-0.3	0.7	1.2	6.1
NH2	300.0	954.0	49.5	16.1	36.3	11.0	4.1	37.5	15.7	10.3	3.5	0.6	3.1	0.7	-0.1	0.7	1.0	7.4
NH3	180.0	726.0	33.3	11.9	15.0	9.1	0.7	5.2	0.7	0.7	2.1	-0.1	0.9	0.4	-0.1	0.4	0.4	1.3
NH4	170.0	654.0	37.5	10.2	20.4	13.2	3.5	23.1	9.5	5.3	2.5	0.4	2.2	0.4	-0.1	0.4	0.6	3.9
NH5	171.0	636.0	46.2	8.1	34.8	11.9	2.8	43.2	18.1	10.6	1.2	0.4	1.7	0.7	-0.1	0.7	0.9	6.4
NH6	190.0	753.0	57.9	11.4	45.3	13.5	3.4	44.4	2.9	13.0	1.4	0.6	1.2	0.8	0.3	0.8	0.9	8.1
NH7	225.0	729.0	51.0	11.4	46.5	17.2	0.6	72.6	30.0	23.8	5.8	1.0	5.1	1.1	-0.3	1.1	1.7	15.2
NH8	195.0	801.0	47.1	12.2	29.4	10.3	5.0	33.6	2.2	8.8	5.6	0.5	6.7	0.7	0.3	0.7	0.9	6.4
NH9	192.0	750.0	62.1	12.2	43.2	12.8	4.0	40.2	2.7	11.3	3.5	0.6	3.0	0.3	-0.3	0.9	1.1	7.4

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	001 -LA	002 -LA	003 -LB	004 -LA	005 -LB	006 -LB	007 -LA	008 -LB	009 -LB	010 -LB	011 -LA	012 -LB	013 -LBA	014 -LB	015 -LAR	016 -LB	017 -LB	018 -LB
NHH1	126.0	549.0	39.6	11.5	15.2	9.2	1.1	7.1	2.9	0.6	1.3	-0.1	0.9	0.5	-0.1	0.5	0.3	1.5
NHH10	223.0	741.0	44.1	13.0	17.7	11.8	0.6	10.8	2.0	2.5	0.0	-0.1	0.9	0.3	-0.1	0.3	0.4	2.0
NHH11	201.0	651.0	36.0	11.9	17.4	11.5	0.6	12.7	5.3	3.1	1.6	-0.1	0.8	0.5	-0.1	0.5	0.3	2.2
NHH11-R	210.0	672.0	37.2	12.3	18.3	12.0	0.7	13.1	2.5	3.2	1.7	-0.1	1.1	0.4	-0.1	0.4	0.3	2.2
NHH12	147.0	417.0	20.1	9.2	12.1	7.7	2.7	16.7	3.2	3.9	1.8	0.5	1.7	0.3	-0.1	0.3	0.5	2.7
NHH2	162.0	539.0	50.4	11.9	25.5	7.4	5.6	27.3	11.3	6.6	5.8	-0.5	5.8	0.6	0.3	0.6	0.8	4.9
NHH3	135.0	525.0	24.4	9.8	13.1	7.8	2.6	13.9	5.7	3.5	1.5	0.2	1.3	0.4	-0.1	0.4	0.5	2.8
NHH4	145.0	660.0	122.0	11.4	66.6	16.9	3.8	52.8	21.7	12.9	2.6	0.5	2.3	4.7	-0.1	4.7	0.9	7.0
NHH5	218.0	648.0	28.0	9.7	20.5	7.6	4.5	24.3	10.1	7.0	3.4	0.5	3.2	0.4	-0.1	0.4	0.6	4.3
NHH6	210.0	747.0	66.9	12.8	53.7	15.4	4.5	44.1	16.4	13.9	2.8	-0.6	6.7	3.8	-0.1	3.8	0.9	6.1
NHH7	254.0	684.0	55.5	10.2	48.6	15.5	4.6	48.0	19.8	13.1	4.5	0.5	4.2	0.6	-0.1	0.6	0.9	6.3
NHH8	149.0	552.0	30.6	10.1	14.5	8.8	1.3	11.8	2.2	2.8	1.4	-0.1	1.5	0.6	-0.1	0.5	0.5	2.3
NHH9	157.0	606.0	39.0	10.4	14.8	8.9	1.5	9.7	4.0	1.6	1.7	-0.1	0.9	0.3	-0.1	0.3	0.3	1.8
NI1	230.0	906.0	84.3	20.4	78.0	20.6	9.4	109.0	44.4	31.5	12.0	1.2	11.4	1.1	0.3	1.1	2.6	24.1
NI10	261.0	831.0	46.8	12.1	28.4	9.8	4.5	26.9	11.3	6.6	4.4	0.5	5.5	0.6	-0.1	0.6	0.9	5.5
NI2	189.0	507.0	60.6	10.2	39.0	10.0	3.1	30.9	12.7	7.3	1.7	-0.3	1.7	0.4	-0.3	0.4	0.7	4.6
NI3	188.0	618.0	28.2	11.4	16.8	10.8	0.8	16.9	7.1	4.3	1.6	0.2	1.3	0.5	-0.1	0.5	0.6	3.4
NI4	266.0	810.0	57.3	17.2	47.7	15.7	6.9	56.4	23.3	15.9	7.2	0.7	2.6	6.6	0.3	6.6	1.6	11.9
NI5	228.0	714.0	41.1	11.5	36.0	13.5	2.4	39.9	16.5	11.4	1.5	0.4	3.0	0.5	1.1	0.5	1.0	7.7
NI5-R	224.0	696.0	42.0	11.2	38.4	14.0	2.1	41.7	17.3	11.2	1.3	-0.3	1.7	4.7	0.3	4.7	1.1	7.6
NI6	111.0	519.0	22.1	8.3	14.0	9.3	0.3	15.8	3.5	3.5	1.6	0.2	1.4	0.3	0.3	0.3	0.5	3.1
NI7	345.0	915.0	76.5	23.7	78.3	25.8	14.8	113.0	46.5	32.4	22.4	1.3	21.9	13.6	-0.3	19.6	2.7	23.6
NI8	146.0	525.0	26.0	7.4	17.2	11.6	0.9	14.2	3.0	3.6	1.1	0.3	0.9	0.4	-0.1	0.4	0.5	2.8
NI9	187.0	618.0	42.0	9.6	39.9	13.5	3.1	62.4	5.1	19.0	3.6	1.1	3.2	1.4	0.3	1.4	1.9	8.9
NII1	121.0	540.0	6.6	8.2	3.8	1.4	-0.1	4.1	0.7	-0.1	0.7	-0.1	1.0	0.3	-0.1	0.3	0.4	1.4
NII10	257.0	621.0	48.3	12.0	28.4	10.1	3.0	29.1	12.2	7.7	2.7	0.6	2.3	0.6	-0.1	0.6	0.8	5.4
NII11	284.0	1150.0	43.5	24.2	22.4	7.5	3.4	21.7	4.4	5.5	2.5	0.3	2.1	0.6	-0.1	0.6	0.7	4.9
NII1-R	286.0	1170.0	45.3	28.0	23.9	8.3	3.3	22.2	9.2	5.8	2.5	-0.3	1.9	0.6	-0.1	0.6	0.7	4.8
NII12	154.0	627.0	52.2	8.5	36.3	12.1	0.6	33.3	7.7	10.0	1.7	0.5	3.3	0.5	-0.1	0.5	1.1	5.3
NII13	516.0	954.0	124.0	43.8	113.0	35.7	9.5	126.0	52.2	40.5	14.0	1.4	15.4	0.9	-0.3	0.9	3.8	20.7
NII14	209.0	615.0	45.3	12.2	22.8	7.2	1.3	17.8	3.9	3.9	1.5	0.2	1.3	0.3	-0.1	0.3	0.6	2.7
NII2	166.0	561.0	45.9	11.0	33.0	9.8	3.1	37.2	8.4	9.0	3.6	0.4	3.0	0.6	0.3	0.6	0.9	6.2
NII3	315.0	990.0	51.6	22.2	31.5	7.4	9.9	33.9	14.2	7.3	11.7	0.9	11.1	4.8	0.3	4.8	1.0	7.2
NII4	564.0	2080.0	54.0	61.8	28.4	7.1	14.8	28.4	12.0	12.6	19.3	0.6	6.8	4.2	-0.3	4.2	1.1	6.4
NII5	164.0	672.0	40.5	9.6	18.8	11.8	0.9	17.5	4.0	3.8	1.8	0.2	1.5	0.4	-0.1	0.4	0.7	3.2
NII6	254.0	828.0	32.1	15.7	13.6	4.6	3.0	12.9	2.9	1.9	2.4	0.2	2.2	0.3	-0.1	0.6	0.5	2.3
NII7	168.0	687.0	20.0	11.3	8.1	4.7	0.9	3.3	1.3	0.2	1.0	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	0.9
NII8	498.0	1560.0	68.3	34.5	44.7	13.4	10.9	42.3	17.7	11.9	8.3	0.7	7.9	5.6	-0.3	5.6	1.2	8.6
NII9	245.0	828.0	33.6	16.5	14.3	4.7	1.0	11.9	2.4	2.1	1.8	0.2	0.3	0.6	-0.1	0.6	0.6	2.4
NJ1	196.0	684.0	42.5	12.1	39.6	12.7	3.0	49.2	17.7	12.8	3.5	-0.6	3.0	0.5	1.1	0.5	1.2	7.2
NJ2	168.0	582.0	32.4	9.1	29.3	10.7	3.6	39.9	16.6	10.9	4.4	0.6	4.4	0.3	-0.1	0.3	1.1	6.9
NJ3	157.0	630.0	41.7	10.8	42.9	18.9	4.2	48.9	20.3	14.8	5.7	0.6	5.4	4.5	-0.1	4.5	1.2	7.8
NJ4	131.0	170.0	29.6	8.6	24.2	8.6	1.0	29.8	12.4	8.1	2.6	0.5	2.5	0.4	-0.1	0.4	0.9	5.2
NJ5	164.0	522.0	25.1	9.0	14.9	9.1	0.9	17.9	1.8	4.9	3.9	-0.3	1.8	0.3	-0.1	0.6	0.5	2.8
NJ6	137.0	177.0	35.1	9.3	23.0	6.7	2.1	10.2	4.6	4.4	2.2	0.2	2.0	0.3	-0.1	0.3	0.5	3.5
NJ7	201.0	771.0	43.7	13.7	43.8	18.3	3.6	54.0	22.3	14.4	7.3	0.6	3.7	0.7	-0.3	0.7	1.4	10.5
NJJ1	156.0	681.0	42.9	9.0	22.2	7.0	1.6	22.5	9.3	5.7	1.3	0.4	1.1	0.3	-0.1	0.3	0.8	4.2
NJJ10	543.0	2390.0	228.0	33.4	432.0	106.0	51.3	594.0	243.0	481.0	69.2	1.7	66.0	2.5	1.2	2.5	8.2	77.7
NJJ11	173.0	666.0	46.5	10.5	19.0	6.2	0.7	15.0	2.1	3.5	1.5	0.2	0.5	0.9	-0.1	0.9	0.9	2.0
NJJ12	276.0	753.0	83.7	17.9	56.7	18.1	4.0	49.5	3.1	13.8	8.9	0.6	4.9	0.7	-0.3	0.7	1.3	6.8
NJJ13	402.0	1370.0	75.6	26.6	54.0	15.0	9.8	48.3	20.2	12.5	9.7	0.9	11.1	0.5	0.3	0.5	1.1	8.7
NJJ14	165.0	726.0	36.3	10.2	12.6	7.8	0.7	10.3	2.2	2.2	1.2	-0.1	1.5	0.3	-0.1	0.6	0.3	2.1
NJJ2	246.0	873.0	46.5	13.8	29.9	11.1	4.3	44.1	7.5	7.9	4.1	0.8	1.0	0.5	0.3	0.5	0.8	5.9
NJJ3	223.0	741.0	55.7	11.7	34.2	10.3	2.7	25.7	5.8	5.6	4.1	0.3	1.2	0.5	-0.1	0.5	0.5	3.6
NJJ4	306.0	1080.0	82.8	21.1	57.6	11.7	9.2	49.2	13.1	13.2	16.1	0.4	15.4	0.7	0.3	0.7	0.9	6.1
NJJ5	267.0	966.0	110.0	17.3	86.7	29.6	10.5	69.3	15.8	15.5	5.8	1.1	2.2	5.0	0.3	5.0	1.1	7.7
NJJ6	167.0	657.0	39.9	10.5	20.7	6.5	2.6	22.4	5.3	4.8	3.1	0.4	3.6	0.3	-0.1	0.3	0.5	3.9
NJJ7	124.0	209.0	19.0	7.4	8.1	5.5	1.5	7.6	1.5	-0.1	0.1	-0.1	0.3	0.2	-0.1	0.3	0.3	2.1
NJJ8	286.0	993.0	40.2	19.5	22.7	6.6	6.6	23.1	5.1	3.9	5.3	0.5	6.3	0.4	0.3	0.4	0.6	3.2
NJJ8-R	278.0	1010.0	39.8	18.9	23.0	6.9	7.8	25.1	10.4	5.7	7.9	-0.6	9.1	0.5	0.3	0.6	0.7	4.3
NJJ9	546.0	1990.0	76.8	52.8	61.2	20.8	29.9	107.0	8.9	25.5	41.1	1.7	40.8	4.6	0.4	4.6	3.4	14.9
NK1	186.0	619.0	25.2	8.1	28.3	9.5	1.0	44.1	18.2	12.8	2.3	0.5	1.9	5.0	-0.1	6.0	1.0	7.6
NK2	168.0	651.0	28.3	11.0	13.6	8.7	0.9	14.6	6.0	3.3	1.7	0.2	1.1	0.3	-0.1	0.3	0.5	3.2
NK3	166.0	576.0	30.6	8.9	34.8	12.2	0.9	30.0	16.3	10.4	2.2	0.6	1.8	3.8	-0.1	0.7	0.7	5.8
NK4	194.0	681.0	30.6	9.9	16.2	10.4	0.9	13.1	2.0	2.9	1.3	0.3	1.4	2.5	-0.1	2.5	0.5	3.0
NK4-R	199.0	678.0	30.0	9.9	15.8	10.7	0.9	12.2	2.3	2.9	4.1	0.2	1.3	0.8	-0.1	0.8	0.4	2.9
NK5	135.0	510.0	21.5	9.3	21.2	7.0	3.4	34.8	14.4	9.2	2.6	0.5	2.3	0.3	-0.1	0.3	0.9	5.9

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	001-LA	002-LA	003-LB	004-LA	005-LB	006-LB	007-LA	008-LB	009-LB	010-LB	011-LA	012-LB	013-LBA	014-LB	015-LAR	016-LB	017-LB	018-LB
NK6	172.0	576.0	26.6	9.5	18.6	6.1	0.4	20.1	3.7	4.9	1.6	0.2	1.9	0.4	-0.1	0.4	0.5	3.8
NK7	184.0	651.0	47.7	11.5	34.2	10.0	2.5	29.0	6.4	6.4	2.3	0.3	2.9	0.4	-0.1	0.4	0.6	4.5
NKK1	247.0	675.0	24.6	9.9	8.0	5.0	0.8	6.9	1.2	1.5	0.9	-0.1	0.3	0.4	-0.1	0.4	0.5	1.4
NKK10	173.0	708.0	43.2	10.3	18.0	11.6	0.8	14.0	2.6	3.0	1.5	-0.1	1.0	0.4	-0.1	0.4	0.5	2.1
NKK11	297.0	816.0	57.0	13.1	35.7	11.2	5.0	36.0	15.2	8.9	5.8	-0.0	5.0	0.4	0.3	0.4	0.8	5.8
NKK2	185.0	735.0	29.6	11.7	12.3	7.5	0.9	11.3	2.1	2.6	2.0	-0.1	0.5	0.4	-0.1	0.4	0.3	2.4
NKK3	248.0	634.0	47.7	14.4	23.6	10.6	5.3	20.6	4.7	4.4	0.5	0.5	5.9	0.5	-0.1	0.5	0.7	3.7
NKK4	354.0	1250.0	63.9	27.1	40.2	10.6	11.4	42.0	17.4	9.5	18.0	0.9	12.8	5.2	0.4	5.2	1.0	7.7
NKK5	463.0	1450.0	57.9	35.1	39.0	11.4	16.8	46.9	18.9	12.8	21.2	1.0	20.4	6.7	0.3	6.7	1.4	9.3
NKK6	297.0	954.0	56.4	16.1	32.1	10.2	5.9	35.1	3.5	7.7	7.2	0.4	8.1	0.4	0.3	0.4	0.6	4.3
NKK7	272.0	604.0	43.6	12.9	27.5	7.4	3.5	19.6	4.1	4.1	8.5	0.3	4.1	0.3	-0.1	0.3	0.5	3.0
NKK8	165.0	723.0	39.9	11.1	16.2	10.5	0.8	13.4	2.9	3.0	1.6	-0.1	2.0	0.3	-0.1	0.3	0.6	2.8
NKK9	179.0	303.0	44.1	9.5	24.6	8.8	2.2	25.7	5.8	6.4	2.8	0.3	1.8	0.4	-0.1	0.4	0.8	4.2
NL1	170.0	645.0	47.7	10.2	19.0	11.9	0.5	6.7	2.7	2.3	1.1	-0.1	0.9	0.3	-0.1	0.3	0.3	1.7
NL2	172.0	669.0	26.0	9.5	9.2	5.6	0.9	10.9	0.9	0.9	0.9	-0.1	1.1	0.2	-0.1	0.2	0.8	1.8
NL3	119.0	183.0	18.7	7.5	6.8	4.4	-0.1	5.9	1.4	0.7	0.3	-0.1	0.4	0.2	-0.1	0.2	0.3	1.6
NL3-R	166.0	753.0	39.9	11.5	16.9	10.5	1.2	14.6	2.9	3.3	0.9	0.2	1.0	0.5	-0.1	0.5	0.5	3.1
NL4	187.0	699.0	48.0	11.5	18.7	5.5	0.6	6.2	2.5	0.3	1.3	-0.1	0.9	0.4	-0.1	0.4	0.2	1.8
NLL1	259.0	921.0	93.8	44.4	59.1	17.9	0.4	39.3	2.9	12.1	0.9	-0.5	0.3	0.3	-0.1	0.3	0.7	5.5
NLL10	209.0	789.0	38.4	11.9	16.0	10.1	2.1	13.0	2.6	2.8	2.1	0.3	2.7	0.3	-0.1	0.3	0.5	2.4
NLL11	169.0	563.0	34.5	8.7	15.4	10.2	2.3	20.5	9.7	6.0	2.7	-0.5	2.4	0.3	-0.1	0.8	1.0	4.7
NLL12	144.0	615.0	41.7	7.7	19.4	6.9	3.1	26.9	11.1	7.0	1.7	0.3	1.3	0.3	-0.1	0.3	0.7	4.2
NLL2	135.0	699.0	35.0	11.1	16.4	5.3	0.7	14.2	1.3	3.4	4.5	-0.1	1.6	1.2	-0.1	1.2	0.8	2.3
NLL3	156.0	726.0	54.3	11.9	28.2	6.7	0.8	12.3	1.3	4.4	1.1	-0.1	0.4	0.3	-0.1	0.3	0.4	2.2
NLL4	200.0	744.0	68.1	10.3	53.1	18.2	1.0	49.5	20.3	13.4	1.4	-0.5	7.1	0.4	-0.1	0.4	1.1	8.3
NLL5	600.0	1670.0	56.7	36.6	36.9	10.5	13.3	40.8	8.9	7.9	15.4	0.9	10.9	0.6	0.3	0.6	0.9	6.8
NLL6	393.0	1400.0	61.2	24.5	34.8	9.4	11.5	29.2	7.1	6.4	10.7	0.6	6.6	3.5	-0.3	3.5	0.8	5.5
NLL7	633.0	2070.0	127.0	66.9	130.0	35.7	32.4	158.0	65.7	47.1	41.4	2.5	43.8	1.5	0.4	1.5	4.7	27.8
NLL8	215.0	837.0	48.0	13.6	21.6	7.3	0.6	16.4	3.3	3.5	2.0	0.2	1.7	0.4	-0.1	0.4	0.3	2.5
NLL9	300.0	924.0	67.8	22.6	55.2	15.3	16.8	60.3	24.8	14.3	21.3	1.4	24.3	1.0	0.4	1.0	2.0	10.4
NLL9-R	315.0	866.0	68.4	24.5	57.0	16.2	19.1	61.2	26.8	17.0	24.9	1.1	23.5	1.1	-0.8	1.1	1.4	11.3
NM1	184.0	675.0	34.8	10.6	32.1	10.7	9.7	62.4	25.8	16.0	9.6	1.0	11.1	0.9	0.4	0.9	1.1	9.2
NM2	127.0	599.0	46.3	7.9	26.7	8.0	0.4	28.4	2.0	7.4	1.7	-0.5	1.8	0.6	-0.1	0.6	0.6	4.0
NMM1	188.0	729.0	47.7	13.1	28.3	9.1	2.4	29.5	3.3	8.9	2.3	0.5	2.8	0.4	-0.1	0.4	0.7	4.4
NMM10	312.0	1450.0	77.1	88.7	80.1	23.6	29.8	103.0	42.3	26.8	34.5	1.3	36.9	1.4	-0.9	4.4	1.7	16.0
NMM2	287.0	1030.0	53.4	14.3	31.5	9.4	3.3	36.9	8.9	7.2	4.0	0.5	1.8	0.8	-0.1	0.8	0.8	6.1
NMM3	245.0	900.0	46.8	14.2	28.4	8.5	3.0	24.4	4.7	5.9	2.6	1.0	2.9	0.2	-0.1	0.2	0.9	4.2
NMM3-R	244.0	891.0	44.1	14.2	24.6	8.9	3.0	20.4	1.4	5.3	2.3	0.3	2.7	0.5	-0.1	0.5	0.5	3.5
NMM4	198.0	732.0	60.0	10.3	29.6	7.6	2.3	9.6	4.2	3.8	4.1	0.2	0.9	0.4	-0.1	0.4	0.3	2.5
NMM5	215.0	732.0	40.2	11.1	23.5	15.5	1.1	16.7	3.6	3.9	1.3	0.2	1.2	1.2	-0.1	1.2	0.3	2.4
NMM6	609.0	1870.0	115.0	51.9	91.2	29.8	10.7	113.0	7.0	28.7	9.3	1.2	10.9	2.2	-1.2	2.2	1.7	13.1
NMM7	154.0	558.0	27.1	7.3	11.5	7.5	-0.1	4.2	2.0	0.2	0.6	-0.1	0.2	0.3	-0.1	0.3	0.2	1.4
NMM8	260.0	1400.0	56.1	33.6	41.1	11.1	29.7	43.5	18.1	8.5	43.8	1.0	42.3	0.8	-0.8	0.8	1.1	6.8
NMM9	221.0	813.0	83.1	14.0	61.8	16.7	6.5	59.7	24.6	15.9	8.6	0.7	8.3	0.7	0.3	0.7	1.2	9.4
NN1	118.0	599.0	33.6	9.1	9.8	6.2	0.6	4.4	1.7	0.2	0.0	-0.1	0.7	0.3	-0.1	0.3	0.3	1.2
NNN1	222.0	765.0	42.0	12.5	19.3	5.4	3.2	12.8	3.0	2.1	1.4	-0.1	1.2	0.5	-0.1	0.5	0.4	2.1
NNN10	573.0	795.0	111.0	40.5	121.0	37.8	0.4	56.4	5.0	29.7	2.3	-0.3	0.8	0.8	-0.1	0.8	0.9	9.2
NNN2	188.0	708.0	35.1	11.1	15.0	4.8	0.8	10.6	2.7	0.3	1.8	-0.1	1.5	0.9	-0.1	0.9	0.3	1.8
NNN3	193.0	669.0	43.2	7.4	16.9	10.5	0.4	5.1	2.1	0.2	0.9	-0.1	0.8	0.1	-0.1	0.1	-0.1	1.0
NNN4	281.0	858.0	66.9	13.9	49.2	16.3	5.7	48.0	19.9	14.7	6.1	0.6	5.7	0.8	0.3	0.8	1.6	9.1
NNN5	247.0	822.0	34.2	12.2	16.1	9.5	2.8	9.1	2.3	0.2	1.4	-0.1	1.3	0.3	-0.1	0.3	0.4	1.8
NNN6	215.0	759.0	31.8	8.8	12.0	6.9	-0.1	3.6	1.5	0.3	0.9	-0.1	0.5	0.3	-0.1	0.3	0.3	1.0
NNN7	161.0	600.0	40.8	7.4	15.4	9.6	-0.1	3.4	2.0	0.3	1.0	-0.1	0.6	0.2	-0.1	0.2	0.3	1.1
NNN8	220.0	741.0	51.6	12.6	47.4	15.8	11.4	85.5	35.1	24.3	13.4	1.1	12.2	1.3	0.4	1.3	1.6	11.9
NNN9	321.0	1100.0	102.0	28.7	85.8	25.1	18.2	81.3	35.4	22.2	18.9	1.3	21.0	1.1	-0.3	1.1	2.2	11.8
NO1	204.0	759.0	44.4	11.0	22.6	6.6	1.1	19.0	7.9	4.2	1.1	0.3	0.8	0.4	-0.1	0.4	0.5	3.2
NO2	101.0	465.0	11.9	8.1	1.8	0.6	-0.1	1.2	0.5	0.5	0.6	-0.1	1.1	-0.1	-0.1	-0.1	-0.1	0.7
NOO1	624.0	1840.0	46.5	49.5	18.4	5.5	7.1	10.1	2.3	3.2	3.4	-0.1	1.0	1.0	-0.1	1.0	0.5	2.0
NOO2	194.0	726.0	35.0	10.4	17.9	7.4	2.8	16.6	6.8	4.3	3.2	0.3	2.7	0.2	-0.1	0.2	0.0	3.6
NOO3	477.0	1550.0	84.3	41.7	46.5	11.1	8.1	29.9	1.8	7.4	3.1	-0.1	3.5	1.1	-0.1	1.1	0.4	3.4
NOO4	287.0	828.0	45.3	12.1	25.2	9.1	4.5	22.8	9.5	6.5	2.9	0.5	0.5	0.5	-0.3	0.5	1.0	4.7
NOO5	252.0	786.0	34.5	12.4	11.2	5.5	1.1	7.7	1.7	-0.1	1.7	-0.1	1.5	0.3	-0.1	0.3	0.1	1.4
NOO6	195.0	744.0	39.6	9.9	12.7	8.0	-0.1	4.6	1.8	0.4	0.9	-0.1	0.6	0.4	-0.1	0.4	0.3	1.5
NOO6-R	197.0	720.0	32.7	10.1	12.2	7.7	0.5	3.5	2.4	0.6	0.9	-0.1	0.6	0.4	-0.1	0.4	0.4	1.3
NOO7	156.0	645.0	33.2	10.3	7.0	4.1	0.8	5.9	2.1	0.5	1.1	-0.1	0.9	0.3	-0.1	0.3	0.3	0.9
NOO8	256.0	804.0	50.4	17.1	36.6	9.5	12.6	38.7	16.1	8.9	18.8	0.6	17.3	0.9	0.3	0.9	1.0	7.7
NP1	197.0	741.0	52.2	17.5	28.1	17.9	2.7	18.2	4.7	4.1	1.1	0.2	0.8	1.3	-0.1	1.3	0.4	2.8

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A14-06865

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	001 -LA	002 -LA	003 -LB	004 -LA	005 -LB	006 -LB	007 -LA	008 -LB	009 -LB	010 -LB	011 -LA	012 -LB	013 -LBA	014 -LB	015 -LAR	016 -LB	017 -LB	018 -LB
NP2	152.0	648.0	48.0	9.4	21.7	14.0	1.0	6.7	3.1	0.4	0.9	-0.1	0.7	0.5	-0.1	0.5	0.3	1.6
NR3	209.0	789.0	43.8	11.0	20.6	13.6	0.7	5.8	0.9	2.6	0.9	-0.1	0.4	0.4	-0.1	0.4	0.4	1.7
NP4	175.0	729.0	59.4	10.6	50.7	21.2	8.6	77.4	32.4	22.8	10.6	1.1	10.2	1.1	1.5	1.1	2.3	15.9
NPP4	200.0	702.0	35.7	8.6	8.5	5.0	0.1	2.5	1.2	0.2	1.3	-0.1	0.7	0.3	-0.1	0.3	0.3	1.0
NPP1-R	190.0	699.0	34.5	7.6	8.1	4.8	-0.1	2.3	0.9	0.2	1.2	-0.1	0.8	0.4	-0.1	0.4	0.4	0.8
NPP2	369.0	1110.0	50.4	19.8	29.3	9.4	3.8	26.2	3.6	6.4	3.0	-0.3	2.8	3.6	-0.1	3.6	0.8	5.1
NPP3	275.0	612.0	45.0	11.5	23.1	6.7	1.0	16.5	6.9	4.1	2.0	-0.1	1.6	0.3	-0.1	0.3	0.3	3.0
NPP4	298.0	1080.0	81.0	24.2	77.1	23.1	11.1	75.0	31.2	23.3	8.5	1.1	7.4	0.8	-0.3	0.8	1.4	10.9
NQ1	148.0	657.0	51.0	8.6	25.8	7.8	2.5	28.4	2.2	7.7	1.1	0.3	0.3	0.4	0.2	0.4	0.4	3.9
NQ2	224.0	921.0	37.2	16.7	17.3	11.1	0.7	13.6	2.9	2.7	1.9	-0.1	2.2	0.8	-0.1	0.8	0.8	2.5
NQ2-R	218.0	876.0	36.9	15.7	17.6	5.3	3.5	15.5	3.4	2.8	2.1	0.2	2.5	0.4	-0.1	0.4	0.4	2.7
NQ3	188.0	723.0	38.1	11.2	12.8	5.1	0.5	5.6	2.5	0.2	1.4	-0.1	0.8	0.4	-0.1	0.4	0.4	1.2
NQ4	140.0	627.0	61.5	8.6	35.4	9.7	1.0	26.6	6.2	6.2	2.2	0.4	1.9	0.4	-0.1	0.4	0.6	3.7
NQ5	202.0	771.0	83.4	11.0	70.5	21.4	3.0	60.6	13.1	15.4	1.3	0.7	0.7	0.5	-0.1	0.5	0.9	7.1
NQ6	137.0	585.0	38.4	9.9	14.9	9.5	-0.1	4.6	1.9	0.4	0.9	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	1.1
NQ7	147.0	627.0	42.3	8.1	20.8	7.7	4.1	35.7	14.7	10.2	3.5	-0.3	3.5	0.7	-0.1	0.7	1.0	6.2
NQ8	179.0	669.0	39.9	9.3	23.9	8.3	2.3	25.0	10.4	6.9	2.2	0.3	2.1	0.4	-0.1	0.4	0.6	3.8
NQQ1	166.0	663.0	36.6	9.3	17.0	10.9	1.2	18.2	7.6	4.5	2.0	0.4	1.7	0.3	-0.1	0.3	0.5	3.3
NR1	164.0	648.0	22.6	7.1	10.4	6.4	0.5	3.8	1.8	0.3	0.9	-0.1	0.4	0.3	-0.1	0.3	0.3	1.1
NR11	218.0	800.0	51.9	13.1	30.3	9.6	5.8	30.9	2.2	8.3	4.2	0.8	5.0	1.3	-0.3	1.3	0.9	5.3
NR12	273.0	843.0	78.3	13.8	63.6	22.1	4.0	55.5	23.3	16.7	2.8	0.7	2.3	0.6	1.1	0.6	1.2	8.8
NR13	627.0	261.0	155.0	20.8	198.0	54.9	7.0	145.0	60.0	44.7	19.6	1.1	13.1	1.0	-0.3	1.0	2.3	18.7
NR14	166.0	633.0	34.8	10.2	13.8	4.3	0.6	16.3	1.5	3.0	1.9	-0.1	1.7	1.5	-0.1	1.5	0.8	2.1
NR2	191.0	690.0	42.0	10.9	22.1	10.6	1.5	17.8	3.7	4.0	2.1	0.3	2.2	0.4	-0.1	0.4	0.5	3.2
NR3	528.0	1630.0	103.0	30.3	102.0	35.7	4.4	87.0	36.3	28.0	2.2	1.0	1.9	8.1	-0.1	8.1	1.6	13.7
NR4	168.0	642.0	34.5	12.0	19.1	6.2	0.6	6.8	3.1	3.5	0.9	0.2	0.8	0.3	-0.1	0.3	0.3	2.6
NR5	208.0	708.0	43.5	9.6	20.3	13.2	1.0	7.1	3.1	2.9	1.0	-0.1	0.8	0.3	-0.1	0.3	0.2	2.0
NR6	229.0	786.0	41.7	11.8	20.1	14.2	0.6	13.7	2.8	3.3	0.9	-0.1	0.8	0.3	-0.1	0.3	1.1	2.1
NR7	224.0	738.0	49.8	11.8	30.9	10.5	4.4	30.0	12.6	8.0	4.6	0.6	1.5	0.5	0.3	0.5	1.0	5.9
NR8	157.0	636.0	40.2	10.8	18.4	12.2	0.6	11.7	1.6	2.9	1.6	-0.1	1.4	0.3	-0.1	0.3	0.3	1.8
NS1	139.0	579.0	72.3	8.0	45.9	13.9	0.8	14.0	2.0	6.7	0.9	0.2	0.2	1.5	-0.1	1.5	0.3	2.8
NS10	199.0	672.0	61.5	9.4	54.3	16.0	6.9	57.8	23.9	14.5	6.2	-0.5	6.3	0.6	-0.4	0.6	1.0	6.9
NS11	207.0	693.0	39.0	10.9	23.1	7.8	1.5	23.6	9.9	6.4	1.9	0.3	2.3	2.3	-0.1	2.3	0.7	4.5
NS12	159.0	183.0	43.5	9.1	32.1	9.9	4.4	39.3	16.3	9.5	0.4	0.4	3.2	4.4	-0.1	4.4	0.9	6.3
NS13	244.0	948.0	47.1	21.0	43.2	15.4	5.6	48.3	20.2	15.0	4.5	0.8	5.4	0.6	1.1	0.6	1.3	8.9
NS14	312.0	866.0	56.3	15.8	63.3	29.7	10.0	86.1	35.7	27.1	8.4	1.3	7.9	1.1	1.3	1.1	3.1	16.2
NS2	143.0	176.0	60.6	9.0	30.9	8.4	0.8	19.1	1.3	4.4	1.6	0.4	1.2	0.4	-0.1	0.4	0.3	2.5
NS3	128.0	200.0	62.4	6.0	61.8	26.6	3.4	69.3	28.5	20.0	0.8	0.3	1.9	6.1	-0.1	6.1	2.0	9.5
NS4	197.0	699.0	46.5	7.6	21.4	13.9	0.8	6.5	3.0	3.2	1.2	-0.1	0.5	1.2	-0.1	1.2	0.3	2.4
NS4-R	177.0	684.0	45.3	7.9	20.2	13.6	0.6	12.0	3.2	2.8	0.9	-0.1	0.5	1.1	-0.1	1.1	0.3	2.2
NS5	137.0	597.0	29.7	10.2	10.2	6.1	0.6	2.9	1.3	0.2	1.2	-0.1	0.4	0.5	-0.1	0.5	0.2	1.5
NS6	193.0	645.0	23.1	10.9	12.3	8.0	0.7	4.6	2.1	2.4	0.9	-0.1	0.7	0.4	-0.1	0.4	0.2	2.1
NS7	242.0	858.0	32.4	15.7	16.9	5.1	0.8	15.5	3.1	3.5	2.2	0.2	0.8	0.4	-0.1	0.4	0.4	3.2
NS8	165.0	660.0	64.8	11.2	47.4	18.2	3.9	40.5	2.9	12.5	3.6	-0.5	4.3	4.1	-0.3	4.1	1.0	7.4
NS9	126.0	561.0	38.4	9.1	16.4	10.0	0.7	5.0	2.4	0.4	0.8	-0.1	0.6	0.4	-0.1	0.4	0.2	1.4
NT1	510.0	483.0	86.4	20.0	64.2	9.0	0.8	44.1	2.8	12.8	5.7	-0.1	1.4	0.5	-0.1	0.5	0.5	4.5
NT10	213.0	654.0	113.0	11.2	80.1	22.8	1.7	48.9	10.3	12.1	1.1	0.4	0.7	0.4	-0.1	0.4	1.1	5.4
NT11	298.0	864.0	52.2	10.1	20.7	13.6	1.0	6.3	3.1	2.9	1.4	-0.1	0.3	0.5	-0.1	0.5	0.3	1.9
NT12	226.0	600.0	41.4	8.3	34.8	10.9	3.2	36.3	14.9	9.2	1.1	0.6	2.3	0.5	1.1	0.5	1.2	5.3
NT13	214.0	711.0	34.6	12.2	22.1	14.9	2.1	17.0	3.4	3.9	1.2	0.2	1.0	0.4	-0.1	0.4	0.4	2.7
NT14	134.0	534.0	24.9	8.4	14.4	8.9	0.3	16.0	3.2	3.6	1.4	0.2	1.4	0.4	-0.1	0.4	0.4	3.0
NT15	226.0	881.0	106.0	10.7	95.7	32.1	9.0	90.8	37.2	28.8	8.4	1.3	13	0.7	-0.3	0.7	2.2	12.6
NT16	162.0	723.0	46.5	12.7	35.4	12.3	2.9	40.5	16.7	12.7	2.9	0.6	2.6	0.6	0.3	0.6	1.1	7.8
NT17	143.0	573.0	24.0	9.5	13.0	8.2	0.5	12.4	2.1	3.3	4.5	-0.1	1.4	0.6	-0.1	0.6	0.4	2.7
NT18	256.0	876.0	51.0	22.1	43.8	10.5	1.6	15.7	1.4	6.1	1.2	0.4	1.9	0.5	-0.1	0.5	0.4	2.5
NT2	480.0	954.0	69.0	13.0	58.2	18.9	0.8	46.5	2.7	13.3	2.0	0.5	1.8	1.3	-0.2	1.3	0.6	5.0
NT3	157.0	600.0	48.3	11.6	17.4	4.7	0.8	4.5	2.1	0.3	0.9	-0.1	0.7	0.4	-0.1	0.4	0.4	1.1
NT4	235.0	732.0	57.6	10.9	23.8	15.8	0.7	2.6	2.6	4.0	-0.1	0.7	0.3	-0.1	0.3	0.3		1.3
NT5	180.0	621.0	33.6	9.7	15.6	10.3	0.7	11.2	2.0	2.6	0.8	-0.1	0.6	0.5	-0.1	0.5	1.0	1.9
NT6-R	176.0	624.0	32.4	10.6	16.5	10.6	1.0	12.4	2.6	2.9	0.9	-0.1	0.7	0.4	-0.1	0.4	0.3	2.3
NT6	236.0	816.0	38.7	13.9	14.8	5.1	0.8	11.1	2.2	2.2	1.0	-0.1	0.6	0.4	-0.1	0.4	-0.1	1.9
NT7	286.0	1090.0	63.6	25.0	61.8	17.0	2.8	42.6	2.7	11.4	4.2	0.3	0.4	3.1	-0.1	3.1	0.7	5.3
NT8	268.0	1000.0	43.8	24.3	31.5	16.1	5.2	37.5	15.8	11.0	4.5	0.3	1.8	5.5	1.2	5.5	1.2	8.2
NT9	208.0	795.0	46.9	10.7	23.1	7.2	2.6	13.9	2.6	3.6	1.8	-0.1	0.3	0.8	-0.1	0.8	0.3	2.1
NU1	209.0	666.0	72.6	8.3	34.5	11.7	1.2	15.7	1.5	3.6	1.0	0.2	0.9	0.6	-0.1	0.6	1.4	1.7
NU10	339.0	879.0	43.6	18.6	24.7	7.2	2.6	23.0	5.2	5.4	2.3	0.4	0.7	0.2	-1.0	0.2	0.9	4.2
NU11	702.0	798.0	46.8	18.3	26.4	8.3	1.0	19.1	4.9	3.8	1.8	0.4	2.2	0.4	-0.1	0.4	0.5	2.8

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	001-LA	002-LA	003-LB	004-LA	005-LB	006-LB	007-LA	008-LB	009-LB	010-LB	011-LA	012-LB	013-LBA	014-LB	015-LAR	016-LB	017-LB	018-LB
NU12	134.0	582.0	30.0	10.6	13.6	7.8	2.9	10.7	0.9	1.7	1.7	-0.1	1.4	0.1	-0.1	0.1	0.7	1.9
NU12-R	126.0	555.0	29.6	10.2	12.8	7.5	0.2	10.0	2.5	1.7	1.5	-0.1	0.8	0.4	-0.1	0.4	0.7	1.6
NU13	211.0	663.0	41.1	14.0	23.3	9.3	5.2	20.6	9.8	8.8	6.2	1.1	2.5	2.7	-0.3	2.7	1.0	5.4
NU14	202.0	588.0	25.7	9.6	16.9	10.3	1.0	20.5	8.6	4.7	1.8	0.4	5.6	0.5	-0.1	0.5	0.7	3.8
NU15	156.0	609.0	42.6	9.8	24.0	7.2	3.1	24.7	16.3	6.7	2.2	0.3	1.8	0.3	-0.1	0.6	0.8	4.6
NU16	224.0	834.0	43.5	19.2	28.1	7.5	5.6	32.1	13.2	8.7	5.0	0.6	4.8	4.7	0.3	4.7	1.5	7.6
NU17	248.0	642.0	34.5	10.2	25.9	9.3	3.5	33.6	14.1	9.8	4.2	0.6	4.0	0.8	-0.3	0.3	1.1	6.6
NU18	188.0	741.0	41.7	14.9	21.1	8.7	1.0	20.5	2.0	4.7	2.2	0.6	2.1	0.9	-0.1	0.9	0.7	2.9
NU19	207.0	780.0	36.3	9.0	28.0	10.5	1.0	36.0	13.2	9.4	2.4	0.5	1.8	0.4	-0.1	0.4	0.8	5.2
NU2	348.0	921.0	168.0	23.3	123.0	32.4	12.7	121.0	6.6	31.5	17.1	1.3	12.8	1.6	0.3	1.6	1.8	13.4
NU3	143.0	597.0	69.6	10.1	33.9	8.9	0.6	13.5	1.5	4.9	1.4	-0.1	0.7	0.3	-0.1	0.3	0.7	2.2
NU4	435.0	1260.0	78.3	26.9	47.4	12.7	6.1	45.6	19.0	12.2	5.5	0.8	6.3	0.6	0.3	0.6	1.3	8.9
NU5	204.0	705.0	39.3	9.5	13.7	9.0	0.7	7.4	1.7	0.3	0.6	-0.1	0.3	0.3	-0.1	0.6	0.4	1.2
NU6	168.0	654.0	27.4	12.8	10.9	4.5	0.7	8.8	1.7	-0.1	1.1	-0.1	0.4	0.4	-0.1	0.4	0.8	1.5
NU7	177.0	645.0	30.0	10.9	13.5	7.8	0.9	11.1	4.6	2.6	4.7	-0.1	2.3	0.7	-0.1	0.7	0.4	2.2
NU8	197.0	813.0	27.2	16.1	14.5	5.8	4.9	19.5	8.0	5.2	3.4	0.7	2.9	0.3	-0.1	0.3	1.0	4.6
NU9	272.0	843.0	49.2	15.6	34.2	11.9	2.5	37.8	2.6	12.1	0.7	0.3	2.0	0.4	-1.1	0.4	1.1	6.4
NV1	405.0	1160.0	89.7	31.8	78.9	19.2	15.2	78.3	18.2	14.9	13.4	1.0	12.3	0.8	0.3	0.8	1.4	9.5
NV10	303.0	978.0	89.2	20.2	61.2	20.5	1.9	31.2	3.2	10.6	4.5	0.4	1.7	3.0	-0.1	0.3	0.6	4.4
NV11	211.0	564.0	37.8	8.4	28.8	8.3	0.5	24.0	4.3	5.5	1.1	0.4	0.8	0.5	-0.1	0.5	0.7	3.5
NV12	232.0	612.0	58.1	10.7	45.3	12.7	2.9	36.3	7.4	7.6	1.2	0.4	2.5	0.4	-0.1	0.4	0.7	4.5
NV13	289.0	657.0	55.8	11.7	60.6	22.9	6.8	74.1	30.3	19.7	6.0	0.7	5.6	5.9	0.3	5.9	1.3	9.0
NV14	198.0	631.0	30.0	8.1	25.0	8.5	3.2	31.6	42.9	8.3	4.3	0.4	1.4	0.5	-0.1	0.5	0.7	5.0
NV15	270.0	453.0	45.3	9.9	33.0	12.6	1.9	29.6	6.0	7.1	2.1	0.5	1.8	0.5	-0.1	0.5	0.7	3.6
NV16	222.0	893.0	64.2	9.2	67.5	24.3	5.3	84.3	36.3	24.9	3.6	0.6	2.9	1.0	0.3	1.0	1.8	13.4
NV17	289.0	825.0	110.0	14.7	91.5	27.7	3.7	66.0	4.9	20.9	2.5	0.3	2.8	0.5	-0.1	0.5	1.2	9.2
NV18	292.0	693.0	85.8	10.3	72.0	19.5	3.2	51.3	9.8	12.7	3.0	0.8	2.7	0.6	-0.1	0.5	1.2	6.3
NV2	429.0	1060.0	81.9	20.1	56.7	19.6	8.3	47.4	10.9	9.8	5.6	0.3	5.0	0.7	0.3	0.7	0.9	6.4
NV3	185.0	521.0	44.7	11.4	23.2	7.4	3.2	17.3	1.4	4.0	2.2	0.3	1.5	0.5	-0.1	0.6	0.3	2.5
NV4	230.0	684.0	44.4	11.4	38.4	12.7	4.7	36.6	15.4	9.6	3.2	0.7	2.9	0.6	0.3	0.6	0.8	6.2
NV5	248.0	765.0	52.2	44.0	46.2	14.6	3.8	41.4	47.0	12.3	3.6	0.6	3.1	0.6	-0.3	0.6	1.0	7.4
NV5-R	255.0	792.0	56.1	14.9	56.4	17.5	4.8	55.2	10.9	15.1	4.4	0.3	3.8	0.7	0.3	0.7	1.2	8.6
NV6	244.0	957.0	168.0	20.0	112.0	24.7	4.1	63.0	3.6	17.6	2.6	1.0	3.2	0.7	0.5	0.7	0.8	6.3
NV7	208.0	615.0	42.6	12.4	25.1	7.8	0.9	19.1	4.4	4.5	1.6	0.3	1.3	0.7	-0.1	0.7	0.5	3.2
NV8	222.0	636.0	59.7	10.2	44.1	12.4	1.0	29.3	6.3	7.0	4.5	0.5	1.1	1.9	-0.1	1.9	0.7	3.9
NV9	233.0	645.0	36.0	12.1	26.0	8.1	3.8	24.1	4.9	5.5	2.3	0.5	2.1	0.4	-0.1	0.4	0.6	3.9
NW1	402.0	966.0	64.2	21.5	34.8	8.2	13.3	33.8	13.9	7.1	17.3	0.7	18.5	0.4	0.3	0.4	0.8	5.2
NW10	209.0	708.0	64.8	12.6	35.7	10.2	0.7	11.7	1.6	5.3	1.7	0.5	0.8	1.5	-0.1	1.5	0.4	2.9
NW11	208.0	660.0	45.3	11.0	27.5	7.3	2.7	16.4	1.1	4.0	4.5	-0.1	1.2	1.1	-0.1	1.1	0.4	2.3
NW12	285.0	660.0	24.9	11.0	51.6	19.5	0.8	55.8	23.3	14.9	1.5	0.9	0.8	0.7	1.2	0.7	1.1	8.1
NW13	296.0	882.0	148.0	16.3	165.0	51.0	7.4	153.0	8.8	53.1	7.6	1.8	9.2	0.9	-0.3	0.9	4.5	20.8
NW14	318.0	894.0	78.9	16.1	43.2	11.8	0.9	34.5	2.7	7.6	2.0	0.2	1.8	1.4	-0.1	1.4	0.5	3.2
NW15	378.0	1060.0	65.1	28.1	67.2	19.5	15.8	79.5	32.7	21.6	16.8	1.1	18.1	1.1	-0.3	1.1	1.5	12.3
NW16	318.0	756.0	101.0	10.9	72.9	20.6	1.8	44.1	10.1	11.2	1.1	0.3	0.9	0.3	0.3	0.3	0.6	4.9
NW17	270.0	591.0	82.2	10.4	25.7	7.2	0.8	16.6	3.9	3.5	1.3	0.2	0.8	1.3	-0.1	1.3	0.3	2.5
NW17-R	273.0	597.0	75.3	10.3	23.7	6.7	0.8	15.2	3.5	3.3	1.3	0.2	0.8	1.2	-0.1	1.2	0.3	2.5
NW18	306.0	1040.0	50.7	28.3	33.3	9.5	3.3	25.4	5.6	5.9	2.5	0.4	3.1	2.0	-0.1	2.0	0.5	3.7
NW2	232.0	648.0	59.1	10.6	36.3	10.1	2.5	29.3	2.2	7.2	2.6	0.5	2.4	0.5	-0.1	0.5	0.5	4.1
NW2-R	200.0	621.0	50.1	10.1	29.5	9.0	3.3	27.8	2.5	7.3	2.3	0.5	2.1	0.5	-0.1	0.5	0.5	3.5
NW3	191.0	456.0	26.5	10.6	11.9	7.0	1.3	10.2	2.1	2.1	1.5	-0.1	1.3	0.4	-0.1	0.4	0.5	1.9
NW4	189.0	489.0	37.2	13.0	18.7	6.0	3.0	19.2	1.7	4.3	2.1	0.4	1.9	0.9	-0.1	0.9	0.3	2.3
NW5	360.0	1040.0	45.0	20.8	28.6	7.6	5.1	30.6	6.6	7.8	6.3	0.6	5.3	1.3	0.3	1.3	1.0	7.1
NW6	234.0	663.0	33.3	12.4	17.1	10.3	2.9	14.7	3.3	2.9	2.1	0.3	1.8	0.3	-0.1	0.3	0.5	2.6
NW7	187.0	546.0	23.2	10.7	11.3	6.5	0.6	9.8	2.0	2.1	1.4	-0.1	1.2	0.4	-0.1	0.4	0.9	1.8
NW8	339.0	1210.0	165.0	27.7	209.0	66.6	9.7	233.0	88.1	76.8	12.2	1.8	7.4	2.0	-0.3	2.0	3.1	33.0
NW9	229.0	720.0	44.1	13.1	21.5	13.2	3.3	15.0	3.1	3.0	2.4	0.3	2.1	0.4	-0.1	0.4	0.4	2.2
NX1	237.0	735.0	80.4	8.9	33.0	9.8	0.5	18.7	1.3	4.7	1.0	-0.1	0.9	1.3	-0.1	1.3	0.3	2.6
NX10	223.0	696.0	51.3	12.1	27.3	7.4	1.5	21.6	5.2	4.5	1.8	0.3	1.5	0.4	-0.1	0.4	0.5	3.5
NX11	126.0	534.0	34.5	9.3	13.4	7.7	0.1	5.5	0.9	0.9	0.1	-0.1	0.7	0.4	-0.1	0.4	0.4	1.1
NX12	127.0	516.0	37.5	9.0	18.7	11.0	0.3	12.0	2.9	2.2	1.1	-0.1	1.2	0.3	-0.1	0.3	0.9	1.8
NX13	174.0	675.0	63.9	12.0	41.7	12.8	2.1	35.4	7.3	9.7	2.4	0.5	2.0	0.5	-0.1	0.5	0.6	5.7
NX14	188.0	588.0	40.2	10.8	26.1	8.8	1.7	27.8	5.7	6.9	2.4	0.4	2.0	0.5	-0.1	0.5	0.6	4.1
NX14-R	200.0	600.0	47.1	10.2	30.6	10.1	0.9	30.0	6.5	7.7	2.0	0.3	1.6	0.4	-0.1	0.4	0.5	4.3
NX15	182.0	555.0	30.9	8.9	16.2	9.9	0.9	14.4	2.5	3.3	1.8	0.2	2.1	0.7	-0.1	0.7	0.3	2.7
NX16	148.0	597.0	37.2	9.9	19.3	6.0	2.9	17.8	3.4	4.0	2.0	0.4	1.0	0.2	-0.1	0.2	0.5	3.1
NX17	186.0	528.0	47.7	12.1	37.8	13.0	5.1	54.0	22.3	16.5	6.8	0.5	6.1	0.8	0.3	0.8	1.5	10.2
NX2	142.0	594.0	49.8	10.1	20.7	12.4	0.7	6.9	3.1	2.9	1.4	-0.1	1.2	0.4	-0.1	0.4	0.2	2.0

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	001 -LA	002 -LA	003 -LB	004 -LA	005 -LB	006 -LB	007 -LA	008 -LB	009 -LB	010 -LB	011 -LA	012 -LB	013 -LBA	014 -LB	015 -LAR	016 -LB	017 -LB	018 -LB
NX3	210.0	630.0	42.9	10.1	19.5	12.6	0.9	11.3	2.4	0.2	1.0	-0.1	1.0	0.5	-0.1	0.5	1.0	1.6
NX4	327.0	1110.0	47.7	21.5	43.5	15.4	7.5	45.3	18.7	11.8	4.6	0.8	1.7	4.8	0.3	4.8	1.4	7.6
NX5	231.0	789.0	30.6	13.3	10.3	5.9	1.9	7.2	1.5	-0.1	1.7	-0.1	1.6	0.3	-0.1	0.3	0.3	1.5
NX6	192.0	624.0	32.1	11.8	18.3	11.1	2.0	17.2	3.7	3.9	2.2	-0.2	1.8	0.3	-0.1	0.3	0.4	2.7
NX7	420.0	738.0	126.0	50.1	127.0	44.1	5.7	145.0	60.0	54.9	9.6	1.4	9.0	0.9	1.2	0.9	3.3	24.2
NX8	160.0	567.0	34.5	11.1	22.1	7.2	3.1	25.2	10.5	6.6	2.5	-0.4	2.2	0.4	-0.1	0.4	0.8	4.6
NX9	186.0	678.0	52.2	10.9	32.7	10.1	3.8	31.8	13.2	8.8	3.0	0.4	2.5	0.4	-0.1	0.4	0.9	5.5
NY1	136.0	471.0	27.0	10.1	14.6	8.9	0.8	11.8	2.8	2.8	1.4	-0.1	0.3	0.5	-0.1	0.5	0.4	2.4
NY10	172.0	576.0	33.9	9.1	40.2	13.7	6.1	61.5	25.3	16.8	5.9	0.9	6.0	6.1	1.2	6.1	2.0	9.2
NY11	190.0	603.0	48.6	8.2	23.9	6.3	0.5	7.0	3.3	3.0	0.8	-0.1	0.6	0.3	-0.1	0.3	0.3	2.0
NY12	167.0	624.0	42.6	9.4	20.3	13.2	0.7	8.6	3.5	3.0	1.0	-0.1	0.5	1.2	-0.1	1.2	0.3	2.6
NY13	196.0	696.0	48.6	11.8	27.2	7.1	0.8	9.1	1.2	4.1	0.8	-0.1	0.3	0.3	-0.1	0.3	0.3	2.2
NY14	205.0	822.0	47.1	12.8	48.3	18.2	7.2	63.9	26.3	19.4	4.8	1.0	4.4	0.9	0.2	0.9	1.6	11.9
NY15	291.0	816.0	66.0	12.2	44.7	17.8	3.8	43.8	18.2	9.3	5.3	0.4	2.8	0.7	-0.3	0.7	0.9	6.1
NY2	238.0	618.0	39.9	8.0	20.9	14.1	0.6	14.7	3.4	3.3	1.0	-0.1	0.4	1.1	-0.1	1.1	0.4	2.4
NY3	164.0	612.0	26.9	9.2	7.8	4.8	1.2	3.2	1.1	0.2	1.1	-0.1	1.0	0.3	-0.1	0.3	0.3	2.1
NY4	744.0	1690.0	184.0	51.3	192.0	53.4	18.4	206.0	85.2	66.3	20.1	1.5	21.7	1.3	0.3	1.3	3.9	32.7
NY4-R	807.0	1570.0	188.0	56.4	215.0	59.1	21.0	244.0	101.0	78.6	26.4	1.6	29.1	1.4	0.3	1.4	4.8	39.9
NY5	145.0	585.0	30.9	10.7	17.3	10.8	2.0	15.7	3.2	3.4	2.0	0.2	1.7	0.5	-0.1	0.5	0.3	2.8
NY6	218.0	729.0	45.9	13.5	40.8	15.3	5.3	42.0	8.5	10.8	4.4	0.3	5.4	6.5	-0.3	6.5	1.1	7.4
NY7	233.0	771.0	64.2	15.1	66.3	21.4	5.9	63.6	12.6	17.2	5.0	1.2	4.4	6.6	0.3	6.6	2.5	11.6
NY8	125.0	567.0	23.1	10.0	11.5	7.0	0.5	13.7	1.6	3.8	1.4	-0.1	0.4	0.6	-0.1	0.6	0.5	2.6
NY9	298.0	468.0	74.7	17.5	71.7	22.9	0.6	39.0	3.5	18.8	2.2	0.6	2.0	0.7	-0.1	0.7	1.2	7.0
NZ1	250.0	471.0	36.1	10.4	19.3	13.6	0.7	16.4	3.8	3.6	1.3	-0.1	1.3	0.3	-0.1	0.3	0.4	3.0
NZ10	303.0	957.0	108.0	22.4	142.0	40.5	13.8	171.0	12.3	50.7	13.5	1.5	12.6	2.0	0.4	2.0	3.2	22.9
NZ11	179.0	654.0	58.2	9.7	35.7	9.8	0.6	27.6	6.8	6.1	1.5	0.3	2.6	0.4	-0.1	0.4	0.6	3.9
NZ12	179.0	594.0	48.6	8.6	48.3	15.4	5.3	51.0	20.9	11.6	3.6	0.8	3.6	0.6	-0.1	0.6	1.2	6.0
NZ13	148.0	567.0	34.2	10.1	15.8	9.2	0.8	6.8	2.9	2.9	1.5	-0.1	0.6	0.5	-0.1	0.5	0.3	2.2
NZ2	271.0	792.0	36.3	11.3	16.4	10.2	0.8	14.6	2.7	3.2	1.3	0.4	1.2	0.7	-0.1	0.7	0.7	3.5
NZ2-R	107.0	516.0	22.1	6.1	6.6	5.5	0.1	7.5	1.3	0.1	0.8	-0.1	0.1	0.3	-0.1	0.3	0.2	1.8
NZ3	247.0	840.0	39.9	13.8	17.5	5.9	2.1	14.6	2.4	3.7	1.9	0.3	1.5	0.7	-0.1	0.7	0.6	3.6
NZ4	342.0	822.0	34.8	12.6	17.3	5.7	1.5	15.1	8.0	5.7	2.6	0.4	2.3	2.7	-0.1	2.7	0.6	3.7
NZ5	194.0	630.0	26.5	9.9	15.4	10.5	1.8	16.8	3.2	4.1	1.6	0.3	0.4	0.4	-0.1	0.4	0.5	3.6
NZ6	137.0	622.0	22.0	9.5	11.7	7.4	2.2	13.1	2.8	2.9	1.5	0.2	1.2	0.3	-0.1	0.3	0.5	2.5
NZ7	357.0	438.0	34.5	10.1	27.8	9.5	0.6	29.2	5.6	8.5	1.8	0.5	1.3	0.7	-0.1	0.7	1.0	5.9
NZ8	249.0	825.0	37.5	13.2	20.0	13.4	3.8	20.1	3.6	4.7	1.9	0.3	1.5	0.4	-0.1	0.4	0.7	4.3
NZ9	224.0	489.0	68.4	11.1	42.3	10.1	0.7	26.6	1.6	5.9	1.3	0.3	1.0	0.4	-0.1	0.4	0.5	3.3
LMB-QA	141.0	462.0	10.9	8.6	-0.1	-0.1	0.5	0.8	-0.1	-0.1	0.8	-0.1	0.6	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	143.0	453.0	14.0	8.2	-0.1	-0.1	0.6	0.3	-0.1	-0.1	0.4	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	137.0	453.0	10.9	1.4	-0.1	-0.1	-0.1	0.9	-0.1	-0.1	0.8	-0.1	0.4	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	129.0	459.0	11.8	8.2	-0.1	-0.1	0.6	0.9	-0.1	-0.1	0.9	-0.1	0.6	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	163.0	558.0	40.2	8.6	-0.1	-0.1	0.6	1.0	-0.1	-0.1	0.9	-0.1	1.4	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	177.0	567.0	48.5	8.6	1.0	-0.1	0.7	0.9	-0.1	-0.1	0.9	-0.1	0.4	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	160.0	564.0	38.1	8.4	-0.1	-0.1	0.7	0.9	-0.1	-0.1	0.8	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	145.0	661.0	35.4	1.5	-0.6	-0.1	0.7	0.9	-0.1	-0.1	0.9	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	133.0	531.0	22.9	8.4	-0.1	-0.1	0.6	0.9	-0.1	-0.1	0.9	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	118.0	492.0	18.2	8.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	0.3	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	143.0	534.0	16.1	8.1	-0.1	0.5	0.6	0.9	-0.1	-0.1	0.9	-0.1	0.6	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	143.0	619.0	15.4	8.0	-0.4	0.3	0.6	0.9	-0.1	-0.1	0.9	-0.1	0.6	-0.1	-0.1	-0.1	-0.1	-0.1

SOIL GAS HYDROCARBONS (SGH) by GC/MS

A14-06865 - Date: November 19, 2014 - Activation Laboratories Ltd.

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Canstar Resources - Alex Pleson
North Survey Area

R=Replicate Sample
-0.1=Reporting Limit of 0.1pg/g (ppt=parts per trillion)
LMB-QA = Laboratory Materials Blank - Quality Assurance

LEGEND FOR COLUMN HEADINGS - SGH COMPOUND CLASSES

Results represent only the material tested. Actlabs is not liable for any claim/damage from use of this report in excess of the test cost. Unless requested samples are discarded in 90 days. This report is only to be reproduced in full.

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

001 LA 002 LA 003 LB 004 LA 005 LB 006 LB 007 LA 008 LB 009 LB 010 LB 011 LA 012 LB 013 LBA 014 LB 015 LAR 016 LB 017 LB 018 LB

LA, HA, LBA, HBA = ALKYL-ALKANES
LB, HB, LPB, HPB = ALKYL-BENZENES
LAR, MAR, HAR = ALKYL-AROMATICS
LBI, MBI, HBI, LPH, MPH, HPH = ALKYL-POLYAROMATICS
THI = ALKYL-DIVINYLENE SULPHIDES
ALK = ALKYL-ALKENES

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NA1	19.9	7.1	0.6	7.1	16.9	2.3	9.0	4.7	19.6	11.0	4.8	2.4	6.8	8.0	7.9	5.3	1.7	4.1
NA2	1.8	1.9	-0.1	0.8	2.0	1.4	1.5	1.1	1.3	2.9	0.4	1.8	0.5	1.7	0.4	1.5	-0.1	0.2
NAA1	7.2	1.3	-0.1	1.6	3.4	2.0	3.5	1.8	5.2	6.0	3.6	2.8	1.4	3.7	3.8	2.8	1.3	2.8
NAA10	7.3	0.4	-0.1	0.6	6.1	2.0	4.3	1.4	7.0	1.4	3.2	1.9	3.6	3.9	3.7	0.3	2.9	
NAA11	30.9	32.1	1.9	28.8	7.0	4.2	11.1	18.3	38.1	18.8	6.7	8.4	9.8	10.6	11.8	9.2	4.1	15.3
NAA2	74.7	104.0	8.6	105.0	30.6	4.9	24.5	27.5	75.3	41.4	4.4	7.9	15.3	25.0	26.0	12.4	7.7	27.4
NAA3	6.8	8.9	0.4	10.2	2.2	1.9	0.2	13.4	3.9	8.0	2.9	2.6	1.2	2.9	3.5	2.8	0.4	10.6
NAA4	3.2	2.1	-0.1	2.4	4.5	0.2	0.1	2.5	3.3	4.4	0.5	2.8	1.2	2.6	2.8	2.3	1.1	1.5
NAA5	4.8	2.8	-0.1	3.2	7.6	2.3	2.5	3.5	5.0	7.9	0.6	2.8	1.3	3.3	3.3	2.5	1.4	2.5
NAA6	6.1	3.3	-0.1	3.7	12.1	2.2	3.1	1.4	6.8	5.8	2.2	3.2	2.8	4.3	4.2	3.2	1.4	2.9
NAA7	3.5	1.9	-0.1	2.1	3.5	0.7	2.3	1.0	3.3	5.2	1.9	2.0	1.8	2.7	3.0	2.3	1.2	1.9
NAA8	3.2	1.1	-0.1	1.1	4.8	0.2	0.3	2.3	3.1	6.2	0.4	2.4	1.8	2.6	2.6	2.3	1.1	1.7
NAA9	8.2	3.6	0.4	3.5	4.0	2.2	0.1	3.3	7.5	7.1	3.8	3.4	1.6	4.0	4.4	3.3	0.4	2.6
NB1	7.2	5.4	-0.1	6.2	5.5	2.0	3.6	1.3	6.3	4.2	4.1	2.8	2.8	4.2	4.4	9.3	1.2	2.9
NB2	3.4	1.0	-0.1	1.0	2.6	1.8	2.1	1.8	3.2	4.4	0.5	1.7	1.5	2.3	2.2	1.8	-0.1	1.3
NB3	2.4	1.1	-0.1	1.1	4.7	0.2	2.3	1.9	3.8	5.4	1.4	1.1	1.1	2.5	2.5	2.0	-0.1	1.3
NB4	4.9	5.2	-0.1	5.9	2.4	0.2	2.4	1.3	5.9	4.4	3.2	1.9	1.3	2.6	2.6	2.1	1.2	3.5
NBB1	8.0	2.2	-0.1	2.2	7.4	2.1	0.6	3.0	6.1	5.6	1.6	3.4	4.7	3.8	4.2	4.0	0.3	2.1
NBB10	5.0	5.9	-0.1	1.5	3.4	1.6	0.1	1.5	4.7	5.4	3.1	2.8	1.3	3.1	3.3	2.6	0.9	2.8
NBB11	14.9	3.5	0.5	5.0	4.6	3.5	0.7	8.4	16.2	7.9	2.5	6.5	6.5	7.7	8.7	7.7	1.1	5.8
NBB2	8.1	10.2	0.8	10.5	6.5	2.5	0.3	6.8	7.7	10.6	1.4	4.3	1.6	4.0	4.6	3.9	1.2	6.9
NBB2-R	7.3	9.8	0.7	10.1	6.0	2.3	0.2	6.3	6.6	9.6	1.5	3.2	1.6	3.5	4.0	3.5	1.1	6.3
NBB3	14.3	16.3	1.1	16.6	3.7	2.9	0.2	5.6	9.4	1.8	2.0	5.0	2.0	5.4	6.1	4.4	1.3	9.2
NBB4	1.4	1.0	-0.1	1.3	3.5	2.1	2.9	1.7	4.7	5.1	3.4	2.5	1.3	3.4	3.6	2.6	1.3	2.5
NBB5	3.4	2.3	-0.1	2.6	5.0	1.9	2.5	3.2	3.8	5.4	1.5	1.9	1.9	2.6	2.7	2.4	1.2	2.8
NBB6	13.6	14.8	1.1	14.0	7.1	3.0	0.2	3.7	14.3	13.7	3.4	4.8	5.5	6.0	6.7	4.8	1.6	7.3
NBB7	2.5	1.0	-0.1	1.2	2.0	0.2	0.4	1.5	1.8	2.8	1.5	1.8	0.8	0.6	1.9	1.7	-0.1	1.3
NBB8	5.6	3.2	-0.1	3.6	4.3	0.9	0.4	6.5	4.5	6.5	2.3	3.1	1.6	3.0	3.4	3.2	0.4	3.0
NBB9	16.7	21.3	1.5	21.7	8.0	3.5	4.2	12.4	13.3	16.3	2.8	6.8	5.4	6.5	8.0	6.0	2.3	10.3
NC1	2.8	1.5	-0.1	1.5	2.0	0.2	0.2	1.6	1.9	3.4	1.7	1.3	0.9	1.9	2.2	1.8	1.1	1.1
NC2	3.2	1.7	-0.1	2.0	2.4	0.2	2.2	2.1	2.1	4.4	2.4	1.6	1.6	1.9	2.6	2.1	1.1	0.2
NC3	2.4	2.1	-0.1	2.3	2.2	0.2	1.7	0.9	1.5	3.7	1.5	1.3	0.8	1.8	1.9	1.7	1.1	1.9
NC4	1.9	0.5	-0.1	0.6	1.9	1.5	0.3	0.7	1.2	3.0	1.1	1.7	0.7	1.6	0.5	1.7	-0.1	0.8
NC5	3.4	3.8	-0.1	3.3	3.0	0.2	1.0	1.0	3.5	4.7	2.7	2.0	1.7	2.4	2.6	2.1	1.1	2.4
NC6	8.0	1.8	-0.1	2.2	2.8	2.1	4.1	3.3	7.1	4.4	3.9	2.8	2.9	4.0	4.2	3.3	1.2	2.2
NC7	2.9	2.4	-0.1	1.9	2.2	0.2	0.3	0.7	2.0	3.3	1.9	2.0	1.0	1.9	2.1	2.0	0.3	1.4
NC8	3.0	1.5	-0.1	1.7	3.8	0.2	0.2	2.0	2.9	4.9	0.4	2.2	1.1	2.2	2.3	2.1	1.1	1.5
NCC1	11.2	8.0	0.6	8.5	2.3	1.6	4.7	3.6	5.7	6.3	3.2	2.9	1.3	3.9	4.1	2.9	1.5	8.0
NCC1-R	10.2	7.4	0.5	7.9	2.2	1.6	4.2	3.5	5.4	5.9	3.1	2.8	1.2	3.6	3.8	2.8	1.6	7.5
NCC2	18.8	2.5	0.5	3.3	4.4	2.8	8.9	6.5	17.6	6.3	6.0	7.5	8.0	8.0	5.7	2.4	4.0	
NCC3	9.8	2.2	-0.1	2.8	3.7	2.3	5.0	4.3	7.2	5.7	4.1	3.5	1.8	4.2	4.5	3.5	1.1	2.6
NCC4	14.9	2.7	0.6	3.8	4.8	8.0	5.9	4.9	11.3	10.2	2.3	5.6	5.1	6.1	6.4	4.4	2.4	5.0
NCC5	6.2	3.2	-0.1	4.1	3.8	2.2	3.1	5.7	6.5	7.4	3.6	3.5	1.6	3.7	3.7	3.1	0.8	4.6
NCC6	3.2	0.6	-0.1	0.9	1.3	2.0	1.7	1.3	2.5	1.1	2.8	1.9	0.3	2.4	2.4	2.0	0.6	2.5
ND1	8.5	1.4	-0.1	1.7	3.9	2.1	4.2	1.0	7.9	6.2	4.3	2.8	1.7	4.0	3.8	2.9	1.4	1.6
ND10	2.8	0.6	-0.1	0.9	2.5	0.2	1.8	1.0	2.1	4.5	2.3	1.4	0.9	2.2	2.1	1.9	-0.1	2.4
ND2	7.0	0.7	-0.1	0.7	3.5	2.2	3.3	1.5	7.3	7.5	4.1	2.8	1.6	3.7	3.7	2.9	1.7	3.1
ND3	2.8	0.8	-0.1	0.9	1.6	1.5	1.6	0.3	0.6	3.0	1.6	0.9	1.1	1.7	1.8	1.5	-0.1	1.5
ND4	5.7	7.2	-0.1	5.6	1.4	1.9	2.6	5.0	4.8	4.6	3.1	2.9	1.3	3.5	3.6	2.7	0.3	7.3
ND5	1.8	2.1	-0.1	0.4	1.8	1.5	1.3	0.6	0.8	3.2	4.3	1.8	0.7	1.6	1.5	1.5	-0.1	2.1
ND6	4.1	3.6	-0.1	2.8	3.6	1.9	2.2	1.0	3.1	5.4	0.5	1.9	1.8	2.6	2.5	2.2	1.1	2.3
ND7	2.9	1.8	-0.1	1.2	1.2	0.6	0.4	1.7	3.2	0.5	1.8	1.0	2.5	2.9	2.4	-0.1	1.3	
ND8	3.3	1.7	-0.1	1.8	3.8	0.2	2.2	0.7	3.5	5.3	0.7	1.6	1.1	2.5	2.5	2.2	1.1	1.8
ND8-R	4.0	2.7	-0.1	2.9	4.7	1.9	2.3	1.1	3.1	6.1	0.7	2.5	1.3	2.5	2.9	2.4	0.3	2.2
ND9	4.2	2.3	-0.1	2.5	4.5	2.0	2.5	0.9	3.2	6.4	1.4	2.1	1.9	2.9	2.8	2.3	1.1	2.5
NDD1	21.8	16.7	1.1	16.7	18.8	2.8	8.6	4.2	22.1	14.2	5.2	2.2	7.4	7.7	7.9	5.8	1.8	8.1
NDD2	15.1	5.1	0.4	5.1	7.8	2.7	7.4	3.8	11.5	8.8	3.2	4.3	2.0	5.2	5.3	3.7	0.3	3.1
NDD3	9.2	4.0	0.4	4.0	12.1	2.2	0.6	3.7	9.3	7.8	1.7	4.2	5.3	4.7	5.2	4.9	0.3	3.0
NDD4	6.8	2.5	-0.1	2.5	7.4	2.2	0.4	3.9	7.1	7.7	1.5	3.6	1.9	4.2	4.5	4.1	0.3	3.0
NDD5	3.3	1.5	-0.1	1.7	4.4	1.8	2.1	3.2	2.7	6.1	0.6	1.8	1.0	2.7	2.5	2.4	1.1	2.4
NE1	22.6	16.7	1.2	16.9	9.0	3.0	9.4	3.5	23.4	16.3	4.3	7.5	7.6	8.6	9.3	6.3	1.8	7.5
NE10	1.8	1.2	-0.1	1.4	0.1	1.1	1.4	0.4	0.9	0.5	4.0	0.6	0.7	1.5	1.5	1.3	-0.1	1.1
NE11	2.1	0.3	-0.1	0.4	1.2	0.3	1.5	0.6	1.9	2.6	1.5	0.8	0.7	1.9	1.8	1.7	-0.1	1.6
NE2	1.8	0.6	-0.1	0.7	0.1	1.8	0.3	1.3	0.7	1.8	0.0	1.4	0.7	1.3	1.3	1.3	-0.1	1.1
NE3	4.8	2.1	-0.1	2.3	2.1	0.7	2.8	2.0	3.4	3.7	2.8	2.0	2.0	2.5	2.9	2.3	1.1	1.4
NE4	5.0	2.5	-0.1	2.0	3.1	0.2	0.2	1.8	3.5	4.4	3.2	2.6	2.4	2.8	3.1	2.8	0.3	1.4
NE4-R	3.5	0.9	-0.1	1.1	2.2	0.2	0.3	1.5	2.4	3.2	2.4	2.1	1.1	2.2	2.3	2.2	-0.1	0.6

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NE5	12.8	1.5	-0.1	1.5	7.1	2.5	6.0	3.2	11.3	8.9	3.2	3.7	2.1	5.6	5.6	3.6	1.4	2.0
NE6	2.4	1.1	-0.1	1.5	2.5	1.6	1.6	2.1	1.8	4.2	2.1	2.1	0.9	2.0	2.0	1.8	0.2	1.7
NE7	7.2	2.0	-0.1	1.6	1.3	0.7	3.5	1.5	3.8	2.2	2.7	2.1	1.2	2.8	3.2	2.3	-0.1	1.2
NE8	1.5	1.4	-0.1	1.2	-0.1	1.4	1.3	0.6	1.3	2.7	1.5	1.6	0.6	1.5	1.4	1.4	-0.1	1.6
NE9	4.0	1.5	-0.1	1.5	3.9	2.0	2.0	2.3	4.4	6.8	1.6	2.9	1.5	3.1	3.2	2.9	-0.1	0.9
NEE1	3.8	2.3	-0.1	3.7	2.8	2.1	2.0	1.5	3.9	5.9	2.8	2.2	1.0	2.7	2.7	2.2	1.3	2.1
NEE2	9.1	8.6	0.6	6.6	3.8	2.2	0.4	5.9	6.2	8.2	1.6	3.5	1.7	3.9	4.4	6.7	1.0	5.7
NEE3	6.2	3.0	-0.1	3.0	6.6	2.0	3.4	3.8	5.9	9.4	1.7	2.9	1.4	3.3	3.8	2.8	0.3	3.0
NEE4	3.2	2.1	-0.1	2.7	1.5	0.2	0.3	2.8	2.1	4.4	1.6	1.7	0.9	1.9	2.1	1.6	0.3	4.4
NEE5	3.0	1.0	-0.1	1.0	2.8	0.2	2.0	2.2	2.4	4.2	2.3	1.6	1.0	1.9	2.1	1.8	1.2	1.7
NEE6	11.0	5.3	0.5	5.8	12.4	2.4	0.1	6.4	11.2	8.4	1.7	4.7	5.9	5.6	6.1	5.5	0.4	5.6
NEE7	9.2	0.9	-0.1	1.3	12.4	1.9	5.4	4.7	14.1	3.8	3.7	4.5	5.8	5.1	5.3	3.8	1.5	3.1
NF1	7.6	1.4	-0.1	1.6	7.3	2.4	4.3	4.4	7.7	8.9	1.6	3.0	4.9	4.3	4.4	4.1	-0.5	3.0
NF10	2.6	1.7	-0.1	1.7	2.3	0.2	1.9	1.8	2.7	3.7	2.1	1.8	1.5	2.3	2.5	2.1	0.3	1.4
NF11	6.1	5.7	-0.1	4.7	1.6	0.7	0.2	1.3	4.1	4.5	8.2	2.6	2.5	3.2	3.8	2.9	1.3	3.2
NF2	1.4	2.2	-0.1	1.5	1.4	1.4	0.3	2.0	1.0	2.8	0.3	1.6	0.5	1.5	1.7	1.5	-0.1	1.7
NF3	2.7	2.9	-0.1	1.0	1.8	0.2	0.4	2.4	1.8	3.6	1.7	1.9	1.0	1.9	2.3	2.0	1.1	1.8
NF4	2.1	3.1	-0.1	1.2	2.9	0.2	0.3	2.5	1.6	4.4	0.5	2.1	1.0	1.9	2.0	1.9	0.3	1.9
NF5	2.1	1.4	-0.1	1.6	2.1	0.5	1.5	0.6	0.4	3.7	0.8	0.6	1.9	1.8	1.7	1.1	1.7
NF6	2.7	3.2	-0.1	3.8	1.5	0.2	1.7	1.0	1.8	3.2	2.0	1.5	0.8	1.8	0.6	1.8	0.2	2.3
NF7	18.0	9.5	0.5	10.3	3.8	3.7	7.0	11.2	18.5	8.3	3.6	7.7	7.2	10.0	10.3	7.0	1.2	6.8
NF8	3.6	0.5	-0.1	0.5	3.2	0.2	2.4	0.7	3.6	5.0	3.0	1.2	1.9	2.7	2.8	2.4	-0.1	1.3
NF8-R	4.8	6.8	-0.1	7.8	3.5	0.6	3.1	1.6	5.0	7.8	8.5	2.6	2.5	3.3	3.6	6.0	0.3	3.6
NF9	3.4	3.8	-0.1	4.4	3.2	0.2	2.1	1.1	3.5	5.3	2.7	1.6	1.8	2.5	2.6	2.2	0.2	2.1
NFF1	4.0	0.8	-0.1	0.8	2.7	1.9	0.1	2.1	2.5	4.6	2.6	2.0	1.7	2.5	2.8	2.1	1.2	1.3
NFF2	11.3	7.5	0.6	8.0	9.8	2.4	4.4	6.0	12.6	11.2	3.5	4.0	2.3	5.3	5.3	3.9	1.8	5.1
NFF3	2.4	2.8	-0.1	3.0	0.1	0.2	0.4	3.3	1.2	0.6	4.2	1.7	0.8	0.9	0.5	4.7	0.3	2.6
NFF4	16.9	6.4	0.6	6.8	12.7	2.9	7.0	4.2	16.2	8.7	4.3	5.7	6.5	7.1	7.3	4.9	1.1	3.2
NFF5	10.2	2.7	-0.1	2.7	11.4	2.2	4.3	3.9	9.9	7.3	2.0	3.6	1.9	4.8	4.7	3.6	1.5	2.8
NFF6	2.8	1.5	-0.1	1.6	3.7	0.2	1.8	1.9	2.7	4.7	2.1	2.2	1.1	2.0	2.4	2.0	1.1	1.6
NFF6-R	3.0	1.6	-0.1	1.7	3.8	0.2	0.2	1.9	2.9	4.7	2.3	2.2	1.1	1.4	2.3	2.2	0.3	1.4
NFF7	5.1	1.6	-0.1	1.6	5.6	2.0	2.9	2.6	5.0	8.0	1.6	2.6	1.2	3.1	3.1	2.4	1.3	2.1
NFF8	4.6	2.4	-0.1	2.4	4.2	2.0	2.8	2.9	4.8	7.2	3.6	2.4	2.3	3.1	3.4	2.6	0.4	2.0
NG1	1.2	0.2	-0.1	0.3	-0.1	0.4	1.1	0.4	0.9	0.8	1.3	0.5	1.3	1.2	1.3	-0.1	1.2
NG10	3.0	0.5	-0.1	0.8	1.4	0.2	1.8	1.0	2.4	3.8	2.2	1.3	0.9	2.3	2.2	4.9	1.2	2.5
NG11	3.8	3.5	-0.1	3.8	1.1	0.3	1.9	0.9	1.9	2.8	1.5	1.4	0.8	2.0	2.1	1.9	-0.1	2.2
NG2	1.6	2.3	-0.1	0.5	0.1	0.2	1.6	0.7	2.5	0.7	2.0	2.0	0.3	1.6	0.3	1.6	-0.1	1.9
NG2-R	1.6	2.3	-0.1	0.4	1.3	0.2	1.6	0.8	2.7	2.8	2.0	1.9	0.7	1.6	0.3	1.4	-0.1	1.7
NG3	1.8	0.2	-0.1	0.3	0.1	0.2	1.4	0.3	1.2	1.7	-0.1	1.6	0.6	1.6	1.5	1.4	-0.1	1.1
NG4	4.9	1.3	-0.1	1.7	2.4	1.8	2.6	0.7	3.9	4.0	3.1	2.2	1.2	2.9	2.8	2.3	1.1	0.5
NG5	13.3	8.2	0.4	2.8	7.9	2.7	5.3	5.5	11.7	10.9	3.3	4.7	2.8	5.9	6.0	4.3	1.8	5.2
NG6	3.8	1.9	-0.1	2.2	-0.1	0.3	1.9	1.7	2.5	0.4	2.1	1.7	0.9	2.5	2.4	2.0	1.2	1.4
NG7	1.9	0.3	-0.1	0.4	0.1	1.4	1.4	0.3	1.0	1.0	1.9	1.3	0.4	1.4	1.3	-0.1	0.9
NG8	2.6	1.2	-0.1	1.6	1.4	1.7	1.6	0.6	2.5	3.2	1.4	0.9	0.7	2.1	2.1	1.7	-0.1	1.5
NG9	6.1	2.4	-0.1	2.7	3.1	1.9	3.3	2.8	5.5	6.0	3.3	2.3	2.3	3.2	3.3	2.7	1.3	2.4
NGG1	3.1	1.1	-0.1	1.1	2.7	1.9	1.4	2.8	3.0	4.8	2.4	1.0	0.9	2.2	2.3	1.9	1.3	2.2
NGG10	5.6	7.6	0.6	8.6	3.0	1.5	0.4	4.1	4.0	6.1	2.4	2.9	1.5	2.7	2.9	2.9	0.5	4.4
NGG2	2.3	1.0	-0.1	1.4	2.7	1.8	1.6	2.1	2.4	4.4	2.1	0.9	0.7	2.1	2.1	1.9	-0.1	1.6
NGG3	14.3	5.6	0.5	6.8	2.1	1.9	6.0	6.9	14.7	2.0	5.3	4.7	5.8	5.6	5.7	3.9	1.5	16.8
NGG4	7.9	3.8	-0.1	3.8	6.7	2.2	3.8	3.9	7.7	7.9	2.2	3.1	1.7	3.9	4.1	3.1	0.4	3.4
NGG5	6.7	2.3	-0.1	2.3	4.4	2.0	3.3	2.9	6.0	6.6	3.5	2.8	1.5	3.1	3.3	2.6	0.3	2.1
NGG6	3.5	1.2	-0.1	1.6	2.7	1.9	2.1	2.8	3.5	5.0	2.6	1.1	1.7	2.7	2.7	2.3	0.2	0.1
NGG7	5.8	2.3	-0.1	2.4	6.2	2.1	2.9	3.9	6.0	8.8	2.0	2.5	1.4	3.2	3.2	2.6	0.3	2.8
NGG8	3.4	4.6	-0.1	5.1	3.2	1.8	1.8	4.6	3.0	6.1	1.7	1.1	0.9	2.4	2.5	2.0	1.3	3.5
NGG9	2.1	1.1	-0.1	1.4	2.1	0.2	0.3	3.3	1.2	1.8	1.5	1.8	0.7	1.6	0.5	1.6	0.3	2.6
NH1	3.5	3.0	-0.1	3.3	3.0	0.2	-0.1	3.9	2.7	5.5	1.9	2.4	1.2	2.3	2.5	2.2	0.3	3.0
NH10	10.7	1.3	-0.1	1.6	3.4	1.9	8.5	2.2	7.2	5.1	4.2	2.9	1.8	3.9	4.1	2.9	1.1	1.5
NH10-R	19.2	1.4	-0.1	1.7	3.4	1.9	8.6	2.9	12.4	5.5	5.5	4.4	6.2	6.2	6.5	4.3	1.2	0.8
NH11	6.3	6.0	-0.1	4.6	7.5	2.1	3.2	1.7	6.4	8.4	1.9	2.8	1.5	3.7	3.7	2.9	1.3	3.2
NH2	7.7	3.1	-0.1	3.5	4.8	2.2	3.4	1.9	7.5	7.4	4.0	3.0	1.5	4.0	4.2	2.9	1.2	3.2
NH3	2.5	1.1	-0.1	1.4	1.3	0.5	0.2	0.5	1.3	2.3	0.3	1.5	0.4	1.5	1.4	1.3	-0.1	1.2
NH4	4.3	1.8	-0.1	1.8	4.6	2.0	2.4	3.0	4.4	7.0	3.1	2.2	2.0	2.9	2.9	2.4	1.2	2.3
NH5	6.6	0.7	-0.1	1.0	3.7	2.0	3.2	2.2	6.1	5.4	3.7	2.7	1.4	3.7	3.8	2.8	1.2	1.5
NH6	8.4	0.3	-0.1	0.3	7.7	2.1	4.9	1.6	7.5	7.6	2.0	3.0	1.8	4.3	4.4	3.4	1.2	2.1
NH7	15.8	5.0	0.5	5.0	18.1	2.3	7.3	3.1	16.3	11.1	4.3	5.3	6.5	7.3	7.5	5.0	1.4	4.7
NH8	6.6	4.1	-0.1	4.6	9.0	2.0	3.6	2.2	5.4	7.1	2.1	2.8	1.5	4.0	4.1	3.2	1.3	3.1
NH9	8.4	4.1	-0.1	4.6	8.9	2.1	4.0	2.2	5.8	8.3	2.2	3.1	1.5	4.3	4.3	3.2	1.3	3.3

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NHH1	1.7	1.3	-0.1	1.8	1.6	1.4	0.3	2.3	0.9	3.1	1.2	1.6	0.6	1.5	1.5	1.5	1.1	1.8
NHH10	2.3	0.8	-0.1	0.9	1.8	1.6	1.5	3.3	1.7	3.7	1.0	0.9	0.7	1.8	1.7	1.6	1.1	2.5
NHH11	2.5	1.3	-0.1	1.4	1.7	1.6	1.6	2.1	2.3	3.1	1.9	1.0	0.7	1.9	1.8	1.6	1.2	1.6
NHH11-R	2.5	1.1	-0.1	1.1	2.1	1.6	1.6	2.0	2.2	3.5	1.6	0.9	0.7	2.0	1.9	1.6	1.2	1.6
NHH12	2.9	1.5	-0.1	2.0	1.7	0.2	1.8	2.7	2.7	2.5	1.7	2.2	0.8	2.2	2.2	2.0	0.3	2.1
NHH2	4.8	7.7	0.5	3.1	4.2	2.1	2.6	3.1	5.1	9.2	3.2	2.6	1.3	3.0	3.1	2.6	1.1	5.7
NHH3	3.1	1.2	-0.1	1.2	3.1	0.2	0.4	2.6	2.6	5.2	2.5	2.2	1.1	2.3	2.4	2.2	0.3	1.9
NHH4	7.1	2.0	-0.1	2.0	2.5	1.7	3.3	2.8	4.9	5.1	3.2	3.0	1.3	3.2	3.5	2.5	0.4	1.9
NHH5	4.5	3.7	-0.1	3.7	5.9	0.7	2.6	4.1	3.4	8.5	1.4	2.6	1.3	2.7	2.8	2.5	0.4	3.9
NHH6	6.5	1.1	-0.1	1.1	3.0	2.0	3.2	3.9	4.2	5.7	3.2	2.1	1.4	2.7	3.0	2.3	0.3	2.6
NHH7	6.6	3.0	-0.1	2.9	6.0	2.1	3.0	4.0	6.0	8.8	1.7	2.5	1.4	3.1	3.0	2.5	0.5	3.4
NHH8	2.6	0.8	-0.1	1.1	2.4	0.2	0.2	2.7	1.8	4.0	1.7	2.0	0.9	1.8	1.9	1.8	0.3	2.3
NHH9	2.0	2.8	-0.1	3.2	1.8	0.2	0.3	3.2	1.5	3.9	1.5	1.8	0.8	1.7	1.8	1.8	0.3	2.7
NI1	24.7	4.8	1.0	19.7	4.8	3.5	9.6	8.2	18.1	15.0	1.9	9.8	6.1	11.8	12.9	8.4	1.3	14.1
NI10	5.8	5.1	-0.1	5.6	4.5	2.0	2.6	2.1	4.6	7.7	3.5	2.8	1.3	3.3	3.3	2.5	1.3	3.7
NI2	5.0	1.5	-0.1	2.0	2.3	1.9	2.9	2.2	3.6	4.3	2.5	2.2	1.0	2.9	2.9	2.2	1.2	1.5
NI3	3.7	1.1	-0.1	1.1	3.4	1.9	2.0	2.6	2.9	5.3	2.5	1.9	1.0	2.7	2.6	2.1	-0.1	2.0
NI4	12.3	3.2	-0.1	4.1	7.8	2.4	4.6	5.9	9.5	14.0	3.1	4.2	5.0	5.8	6.0	4.2	1.7	8.0
NI5	8.2	1.1	-0.1	1.6	9.0	2.3	4.0	2.4	8.7	7.8	2.1	3.2	1.7	4.5	4.4	3.4	1.3	3.8
NI5-R	8.2	0.8	-0.1	0.9	8.6	2.3	3.8	2.0	8.3	6.9	2.0	3.0	1.7	4.3	4.3	3.3	1.3	3.1
NI6	3.2	1.8	-0.1	2.3	1.9	0.2	2.0	3.1	3.1	2.9	2.3	2.4	1.0	2.5	2.8	2.2	1.1	1.9
NI7	24.2	9.7	1.8	54.8	6.1	3.7	7.7	29.0	16.7	20.4	0.3	9.0	5.8	9.5	10.5	7.3	1.5	28.3
NI8	3.1	2.0	-0.1	1.9	1.2	0.2	1.9	1.0	2.3	3.1	2.2	1.3	0.8	2.1	2.1	1.8	-0.1	1.8
NI9	8.9	3.7	-0.1	3.9	8.9	2.9	4.5	1.9	9.4	8.5	2.1	3.4	1.9	4.8	5.0	3.5	1.2	3.5
NII1	1.6	0.6	-0.1	0.7	3.1	1.5	1.6	1.2	1.9	3.0	0.2	1.9	0.7	1.9	1.8	1.6	-0.1	0.9
NII10	5.7	5.0	-0.1	4.2	8.5	2.1	2.8	1.3	5.9	7.2	1.8	2.8	1.5	3.4	3.5	2.8	1.4	3.0
NII11	5.1	3.7	-0.1	3.2	3.7	1.9	2.7	0.8	5.2	5.5	3.3	2.4	2.2	3.2	3.5	2.5	1.2	1.9
NII11-R	5.0	2.5	-0.1	2.6	3.5	2.0	2.7	0.8	5.2	5.1	3.4	2.9	1.3	3.3	3.5	2.6	1.2	2.3
NII12	5.5	1.2	-0.1	1.2	4.3	1.9	3.5	1.9	5.2	5.7	3.4	2.4	1.3	3.2	3.2	2.6	1.2	1.6
NII13	21.2	16.3	1.3	15.8	3.3	2.8	8.6	2.7	15.1	1.4	9.2	0.2	5.1	8.1	8.6	5.6	1.2	8.3
NII14	2.9	2.3	-0.1	1.7	2.5	0.2	1.9	1.6	2.7	3.5	1.7	1.2	0.8	2.1	2.0	1.8	-0.1	1.4
NII2	6.4	3.2	-0.1	3.2	4.1	2.0	3.2	1.3	6.0	7.5	3.4	2.7	2.3	3.6	3.8	2.6	1.3	2.7
NII3	7.5	17.4	0.8	18.7	3.0	2.3	2.9	11.8	5.5	12.8	3.2	3.8	1.2	4.4	4.6	3.3	1.2	15.0
NII4	6.7	6.6	1.1	10.6	2.6	2.2	2.8	13.6	6.3	15.7	8.1	3.5	1.1	4.2	4.3	6.2	2.2	21.7
NII5	3.1	3.8	-0.1	2.9	3.5	1.9	2.0	1.1	3.2	5.8	0.5	2.0	0.9	2.4	2.5	2.1	1.2	2.2
NII6	2.5	0.8	-0.1	1.0	2.6	1.8	1.6	0.9	2.6	4.6	2.2	1.1	0.8	2.1	2.0	1.7	1.2	1.8
NII7	0.9	1.8	-0.1	2.0	-0.1	0.4	1.1	1.6	0.7	2.3	-0.1	1.4	0.3	1.3	1.3	1.2	-0.1	1.3
NII8	8.9	15.4	0.6	4.3	3.5	2.3	3.5	7.1	7.2	10.6	5.6	3.8	1.5	4.7	4.9	9.5	0.6	11.2
NII9	2.4	3.5	-0.1	2.8	2.9	1.8	1.6	0.9	2.7	4.8	2.3	1.0	0.8	2.1	2.0	1.8	1.2	2.1
NJ1	7.7	2.1	-0.1	2.1	5.1	2.2	3.9	0.2	7.7	8.1	3.6	2.9	1.4	4.2	4.4	3.2	1.3	2.6
NJ2	7.1	5.4	-0.1	5.9	4.8	2.1	3.5	4.5	7.1	8.0	3.8	2.9	1.6	3.7	3.9	3.0	0.3	3.4
NJ3	8.4	6.9	0.4	7.5	4.6	2.2	4.1	3.5	7.7	8.8	8.9	2.9	1.6	3.7	4.2	6.2	0.3	4.7
NJ4	5.4	3.5	-0.1	3.5	5.3	2.0	3.1	4.0	5.5	8.1	1.1	2.1	1.5	3.3	3.4	2.8	0.3	3.0
NJ5	3.0	1.8	-0.1	2.0	2.9	0.2	0.2	2.1	3.0	4.1	0.5	2.2	1.1	2.3	2.6	2.2	1.1	1.8
NJ6	3.8	1.7	-0.1	1.7	2.7	0.2	2.4	3.0	2.8	4.9	2.6	1.4	1.2	2.3	2.4	2.2	0.2	2.2
NJ7	10.9	7.3	0.4	7.9	7.5	2.3	0.6	10.9	8.3	9.3	4.2	4.7	4.9	4.9	5.4	6.3	0.4	4.9
NJJ1	4.4	0.9	-0.1	1.1	3.0	2.0	2.5	0.8	3.3	4.7	3.0	2.2	2.0	2.9	2.9	2.4	1.2	2.1
NJJ10	80.4	51.0	4.1	48.0	33.6	3.8	26.9	12.8	30.7	23.2	13.4	7.8	16.8	21.3	21.7	10.7	3.3	17.1
NJJ11	2.3	0.6	-0.1	0.8	2.0	0.2	1.6	0.7	1.6	3.2	1.3	0.8	0.7	1.9	1.8	1.6	-0.1	1.5
NJJ12	7.1	2.4	-0.1	2.5	5.3	1.9	4.2	2.6	6.5	6.9	2.2	2.7	1.5	3.6	3.6	2.9	1.2	2.3
NJJ13	9.0	10.6	0.7	11.2	6.2	2.2	3.6	4.0	7.4	11.0	4.0	3.8	1.5	4.6	4.8	3.4	1.8	9.8
NJJ14	2.2	1.0	-0.1	1.2	2.8	1.6	1.6	0.5	2.3	3.8	1.0	1.0	0.8	2.1	2.0	1.7	-0.1	1.3
NJJ2	5.7	3.3	-0.1	3.5	5.4	2.1	2.8	2.9	5.6	8.6	4.0	2.9	1.6	3.7	3.6	2.8	1.3	3.3
NJJ3	3.8	1.4	-0.1	1.8	2.0	0.2	2.0	0.7	2.7	3.5	2.3	1.1	0.9	2.3	2.2	4.1	1.1	1.4
NJJ4	5.8	24.4	1.6	23.0	2.6	2.2	2.6	8.2	6.1	5.3	2.8	2.8	0.9	3.3	3.6	2.7	2.4	13.8
NJJ5	8.0	1.2	-0.1	1.5	2.5	2.1	3.5	1.3	5.3	5.3	3.6	2.6	1.5	3.5	3.6	2.6	1.4	2.3
NJJ6	4.1	3.0	-0.1	3.0	3.4	2.0	2.4	1.1	3.3	5.7	1.2	2.3	1.1	2.8	2.8	2.3	1.2	2.4
NJJ7	2.4	0.6	-0.1	0.8	4.0	0.2	1.4	2.8	2.8	6.4	0.5	0.9	0.9	2.4	2.3	4.9	-0.1	1.1
NJJ8	3.4	4.4	-0.1	5.1	2.9	2.0	1.9	1.3	3.5	5.4	2.5	1.5	0.9	2.5	2.5	1.9	1.3	2.4
NJJ8-R	4.5	1.7	-0.1	2.3	4.1	1.9	2.2	2.1	4.6	7.4	3.4	2.2	1.1	2.9	2.8	2.8	1.5	3.5
NJJ9	16.8	45.0	3.0	42.3	11.4	3.1	5.4	14.1	14.2	23.6	2.0	6.4	5.8	7.9	8.1	5.5	1.6	23.0
NK1	7.8	1.7	-0.1	1.7	3.4	2.1	0.3	2.6	6.2	5.9	8.1	3.4	1.8	3.8	4.1	0.7	0.3	1.7
NK2	3.5	2.5	-0.1	1.5	2.8	0.2	1.9	2.0	2.6	4.9	1.7	1.9	0.9	2.5	2.4	2.0	1.1	3.1
NK3	6.0	2.4	-0.1	2.7	4.7	2.1	3.4	3.5	5.8	6.4	2.3	3.0	1.6	3.3	3.8	2.5	1.1	2.5
NK4	3.2	2.0	-0.1	2.5	1.1	1.7	2.0	2.8	3.5	0.9	2.5	1.9	0.9	2.6	2.5	2.1	1.2	4.7
NK4-R	2.9	0.4	-0.1	0.2	1.1	0.2	1.8	2.2	2.8	0.8	2.2	0.9	0.9	2.4	2.5	4.9	1.2	3.6
NK5	6.1	2.8	-0.1	3.4	1.9	2.0	3.3	3.7	4.9	3.2	2.8	2.5	1.6	3.5	3.9	3.2	0.2	2.0

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NK6	4.2	1.1	-0.1	1.1	3.5	2.0	2.4	3.2	4.3	5.9	3.0	2.0	2.0	3.0	2.9	2.4	1.1	2.5
NK7	4.9	3.2	-0.1	3.5	3.0	1.9	2.6	1.5	3.4	4.9	2.9	2.1	1.1	2.8	2.8	2.3	1.2	2.7
NKK1	1.5	0.2	-0.1	0.3	-0.1	1.4	1.3	0.4	1.5	0.5	1.0	1.7	0.6	1.6	1.7	1.4	-0.1	1.2
NKK10	2.3	1.1	-0.1	1.6	2.0	1.7	1.5	1.7	2.3	3.6	2.0	0.9	0.7	1.9	1.9	1.7	-0.1	1.6
NKK11	6.1	5.5	0.4	4.8	7.0	2.1	3.1	6.2	6.1	7.5	2.0	2.8	1.4	3.4	3.5	2.7	0.3	2.5
NKK2	2.7	2.8	-0.1	3.0	2.5	1.9	1.7	0.8	2.3	4.2	2.3	1.3	0.9	2.2	2.2	1.8	1.1	1.6
NKK3	3.9	5.8	-0.1	4.6	3.1	0.8	2.0	1.4	3.2	5.9	2.3	2.1	1.0	2.7	2.6	2.1	0.3	3.2
NKK4	8.0	25.0	1.7	25.6	3.3	2.4	3.1	10.2	7.4	8.8	3.3	3.8	1.2	4.4	4.5	3.3	2.7	18.9
NKK5	9.6	23.8	1.4	24.4	3.2	2.7	3.1	14.9	9.5	9.8	3.6	4.1	1.5	4.9	5.0	3.5	2.5	21.9
NKK6	3.9	5.7	-0.1	4.2	3.2	2.2	2.2	1.5	3.4	6.5	3.0	2.3	1.1	2.9	2.8	2.3	0.4	3.3
NKK7	3.3	3.1	-0.1	2.1	2.2	0.7	1.7	0.9	2.2	4.1	1.7	1.6	0.9	2.2	2.2	1.9	1.5	2.4
NKK8	2.9	1.2	-0.1	1.2	5.4	0.2	1.9	0.7	3.2	6.9	0.5	1.7	1.0	2.5	2.7	2.1	1.1	1.8
NKK9	4.4	0.9	-0.1	0.9	4.7	1.8	2.4	2.5	4.6	7.0	1.5	2.1	2.0	2.9	2.8	2.6	1.1	1.9
NL1	1.9	1.1	-0.1	1.4	1.8	1.5	1.5	1.6	1.3	2.9	1.4	0.8	0.8	1.7	1.8	1.5	-0.1	1.1
NL2	2.1	0.3	-0.1	0.5	1.9	1.6	1.4	2.7	1.4	3.8	4.5	0.9	0.7	1.8	1.7	1.5	1.1	1.8
NL3	1.8	0.5	-0.1	0.7	2.0	1.5	1.4	1.4	1.9	2.8	1.5	0.9	0.7	1.8	1.7	1.5	-0.1	1.2
NL3-R	3.4	0.6	-0.1	1.0	3.5	1.9	2.0	1.4	2.7	5.5	0.6	1.5	1.0	2.6	2.4	2.0	1.2	2.2
NL4	1.9	0.4	-0.1	0.8	1.5	1.5	1.4	1.0	1.3	3.2	1.4	1.8	0.6	1.7	1.6	1.4	1.1	1.8
NLL1	5.7	0.3	-0.1	0.4	1.3	0.6	2.7	0.6	4.1	0.5	2.8	2.2	1.1	2.8	3.0	2.1	1.1	1.3
NLL10	2.7	2.1	-0.1	2.3	3.2	1.9	1.7	0.8	2.8	4.5	2.4	1.1	0.8	2.2	2.3	1.8	1.1	2.0
NLL11	4.7	2.1	-0.1	2.2	5.4	2.0	2.5	2.7	5.2	7.7	0.8	2.6	1.2	3.3	3.3	2.7	1.2	2.1
NLL12	4.1	1.6	-0.1	1.9	5.0	2.0	2.4	2.5	4.5	6.0	0.8	2.1	1.9	3.1	3.1	2.4	-0.1	0.5
NLL3	2.3	2.2	-0.1	0.8	1.2	1.5	1.6	0.6	1.3	2.6	1.4	1.1	0.6	1.7	1.7	1.6	-0.1	1.9
NLL4	8.9	1.2	-0.1	1.5	5.4	2.2	3.7	2.9	7.0	6.8	2.1	3.8	1.5	4.6	4.6	3.0	1.6	3.7
NLL5	7.0	22.0	1.3	20.4	3.0	3.0	2.7	7.8	6.1	11.3	3.2	3.7	1.2	4.3	4.3	3.2	1.5	14.9
NLL6	5.7	13.7	1.0	14.1	2.2	2.1	2.5	5.2	4.3	9.2	2.9	2.8	1.0	3.4	3.4	2.5	1.2	12.4
NLL7	29.3	50.1	3.5	47.4	8.6	4.1	9.5	14.2	25.9	22.6	0.3	9.3	6.7	11.5	12.0	7.3	2.9	25.9
NLL8	2.8	2.9	-0.1	2.4	2.8	0.2	1.8	0.8	2.7	4.1	1.6	1.3	0.9	2.2	2.1	1.9	1.1	1.8
NLL9	10.8	17.7	1.2	16.1	7.9	2.8	4.1	3.2	11.3	14.3	2.8	4.5	2.4	5.3	5.6	4.0	1.9	7.6
NLL9-R	11.7	22.5	1.6	21.1	8.9	2.9	4.4	3.6	12.4	16.1	2.9	4.9	5.2	5.5	5.7	3.9	2.0	7.9
NM1	9.5	5.6	0.5	5.6	8.7	2.5	3.8	2.0	9.0	10.2	2.0	3.7	1.6	4.4	4.4	3.2	0.6	3.5
NM2	4.2	1.0	-0.1	1.0	2.5	1.9	2.3	1.9	3.9	4.2	2.7	2.1	1.7	2.6	2.6	2.4	1.1	1.4
NMM1	4.4	3.1	-0.1	2.5	4.3	1.7	2.6	0.8	3.4	5.1	1.3	2.1	2.1	3.0	3.1	2.4	-0.1	1.9
NMM10	16.5	21.0	1.8	20.8	12.0	3.2	5.9	10.3	15.7	22.2	3.7	6.1	5.6	7.9	7.1	4.9	2.1	12.1
NMM2	5.9	5.3	-0.1	5.7	4.9	1.9	2.9	2.4	4.6	8.0	0.8	2.9	1.1	3.5	3.8	2.7	1.3	5.2
NMM3	4.4	3.8	-0.1	3.0	3.4	2.0	2.2	1.1	3.1	5.3	2.2	2.1	1.9	2.7	2.8	2.2	1.1	2.5
NMM3-R	3.5	1.6	-0.1	1.6	3.0	0.2	2.1	0.9	2.7	4.8	2.6	1.9	1.7	2.5	2.6	2.1	-0.1	1.9
NMM4	2.6	1.1	-0.1	1.5	2.3	0.2	1.8	0.7	2.6	3.8	2.2	1.5	0.8	2.1	2.1	1.8	-0.1	1.5
NMM5	2.7	0.4	-0.1	0.6	2.1	0.2	1.8	0.6	1.8	3.1	1.6	1.2	0.8	2.0	1.9	1.7	-0.1	1.4
NMM6	14.3	10.1	0.6	2.5	2.8	2.5	5.2	2.2	9.5	7.3	4.4	4.4	1.8	5.7	5.9	4.0	1.5	5.3
NMM7	1.6	0.2	-0.1	0.3	-0.1	0.5	1.3	0.3	0.9	1.7	1.1	0.7	0.4	1.5	1.4	1.3	-0.1	1.0
NMM8	7.0	25.1	1.7	28.6	5.9	2.5	3.6	11.1	6.6	12.3	3.6	3.6	1.3	4.0	4.2	3.1	2.3	13.6
NMM9	9.4	8.3	0.6	8.7	4.0	2.2	4.2	1.9	8.9	8.2	3.8	3.4	1.5	4.4	4.7	3.4	1.6	4.5
NN1	1.3	0.2	-0.1	0.3	1.2	1.3	1.3	1.2	0.7	1.8	0.4	1.4	0.4	1.3	1.4	1.3	-0.1	1.0
NNN1	2.4	1.1	-0.1	2.0	1.9	1.7	1.5	2.4	2.4	4.4	2.1	0.9	0.7	1.9	2.0	1.6	-0.1	1.8
NNN10	9.5	2.3	0.4	1.8	1.5	1.8	3.5	0.6	4.9	2.8	3.2	2.2	1.3	3.1	3.3	2.6	1.2	1.6
NNN2	2.1	1.1	-0.1	1.2	1.8	1.6	1.5	0.8	1.6	3.6	1.9	0.8	0.7	1.9	1.8	1.6	-0.1	1.9
NNN3	1.1	0.4	-0.1	0.8	-0.1	0.4	1.1	1.3	0.4	0.4	1.2	0.5	1.2	1.2	1.2	1.2	-0.1	1.1
NNN4	9.5	7.6	0.5	6.2	10.4	2.3	4.7	2.2	10.1	8.9	2.8	3.7	1.9	4.7	4.8	3.6	0.3	4.3
NNN5	2.0	2.9	-0.1	3.4	1.2	1.6	1.5	0.9	1.6	3.5	1.8	0.8	0.6	1.7	1.7	1.4	-0.1	1.8
NNN6	1.2	0.4	-0.1	0.5	-0.1	0.4	1.2	0.5	0.6	2.0	0.9	1.4	0.5	1.4	1.3	1.2	-0.1	1.4
NNN7	1.2	1.4	-0.1	1.8	-0.1	1.3	1.2	1.1	0.6	0.4	0.9	1.4	0.5	1.3	1.2	1.2	-0.1	1.1
NNN8	12.4	13.4	0.9	12.2	10.0	2.8	5.2	6.2	13.1	12.5	3.1	4.5	4.0	5.7	5.6	4.1	2.0	5.8
NNN9	12.2	12.6	0.9	12.6	8.4	2.7	4.7	3.2	11.6	13.1	2.9	4.6	5.0	5.6	6.0	3.8	2.0	6.9
NO1	3.4	2.5	-0.1	1.1	2.9	0.7	1.9	0.9	2.5	4.8	2.5	1.5	0.9	2.4	2.5	1.9	1.2	1.9
NO2	0.7	0.2	-0.1	0.2	-0.1	0.4	1.1	1.1	0.4	0.3	0.1	1.2	0.4	1.2	1.2	1.1	-0.1	0.8
NOO1	1.9	3.6	-0.1	2.9	1.3	0.2	1.4	1.1	2.0	3.4	1.8	0.9	0.6	1.8	1.7	1.4	-0.1	2.5
NOO2	3.8	5.5	-0.1	4.1	4.0	1.8	2.0	1.6	3.4	6.6	2.9	2.1	1.1	2.7	2.6	2.2	1.3	3.9
NOO3	3.5	2.9	-0.1	2.4	1.4	0.3	2.1	0.6	2.1	2.5	2.2	1.3	0.8	1.7	2.0	1.8	-0.1	1.8
NOO4	4.9	4.6	-0.1	3.8	5.0	1.9	2.7	1.5	5.2	5.7	1.9	2.3	1.4	3.0	3.0	2.3	1.2	3.4
NOO5	1.7	12.7	-0.1	13.3	1.6	1.8	1.3	2.4	0.7	3.7	1.7	1.9	0.6	1.7	1.6	1.5	1.1	1.9
NOO6	1.6	0.3	-0.1	0.4	-0.1	1.4	1.3	0.5	1.0	0.5	1.1	1.0	0.6	1.6	1.5	1.4	-0.1	1.4
NOO6-R	1.5	0.6	-0.1	0.7	-0.1	1.4	1.4	0.6	1.0	0.6	1.2	1.6	0.7	1.5	1.4	1.3	-0.1	1.4
NOO7	1.1	0.7	-0.1	0.8	1.2	1.3	1.1	0.6	0.3	2.2	0.9	1.4	0.3	1.4	1.3	1.3	-0.1	1.5
NOO8	8.0	15.7	1.4	15.4	6.5	2.5	3.4	4.1	8.1	15.1	2.0	3.9	1.4	4.6	4.7	3.5	2.5	9.7
NP1	2.9	1.3	-0.1	1.8	2.6	1.7	2.0	0.7	2.3	4.2	2.4	1.4	0.8	2.3	2.3	1.9	-0.1	1.5

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A14-06865

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NP2	1.8	1.1	-0.1	1.3	1.3	1.5	1.4	0.5	1.2	2.6	1.1	1.7	0.5	1.6	1.6	1.5	-0.1	1.3
NP3	1.9	1.6	-0.1	1.9	1.2	1.4	1.4	0.6	1.2	2.5	-0.1	1.7	0.6	1.6	1.7	1.4	-0.1	1.3
NP4	16.2	2.8	0.8	3.7	8.1	3.2	6.1	2.2	16.7	14.1	4.0	6.1	5.7	7.7	7.9	5.3	1.9	5.9
NPP4	1.3	3.5	-0.1	4.1	-0.1	1.3	1.1	0.7	0.7	2.3	0.9	1.4	0.5	1.2	1.2	1.2	-0.1	1.6
NPP1-R	1.0	2.9	-0.1	3.3	-0.1	0.4	1.1	0.7	0.6	2.3	0.7	1.3	0.5	1.3	1.2	1.2	-0.1	1.4
NPP2	5.5	6.5	-0.1	5.4	2.4	1.8	2.7	1.9	5.3	5.5	3.4	2.7	1.3	3.1	3.4	2.4	1.3	4.9
NPP3	3.3	5.9	-0.1	6.4	3.5	2.0	2.0	1.2	2.9	5.7	2.7	1.5	1.0	2.5	2.4	2.0	1.2	2.7
NPP4	11.3	4.1	0.4	4.1	10.2	2.4	4.6	3.8	11.5	9.5	2.8	4.1	2.0	5.9	5.0	3.5	1.5	3.1
NQ1	4.1	0.8	-0.1	1.0	2.8	0.3	2.6	0.5	3.4	3.9	3.0	2.1	1.1	2.9	3.0	2.2	1.2	1.2
NQ2	2.6	3.0	-0.1	2.1	2.8	0.2	1.7	0.8	2.8	4.5	2.4	1.3	0.9	2.2	2.2	1.8	1.2	2.1
NQ2-R	2.6	2.1	-0.1	2.2	3.3	0.2	1.8	1.0	2.4	5.3	2.5	1.9	1.6	2.4	2.4	1.9	1.2	2.3
NQ3	1.4	2.8	-0.1	2.4	1.3	0.4	1.3	0.7	1.1	3.0	1.0	1.6	0.5	1.5	1.5	1.1	-0.1	1.9
NQ4	3.8	2.1	-0.1	2.1	3.1	1.9	2.3	0.7	3.5	4.6	2.7	2.0	1.7	2.5	2.5	2.1	1.2	1.7
NQ5	7.6	0.4	-0.1	0.6	3.5	2.0	4.0	2.2	5.1	5.1	3.7	2.4	1.5	3.4	3.4	2.7	1.3	1.3
NQ6	1.2	1.5	-0.1	1.8	-0.1	1.3	1.3	0.5	0.7	2.3	-0.1	1.3	0.5	1.3	1.3	1.3	-0.1	1.2
NQ7	2.9	1.7	-0.1	2.4	4.4	2.2	3.2	3.6	6.8	5.9	3.7	2.7	2.6	3.7	3.7	3.0	1.2	2.5
NQ8	4.0	1.5	-0.1	1.5	4.0	2.0	2.4	2.0	4.1	5.9	2.9	1.4	1.1	2.8	2.6	2.3	1.1	0.3
NQQ1	3.5	1.5	-0.1	2.1	5.4	2.0	2.2	2.1	3.8	7.1	1.7	1.7	1.3	2.5	2.5	2.1	1.2	1.4
NR1	1.2	1.6	-0.1	0.3	-0.1	0.5	1.4	0.5	0.7	1.7	0.9	1.4	0.5	1.4	1.4	1.3	-0.1	1.7
NR11	5.4	7.6	-0.1	8.6	4.4	1.9	2.7	1.7	5.2	7.9	3.4	2.4	1.2	3.4	3.4	2.7	1.2	3.7
NR12	9.2	2.7	-0.1	2.7	6.9	2.2	4.3	3.2	7.0	8.3	2.5	3.1	1.7	4.3	4.3	3.3	1.3	2.8
NR13	19.5	14.0	1.0	14.2	5.8	2.9	7.0	3.8	14.7	12.4	9.4	8.3	5.0	6.8	7.4	4.7	1.9	7.1
NR14	2.0	3.8	-0.1	4.7	1.5	0.3	1.5	1.2	2.3	1.0	1.6	1.0	0.8	2.1	2.0	1.9	1.2	3.1
NR2	3.6	2.0	-0.1	2.0	3.4	0.2	2.0	1.0	2.8	5.6	2.7	1.5	1.0	2.5	2.4	2.0	1.2	2.5
NR3	14.5	4.4	-0.1	3.0	2.4	2.0	5.3	1.5	10.0	5.4	4.4	4.0	5.1	5.5	5.8	3.8	1.3	3.2
NR4	2.9	1.0	-0.1	1.3	1.4	0.2	1.9	1.6	2.0	2.9	1.6	1.0	0.8	2.0	1.9	1.8	-0.1	1.3
NR5	2.3	1.6	-0.1	1.2	1.5	1.6	1.5	0.6	2.0	2.6	1.1	0.9	0.7	1.8	1.8	1.5	-0.1	1.4
NR6	2.3	1.9	-0.1	0.6	1.7	1.6	1.6	1.6	1.6	2.9	1.6	0.9	1.8	1.7	1.7	1.4	-0.1	1.2
NR7	6.5	6.4	-0.1	5.1	4.1	2.1	3.0	3.8	7.3	9.3	3.8	3.0	1.4	3.8	4.0	2.9	1.6	4.7
NR8	2.0	0.3	-0.1	0.8	1.4	1.5	1.4	0.6	1.4	2.7	1.4	0.8	0.6	1.7	1.7	1.4	-0.1	1.5
NS1	3.1	0.3	-0.1	0.3	-0.1	0.2	0.2	1.9	0.5	1.4	1.4	1.7	1.0	0.7	0.4	0.3	-0.1	1.8
NS10	7.2	6.0	0.5	6.0	2.7	1.4	0.3	3.6	7.1	6.1	3.0	3.1	2.9	3.5	3.8	3.9	0.4	3.2
NS11	4.9	4.6	-0.1	5.4	8.7	1.8	2.9	3.8	4.4	6.2	1.8	2.6	1.4	3.4	3.2	2.5	1.2	3.4
NS12	6.5	3.9	-0.1	4.3	8.0	2.2	0.2	3.5	6.3	7.5	1.5	3.5	1.7	3.6	3.9	5.5	0.4	3.2
NS13	9.4	7.4	-0.1	8.6	7.8	2.2	4.4	2.2	7.8	9.0	1.9	3.1	1.9	4.5	4.6	3.3	0.3	4.6
NS14	16.9	16.7	0.6	19.4	18.4	2.1	7.4	9.7	22.4	0.5	5.4	2.5	7.5	8.6	8.5	5.7	1.2	8.7
NS2	2.7	1.4	-0.1	1.4	1.8	0.2	1.7	2.0	1.7	3.2	1.5	1.3	0.7	1.7	1.8	1.7	0.2	1.8
NS3	9.7	2.3	-0.1	3.2	2.9	2.2	4.4	3.6	6.7	3.8	5.8	3.2	1.6	3.8	3.9	9.0	0.3	0.4
NS4	2.6	5.0	-0.1	5.7	-0.1	0.2	1.6	2.5	1.7	3.7	1.4	0.9	0.8	2.2	2.0	1.8	1.2	3.3
NS4-R	2.4	4.3	-0.1	4.9	-0.1	1.5	1.6	2.1	1.5	3.3	1.1	0.8	0.7	2.0	1.9	1.6	1.2	2.8
NS5	1.7	4.3	-0.1	5.0	1.8	1.6	1.4	1.9	1.4	4.6	1.5	1.9	0.9	1.7	0.5	1.7	0.3	3.1
NS6	2.4	1.4	-0.1	1.7	1.2	1.7	1.6	0.7	2.3	2.9	1.4	0.9	0.7	2.0	2.0	1.6	1.1	1.6
NS7	3.4	7.0	-0.1	8.0	2.3	0.2	2.0	2.4	2.5	5.6	2.4	1.9	1.1	2.3	2.4	2.1	0.3	4.5
NS8	7.7	6.4	-0.1	5.0	3.6	1.9	4.2	2.1	6.1	7.4	3.9	3.0	1.7	3.7	4.0	3.2	0.4	3.6
NS9	1.5	2.1	-0.1	0.7	-0.1	1.3	0.3	0.8	0.7	2.5	0.9	1.4	0.9	1.4	1.4	1.4	-0.1	1.6
NT1	4.8	1.4	-0.1	1.6	1.7	0.2	0.3	1.5	2.6	2.6	2.0	2.0	1.3	1.3	2.5	2.3	-0.1	1.2
NT10	5.8	0.7	-0.1	0.8	1.6	1.8	2.7	0.7	3.8	3.0	2.6	2.0	1.1	2.6	2.8	2.0	1.3	1.6
NT11	2.2	2.4	-0.1	2.7	-0.1	1.5	1.4	0.6	1.1	0.5	1.4	0.7	0.7	1.7	1.6	1.5	-0.1	1.8
NT12	5.6	1.5	-0.1	0.4	2.8	2.0	2.6	0.8	5.6	4.7	3.1	2.3	1.2	3.0	2.9	2.4	1.6	1.5
NT13	2.9	1.6	-0.1	2.4	1.7	0.2	1.7	0.6	2.1	3.5	2.1	1.3	0.6	2.0	2.0	1.7	1.1	1.9
NT14	3.1	1.0	-0.1	1.0	3.4	0.2	0.4	2.0	3.0	5.0	0.5	2.2	1.2	2.3	2.5	2.4	0.3	1.4
NT15	13.1	4.0	-0.1	4.1	7.0	2.4	-0.1	6.3	9.3	7.6	0.9	4.4	2.5	4.3	5.5	3.2	0.3	3.7
NT16	8.1	3.6	-0.1	3.6	5.8	2.1	4.0	3.6	7.7	9.0	4.1	3.0	1.6	3.7	4.0	3.1	0.3	2.6
NT17	3.0	3.6	-0.1	4.1	3.5	0.2	2.0	2.7	2.9	5.1	0.5	2.2	1.1	2.3	2.6	2.2	1.2	2.5
NT18	3.0	0.2	-0.1	0.3	-0.1	0.2	0.2	1.2	1.4	1.3	1.3	1.0	0.5	1.7	1.8	1.6	-0.1	1.0
NT2	5.4	1.8	-0.1	1.9	1.4	0.6	0.6	1.7	3.1	2.7	1.8	2.1	1.3	2.2	2.5	2.4	0.2	0.8
NT3	1.1	1.6	-0.1	1.9	-0.1	1.3	0.2	0.5	0.4	2.0	-0.1	1.3	0.5	1.3	1.2	1.2	-0.1	1.3
NT4	1.5	1.0	-0.1	0.5	-0.1	1.3	0.6	0.9	0.9	2.3	-0.1	1.5	0.6	1.4	1.3	1.3	-0.1	1.4
NT5	2.2	1.1	-0.1	1.2	-0.1	0.2	1.5	0.4	1.3	0.5	1.4	0.8	0.6	1.7	1.6	1.5	-0.1	1.2
NT6-R	2.6	1.2	-0.1	1.6	1.2	0.2	1.7	0.6	1.9	2.8	0.9	0.8	0.7	2.0	1.9	1.7	1.1	1.4
NT6	2.0	0.3	-0.1	0.5	1.6	1.6	1.4	0.7	1.5	3.2	1.4	0.8	0.6	1.9	1.8	1.6	-0.1	1.7
NT7	5.5	0.4	-0.1	0.6	1.2	0.6	2.6	2.0	3.3	0.7	2.5	2.0	1.0	2.5	2.7	2.1	1.3	3.1
NT8	8.0	7.7	-0.1	6.5	4.4	2.4	3.4	2.3	6.6	8.7	4.0	3.2	1.5	4.5	4.8	3.4	1.7	5.2
NT9	2.4	0.8	-0.1	1.4	1.2	0.2	1.5	0.5	1.5	2.1	1.1	0.7	0.8	1.8	1.6	1.4	-0.1	1.4
NU1	2.0	4.2	-0.1	4.8	-0.1	0.2	1.6	1.0	2.2	3.2	1.8	0.9	0.7	1.8	1.8	1.5	1.2	1.9
NU10	4.3	4.3	-0.1	1.1	3.3	0.2	2.4	1.4	3.2	6.1	8.0	2.1	2.0	2.8	2.8	2.3	1.3	2.9
NU11	3.1	2.4	-0.1	1.9	0.8	0.2	0.4	1.8	1.9	2.7	1.6	1.8	0.9	1.8	2.0	2.0	0.3	1.6

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NU12	1.9	4.5	-0.1	3.7	2.2	1.6	1.9	2.2	2.2	4.3	0.5	2.0	0.3	1.9	1.9	1.7	0.2	1.5
NU12-R	1.9	4.7	-0.1	4.1	2.0	1.6	1.4	0.8	1.9	3.6	1.7	1.8	0.6	1.8	1.8	1.7	0.2	1.4
NU13	6.0	8.5	0.5	1.9	5.8	2.3	2.9	1.9	5.7	7.7	5.1	2.6	3.4	3.8	3.8	6.3	1.1	4.9
NU14	4.0	0.8	-0.1	0.8	5.2	2.0	2.1	2.5	4.3	8.0	0.5	1.4	1.9	3.0	2.9	2.4	1.2	1.7
NU15	4.8	4.4	-0.1	1.6	4.6	1.9	2.0	2.9	4.8	6.8	3.5	2.5	2.3	2.9	3.2	2.4	0.3	2.1
NU16	7.9	16.8	0.5	19.6	4.3	2.2	3.5	9.4	7.8	10.4	3.8	3.6	1.5	4.2	4.6	3.4	0.4	9.0
NU17	7.2	6.9	0.4	5.0	10.7	2.3	3.5	4.6	7.6	8.3	2.1	2.8	1.8	4.0	4.3	6.3	0.4	4.3
NU18	2.8	5.9	-0.1	7.0	3.7	0.2	2.2	3.8	4.1	6.3	3.0	1.8	1.9	2.3	2.5	2.1	1.2	3.2
NU19	5.4	1.4	-0.1	1.4	5.4	2.0	3.2	2.8	5.4	8.1	1.7	2.8	1.6	3.1	3.2	2.9	0.3	1.1
NU2	13.7	17.5	1.0	16.1	3.9	2.9	0.5	10.8	8.8	2.9	1.7	4.7	4.9	4.9	5.7	4.8	2.0	11.5
NU3	2.4	1.9	-0.1	1.3	0.1	0.2	0.3	1.6	1.2	2.1	1.3	1.6	0.9	1.5	0.5	1.6	0.1	1.3
NU4	9.3	13.9	0.5	15.6	5.3	2.5	3.6	6.5	7.1	12.1	4.0	3.8	1.6	4.8	4.9	3.4	1.6	9.5
NU5	1.3	2.7	-0.1	3.0	0.1	1.8	1.2	0.5	0.8	2.2	0.9	1.4	0.5	1.4	1.3	1.6	0.1	1.7
NU6	1.8	3.1	-0.1	3.5	1.3	1.4	1.4	1.1	1.2	3.0	1.1	1.6	0.7	1.6	1.6	1.5	1.1	2.1
NU7	2.5	4.9	-0.1	5.5	2.0	0.2	1.6	1.6	1.8	4.7	2.1	2.1	0.9	1.9	0.6	1.8	0.3	2.6
NU8	4.8	5.8	-0.1	4.3	6.6	2.1	2.5	2.2	3.8	9.3	1.5	2.9	2.4	3.0	3.2	2.9	0.3	4.0
NU9	6.3	2.6	-0.1	2.9	5.8	2.1	3.8	2.3	4.7	7.0	2.2	2.0	2.7	3.6	3.7	3.0	1.2	1.0
NV1	9.7	4.8	0.5	4.7	11.4	2.3	4.2	5.0	8.7	11.5	2.1	3.9	1.5	4.6	4.7	3.4	1.7	3.7
NV10	4.7	2.2	-0.1	1.5	1.3	0.2	2.7	1.6	3.7	2.6	2.5	1.8	1.1	2.3	2.4	2.0	0.1	1.5
NV11	3.7	2.8	-0.1	1.7	3.5	1.9	2.2	2.1	2.8	5.2	0.7	1.1	1.0	2.5	2.4	0.8	1.2	1.4
NV12	4.9	1.4	-0.1	1.9	4.4	1.9	2.5	0.8	4.4	6.1	0.8	2.2	1.9	2.9	3.0	2.2	1.3	1.6
NV13	9.3	5.0	0.5	5.0	4.5	2.2	4.8	3.7	8.6	8.3	3.6	3.1	1.9	3.9	4.2	3.5	0.4	2.6
NV14	5.1	1.4	-0.1	1.4	5.3	2.0	0.5	2.7	5.0	7.5	0.4	2.9	1.7	3.1	3.2	6.2	1.2	1.8
NV15	3.8	1.1	-0.1	1.2	3.6	0.2	2.3	1.9	3.6	4.9	0.7	1.6	0.9	2.4	2.4	2.1	1.2	1.4
NV16	13.9	2.6	-0.1	3.4	11.0	2.8	6.0	4.4	12.7	12.0	3.6	4.9	2.3	6.2	6.2	3.4	1.8	0.8
NV17	9.8	2.2	-0.1	2.7	6.8	2.0	5.1	3.3	7.6	7.2	1.7	3.1	2.0	4.1	4.1	3.0	1.3	2.5
NV18	6.5	2.2	-0.1	2.2	7.1	2.0	0.2	2.0	4.9	9.6	4.3	3.0	1.8	2.9	3.1	2.8	0.3	2.6
NV2	6.7	2.4	-0.1	2.6	4.8	2.0	3.3	1.4	6.1	7.8	3.7	2.7	1.4	3.5	3.5	2.7	1.4	3.0
NV3	2.7	1.5	-0.1	1.5	2.8	0.2	0.2	2.4	1.9	4.7	0.6	2.1	1.0	0.9	2.2	1.0	1.1	1.7
NV4	6.5	1.8	-0.1	1.8	8.6	2.0	0.2	3.6	5.3	11.5	1.6	3.4	2.0	3.5	3.8	3.4	0.3	2.8
NV5	7.7	2.3	-0.1	2.3	7.5	2.2	3.8	4.0	7.6	11.9	1.9	3.0	1.9	4.2	4.2	3.2	1.3	3.4
NV5-R	9.0	3.5	-0.1	3.5	12.2	2.2	4.6	1.7	8.7	8.3	1.9	3.4	1.7	4.6	4.9	3.4	1.3	3.6
NV6	6.4	3.1	-0.1	2.6	0.1	0.7	0.1	1.9	3.0	0.6	2.6	2.0	1.0	2.5	2.7	2.2	1.1	1.5
NV7	3.3	1.3	-0.1	1.5	4.6	0.2	0.3	2.8	6.5	6.5	0.5	2.6	1.4	2.6	2.7	2.6	1.1	2.5
NV8	4.2	1.0	-0.1	1.0	2.5	0.2	0.4	1.8	3.9	4.1	4.5	2.2	1.3	1.6	2.4	2.3	0.3	1.4
NV9	4.2	2.1	-0.1	2.1	5.3	1.8	0.2	3.1	3.3	7.5	1.3	2.5	1.4	2.5	2.7	2.6	0.3	2.6
NW1	5.6	2.4	1.3	4.0	8.2	2.2	2.3	3.8	5.8	15.1	0.6	3.0	1.2	3.5	3.5	2.7	1.5	8.7
NW10	3.2	3.0	-0.1	3.6	1.5	0.6	0.5	2.3	2.1	3.0	1.7	1.9	1.2	1.0	2.2	2.2	1.1	2.3
NW11	2.6	1.6	-0.1	1.9	2.4	1.6	1.7	1.8	2.4	3.8	2.1	1.0	0.8	1.9	1.8	1.5	0.1	1.5
NW12	8.5	0.3	-0.1	0.3	2.3	2.2	3.9	1.1	6.3	4.9	3.8	2.8	1.6	3.9	3.9	2.9	1.5	1.9
NW13	21.7	9.7	0.7	9.1	5.4	2.1	8.7	1.9	13.6	8.4	5.5	4.9	5.9	6.7	6.9	4.5	0.3	5.0
NW14	3.1	3.3	-0.1	3.6	1.6	0.2	1.9	0.8	2.3	3.3	2.2	1.2	0.8	2.1	2.2	1.8	1.1	1.6
NW15	12.9	12.4	0.7	3.5	12.1	2.8	5.7	2.9	12.4	11.1	3.1	4.4	5.7	5.8	5.8	4.1	0.7	5.7
NW16	5.1	1.1	-0.1	1.3	2.5	0.6	2.7	1.8	3.2	3.9	2.0	1.9	1.0	2.5	2.6	2.0	1.1	0.4
NW17	2.8	2.3	-0.1	0.5	2.4	0.2	0.3	1.9	2.0	3.7	1.0	2.1	1.1	2.0	2.4	2.0	0.3	1.6
NW17-R	2.7	2.3	-0.1	0.5	2.7	0.2	0.3	1.9	2.0	3.8	1.9	2.1	1.1	2.0	2.4	2.1	1.2	1.7
NW18	4.0	6.5	-0.1	7.6	1.4	0.5	2.3	3.0	3.4	3.7	2.5	3.0	1.1	2.3	2.5	2.1	1.1	3.6
NW2	4.2	2.1	-0.1	2.1	3.4	0.2	0.2	2.6	3.1	5.3	2.4	2.5	1.5	2.5	2.8	2.6	0.3	2.2
NW2-R	3.7	3.4	-0.1	2.8	3.5	0.2	0.3	2.4	2.8	5.0	2.9	2.4	1.3	2.6	2.9	2.4	1.1	2.1
NW3	2.1	2.5	-0.1	1.8	3.6	2.1	1.7	2.0	1.8	4.8	0.4	2.1	0.9	1.9	2.0	1.8	1.1	1.9
NW4	2.6	3.0	-0.1	2.4	3.1	0.2	1.8	2.1	1.9	4.3	2.2	1.3	0.8	2.0	2.2	1.9	1.1	1.7
NW5	6.7	26.9	0.6	8.9	5.4	2.2	3.3	8.6	5.9	16.0	3.9	3.4	1.6	4.0	4.4	3.4	0.5	15.1
NW6	2.8	1.9	-0.1	1.9	3.8	0.2	1.8	2.2	2.6	5.4	0.3	1.5	1.6	2.1	2.4	1.9	1.2	2.0
NW7	2.1	0.8	-0.1	0.8	3.5	1.6	0.3	2.0	2.3	5.2	0.3	2.1	0.9	1.9	2.0	1.9	1.1	1.5
NW8	33.9	17.2	1.1	18.0	2.5	3.2	4.7	4.7	19.8	5.6	1.5	8.2	5.9	9.6	10.1	6.4	1.1	11.7
NW9	2.5	4.8	-0.1	3.5	3.0	0.2	1.7	3.4	1.9	5.6	2.3	2.1	1.0	1.9	2.0	1.8	0.3	2.9
NX1	2.9	2.1	-0.1	0.2	0.1	0.2	1.9	0.6	1.9	0.6	1.6	1.1	0.8	2.1	2.2	1.7	0.1	1.4
NX10	3.7	2.7	-0.1	0.7	4.4	0.8	2.2	1.0	3.5	6.4	0.5	2.2	1.1	2.8	2.7	2.2	1.3	2.1
NX11	1.2	1.2	-0.1	1.5	0.6	1.3	0.3	1.4	0.5	2.3	0.2	1.4	0.4	1.4	0.4	1.4	0.1	1.1
NX12	1.9	0.5	-0.1	1.2	1.9	0.2	0.3	1.5	1.2	2.9	1.0	1.8	0.7	1.7	0.5	1.6	0.1	1.1
NX13	6.1	5.5	-0.1	6.2	5.0	1.9	3.5	1.6	4.6	8.1	1.7	2.5	2.6	3.0	3.2	2.6	0.3	2.1
NX14	4.1	2.4	-0.1	2.7	2.3	0.8	2.5	2.4	3.1	4.4	1.5	2.3	1.2	2.3	2.5	2.3	0.3	1.5
NX14-R	4.5	1.8	-0.1	1.9	3.0	0.2	0.1	2.2	3.2	5.0	2.9	2.5	1.3	2.5	2.7	2.4	0.3	1.3
NX15	3.0	4.1	-0.1	4.8	4.7	0.2	0.1	3.4	2.8	7.6	0.5	2.5	1.3	2.5	2.6	2.4	0.3	2.6
NX16	3.4	5.2	-0.1	3.7	5.0	1.9	2.1	3.5	3.7	7.8	0.5	1.8	1.1	2.5	2.5	2.1	0.3	2.4
NX17	10.6	11.5	0.5	10.0	15.7	2.7	4.5	2.9	10.8	10.8	2.6	4.1	3.6	5.0	5.1	3.9	1.1	5.3
NX2	2.1	2.6	-0.1	0.6	2.0	0.2	0.4	1.9	1.4	3.2	1.5	1.9	0.8	1.8	1.9	1.7	0.3	1.6

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019-LB	020-LA	021-LPH	022-LBA	023-LAR	024-LB	025-LAR	026-LBA	027-LB	028-ALK	029-HB	030-HB	031-HB	032-HB	033-HB	034-HB	035-LAR	036-LBA
NX3	1.9	1.2	-0.1	1.5	1.4	0.2	1.4	1.6	1.4	2.7	1.1	0.8	0.7	1.7	1.7	1.5	1.2	1.3
NX4	8.0	10.2	-0.1	3.0	1.9	2.0	3.2	2.9	7.2	6.9	3.4	3.2	1.4	4.1	4.2	3.1	1.7	6.0
NX5	1.8	4.0	-0.1	3.1	1.9	1.5	1.3	1.0	1.7	4.1	1.3	1.8	0.6	1.6	1.5	1.5	1.1	2.3
NX6	2.9	3.8	-0.1	2.8	3.4	0.2	2.0	2.6	2.9	5.5	0.4	2.3	1.0	2.3	2.4	2.2	0.3	2.0
NX7	25.2	19.1	1.0	17.2	4.5	2.3	9.9	6.1	18.0	13.3	4.0	6.3	7.0	7.7	8.7	6.2	1.8	14.6
NX8	4.7	7.2	-0.1	2.9	4.6	1.9	2.7	1.8	4.7	8.2	3.2	2.5	1.3	3.0	3.3	2.6	0.3	0.2
NX9	5.6	3.4	-0.1	3.4	3.8	2.0	3.0	1.4	5.5	7.5	3.4	2.7	2.4	3.1	3.4	2.7	0.4	2.4
NY1	2.6	0.8	-0.1	0.8	2.4	0.2	0.3	2.1	1.8	3.8	1.8	2.0	1.1	1.9	2.1	2.0	1.1	1.7
NY10	9.4	5.6	0.5	5.6	5.7	2.3	4.9	4.6	9.5	10.1	3.9	3.3	2.0	4.6	4.7	3.8	0.3	4.1
NY11	2.2	1.0	-0.1	1.3	1.6	1.6	1.5	1.6	1.4	3.0	1.5	0.8	0.7	1.9	1.7	1.6	-0.1	1.3
NY12	2.7	0.7	-0.1	0.9	1.2	0.2	1.8	1.0	1.9	2.6	1.4	0.9	0.8	2.3	2.2	1.8	1.1	1.9
NY13	2.5	0.9	-0.1	1.0	1.7	1.6	1.7	1.5	1.7	2.6	1.8	0.9	0.7	1.9	1.9	1.7	-0.1	1.0
NY14	12.3	2.3	-0.1	2.8	8.6	2.4	5.4	4.1	11.3	7.7	3.2	4.0	2.2	5.4	5.5	3.9	0.3	2.4
NY15	6.5	3.0	-0.1	3.0	10.4	2.0	3.9	3.9	10.2	7.6	2.6	3.5	1.8	3.9	3.9	5.0	1.3	3.0
NY2	2.7	0.6	-0.1	0.8	1.8	1.6	1.7	1.6	1.8	2.7	1.5	0.8	0.7	2.0	1.9	1.6	-0.1	1.2
NY3	1.4	0.9	-0.1	1.1	1.3	0.5	1.4	2.3	0.8	2.5	1.0	1.9	0.4	1.4	1.4	-0.1	-0.1	1.0
NY4	33.3	20.6	1.4	21.4	9.3	2.7	11.1	13.6	22.8	16.7	2.0	9.1	7.1	11.5	11.6	7.0	1.4	14.8
NY4-R	41.1	29.2	1.9	29.8	10.9	3.0	13.3	8.6	28.2	20.6	2.2	11.1	7.9	13.6	13.8	8.0	1.6	19.0
NY5	3.1	2.2	-0.1	2.5	3.0	0.2	0.4	2.4	2.2	4.5	2.0	2.2	1.1	2.2	2.3	2.2	0.2	1.8
NY6	7.8	4.7	-0.1	5.2	5.1	2.1	3.4	4.5	6.0	7.8	3.9	2.7	1.5	4.0	3.8	2.9	1.3	4.2
NY7	12.1	1.7	-0.1	2.7	8.1	2.3	5.0	8.1	10.4	10.1	3.0	3.6	2.1	5.0	5.1	3.6	1.5	5.9
NY8	2.5	1.5	-0.1	2.4	2.5	0.2	0.4	2.1	2.1	3.9	2.4	2.2	1.1	2.2	2.4	2.3	1.1	1.6
NY9	6.8	2.4	-0.1	1.9	-0.1	1.8	0.3	1.8	4.0	0.6	2.4	2.0	1.3	2.6	2.8	2.4	1.1	1.2
NZ1	3.2	1.4	-0.1	1.8	3.4	2.1	2.1	2.2	2.9	5.2	0.4	1.2	1.6	2.5	2.4	2.0	1.2	1.5
NZ10	24.1	11.9	0.8	12.2	14.9	2.8	10.5	6.6	22.7	14.2	5.7	7.8	7.4	10.0	10.6	6.5	0.6	5.2
NZ11	4.1	1.0	-0.1	1.1	2.6	1.9	2.4	2.1	2.9	3.8	2.6	1.9	1.0	2.6	2.6	2.1	-0.1	1.7
NZ12	5.9	3.2	-0.1	3.2	4.2	1.9	0.3	2.9	5.8	3.1	3.0	3.0	1.7	3.0	3.4	3.0	0.3	2.0
NZ13	2.5	1.6	-0.1	1.6	1.9	0.2	1.3	1.8	1.6	2.9	1.0	1.9	0.9	1.0	1.8	1.7	-0.1	1.5
NZ2	3.7	1.3	-0.1	1.8	-0.1	0.7	1.7	1.7	2.5	0.8	1.8	1.9	0.9	2.5	2.6	1.9	1.4	3.4
NZ2-R	2.0	0.2	-0.1	0.3	-0.1	0.1	1.4	1.5	1.2	0.4	1.3	1.0	0.7	1.7	1.7	1.5	-0.1	1.2
NZ3	3.8	1.5	-0.1	1.8	3.9	1.9	2.1	4.0	2.1	5.3	2.8	1.2	1.1	2.7	2.5	2.1	1.3	3.3
NZ4	4.4	2.5	-0.1	2.1	5.0	2.1	2.2	2.7	4.0	5.9	0.5	1.2	1.9	2.9	2.7	2.6	1.2	2.0
NZ5	4.0	1.2	-0.1	1.2	4.3	2.1	2.4	2.4	4.3	5.9	0.8	2.3	2.0	2.8	2.9	2.5	1.2	1.7
NZ6	2.6	1.5	-0.1	1.8	2.2	0.2	1.9	2.1	2.8	3.9	1.6	1.5	1.4	2.3	2.4	2.1	1.1	0.3
NZ7	6.4	1.2	-0.1	1.2	7.7	2.0	3.7	2.6	6.8	9.2	1.8	2.5	2.8	4.0	3.9	3.0	1.1	0.4
NZ8	4.5	1.8	-0.1	0.8	8.2	1.8	2.4	6.2	3.7	6.2	1.9	2.3	2.2	3.0	3.0	2.4	1.2	2.0
NZ9	3.6	1.2	-0.1	1.5	3.1	0.2	2.4	1.8	2.3	4.2	1.9	1.8	1.6	2.4	2.4	1.9	-0.1	1.3
LMB-QA	-0.1	1.1	-0.1	1.3	-0.1	1.0	-0.1	1.3	-0.1	0.6	-0.1	1.1	-0.1	1.1	1.2	1.1	-0.1	0.3
LMB-QA	-0.1	1.0	-0.1	1.2	-0.1	1.0	-0.1	1.3	-0.1	1.0	-0.1	1.0	-0.1	0.1	1.0	1.1	-0.1	1.1
LMB-QA	-0.1	0.7	-0.1	0.9	-0.1	1.0	-0.1	1.2	-0.1	1.6	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	1.0
LMB-QA	-0.1	0.3	-0.1	0.4	-0.1	1.0	-0.1	1.0	-0.1	0.5	-0.1	1.0	-0.1	1.1	-0.1	1.1	-0.1	1.1
LMB-QA	-0.1	0.2	-0.1	0.4	-0.1	1.0	-0.1	1.2	-0.1	0.4	-0.1	1.1	-0.1	0.2	1.2	1.1	-0.1	0.3
LMB-QA	-0.1	0.3	-0.1	0.5	-0.1	1.0	-0.1	1.2	-0.1	0.4	-0.1	1.0	-0.1	1.1	1.0	1.1	-0.1	0.3
LMB-QA	-0.1	0.6	-0.1	0.8	-0.1	1.0	-0.1	1.2	-0.1	0.5	-0.1	1.0	-0.1	1.1	1.0	1.1	-0.1	1.0
LMB-QA	-0.1	0.4	-0.1	0.6	-0.1	1.1	-0.1	1.3	-0.1	0.4	-0.1	1.1	-0.1	1.1	1.1	1.1	-0.1	1.1
LMB-QA	-0.1	0.3	-0.1	0.4	-0.1	1.1	-0.1	1.2	-0.1	0.4	-0.1	1.1	-0.1	1.1	1.1	1.1	-0.1	1.0
LMB-QA	-0.1	0.2	-0.1	0.2	-0.1	1.0	-0.1	1.1	-0.1	0.3	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	0.3
LMB-QA	-0.1	0.2	-0.1	0.4	-0.1	1.0	-0.1	1.1	-0.1	0.4	-0.1	1.0	-0.1	1.1	0.2	-0.1	-0.1	1.0
LMB-QA	-0.1	0.2	-0.1	0.3	-0.1	1.0	-0.1	1.1	-0.1	0.4	-0.1	1.0	-0.1	0.1	1.1	0.2	-0.1	1.0

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	019 LB	020 LA	021 LPH	022 LBA	023 LAR	024 LB	025 LAR	026 LBA	027 LB	028 ALK	029 HB	030 HB	031 HB	032 HB	033 HB	034 HB	035 LAR	036 LBA

A14-06865

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SOIL GAS HYDROCARBONS
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Activation Laboratories Ltd.
Date: November 25, 2014
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	037-HB	038-LBA	039-LAR	040-LRB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-NB	050-LBA	051-LB	052-LPB	053-LPB	054-HB
NA1	2.5	4.2	2.5	4.3	11.0	2.1	1.4	0.3	10.6	1.2	5.8	1.9	1.5	3.8	0.7	0.9	1.8	1.5
NA2	1.1	1.1	0.8	0.9	1.8	0.9	0.4	1.0	1.6	-0.1	1.4	0.9	0.7	1.2	-0.1	-0.1	-0.1	0.8
NAA1	0.5	2.5	1.3	1.9	2.6	1.3	0.5	0.9	2.2	-0.1	2.8	1.2	1.2	2.9	0.8	0.7	1.0	1.2
NAA10	1.9	2.7	1.3	1.9	0.4	0.9	1.9	2.0	0.4	-0.1	2.5	-0.3	1.0	2.4	-0.7	0.7	1.0	1.0
NAA11	2.7	13.1	2.9	5.6	6.4	1.4	1.6	0.4	3.5	0.8	5.5	1.7	1.3	10.5	1.2	0.2	0.4	1.3
NAA2	3.9	40.2	6.9	13.1	137.0	4.9	2.3	2.5	142.0	10.2	17.5	2.5	14	31.5	2.1	0.6	4.1	0.6
NAA3	1.6	10.2	1.3	1.6	12.2	1.3	1.3	0.7	11.1	-0.1	8.0	1.4	1.2	8.7	1.1	0.7	0.9	1.2
NAA4	1.8	1.9	1.0	1.3	2.1	1.2	1.4	0.6	2.5	-0.1	1.0	1.3	1.1	2.0	-0.1	0.7	0.9	1.0
NAA5	1.7	2.5	1.1	1.6	3.3	1.3	1.1	0.5	3.2	-0.1	0.8	1.5	1.2	2.6	0.7	0.7	0.9	1.2
NAA6	1.7	2.9	1.2	1.9	5.6	1.3	0.9	1.0	5.3	-0.8	2.6	1.4	1.2	2.8	-0.7	0.8	1.0	1.3
NAA7	0.3	1.9	1.1	1.3	3.9	1.0	0.4	1.4	3.6	0.8	3.3	1.1	1.0	2.2	0.7	0.7	0.8	1.0
NAA8	1.5	1.6	0.9	1.2	2.5	0.9	0.3	1.4	2.3	-0.1	0.7	1.0	0.9	1.8	-0.1	-0.1	0.8	0.9
NAA9	1.7	2.9	1.6	2.3	6.4	1.4	0.6	0.4	6.2	0.9	3.5	1.3	1.2	2.9	0.7	0.8	1.2	1.3
NB1	1.7	3.1	1.4	1.8	7.3	1.4	0.9	0.3	6.5	-1.0	5.2	1.6	1.3	2.8	-0.7	0.7	1.0	1.3
NB2	1.3	1.3	0.9	1.2	2.7	0.9	0.4	1.2	2.5	0.7	2.4	0.9	0.8	1.4	-0.1	-0.1	0.8	0.8
NB3	1.1	1.4	0.9	1.2	2.4	1.0	0.6	1.1	2.3	0.7	1.0	1.0	0.8	1.5	-0.1	-0.1	0.8	0.9
NB4	1.2	3.5	1.0	1.4	7.9	1.1	0.4	0.7	7.5	1.0	4.4	1.2	1.0	3.4	0.8	0.7	0.9	1.0
NBB1	2.0	2.0	1.4	2.3	3.5	0.7	1.9	1.9	3.2	-0.7	5.7	1.2	1.0	2.6	-0.1	0.7	1.1	1.1
NBB10	1.7	3.0	1.2	1.7	7.0	1.2	1.3	0.6	6.7	0.9	3.9	1.2	1.1	2.7	0.7	0.8	1.0	1.2
NBB11	3.3	5.1	2.1	4.1	7.8	1.1	2.4	1.7	7.4	1.1	9.8	1.9	1.7	5.6	0.8	0.9	0.3	1.6
NBB2	2.1	6.1	1.4	2.3	11.9	1.4	1.6	0.8	11.8	1.2	7.3	1.5	1.3	5.6	0.9	0.8	1.1	1.3
NBB2-R	2.0	5.3	1.6	2.0	9.7	1.3	1.6	0.6	9.4	1.0	8.0	1.3	1.2	4.9	-0.8	0.7	1.1	1.2
NBB3	1.9	9.4	1.8	3.0	15.9	1.6	1.2	0.3	15.3	1.3	4.1	1.5	1.4	7.1	1.0	0.8	1.4	1.4
NBB4	1.6	2.5	1.2	1.6	2.2	1.1	0.4	1.3	1.8	-0.1	2.0	1.1	1.0	1.9	-0.1	0.7	0.9	1.0
NBB5	1.6	2.7	1.0	1.3	3.1	1.1	0.5	1.3	3.0	-0.1	1.0	1.1	1.0	2.1	-0.1	0.6	0.8	0.9
NBB6	2.5	7.4	2.1	3.2	15.0	1.7	1.9	0.9	16.2	1.5	9.4	1.7	1.4	6.4	0.8	0.2	1.3	1.4
NBB7	1.3	1.3	0.8	1.0	1.9	-0.1	1.3	1.2	1.7	-0.1	1.7	0.9	0.7	1.2	-0.1	-0.1	0.7	0.8
NBB8	1.5	2.7	1.2	1.7	0.7	0.2	1.7	1.7	0.2	-0.1	0.7	-0.3	0.3	2.7	-0.7	0.7	0.8	1.0
NBB9	2.6	10.1	2.0	3.4	14.1	1.7	2.0	1.3	16.6	1.5	13.8	1.8	1.4	9.5	1.0	0.2	0.3	1.4
NC1	1.3	1.4	0.9	1.1	2.3	0.9	0.3	1.2	1.8	-0.1	0.7	0.9	0.8	1.5	-0.1	-0.1	0.8	0.8
NC2	1.4	1.6	1.0	1.2	3.0	1.0	0.5	1.4	2.7	0.7	1.6	1.1	0.9	2.4	-0.1	0.6	0.8	0.9
NC3	1.1	2.0	0.8	1.1	3.0	0.9	0.6	1.2	2.7	-0.1	2.5	1.0	0.9	2.6	-0.7	0.4	0.7	0.9
NC4	1.2	1.3	0.8	1.0	1.7	0.9	0.2	1.2	1.4	-0.1	2.0	0.9	0.8	1.6	0.7	-0.1	0.7	0.8
NC5	1.5	2.4	1.0	1.3	5.0	1.0	0.5	1.4	4.4	-0.8	4.9	1.1	1.0	3.8	-0.8	0.7	0.8	1.0
NC6	1.8	2.7	1.4	2.1	3.9	1.3	1.1	1.4	4.8	0.8	2.0	1.4	1.3	2.5	-0.1	0.7	1.1	1.2
NC7	1.4	1.6	0.9	1.1	2.5	1.0	1.4	1.5	2.2	-0.1	1.0	1.0	0.8	2.0	-0.1	0.4	0.8	0.8
NC8	1.3	1.6	0.9	1.2	2.4	1.0	0.5	1.4	2.1	-0.1	3.0	1.0	0.9	2.6	0.7	-0.1	0.8	0.9
NCC1	0.4	8.0	1.5	2.1	13.5	1.3	-0.1	0.7	12.7	1.1	8.5	1.4	1.2	6.9	-1.1	0.2	1.2	1.3
NCC1-R	0.4	7.4	1.5	2.0	13.0	1.4	-0.1	0.7	11.9	1.2	8.5	1.4	1.2	11.7	1.0	0.1	1.2	1.3
NCC2	2.6	4.4	2.7	4.4	2.6	2.0	1.5	0.9	12.3	1.2	8.1	1.8	1.6	5.4	-0.9	0.2	2.0	1.6
NCC3	1.4	3.4	1.6	2.6	5.6	2.4	0.7	0.7	7.4	0.9	6.7	2.1	1.6	3.6	0.7	0.8	1.3	1.6
NCC4	2.0	5.0	2.2	3.5	12.8	1.8	1.2	1.2	12.7	1.2	7.7	1.7	1.5	6.9	-0.9	0.2	1.5	1.5
NCC5	0.3	4.1	1.2	1.8	7.3	1.4	0.9	0.6	7.5	0.8	6.7	1.8	1.4	3.7	0.7	1.0		1.4
NCC6	1.7	2.3	0.9	1.2	2.8	0.9	1.3	0.3	2.5	-0.1	2.6	1.1	1.1	2.5	0.8	0.7	0.8	1.1
ND1	1.9	1.8	1.4	2.1	2.3	1.2	0.9	1.5	2.1	-0.1	2.1	1.2	1.1	2.1	-0.1	0.7	1.0	1.1
ND10	1.6	2.2	1.0	1.4	2.7	1.0	0.8	1.4	2.4	-0.1	2.5	1.1	1.0	3.0	-0.9	-0.1	0.8	1.0
ND2	1.9	3.1	1.3	1.9	3.8	1.3	0.8	1.2	3.4	-0.1	5.7	1.4	1.3	3.6	0.8	0.7	1.0	1.3
ND3	1.1	1.5	0.8	0.9	1.7	0.9	0.2	1.1	1.4	-0.1	1.9	0.9	0.7	2.0	0.7	0.4	0.7	0.8
ND4	1.5	7.5	1.2	1.8	9.8	1.4	0.8	0.3	9.0	-0.1	9.5	1.6	1.3	8.6	1.3	0.2	0.4	1.4
ND5	1.1	2.0	0.7	0.9	2.4	0.9	0.5	1.2	2.1	-0.1	2.5	1.0	0.8	2.7	-0.8	-0.1	0.1	0.8
ND6	1.3	2.4	1.0	1.3	3.8	1.0	0.7	1.3	3.5	0.7	5.1	1.0	1.0	2.7	0.8	-0.1	0.8	1.0
ND7	0.3	1.3	1.2	1.4	2.0	1.0	0.3	1.1	1.6	-0.1	0.6	0.9	0.8	1.5	-0.1	0.4	0.8	1.0
ND8	1.3	1.8	0.9	1.3	2.8	1.2	0.7	0.7	2.6	-0.1	1.7	1.3	1.0	2.6	0.7	0.7	0.8	1.0
ND8-R	1.6	2.4	1.0	1.3	3.8	1.2	1.3	0.7	3.5	-0.1	2.3	1.4	1.1	4.9	-0.7	0.2	0.8	1.1
ND9	1.5	2.5	1.0	1.6	4.7	1.4	0.7	0.2	4.4	0.8	3.3	1.5	1.2	4.4	0.8	0.7	0.9	1.2
NDD1	2.3	8.5	2.3	4.2	19.2	2.2	1.2	0.4	18.4	1.7	5.6	1.9	1.6	8.2	-0.9	0.8	1.8	1.6
NDD2	2.3	3.2	1.8	2.9	8.3	1.6	1.4	0.3	8.0	1.0	3.4	1.6	1.3	3.4	0.7	0.8	1.3	1.3
NDD3	2.1	2.7	1.6	2.7	6.7	0.8	1.8	0.9	6.3	-0.1	6.3	1.6	1.2	2.8	-0.1	0.2	1.2	1.2
NDD4	1.9	2.5	1.3	2.0	3.9	1.3	0.6	2.0	3.5	-0.1	3.9	1.4	1.2	2.4	-0.1	0.7	0.3	1.2
NDD5	1.4	2.3	0.9	1.2	2.4	1.0	0.7	1.3	2.1	-0.1	2.4	1.0	0.9	2.3	-0.1	0.4	0.8	0.9
NE1	2.5	8.0	2.7	4.8	23.4	2.2	1.4	0.2	22.9	1.8	4.7	1.8	1.6	7.9	1.0	1.0	2.0	1.6
NE10	0.2	1.0	0.7	0.9	1.2	-0.1	0.9	0.0	1.0	-0.1	1.2	0.7	0.7	1.0	-0.1	-0.1	-0.1	0.7
NE11	1.2	1.4	0.8	1.0	1.9	0.9	0.4	1.2	1.7	-0.1	2.6	0.9	0.8	2.2	0.7	-0.1	0.8	0.8
NE2	1.0	1.0	0.7	0.8	1.3	-0.1	0.9	0.9	1.1	-0.1	1.8	0.8	0.7	1.2	0.7	0.4	-0.1	0.7
NE3	0.2	1.7	1.1	1.4	2.8	1.0	0.4	1.3	2.3	-0.1	0.9	1.0	0.9	1.8	-0.1	0.6	0.8	0.9
NE4	0.1	1.4	1.2	1.6	2.5	1.0	1.4	1.4	2.3	-0.1	2.5	0.9	0.9	1.6	-0.1	0.7	0.9	0.9
NE4-R	1.3	1.2	1.0	1.2	1.7	0.9	0.4	1.2	1.5	-0.1	0.5	0.9	0.8	1.2	-0.1	-0.1	-0.1	0.8

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	D37-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	D43-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBA	052-LPB	053-LPB	054-HB
NE5	1.9	2.2	1.8	2.7	2.9	1.5	1.0	0.3	2.2	-0.1	2.5	1.4	1.3	2.4	-0.1	0.6	1.2	1.3
NE6	1.2	1.6	0.8	1.0	2.0	1.0	1.2	1.2	1.7	-0.1	2.1	1.0	0.8	1.9	-0.1	-0.1	0.7	0.8
NE7	1.3	1.2	1.2	1.5	2.3	1.1	0.5	1.2	2.1	-0.7	0.5	1.0	0.9	1.2	-0.1	0.7	0.8	0.9
NE8	0.2	1.4	0.7	-0.1	1.8	0.8	1.1	1.1	1.6	-0.1	1.8	0.9	0.8	-0.1	0.7	-0.1	-0.1	0.8
NE9	1.9	1.5	1.0	1.5	2.5	0.2	1.8	1.7	2.2	-0.1	2.6	1.1	1.0	1.9	-0.1	0.7	0.8	0.9
NEE1	1.5	2.1	1.0	1.3	2.4	1.0	0.7	1.6	2.2	-0.1	2.3	1.1	1.0	2.2	0.7	0.7	0.8	1.0
NEE2	1.7	5.4	3.5	2.3	10.3	1.3	0.6	0.9	9.9	1.1	1.2	1.2	1.2	4.4	0.8	0.3	1.2	1.2
NEE3	1.8	3.5	1.3	1.9	8.0	1.2	0.8	0.6	7.7	1.0	7.0	1.4	1.2	3.4	0.7	0.7	0.9	1.2
NEE4	1.2	4.0	0.9	1.2	4.4	1.0	0.4	1.3	4.0	0.8	1.7	1.1	0.9	2.7	0.7	0.7	0.8	1.0
NEE5	1.2	1.7	0.8	1.2	2.6	1.0	0.5	1.2	2.4	-0.1	0.9	1.0	0.9	1.7	-0.1	0.6	0.8	0.9
NEE6	2.4	4.7	1.7	2.8	6.5	0.4	2.2	1.3	7.3	-1.0	4.5	1.5	1.4	4.4	0.8	0.2	1.2	1.4
NEE7	2.1	2.9	1.9	2.9	2.8	1.6	0.9	1.6	2.0	-0.1	2.4	1.4	1.3	2.4	0.7	0.8	1.2	1.3
NF1	2.3	2.8	1.3	2.2	0.9	0.8	2.0	1.4	3.1	-0.1	3.5	1.6	1.2	3.5	0.8	0.7	1.1	1.3
NF10	1.4	1.5	0.9	1.2	2.9	0.9	0.3	1.3	2.5	-0.1	1.1	1.1	1.0	1.6	-0.1	0.7	0.8	1.0
NF11	0.3	3.7	1.6	1.8	9.9	1.2	0.5	0.8	9.4	1.1	4.3	1.3	1.1	3.8	0.7	0.7	1.0	1.2
NF2	1.1	1.5	0.7	0.8	2.0	0.8	1.1	1.2	1.6	-0.1	2.4	0.8	0.8	2.3	0.7	-0.1	-0.1	0.8
NF3	1.3	1.6	0.9	1.1	1.9	0.9	1.4	1.4	1.6	-0.1	1.9	0.9	0.8	2.0	-0.7	-0.1	0.7	0.8
NF4	1.4	2.0	0.8	1.0	3.0	0.9	1.5	1.5	2.7	-0.1	1.8	1.0	0.9	2.9	0.7	-0.1	0.7	0.9
NF5	0.2	1.6	0.6	1.0	3.2	0.9	0.2	1.4	2.9	-0.7	4.5	1.1	1.0	2.4	-0.7	-0.1	0.7	1.2
NF6	1.3	2.5	0.8	1.1	5.1	1.0	0.6	0.6	4.8	0.8	2.4	1.1	1.0	3.5	0.9	0.6	0.8	1.0
NF7	2.4	7.3	2.8	5.1	15.2	2.7	1.4	0.3	18.4	1.4	7.8	2.5	2.2	8.8	1.0	1.2	2.1	2.0
NF8	1.6	1.5	1.0	1.3	2.4	1.0	0.4	1.2	2.2	-0.1	2.4	1.0	1.0	2.0	0.7	-0.1	0.8	1.0
NF8-R	1.6	3.8	1.2	1.8	9.2	1.2	0.9	0.9	8.7	1.0	5.3	1.4	1.4	4.3	1.0	0.7	1.0	1.4
NF9	1.5	2.3	1.0	1.2	4.1	1.0	0.5	1.3	3.7	0.7	2.3	1.0	0.9	3.4	0.7	0.7	0.8	0.9
NFF1	0.4	1.7	1.1	1.3	2.7	1.0	0.4	1.4	2.4	-0.1	0.8	1.1	0.9	1.8	-0.1	0.7	0.8	1.0
NFF2	2.2	4.8	1.6	2.5	9.7	1.5	0.2	1.6	9.3	1.1	2.9	1.5	1.4	4.8	0.8	0.8	1.2	1.3
NFF3	1.2	2.8	0.8	1.0	2.2	0.9	1.4	1.6	2.7	-0.1	4.3	1.1	0.9	2.3	-0.1	-0.1	0.7	1.0
NFF4	2.6	3.9	2.1	3.5	11.1	1.8	1.7	0.8	10.9	1.2	2.1	1.6	1.5	3.8	0.8	0.9	1.6	1.5
NFF5	1.8	2.9	1.5	2.2	3.9	1.4	0.9	1.0	3.4	0.7	3.0	1.5	1.3	3.2	0.7	0.8	1.1	1.3
NFF6	1.4	1.5	0.9	1.1	0.9	1.0	1.5	1.5	1.8	-0.1	2.0	1.0	0.9	1.4	-0.1	-0.1	0.7	0.9
NFF6-R	1.4	1.6	0.8	1.2	2.1	1.0	0.5	1.6	1.9	-0.1	2.1	1.1	0.9	1.4	-0.1	-0.1	0.8	0.9
NFF7	1.8	2.1	1.1	1.6	4.9	1.1	0.7	1.5	4.7	0.8	1.9	1.2	0.1	2.3	-0.1	0.7	0.9	1.1
NFF8	1.5	2.2	1.2	1.5	2.7	1.4	0.3	1.3	2.0	-0.1	2.4	1.1	1.1	2.3	0.7	0.7	0.9	1.1
NG1	1.0	1.1	0.6	0.8	1.3	-0.1	0.2	0.9	1.1	-0.1	1.5	0.8	0.7	-0.1	-0.1	-0.1	-0.1	0.7
NG10	1.4	2.2	0.9	1.2	2.5	0.9	0.6	1.2	2.2	-0.1	2.8	1.0	0.9	2.5	0.8	0.7	0.8	0.9
NG11	1.2	2.3	0.8	1.1	4.0	0.9	0.4	1.2	3.6	0.7	6.3	1.0	0.9	3.2	0.9	-0.1	0.8	0.9
NG2	1.0	1.7	0.8	1.1	2.1	0.9	0.4	1.0	1.8	-0.1	2.4	0.9	0.8	2.3	0.8	-0.1	0.7	0.8
NG2-R	1.1	1.7	0.8	1.0	2.0	0.9	0.4	1.0	1.6	-0.1	2.1	0.8	0.8	0.1	0.7	-0.1	-0.1	0.8
NG3	0.9	1.0	0.7	0.9	1.3	-0.1	0.3	0.9	1.1	-0.1	4.6	0.9	0.8	1.2	-0.1	-0.1	-0.1	0.8
NG4	1.4	1.4	1.1	1.4	1.9	1.0	0.7	1.3	1.7	-0.1	2.4	1.1	1.1	1.9	-0.1	0.7	0.9	1.1
NG5	2.5	5.1	1.9	3.1	11.6	1.7	1.3	0.4	10.8	1.1	3.4	1.8	1.4	5.8	0.9	0.8	1.2	1.4
NG6	1.5	1.5	1.0	1.4	1.9	1.0	0.8	1.4	1.7	-0.1	1.9	1.1	0.9	2.2	0.7	0.7	0.8	0.9
NG7	1.0	0.8	0.7	0.9	1.0	-0.1	0.2	1.0	0.9	-0.1	1.1	0.9	0.8	1.1	-0.1	-0.1	-0.1	0.8
NG8	1.2	1.4	0.8	1.0	1.7	1.0	0.7	1.2	1.4	-0.1	2.1	1.0	0.8	1.7	-0.1	-0.1	-0.1	0.8
NG9	1.8	2.4	1.3	1.7	4.8	1.4	0.8	1.3	4.2	0.8	1.8	1.1	1.0	2.8	-0.1	0.7	0.9	1.0
NGG1	1.5	2.1	0.9	1.0	2.9	0.9	0.3	1.4	2.3	0.7	0.7	0.9	0.9	1.9	-0.1	0.6	0.8	0.9
NGG10	2.0	3.8	1.1	1.6	6.2	1.1	1.6	0.9	7.0	1.0	7.9	1.1	1.0	3.9	0.8	0.7	0.9	1.0
NGG2	0.2	1.5	0.8	1.0	2.5	1.0	0.6	0.3	2.0	-0.1	1.9	1.1	0.9	1.8	-0.1	-0.1	0.7	0.9
NGG3	2.1	15.9	1.9	3.1	13.6	1.9	1.3	0.7	12.7	1.1	4.7	0.3	1.4	8.5	0.9	0.2	0.3	1.4
NGG4	1.7	3.4	1.4	1.9	5.9	1.2	1.1	0.6	5.5	0.8	2.8	1.3	1.1	3.2	0.7	0.7	1.0	1.2
NGG5	1.7	2.4	1.2	1.7	4.3	1.2	0.6	0.3	4.1	0.8	1.7	1.2	1.1	2.2	-0.1	0.7	1.0	1.1
NGG6	1.6	2.1	1.0	1.3	3.2	1.1	0.5	0.4	3.0	-0.1	0.8	1.1	1.0	2.1	-0.1	0.7	0.8	1.0
NGG7	1.7	2.8	1.1	1.8	4.5	1.1	0.5	1.3	4.2	0.8	1.5	1.0	1.1	2.6	-0.1	0.7	0.9	1.1
NGG8	1.5	3.4	0.9	1.1	5.6	1.0	0.3	1.5	5.2	0.8	1.8	1.1	0.9	3.0	0.7	-0.1	0.7	0.9
NGG9	1.2	2.3	0.7	0.8	2.0	0.8	1.3	1.3	1.8	-0.1	1.9	0.9	0.8	1.7	-0.1	-0.1	0.7	0.8
NH1	1.5	2.8	0.9	1.2	3.0	1.2	1.2	0.7	2.6	-0.1	0.9	1.2	1.0	2.8	0.7	-0.1	0.8	1.0
NH10	1.3	1.6	1.6	2.2	1.9	1.3	0.7	1.2	1.6	-0.1	1.7	1.2	1.0	1.6	-0.1	0.7	1.1	1.0
NH10-R	1.6	1.7	2.3	3.4	2.1	1.8	0.9	1.4	1.9	-0.1	1.9	1.4	1.3	1.8	-0.1	0.8	1.4	1.3
NH11	1.8	3.1	1.2	1.8	6.2	1.2	0.8	1.4	5.9	0.9	2.0	1.2	1.1	3.2	0.7	0.7	1.0	1.1
NH2	1.5	3.1	1.4	2.0	3.1	1.4	0.8	0.3	2.7	-0.1	3.0	1.4	1.2	3.0	0.7	0.7	1.0	1.2
NH3	1.2	1.1	0.7	0.8	1.2	-0.1	1.0	0.7	1.1	-0.1	1.2	0.8	0.7	1.2	-0.1	-0.1	-0.1	0.7
NH4	1.5	2.4	1.0	1.6	5.5	1.3	0.7	0.2	5.3	0.8	1.6	1.4	1.2	2.7	0.7	0.7	0.9	1.1
NH5	1.9	1.7	1.4	1.9	2.1	1.2	0.8	1.4	1.6	-0.1	1.8	1.1	1.1	1.8	-0.1	0.7	1.0	1.1
NH6	1.8	2.3	1.6	2.3	2.4	1.4	0.8	1.4	2.1	-0.1	2.5	1.3	1.2	2.5	0.7	0.8	1.2	1.2
NH7	2.1	5.0	2.3	3.8	10.6	2.2	1.4	0.3	10.0	1.1	5.5	2.0	1.6	5.5	0.8	0.9	1.8	1.6
NH8	1.9	2.9	1.3	1.9	2.7	1.2	0.9	1.1	2.4	-0.1	2.7	1.2	1.2	2.6	0.7	0.7	1.1	1.2
NH9	1.5	3.1	1.4	2.0	2.9	1.3	0.7	1.0	2.5	-0.1	3.3	1.3	1.1	3.4	0.7	0.7	1.1	1.2

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBH	052-LPB	053-LPB	054-HB
NHH1	1.3	1.6	0.7	0.9	2.1	0.8	1.2	1.3	1.9	-0.1	0.5	0.9	0.8	1.6	-0.1	-0.1	-0.1	0.8
NHH10	1.4	2.3	0.8	0.9	1.9	0.9	0.3	1.3	1.3	-0.1	1.6	0.9	0.8	1.7	-0.1	-0.1	-0.1	0.8
NHH11	1.4	1.6	0.8	1.0	2.3	0.9	0.3	1.4	2.1	-0.1	0.6	0.9	0.8	1.7	-0.1	-0.1	-0.1	0.8
NHH11-R	1.3	1.5	0.8	1.0	2.2	0.9	0.3	1.4	1.9	-0.1	0.5	0.9	0.8	1.6	-0.1	-0.1	0.7	0.8
NHH12	1.3	1.7	0.8	1.1	3.3	1.0	0.5	0.3	3.2	-0.1	1.1	1.1	0.9	2.4	-0.1	0.6	0.8	0.9
NHH2	1.5	5.4	4.1	1.6	9.7	1.3	0.8	0.5	9.4	-1.0	5.8	1.4	1.2	5.3	0.9	0.7	0.2	1.2
NHH3	1.3	1.9	0.9	0.9	2.5	0.9	0.3	1.5	2.4	-0.1	0.6	1.0	0.9	2.0	-0.1	-0.1	0.8	0.9
NHH4	1.3	2.4	1.2	1.8	5.5	1.4	0.7	0.5	5.0	-0.8	2.6	1.5	1.3	2.9	-0.7	0.8	1.1	1.3
NHH5	1.5	3.8	1.0	1.4	5.6	1.1	1.0	0.6	5.1	0.8	2.5	1.2	1.0	3.7	0.8	0.7	0.8	1.0
NHH6	1.5	2.8	1.2	1.4	3.5	1.1	0.6	0.6	3.1	-0.1	1.2	1.2	1.0	2.2	-0.1	0.7	0.9	1.0
NHH7	1.8	3.4	1.1	1.5	5.9	1.1	0.6	1.3	5.5	0.8	2.1	1.2	1.1	3.4	0.8	0.7	0.9	1.0
NHH8	1.5	2.2	0.8	1.0	2.3	0.9	1.4	1.4	2.1	-0.1	2.3	0.8	0.8	1.7	-0.1	-0.1	0.7	0.8
NHH9	1.3	2.5	0.8	1.0	3.2	1.0	1.4	0.6	2.9	-0.1	3.3	1.1	0.9	2.3	-0.1	-0.1	0.7	0.9
NI1	2.8	14.7	3.2	6.0	35.4	2.6	1.6	0.3	34.5	2.4	10.0	1.9	1.5	15.2	1.3	1.3	2.3	1.5
NI10	1.5	3.8	1.1	1.6	5.5	1.1	0.7	1.3	5.1	0.8	3.0	1.1	1.0	4.4	0.7	0.7	0.9	1.0
NI2	1.2	1.5	1.0	1.4	1.8	1.0	0.4	0.9	1.6	-0.1	1.2	1.0	0.9	1.6	-0.1	0.7	0.8	0.9
NI3	1.3	2.0	0.9	1.3	3.5	1.0	0.6	0.9	3.3	0.7	1.2	1.0	0.9	2.1	-0.1	-0.1	0.8	0.9
NI4	1.9	7.6	1.7	2.7	7.0	1.7	1.0	1.0	6.3	0.7	7.8	1.7	1.4	7.5	0.9	0.8	1.3	1.4
NI5	1.7	3.4	1.4	2.2	3.6	1.8	0.9	0.3	3.3	-0.1	3.3	1.7	1.4	3.2	0.8	0.8	1.2	1.4
NI5-R	1.7	2.9	1.4	2.1	3.0	1.5	0.9	0.3	2.7	-0.1	2.8	1.5	1.3	2.7	-0.7	0.7	1.1	1.2
NI6	1.4	1.9	1.0	1.2	3.0	0.9	0.5	1.3	3.7	0.8	1.3	1.0	0.9	2.4	-0.1	0.6	0.8	0.9
NI7	2.4	25.6	2.7	4.7	46.5	2.1	1.4	0.3	45.0	2.9	15.2	1.8	1.5	23.2	1.4	1.0	1.7	1.5
NI8	1.2	1.6	0.9	1.1	1.7	0.9	0.4	0.9	1.5	-0.1	2.0	0.9	0.8	2.0	-0.1	-0.1	-0.1	0.8
NI9	2.0	3.4	1.6	2.3	6.2	1.3	1.0	1.5	5.5	0.9	2.0	1.3	1.3	3.2	0.7	0.7	1.1	1.2
NII1	1.1	0.9	0.8	1.0	1.2	0.8	0.4	-0.1	1.4	-0.1	1.1	0.8	0.7	1.0	-0.1	-0.1	-0.1	0.7
NII10	1.8	3.1	1.1	1.7	6.8	1.1	0.8	1.3	6.5	0.9	5.6	1.3	1.2	5.0	0.8	0.7	0.9	1.1
NII11	1.3	2.1	1.2	1.6	4.5	1.3	0.7	0.3	4.1	0.8	1.6	1.4	1.2	2.6	0.7	0.7	1.0	1.2
NII11-R	1.3	2.2	1.2	1.6	5.5	1.2	0.8	1.5	4.9	0.9	1.8	1.3	1.1	2.9	0.7	0.7	0.9	1.1
NII12	1.9	1.7	1.2	1.7	3.2	1.1	0.8	1.4	3.0	0.7	0.9	1.1	1.0	2.0	-0.1	0.7	0.9	1.0
NII13	2.2	8.7	2.4	4.3	41.1	2.3	1.2	0.3	40.5	3.5	7.7	2.0	1.6	10.7	0.9	1.0	1.8	1.6
NII14	1.4	1.3	0.9	1.1	2.1	0.9	0.3	1.1	1.4	-0.1	0.5	0.9	0.8	1.4	-0.1	-0.1	-0.1	0.8
NII2	1.7	2.9	1.3	1.8	5.1	1.2	0.8	1.5	4.7	0.8	2.2	1.3	1.1	3.4	0.7	0.7	1.0	1.1
NII3	1.7	14.4	1.4	2.1	17.8	1.3	0.9	1.0	16.5	1.1	13.2	1.5	1.3	16.7	1.4	0.8	1.1	1.4
NII4	1.9	21.5	1.9	2.1	44.4	1.7	1.1	0.3	42.3	2.6	39.0	1.7	1.4	18.8	2.1	1.1	1.4	1.4
NII5	1.4	2.3	0.9	1.2	4.6	0.9	0.5	1.2	4.2	0.8	3.0	0.9	0.9	4.0	0.8	-0.1	0.8	0.9
NII6	1.4	1.8	0.8	1.0	2.1	0.9	0.4	1.2	1.9	-0.1	2.4	0.9	0.8	2.2	-0.1	-0.1	-0.1	0.8
NII7	1.4	1.2	0.6	0.8	1.5	0.8	1.1	1.0	1.3	-0.1	1.8	0.8	0.7	1.8	-0.1	-0.1	-0.1	0.7
NII8	2.0	10.9	1.4	2.3	19.9	1.5	0.9	0.3	19.0	1.4	14.7	1.6	1.4	9.9	1.3	0.8	1.1	1.4
NII9	1.4	2.1	0.8	1.1	2.8	0.9	0.6	1.3	2.5	-0.1	3.7	1.1	1.0	3.5	0.7	-0.1	-0.1	1.0
NJ1	1.8	2.6	1.5	2.0	6.8	1.4	0.8	0.3	6.2	0.9	5.5	1.4	1.2	2.5	0.7	0.7	1.0	1.2
NJ2	1.6	3.7	1.3	1.9	7.2	1.3	0.9	0.3	7.1	0.9	3.6	1.4	1.2	3.6	0.7	0.7	1.0	1.2
NJ3	1.8	5.0	1.4	1.9	8.2	1.5	0.9	0.7	7.7	0.9	2.5	1.6	1.2	4.2	0.7	0.7	1.1	1.2
NJ4	1.5	3.3	1.2	1.7	6.2	1.3	1.1	0.4	5.9	0.8	2.8	1.3	1.1	3.1	0.7	0.7	1.0	1.1
NJ5	1.4	1.7	0.9	1.2	2.9	1.0	0.4	1.3	2.7	-0.1	1.1	1.0	0.9	1.8	-0.1	-0.1	0.8	0.9
NJ6	1.3	2.2	1.0	1.3	3.2	1.0	0.4	1.1	2.9	-0.1	1.2	0.9	0.9	2.0	-0.1	-0.1	0.8	0.9
NJ7	2.2	4.5	1.6	2.8	6.3	0.8	0.9	1.7	5.6	0.9	6.7	1.2	1.2	3.9	0.8	0.2	1.2	1.2
NJJ1	1.7	1.9	1.1	1.5	2.5	1.1	0.7	1.3	2.1	-0.1	3.9	1.2	1.1	3.4	0.8	0.7	0.9	1.1
NJJ10	6.7	17.9	6.4	13.7	45.6	3.7	3.0	4.3	44.7	3.5	12.4	0.8	2.1	16.9	1.3	1.9	4.4	-0.4
NJJ11	1.2	1.5	0.8	1.0	2.1	0.9	0.2	1.3	1.9	-0.1	2.3	1.1	0.9	2.1	-0.1	-0.1	0.7	0.9
NJJ12	1.8	2.2	1.4	2.0	2.5	1.3	0.8	1.0	2.3	-0.1	2.7	0.2	1.1	2.5	-0.7	0.7	1.0	1.1
NJJ13	2.2	9.8	1.4	2.4	21.6	1.5	1.3	0.3	20.8	1.8	10.0	1.8	1.4	13.4	1.0	0.9	1.2	1.4
NJJ14	1.3	1.3	0.8	1.0	1.6	0.9	0.5	1.3	1.4	-0.1	1.7	1.0	0.8	1.7	-0.1	-0.1	-0.1	0.8
NJJ2	1.5	3.2	1.1	1.8	3.6	1.2	0.8	0.9	3.2	-0.1	3.8	1.4	1.2	3.5	0.9	0.7	0.9	1.3
NJJ3	1.2	1.3	0.9	1.2	1.7	0.9	0.6	1.1	1.4	-0.1	4.8	0.9	0.8	1.8	-0.1	-0.1	0.8	0.8
NJJ4	1.8	14.2	1.2	1.7	53.7	1.1	0.8	0.9	51.3	4.1	14.2	1.4	1.2	18.5	1.2	0.7	0.9	1.3
NJJ5	1.5	2.1	1.3	1.7	2.4	1.4	0.7	1.4	2.0	-0.1	2.8	1.1	1.0	2.5	-0.7	0.7	0.9	1.1
NJJ6	1.5	2.5	1.0	1.5	5.6	1.2	0.7	0.9	5.3	0.8	3.4	1.2	1.1	3.0	0.8	0.7	0.9	1.1
NJJ7	1.2	1.1	0.9	1.2	1.7	0.9	0.4	1.2	1.5	-0.1	1.4	1.0	0.8	1.3	-0.1	-0.1	-0.1	0.8
NJJ8	0.2	2.2	0.9	1.1	2.2	0.9	0.4	1.2	1.9	-0.1	2.3	1.0	0.8	2.5	-0.1	0.7	-0.1	0.9
NJJ8-R	1.7	3.1	4.0	1.4	2.9	1.0	0.7	1.3	2.6	-0.1	3.7	1.1	1.0	3.8	0.8	0.7	0.8	1.0
NJJ9	2.5	23.2	2.0	3.4	87.6	1.7	1.3	1.3	86.1	6.5	17.5	1.8	1.6	24.7	1.4	1.0	1.5	1.6
NK1	1.8	2.0	1.8	2.1	3.6	1.2	0.1	1.7	3.4	0.9	8.5	1.2	1.0	2.4	0.7	0.7	0.2	1.0
NK2	1.3	2.7	0.9	1.3	2.3	1.0	0.8	1.3	2.0	-0.1	2.5	1.0	0.9	2.6	-0.1	0.7	0.8	0.9
NK3	1.7	2.7	4.3	1.7	3.6	1.4	0.5	1.8	3.3	-0.1	1.8	1.2	1.0	2.4	-0.7	0.7	1.0	1.0
NK4	1.2	4.1	1.0	1.4	3.7	1.1	0.8	1.3	3.2	-0.1	3.5	1.3	1.2	3.1	0.9	0.7	0.9	1.2
NK4-R	1.3	3.3	1.0	1.2	3.1	1.0	0.6	1.4	2.3	-0.1	2.9	1.2	1.1	2.6	-0.8	0.7	0.8	1.1
NK5	0.4	2.4	1.3	1.9	2.7	1.3	1.3	0.7	2.8	-0.1	1.3	1.4	1.2	2.5	-0.1	0.7	1.0	1.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBH	052-LPB	053-LPB	054-HB
NK6	1.6	2.5	1.0	1.6	3.8	1.4	0.7	1.0	3.5	0.7	1.6	1.5	1.1	2.6	0.7	0.7	0.9	1.1
NK7	1.4	2.6	1.1	1.5	4.4	1.2	0.7	0.8	4.1	0.8	1.9	1.1	1.0	2.9	0.7	0.7	0.9	1.0
NKK1	1.0	1.1	0.7	0.9	1.4	0.8	0.3	1.1	1.1	-0.1	1.3	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.8
NKK10	0.2	1.4	0.8	0.9	2.5	0.9	1.4	1.4	2.1	0.7	2.1	0.9	0.8	1.7	-0.1	-0.1	-0.1	0.8
NKK11	1.8	2.7	1.2	1.8	6.6	1.2	1.1	0.3	6.4	0.9	2.1	1.3	1.2	3.1	0.7	0.7	1.0	1.2
NKK2	1.3	1.7	0.8	1.1	2.5	1.0	0.6	1.3	2.1	-0.1	2.6	1.0	0.9	2.2	0.7	0.7	0.8	0.9
NKK3	1.5	3.3	0.9	1.4	3.8	1.0	0.4	1.4	3.4	-0.1	5.2	1.1	1.1	5.2	-0.8	0.7	0.8	1.1
NKK4	0.3	19.3	1.5	2.1	60.0	1.3	-0.1	0.5	57.6	4.0	22.8	1.6	1.4	16.5	2.1	0.9	1.1	1.5
NKK5	1.1	21.1	1.5	2.2	47.1	1.8	0.4	0.8	45.0	2.9	40.6	1.6	1.4	16.9	2.1	0.6	1.1	1.4
NKK6	1.6	3.4	1.0	1.4	3.5	1.0	0.4	1.4	3.1	-0.1	4.6	1.3	1.3	4.4	0.8	0.7	0.9	1.3
NKK7	1.7	2.2	0.6	1.1	2.4	0.9	0.4	1.5	2.1	-0.1	2.5	1.0	0.9	2.6	-0.7	0.7	0.8	0.9
NKK8	1.6	1.8	0.9	1.2	3.4	1.2	1.0	0.5	2.9	0.7	1.6	1.2	1.0	2.4	0.7	0.7	0.8	1.0
NKK9	1.6	1.9	1.0	1.4	3.8	1.0	0.5	1.5	3.4	0.7	1.5	1.1	0.9	2.3	-0.1	0.7	0.8	0.9
NL1	1.2	1.1	0.8	0.9	1.4	-0.1	0.3	1.1	1.0	-0.1	1.3	0.8	0.7	1.2	-0.1	-0.1	-0.1	0.7
NL2	1.3	1.6	0.7	0.9	1.7	0.8	0.3	1.3	1.4	-0.1	1.6	0.9	0.8	1.6	-0.1	-0.1	-0.1	0.8
NL3	1.1	1.0	0.7	1.0	1.3	0.9	0.4	1.2	1.1	-0.1	1.2	0.8	0.8	1.2	-0.1	-0.1	-0.1	0.8
NL3-R	1.4	2.0	0.9	1.2	2.2	1.0	0.6	1.5	1.9	-0.1	2.4	1.0	0.9	2.1	-0.7	0.7	0.8	0.9
NL4	1.1	1.7	0.7	0.9	1.8	0.8	0.4	1.2	1.5	-0.1	2.0	0.9	0.8	2.0	-0.1	-0.1	-0.1	0.8
NLL1	1.3	1.3	1.4	1.5	1.7	1.1	0.7	1.4	1.4	-0.1	1.9	1.1	0.9	1.8	-0.1	0.7	0.7	0.9
NLL10	0.2	2.0	0.8	1.0	3.8	1.2	0.9	0.5	3.2	0.8	1.9	0.3	1.1	2.7	0.7	0.6	0.8	1.1
NLL11	1.5	2.1	1.1	1.5	5.2	1.8	0.8	0.3	4.6	0.8	1.8	1.5	1.2	2.4	0.7	0.7	0.9	1.2
NLL12	1.6	1.8	1.1	1.3	3.7	1.0	0.8	1.4	3.1	0.7	0.7	1.1	1.0	2.0	-0.1	0.7	0.8	1.0
NLL2	1.2	1.8	0.6	1.1	3.3	1.0	0.5	1.0	3.0	0.7	4.5	1.1	0.9	2.2	0.8	-0.1	0.7	0.9
NLL3	1.3	1.7	0.8	1.0	2.1	0.8	0.2	1.2	1.8	-0.1	2.1	0.8	0.8	2.2	0.8	-0.1	-0.1	0.8
NLL4	2.0	3.4	1.1	2.3	4.7	1.5	1.2	0.7	4.1	-0.1	3.4	1.7	1.4	4.9	1.0	0.8	1.1	1.4
NLL5	2.1	14.5	0.3	0.3	41.4	1.3	1.3	0.8	39.6	3.0	20.4	1.8	1.6	20.4	1.6	1.0	1.1	1.4
NLL6	1.9	12.0	1.2	1.8	32.1	1.4	1.3	0.7	30.9	2.3	14.3	1.7	1.3	11.2	1.4	0.8	1.0	1.4
NLL7	2.5	25.2	3.0	5.3	123.0	2.5	1.1	1.3	119.0	9.6	19.0	2.2	1.8	28.8	1.8	1.2	1.9	1.7
NLL8	0.2	1.7	0.8	1.1	3.2	0.9	0.5	1.2	3.0	0.7	1.6	0.9	0.9	2.2	-0.1	0.7	0.8	0.9
NLL9	1.9	7.4	1.6	2.6	8.7	1.5	0.9	0.7	10.5	1.0	3.9	1.6	1.4	6.6	0.9	0.2	1.2	1.4
NLL9-R	1.8	8.4	1.7	2.9	24.9	1.7	0.9	0.9	24.4	2.0	6.2	1.9	1.5	8.2	1.0	0.9	1.1	1.5
NM1	1.8	3.8	1.4	2.2	4.5	1.3	0.9	0.3	4.0	0.7	2.5	1.5	1.3	3.7	0.7	0.8	1.1	1.3
NM2	0.3	1.4	1.0	1.3	3.0	1.0	0.4	1.3	2.7	0.7	0.8	1.0	0.9	1.8	-0.1	-0.1	0.8	0.9
NMM1	1.5	1.8	1.1	1.4	2.7	1.0	0.4	1.2	2.3	0.7	2.4	1.1	0.9	2.1	-0.1	0.7	0.8	1.0
NMM10	2.6	12.4	2.0	3.6	9.0	1.6	1.2	1.7	8.3	0.8	10.5	1.8	1.5	12.7	1.2	0.9	1.5	1.5
NMM2	1.6	5.3	1.2	1.7	11.6	1.1	0.8	1.3	10.8	1.2	6.3	1.3	1.2	8.5	0.9	0.7	1.0	1.2
NMM3	1.5	2.4	1.0	1.2	3.2	1.4	0.8	1.3	2.8	0.7	3.8	1.1	1.1	2.3	0.8	0.7	0.9	1.0
NMM3-R	1.4	2.0	1.0	1.2	2.8	1.0	0.7	1.2	2.3	0.7	3.6	1.1	0.9	2.0	0.7	-0.1	0.8	1.0
NMM4	1.2	1.3	0.8	1.1	1.6	0.9	0.5	1.1	1.4	-0.1	1.7	1.0	0.9	1.7	-0.1	-0.1	0.8	0.9
NMM5	1.2	1.3	0.8	1.0	1.6	0.9	0.4	1.0	1.2	-0.1	1.8	0.9	0.8	1.9	-0.1	-0.1	-0.1	0.8
NMM6	1.9	5.5	1.7	2.8	11.4	1.5	1.0	1.1	10.9	1.1	4.7	1.6	1.4	6.9	0.9	0.8	1.3	1.4
NMM7	1.1	0.9	0.7	0.9	1.1	-0.1	0.4	-0.1	1.0	-0.1	1.1	0.8	0.8	1.1	-0.1	-0.1	-0.1	0.8
NMM8	1.7	14.0	1.3	1.6	6.5	1.2	0.8	0.3	8.5	0.7	13.3	1.6	1.3	13.1	1.2	0.8	0.9	1.4
NMM9	1.8	4.4	1.5	2.5	10.0	1.4	0.9	0.9	9.6	1.1	2.5	1.5	1.3	4.4	0.8	0.8	1.1	1.3
NN1	1.1	0.9	0.7	0.8	1.2	-0.1	1.0	1.0	0.8	-0.1	1.0	0.8	0.7	1.0	-0.1	-0.1	-0.1	0.7
NNN1	0.2	1.6	0.8	1.0	1.9	0.9	0.3	1.1	1.4	-0.1	2.0	0.8	0.8	2.1	-0.1	-0.1	-0.1	0.8
NNN10	0.4	1.7	1.4	1.9	3.2	1.1	0.7	1.1	2.9	0.7	4.2	1.1	1.0	1.9	-0.1	0.7	1.1	1.0
NNN2	1.2	1.8	0.8	0.9	3.2	0.9	0.2	1.2	2.7	0.7	3.3	0.9	0.8	2.7	-0.1	-0.1	-0.1	0.8
NNN3	0.9	1.0	0.6	0.8	1.1	-0.1	0.8	0.8	0.9	-0.1	1.0	0.7	-0.1	1.0	-0.1	-0.1	-0.1	0.1
NNN4	1.8	4.4	1.5	2.5	12.5	1.7	1.0	0.3	12.2	1.3	3.8	0.3	1.4	5.6	0.8	0.8	1.2	1.4
NNN5	1.0	1.7	0.8	0.9	1.8	0.8	0.4	0.3	1.3	-0.1	2.1	0.9	0.8	2.0	-0.1	-0.1	-0.1	0.8
NNN6	0.2	1.2	0.7	0.8	1.4	-0.1	0.9	0.9	1.0	-0.1	1.5	0.8	0.7	-0.1	-0.1	-0.1	-0.1	0.7
NNN7	0.2	1.0	0.6	0.8	1.2	-0.1	0.9	1.0	1.0	-0.1	1.1	0.8	0.7	1.2	-0.1	-0.1	-0.1	0.7
NNN8	2.4	5.7	1.7	3.3	22.2	2.4	1.6	0.3	22.0	1.9	3.2	2.2	1.8	6.3	0.8	0.9	1.5	1.7
NNN9	2.2	6.6	1.8	2.8	13.1	1.3	1.0	1.7	12.0	1.1	3.8	1.4	1.3	5.7	-0.8	0.9	1.4	1.3
NO1	1.3	1.8	0.9	1.2	2.2	1.0	0.6	1.4	1.8	-0.1	2.4	1.0	0.9	2.5	0.7	0.7	0.8	0.9
NO2	1.0	0.8	0.7	0.8	1.0	-0.1	0.2	0.8	0.8	-0.1	0.8	-0.1	-0.1	0.7	-0.1	-0.1	-0.1	0.1
NOO1	1.4	2.3	0.7	0.9	2.7	-0.1	0.3	1.1	2.5	0.7	3.2	0.8	0.8	2.7	-0.1	-0.1	-0.1	0.8
NOO2	1.3	3.7	0.9	1.3	4.0	1.0	0.7	1.3	3.7	0.7	4.4	1.1	1.0	3.7	0.8	0.7	0.8	1.0
NOO3	1.2	1.7	0.9	1.1	3.0	0.9	0.3	1.1	2.8	0.7	1.2	1.0	0.9	1.8	-0.1	-0.1	0.8	0.9
NOO4	1.5	3.2	1.0	1.3	5.4	1.2	0.7	1.3	5.1	0.9	5.6	1.2	1.0	2.7	-0.7	0.7	0.8	1.0
NOO5	0.2	1.8	0.7	0.9	2.0	1.0	0.3	1.2	1.6	-0.1	2.0	1.0	0.9	1.9	-0.1	-0.1	-0.1	0.8
NOO6	1.2	1.3	0.7	0.9	1.4	-0.1	0.3	1.0	1.2	-0.1	1.9	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.7
NOO6-R	1.0	1.3	0.7	0.9	1.4	-0.1	0.3	1.0	1.1	-0.1	1.4	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.7
NOO7	0.2	1.4	0.6	0.1	1.5	0.8	1.1	1.1	1.3	-0.1	1.7	0.8	0.7	1.7	-0.1	-0.1	-0.1	0.7
NOO8	2.4	9.5	1.4	2.4	39.3	1.4	1.0	0.3	38.7	3.1	6.4	1.7	1.4	9.9	1.0	0.8	1.2	1.4
NP1	1.4	1.3	0.9	1.1	1.6	0.9	0.3	1.3	1.3	-0.1	1.7	0.9	0.8	1.8	-0.1	-0.1	-0.1	0.8

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBM	052-LPB	053-LPB	054-HB
NP2	1.3	1.1	0.7	0.9	1.4	0.8	0.2	1.2	1.2	-0.1	1.4	0.9	0.8	1.4	-0.1	-0.1	-0.1	0.8
NP3	1.1	1.2	0.7	0.8	1.4	0.8	1.2	1.2	1.2	-0.1	1.6	0.8	0.7	1.7	-0.1	-0.1	-0.1	0.7
NP4	2.2	5.9	2.3	4.0	15.1	1.9	1.1	0.9	14.9	1.4	9.5	1.8	1.6	5.8	0.9	0.3	0.3	1.5
NPP4	1.1	1.5	0.7	0.8	1.5	-0.1	0.2	1.0	1.3	-0.1	1.7	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.7
NPP1-R	1.0	1.3	0.6	0.8	1.4	-0.1	0.2	0.9	1.2	-0.1	1.4	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.7
NPP2	1.5	4.9	1.2	1.6	5.3	1.2	0.8	0.3	4.3	0.7	3.6	1.3	1.1	5.3	0.8	0.7	0.9	1.2
NPP3	1.5	2.7	0.9	1.2	4.6	0.9	0.4	1.4	4.2	0.8	1.7	1.0	0.9	3.0	0.7	0.7	0.8	0.9
NPP4	2.1	3.0	1.6	2.4	7.2	1.3	1.0	1.8	6.8	1.0	4.9	1.4	1.2	3.2	-0.1	0.8	1.2	1.2
NQ1	0.3	1.3	1.1	1.4	1.8	1.1	0.6	1.3	1.3	-0.1	1.6	1.1	1.0	1.5	-0.1	0.7	0.9	1.0
NQ2	1.5	2.1	0.8	1.1	3.0	1.0	0.4	1.4	2.5	-0.1	3.2	1.1	0.9	3.4	-0.7	0.7	0.8	1.0
NQ2-R	1.4	2.3	0.9	1.2	5.3	1.1	0.4	0.3	4.9	0.8	3.1	1.2	1.0	4.2	0.7	0.7	0.8	1.0
NQ3	1.3	1.8	0.7	0.9	2.4	0.8	0.2	1.2	2.1	-0.1	2.3	0.8	0.8	2.2	0.8	-0.1	-0.1	0.8
NQ4	1.6	1.8	1.0	1.3	4.6	1.0	0.5	1.4	4.2	0.8	1.8	1.0	0.9	2.2	-0.1	0.7	0.8	0.9
NQ5	1.9	1.4	1.3	1.8	1.8	1.2	0.8	1.5	1.6	-0.1	1.6	1.1	1.0	1.5	-0.1	0.7	1.0	1.0
NQ6	1.1	1.1	0.7	0.8	1.3	-0.1	0.2	0.9	1.0	-0.1	1.3	0.8	0.7	1.3	-0.1	-0.1	-0.1	0.7
NQ7	2.0	2.9	1.3	2.3	9.2	2.0	1.2	0.3	8.9	1.1	2.0	1.9	1.6	3.7	0.7	-0.1	0.8	1.5
NQ8	1.6	1.6	1.0	1.3	3.4	1.1	0.6	0.3	3.2	0.7	1.2	1.2	0.9	1.8	-0.1	-0.1	0.8	0.9
NQQ1	1.6	1.4	0.9	1.2	1.7	0.9	0.5	1.2	1.5	-0.1	1.7	0.9	0.9	1.6	-0.1	-0.1	0.8	0.8
NR1	0.2	1.5	0.7	0.8	1.6	-0.1	1.1	1.0	1.2	-0.1	2.4	0.8	0.7	1.6	-0.1	-0.1	-0.1	0.7
NR11	1.7	3.8	1.1	0.4	7.7	2.5	1.0	0.2	7.2	0.9	3.5	2.1	1.6	5.0	0.8	0.7	1.1	1.6
NR12	1.7	2.8	1.4	2.1	4.7	1.2	0.9	1.1	4.3	0.8	1.7	1.2	1.1	2.7	0.7	0.7	1.0	1.1
NR13	2.0	9.1	2.1	3.3	30.9	1.7	1.1	0.4	30.0	2.7	14.2	1.7	1.4	7.9	0.9	0.8	1.3	1.4
NR14	1.5	2.9	0.8	1.0	2.6	0.9	0.4	1.2	2.2	-0.1	3.1	0.9	0.9	3.2	0.7	-0.1	-0.1	0.9
NR2	1.2	2.5	0.9	1.2	3.0	1.0	0.7	1.2	2.6	-0.1	2.8	1.1	0.9	3.0	0.8	0.7	0.8	1.0
NR3	1.5	3.1	1.6	2.6	5.2	1.8	0.9	0.3	4.4	0.8	6.2	1.8	1.4	5.7	0.8	0.8	1.1	1.4
NR4	1.1	1.1	0.8	1.1	1.5	0.9	0.4	0.7	1.3	-0.1	1.6	0.9	0.8	1.6	-0.1	-0.1	-0.1	0.7
NR5	1.2	1.2	0.8	0.9	1.5	0.8	0.4	1.1	1.1	-0.1	1.4	0.8	0.8	1.5	-0.1	-0.1	-0.1	0.8
NR6	1.1	1.1	0.8	1.0	1.4	0.9	0.4	1.1	1.2	-0.1	1.4	0.8	0.7	1.4	-0.1	-0.1	-0.1	0.7
NR7	1.6	4.7	1.3	1.9	6.1	1.5	0.8	0.3	5.2	-0.1	5.3	1.7	1.4	5.5	1.1	0.8	1.0	1.4
NR8	0.2	1.4	0.8	0.9	1.7	0.8	1.2	1.2	1.4	-0.1	2.2	0.9	0.8	2.2	-0.1	-0.1	-0.1	0.8
NS1	1.3	1.6	0.8	1.1	1.8	0.9	1.2	1.3	1.3	-0.1	2.2	0.8	0.8	2.4	0.7	-0.1	0.7	0.8
NS10	0.4	3.4	1.4	2.0	6.2	1.2	1.4	0.8	5.9	0.9	6.4	1.3	1.1	3.2	0.7	0.7	1.0	1.1
NS11	1.4	3.3	1.1	1.6	3.4	1.1	0.8	1.3	3.0	-0.1	3.4	1.2	1.0	3.1	-0.1	0.7	0.9	1.1
NS12	2.1	3.0	1.9	1.9	5.4	1.2	1.8	0.4	5.0	0.8	5.2	1.3	1.2	2.9	0.7	0.7	0.2	1.2
NS13	1.8	4.9	1.5	2.3	7.1	1.6	0.9	0.3	6.8	0.8	4.1	1.6	1.3	5.8	0.8	0.8	1.2	1.3
NS14	2.1	8.5	2.3	4.4	16.0	2.6	1.0	0.2	15.5	1.3	4.9	2.2	1.7	7.9	1.0	1.0	1.8	1.6
NS2	1.3	1.6	0.8	1.0	3.0	0.9	0.4	1.2	2.6	-0.1	1.0	0.9	0.8	2.0	-0.1	-0.1	0.7	0.8
NS3	1.6	2.3	1.4	2.1	3.2	1.3	0.9	0.5	4.2	0.7	1.5	1.3	1.2	3.2	-0.7	0.7	1.1	1.2
NS4	1.3	3.2	0.8	1.1	3.9	0.9	0.5	1.3	3.4	-0.1	3.2	1.0	0.9	4.0	0.9	0.7	0.8	0.9
NS4-R	1.4	2.6	0.8	1.0	3.1	0.9	0.4	1.2	2.7	-0.1	2.9	0.9	0.9	3.4	0.9	0.7	0.7	0.9
NS5	1.4	2.9	0.8	0.9	3.9	1.0	1.2	0.6	3.2	-0.1	4.8	1.1	1.0	5.7	1.0	-0.1	0.7	1.0
NS6	1.4	1.4	0.8	1.0	1.9	0.9	1.5	1.4	1.4	-0.1	2.2	0.9	0.8	2.3	-0.1	-0.1	0.7	0.8
NS7	1.2	4.7	0.9	1.2	5.6	1.0	0.5	1.3	5.0	-0.1	5.6	1.1	1.0	5.4	1.0	0.7	0.8	1.0
NS8	1.6	3.8	1.4	2.1	4.7	1.8	0.4	0.4	4.1	-0.1	3.7	1.3	1.3	4.0	1.0	0.6	1.1	1.3
NS9	1.1	1.5	0.7	0.9	1.8	-0.1	1.1	1.1	1.4	-0.1	1.7	0.8	0.7	2.0	0.7	-0.1	-0.1	0.7
NT1	0.5	1.3	1.4	1.3	2.0	1.0	1.2	1.3	1.8	-0.1	4.3	0.9	0.8	1.2	-0.1	-0.1	0.8	0.8
NT10	1.3	1.4	1.0	1.3	1.8	1.0	0.3	1.2	1.3	-0.1	1.8	0.9	0.9	1.8	-0.1	0.7	0.8	0.9
NT11	1.1	1.6	0.7	0.9	1.9	0.8	0.4	1.2	1.6	-0.1	2.8	0.9	0.8	1.9	0.7	-0.1	-0.1	0.8
NT12	1.9	1.6	1.1	1.5	2.1	1.0	1.0	1.6	2.0	-0.1	1.8	1.1	1.0	1.8	0.7	0.7	0.9	1.0
NT13	1.3	1.7	0.6	1.1	1.9	0.9	0.4	1.4	1.7	-0.1	1.8	0.9	0.8	1.8	-0.1	-0.1	-0.1	0.8
NT14	1.7	1.6	0.9	1.2	1.9	1.0	1.6	1.5	2.5	-0.1	2.7	0.9	0.9	1.8	-0.1	0.6	0.8	0.9
NT15	2.2	3.1	4.7	2.7	3.7	0.8	2.1	0.7	4.3	-0.1	9.5	1.2	1.2	2.8	-0.1	0.7	0.2	1.2
NT16	1.6	2.9	1.3	2.1	5.2	1.5	0.9	0.4	4.8	0.8	2.1	1.5	1.3	2.8	0.7	0.7	1.1	1.3
NT17	1.5	2.4	0.9	1.1	3.7	1.0	1.5	1.6	3.1	-0.1	8.3	1.0	0.8	2.1	-0.1	0.7	0.7	0.8
NT18	0.2	1.0	0.8	0.9	1.2	-0.1	1.0	1.0	1.0	-0.1	1.0	0.8	0.7	0.9	-0.1	-0.1	-0.1	0.7
NT2	0.6	1.4	1.1	1.4	0.9	1.0	1.3	1.5	1.8	-0.1	0.6	1.0	0.8	1.5	-0.1	-0.1	0.9	0.8
NT3	1.0	1.2	0.7	0.8	1.4	-0.1	0.2	0.9	1.2	-0.1	1.6	0.8	0.7	1.7	-0.1	-0.1	-0.1	0.7
NT4	1.1	1.3	0.7	0.8	1.6	-0.1	0.2	1.0	1.4	-0.1	2.3	0.8	0.7	1.7	-0.1	-0.1	-0.1	0.7
NT5	1.1	1.2	0.8	1.0	1.5	0.8	0.4	1.2	1.3	-0.1	1.6	0.9	0.8	1.6	-0.1	-0.1	-0.1	0.8
NT6-R	1.3	1.3	0.8	1.0	1.7	0.9	0.4	1.2	1.4	-0.1	2.0	0.9	0.8	2.1	-0.1	-0.1	0.7	0.8
NT6	1.2	1.5	0.7	1.0	2.0	0.9	0.4	1.3	1.7	-0.1	1.9	0.9	0.8	0.3	0.7	-0.1	-0.1	0.8
NT7	1.4	2.9	1.0	1.4	3.4	1.1	0.6	0.9	3.0	-0.1	8.7	1.2	1.1	4.2	0.9	0.7	0.9	1.1
NT8	1.6	5.2	1.4	2.1	10.1	1.9	0.9	0.8	9.6	1.0	5.6	1.9	1.5	8.4	1.0	0.8	1.0	1.5
NT9	1.2	2.3	0.8	1.0	1.5	0.9	0.4	1.2	1.3	-0.1	2.3	0.8	0.8	1.7	-0.1	-0.1	-0.1	0.8
NU1	1.1	1.8	0.9	1.0	2.1	0.9	0.4	1.2	1.7	-0.1	2.1	1.0	0.8	2.1	0.7	-0.1	0.7	0.9
NU10	1.6	2.6	1.0	1.4	8.4	1.1	0.8	0.3	8.1	1.0	2.3	1.3	1.2	4.2	0.8	0.7	0.9	1.2
NU11	1.4	1.7	0.8	1.1	1.9	0.9	0.3	1.3	1.7	-0.1	2.1	0.8	0.8	1.8	-0.1	-0.1	0.7	0.8

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBM	052-LPB	053-LPB	054-HB
NU12	0.3	1.7	0.8	1.0	3.0	0.9	0.3	1.4	2.7	-0.1	1.8	1.0	0.8	2.7	-0.1	-0.1	0.7	0.8
NU12-R	1.4	1.6	0.7	0.9	2.8	0.8	0.3	1.3	2.5	-0.1	1.9	0.9	0.8	2.0	-0.1	-0.1	-0.1	0.8
NU13	1.9	4.7	1.2	1.7	13.1	1.2	1.3	0.7	12.5	-1.2	2.6	1.4	1.4	4.7	-0.8	0.7	0.9	1.3
NU14	1.7	1.9	1.0	1.4	2.8	1.1	0.7	0.3	2.6	-0.1	2.8	1.3	1.1	2.8	0.7	0.7	0.8	1.1
NU15	1.6	2.7	1.1	1.4	6.8	1.1	0.3	0.8	6.4	1.0	5.2	1.1	1.0	2.6	0.7	0.7	0.8	1.1
NU16	1.7	9.6	1.5	2.4	20.2	1.7	0.9	0.8	19.4	1.6	6.8	1.7	1.5	10.3	1.0	0.8	1.2	1.5
NU17	1.9	4.1	1.6	1.9	12.5	1.5	1.3	0.7	12.1	1.3	10.8	1.7	1.3	4.4	0.7	0.7	1.0	1.3
NU18	1.5	3.0	1.0	1.2	3.7	1.1	0.3	1.5	3.0	-0.1	1.6	1.1	1.0	2.8	0.7	0.7	0.8	1.0
NU19	1.8	0.8	1.1	1.6	2.5	1.1	1.7	-0.1	2.4	-0.1	3.4	1.2	1.0	2.3	-0.1	0.7	0.8	1.0
NU2	1.9	10.2	1.6	2.7	9.0	0.8	1.5	0.9	10.8	1.1	8.1	0.4	1.4	13.4	1.1	0.2	1.2	1.4
NU3	1.2	1.2	0.6	0.9	1.4	-0.1	1.1	1.2	1.1	-0.1	1.3	0.8	0.7	1.3	-0.1	-0.1	0.7	0.7
NU4	2.2	9.8	1.5	2.3	13.5	1.4	1.3	0.8	12.2	0.9	13.2	1.6	1.4	11.6	1.5	0.9	1.1	1.4
NU5	1.1	1.6	0.7	0.8	1.9	-0.1	0.2	1.1	1.6	-0.1	1.9	0.9	0.7	2.1	0.7	-0.1	-0.1	0.7
NU6	1.1	2.0	0.7	0.9	2.4	0.8	1.2	1.3	2.1	-0.1	2.4	0.9	0.8	2.6	0.8	-0.1	-0.1	0.8
NU7	1.3	2.6	0.6	1.0	3.2	1.0	1.2	1.3	2.6	-0.1	4.9	1.1	0.9	3.1	-0.8	-0.1	0.7	0.9
NU8	1.8	3.4	1.0	1.5	6.3	1.1	1.5	0.2	6.0	0.8	3.4	1.3	1.2	5.2	0.8	0.7	0.9	1.1
NU9	1.7	1.1	1.2	1.8	4.7	1.5	0.9	0.3	4.4	0.8	1.6	1.6	1.5	2.6	-0.1	0.7	1.0	1.4
NV1	1.9	3.9	1.5	2.3	3.5	1.2	0.9	1.3	2.9	-0.1	3.7	1.3	1.2	3.4	0.7	0.8	1.1	1.2
NV10	1.3	1.5	1.0	1.4	2.0	1.0	0.6	1.0	1.8	-0.1	0.5	0.9	0.9	1.6	-0.1	0.7	0.8	0.9
NV11	1.5	1.6	1.0	1.3	2.0	1.0	0.4	1.2	1.5	-0.1	1.7	1.0	0.9	1.9	0.7	-0.1	0.8	0.9
NV12	1.4	1.6	1.0	1.3	2.0	1.0	0.3	1.2	1.5	-0.1	1.7	1.0	0.9	1.7	-0.1	0.7	0.8	0.9
NV13	1.9	3.5	1.5	2.3	8.2	1.7	1.2	0.4	8.0	1.0	7.5	1.6	1.4	3.5	-0.1	0.7	1.1	1.3
NV14	1.9	1.8	1.1	1.6	3.5	1.2	1.7	0.7	4.0	-0.8	4.2	1.4	1.1	2.1	-0.1	0.7	0.9	1.0
NV15	1.7	1.6	0.9	1.3	2.5	1.0	0.8	1.3	2.3	0.7	0.5	0.9	0.9	1.6	-0.1	-0.1	0.8	0.9
NV16	2.3	3.8	1.9	3.1	7.9	1.5	0.8	1.1	7.5	0.9	3.6	1.7	1.4	4.1	0.7	0.8	1.3	1.4
NV17	1.6	2.5	1.5	2.2	2.7	1.4	0.7	1.2	2.3	-0.1	2.7	1.3	1.2	2.6	0.7	0.7	1.1	1.2
NV18	1.8	2.4	1.1	1.6	3.5	1.1	0.3	1.4	3.2	-0.1	3.5	1.1	1.0	2.8	-0.1	0.7	0.9	1.0
NV2	1.8	2.9	1.2	1.8	2.8	1.2	0.7	1.4	2.4	-0.1	3.1	1.2	1.1	3.1	0.7	0.7	1.0	1.1
NV3	1.3	1.8	0.9	1.0	2.4	0.9	1.3	1.4	1.9	-0.1	0.6	1.0	0.8	1.9	-0.1	-0.1	0.8	0.8
NV4	2.1	2.6	1.2	1.9	4.6	1.4	1.6	1.0	4.2	0.8	4.9	1.4	1.2	3.0	0.7	0.7	1.0	1.2
NV5	1.8	3.4	1.3	2.2	7.4	1.7	0.4	0.3	6.9	-0.9	3.0	1.6	1.4	4.5	-0.8	0.8	1.1	1.4
NV5-R	2.0	3.7	1.6	2.2	7.1	1.3	1.0	1.5	6.2	0.9	2.6	1.4	1.3	4.1	0.7	0.8	1.2	1.2
NV6	1.4	1.6	1.0	1.4	2.7	1.4	0.4	1.3	2.4	0.7	1.4	1.1	1.0	1.6	-0.1	0.7	0.9	0.9
NV7	1.7	2.5	0.9	1.3	3.5	1.2	1.4	0.7	3.1	-0.1	1.2	1.2	1.0	2.5	-0.1	-0.1	0.8	1.0
NV8	1.5	1.4	0.8	1.3	0.9	1.1	0.4	1.4	1.8	-0.1	2.0	0.9	0.9	1.4	-0.1	-0.1	0.8	0.9
NV9	1.5	2.5	1.0	1.4	4.0	1.0	0.5	1.6	3.7	-0.1	1.7	1.0	0.9	2.7	-0.1	0.7	0.8	0.9
NW1	1.8	8.9	1.1	1.1	28.6	1.4	0.8	1.4	27.7	2.3	5.6	1.4	1.3	8.5	0.9	0.8	0.9	1.3
NW10	1.4	2.0	0.9	1.1	0.4	0.2	1.2	1.3	-0.1	-0.1	2.2	0.9	0.8	2.2	-0.1	-0.1	0.7	0.9
NW11	1.1	1.4	0.8	1.1	1.8	1.0	0.5	1.0	1.4	-0.1	4.9	1.1	0.8	1.9	-0.1	-0.1	-0.1	0.8
NW12	1.8	1.9	1.4	2.1	2.4	1.2	0.8	1.4	2.2	-0.1	2.0	1.1	1.1	1.9	-0.1	0.7	1.0	1.1
NW13	2.0	5.2	2.1	3.5	11.9	1.7	1.1	1.1	11.4	1.2	2.6	1.4	1.3	4.5	-0.8	0.8	1.5	1.3
NW14	0.1	1.6	0.9	1.0	1.7	0.9	0.3	1.2	1.4	-0.1	1.8	0.9	0.9	1.7	-0.1	-0.1	-0.1	0.9
NW15	2.1	5.5	1.7	2.8	5.6	1.5	1.1	1.1	5.3	-0.7	2.8	1.6	1.4	4.5	-0.8	0.8	1.2	1.4
NW16	-0.1	1.3	1.0	1.3	1.6	1.0	0.6	1.1	1.1	-0.1	1.4	1.0	0.9	1.4	-0.1	-0.1	0.8	0.9
NW17	1.3	1.5	0.9	1.1	0.3	0.2	1.4	1.5	0.2	-0.1	1.7	1.0	0.8	1.5	-0.1	0.7	0.7	0.9
NW17-R	1.4	1.5	0.9	1.2	0.3	0.2	1.4	1.5	0.2	-0.1	0.4	0.2	0.8	1.4	-0.1	0.7	0.7	0.9
NW18	1.1	3.6	1.0	1.4	2.6	1.0	0.4	1.2	3.5	-0.1	3.5	1.1	1.1	2.3	-0.1	0.7	0.8	1.1
NW2	1.7	2.3	1.0	1.3	1.0	0.2	1.5	1.5	2.2	-0.1	0.7	0.9	0.9	2.1	-0.1	0.7	0.8	0.9
NW2-R	1.4	2.0	1.0	1.2	1.0	1.0	1.4	1.4	2.1	-0.1	0.7	0.2	0.9	2.2	-0.1	-0.1	0.7	1.0
NW3	1.2	1.8	0.8	1.1	2.6	0.9	0.4	1.2	2.3	-0.1	0.8	1.0	0.9	2.1	-0.1	0.6	0.7	0.9
NW4	1.3	1.8	0.8	1.0	2.8	0.8	1.3	1.3	2.5	-0.1	1.1	0.9	0.8	1.8	-0.1	-0.1	0.8	0.9
NW5	1.7	15.7	1.3	1.9	24.0	1.3	0.8	1.0	22.5	1.6	10.6	1.6	1.3	15.7	1.1	0.8	1.0	1.4
NW6	1.4	2.0	0.9	1.1	3.9	0.9	0.3	1.4	3.4	-0.7	1.6	1.0	0.9	2.2	-0.1	-0.1	0.8	0.9
NW7	1.4	1.6	0.8	1.0	2.4	1.0	0.2	1.5	2.1	-0.1	0.8	1.0	0.9	1.8	-0.1	-0.1	0.7	0.8
NW8	2.7	12.2	2.6	4.9	35.4	2.1	1.8	0.2	34.2	2.7	1.8	1.8	1.5	12.8	1.1	0.8	0.9	1.5
NW9	0.3	2.8	0.8	1.0	3.8	1.0	1.2	0.8	3.5	-0.1	1.4	1.0	0.9	3.3	0.7	-0.1	0.7	0.9
NX1	0.3	1.4	0.9	1.1	1.5	0.9	0.3	1.1	1.3	-0.1	1.5	0.9	0.8	1.5	-0.1	0.6	0.8	0.8
NX10	1.6	2.0	1.0	1.3	2.4	1.0	0.5	1.3	2.1	-0.1	2.5	1.1	1.0	2.6	0.7	0.7	0.8	1.0
NX11	1.1	1.0	0.7	0.8	1.2	-0.1	1.0	1.0	1.0	-0.1	1.0	0.7	0.7	1.0	-0.1	-0.1	-0.1	0.7
NX12	1.3	1.2	0.8	0.9	1.6	0.8	1.2	1.3	1.4	-0.1	1.6	0.8	0.7	1.3	-0.1	-0.1	-0.1	0.7
NX13	1.5	2.6	1.2	1.7	4.1	1.2	0.6	0.4	3.6	0.7	1.9	1.3	1.1	3.1	-0.7	0.7	1.1	1.2
NX14	1.6	2.0	1.0	1.4	3.3	1.3	1.2	0.7	3.0	-0.1	1.2	1.5	1.1	2.4	-0.1	0.7	0.9	1.1
NX14-R	1.4	1.8	1.1	1.4	2.6	1.0	0.5	1.5	2.4	-0.1	0.8	1.1	0.9	2.0	-0.1	0.7	0.8	0.9
NX15	1.6	2.4	0.9	1.2	3.3	0.2	1.6	1.6	3.0	-0.1	1.6	1.0	0.9	2.5	0.7	0.6	0.8	0.9
NX16	1.5	2.7	0.9	1.3	4.5	1.1	0.7	0.5	4.1	-0.7	2.1	1.2	1.0	3.4	-0.7	0.7	0.8	1.0
NX17	1.9	5.3	1.6	2.6	10.0	1.8	1.0	0.8	9.6	1.0	3.5	1.9	1.6	5.8	0.9	0.8	1.2	1.5
NX2	1.1	1.5	0.8	1.0	1.6	0.9	1.2	1.3	1.4	-0.1	1.7	0.9	0.8	1.6	-0.1	-0.1	0.7	0.8

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBH	052-LPB	053-LPB	054-HB
NX3	1.3	1.2	0.8	1.0	1.6	0.9	0.3	1.2	1.4	-0.1	1.5	0.9	0.8	1.4	-0.1	-0.1	0.7	0.8
NX4	1.7	5.6	1.3	2.1	6.8	1.6	1.1	0.2	6.4	0.8	8.5	1.6	1.4	7.5	0.9	0.8	1.1	1.3
NX5	1.3	2.1	0.7	1.0	2.5	0.9	0.4	1.1	2.2	-0.1	2.5	1.0	0.8	2.2	-0.1	-0.1	-0.1	0.8
NX6	0.3	2.3	0.9	1.1	3.3	1.1	1.3	0.8	3.0	-0.1	1.5	1.2	1.0	2.7	0.7	0.7	0.8	1.0
NX7	2.2	15.4	2.6	4.6	28.2	2.4	1.2	0.2	26.8	2.1	9.9	2.0	1.6	16.8	1.2	1.0	0.4	1.5
NX8	1.6	3.0	1.2	1.6	5.0	1.3	0.9	0.4	4.9	0.8	2.0	1.4	1.2	3.4	0.7	0.7	0.9	1.2
NX9	1.6	3.3	1.2	1.8	6.7	1.6	0.9	0.7	6.4	0.7	2.5	1.7	1.4	4.2	0.8	0.8	1.0	1.4
NY1	1.4	1.6	0.8	1.1	2.0	0.9	0.2	1.2	1.8	-0.1	2.0	0.9	0.8	1.4	-0.1	-0.1	0.7	0.8
NY10	2.1	4.0	1.6	2.5	9.7	1.3	0.6	1.6	9.5	1.1	5.5	1.3	1.3	3.8	0.7	0.7	1.2	1.2
NY11	1.0	1.1	0.7	1.1	1.4	0.9	0.4	0.1	1.2	-0.1	1.2	0.8	0.7	1.2	-0.1	-0.1	-0.1	0.7
NY12	1.2	1.7	0.9	1.1	1.8	1.0	0.5	1.2	1.3	-0.1	1.5	1.0	0.8	1.7	-0.1	0.7	0.7	0.8
NY13	1.1	1.0	0.8	1.1	1.4	1.1	0.4	0.4	1.2	-0.1	1.5	1.0	0.8	1.2	-0.1	-0.1	0.8	0.8
NY14	1.9	2.7	1.7	2.9	3.2	1.6	1.0	0.3	3.0	-0.1	4.0	1.6	1.4	4.2	0.8	0.8	1.3	1.4
NY15	1.6	3.2	1.4	2.2	4.9	1.4	0.9	1.5	4.6	0.8	1.9	1.3	1.2	3.0	0.7	0.7	0.2	1.2
NY2	1.0	1.2	0.8	1.1	1.3	0.9	0.4	1.0	1.1	-0.1	1.2	0.9	0.7	1.1	-0.1	-0.1	0.8	0.8
NY3	1.1	1.5	0.7	0.8	1.4	-0.1	0.4	0.9	1.1	-0.1	1.5	0.8	0.7	1.3	-0.1	-0.1	0.7	0.7
NY4	3.0	15.1	2.8	5.3	41.1	2.3	1.2	1.8	39.9	3.2	9.8	2.0	1.6	14.0	1.0	1.0	1.9	1.6
NY4-R	3.4	19.6	3.3	6.3	63.0	2.6	1.2	2.4	61.5	4.7	13.1	2.2	1.7	18.8	1.2	1.2	2.2	1.7
NY5	1.3	1.8	0.9	1.2	1.2	0.9	0.4	1.2	2.1	-0.1	2.3	0.9	0.8	1.5	-0.1	-0.1	0.8	0.8
NY6	1.5	3.9	1.3	1.8	5.5	1.3	0.4	0.9	5.1	0.8	2.1	1.2	1.1	2.9	-0.1	0.7	1.0	1.1
NY7	2.0	5.6	1.6	2.4	3.9	1.5	0.9	1.5	3.2	-0.1	4.3	1.3	1.2	4.4	0.8	0.8	1.2	1.2
NY8	1.6	1.7	0.9	1.1	2.0	1.1	1.4	1.4	2.4	-0.1	2.8	1.2	0.9	1.8	-0.1	0.8	0.8	0.9
NY9	0.5	1.5	1.3	1.6	2.6	1.1	0.3	1.1	2.4	-0.1	1.1	1.0	0.9	1.5	-0.1	-0.1	1.0	0.9
NZ1	1.4	1.5	0.9	1.2	1.8	1.0	0.4	1.2	1.6	-0.1	1.9	1.0	0.9	1.8	-0.1	-0.1	0.8	0.9
NZ10	2.7	5.6	3.2	5.4	16.1	3.1	1.4	0.4	15.8	1.5	8.9	2.4	1.8	6.1	0.8	1.1	2.3	1.8
NZ11	1.2	1.6	1.0	1.3	2.0	1.0	0.4	1.1	1.7	-0.1	1.7	0.9	0.8	1.3	-0.1	0.7	0.8	0.8
NZ12	1.9	2.0	1.2	1.7	4.2	1.3	1.4	0.7	4.2	-0.1	1.8	1.3	1.2	2.6	-0.1	0.7	1.0	1.1
NZ13	1.2	1.4	0.8	1.0	1.7	0.9	0.4	1.1	1.5	-0.1	1.5	0.9	0.7	1.4	-0.1	-0.1	0.7	0.8
NZ2	1.3	3.0	0.9	1.2	2.8	1.0	0.4	1.2	2.0	-0.1	2.6	1.0	1.0	2.7	0.7	0.7	0.8	1.0
NZ2-R	1.0	1.1	0.6	1.0	1.3	0.8	0.3	0.7	1.6	-0.1	1.1	0.8	0.7	1.1	-0.1	-0.1	-0.1	0.8
NZ3	1.2	3.1	0.9	1.3	2.5	1.0	0.4	1.2	2.2	-0.1	2.4	1.0	0.9	2.3	0.7	0.7	0.8	0.9
NZ4	1.7	2.1	0.9	1.2	3.2	1.0	0.5	1.4	2.8	0.7	3.0	1.2	1.1	1.9	-0.1	-0.1	0.8	1.1
NZ5	1.6	1.9	1.1	1.4	2.5	1.1	0.7	1.3	2.1	-0.1	2.3	1.1	1.0	1.8	-0.1	0.7	0.9	1.0
NZ6	1.5	1.7	0.9	1.2	2.9	1.0	0.6	1.3	2.7	-0.1	2.0	1.0	0.9	1.8	-0.1	0.3	0.8	0.9
NZ7	1.9	2.0	1.3	2.0	4.0	1.4	0.9	1.1	3.7	0.8	3.5	1.5	1.3	2.2	-0.1	0.7	1.1	1.2
NZ8	1.6	2.0	4.0	1.4	2.2	1.4	0.9	1.5	1.9	-0.1	2.4	1.2	1.0	2.0	-0.1	0.7	0.9	1.0
NZ9	1.2	1.2	1.0	1.2	1.4	1.0	0.4	1.0	1.1	-0.1	1.3	0.9	0.8	1.2	-0.1	-0.1	0.8	0.8
LMB-QA	1.0	0.2	-0.1	-0.1	1.4	-0.1	0.8	0.8	1.1	-0.1	1.6	0.8	-0.1	1.8	-0.1	-0.1	-0.1	0.7
LMB-QA	1.0	1.0	-0.1	-0.1	1.3	-0.1	0.8	-0.1	1.1	-0.1	1.3	-0.1	-0.1	1.2	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.9	0.2	-0.1	-0.1	0.2	-0.1	0.8	-0.1	0.2	-0.1	1.2	-0.1	-0.1	1.2	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.9	1.0	-0.1	-0.1	0.3	-0.1	0.8	0.8	0.3	-0.1	1.3	-0.1	-0.1	1.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	1.0	1.0	-0.1	-0.1	1.3	-0.1	0.8	0.8	1.0	-0.1	1.4	0.7	-0.1	1.5	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.2	1.0	-0.1	-0.1	1.4	-0.1	0.8	0.7	1.2	-0.1	1.4	-0.1	-0.1	1.4	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.9	1.0	-0.1	-0.1	0.5	-0.1	0.8	0.8	1.3	-0.1	1.4	-0.1	-0.1	1.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	1.0	1.0	-0.1	-0.1	0.5	-0.1	0.8	0.8	1.4	-0.1	1.3	0.7	-0.1	1.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	1.0	0.9	-0.1	-0.1	0.3	-0.1	0.8	0.8	1.1	-0.1	1.1	0.7	-0.1	1.2	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.9	0.8	-0.1	-0.1	1.0	-0.1	0.8	-0.1	0.9	-0.1	0.8	-0.1	-0.1	0.8	-0.1	-0.1	-0.1	-0.1
LMB-QA	1.0	0.2	-0.1	-0.1	1.3	-0.1	0.8	0.7	1.1	-0.1	1.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1
LMB-QA	0.9	0.9	-0.1	-0.1	0.2	-0.1	0.8	0.7	1.1	-0.1	1.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	037-HB	038-LBA	039-LAR	040-LPB	041-LBA	042-LPB	043-HB	044-HB	045-LA	046-LPH	047-LBA	048-HB	049-HB	050-LBA	051-LBJ	052-LPB	053-LPB	054-HB

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055-LRB	056-LBL	057-ALK	058-LRB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HRB	071-HPB	072-HPB
NA1	0.9	-0.1	1.2	1.1	1.7	5.5	1.3	5.6	3.2	8.5	1.9	11.0	1.5	2.7	11.0	4.1	7.1	10.0
NA2	-0.1	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	1.7	0.2	2.3	1.1	2.5	-0.1	-0.1	2.6	-0.1	1.7	2.1
NAA1	0.8	-0.1	0.8	1.1	0.8	3.7	-0.1	3.3	2.3	4.8	1.4	4.8	1.1	1.6	5.0	2.1	2.7	1.1
NAA10	0.7	-0.1	0.7	0.9	0.4	2.4	-0.1	2.7	1.6	5.9	1.4	4.9	1.1	1.2	4.7	1.8	2.4	3.2
NAA11	1.1	0.8	4.0	1.4	2.0	2.7	3.1	10.2	1.8	6.2	2.4	13.0	3.9	2.7	13.5	4.9	9.2	12.6
NAA2	1.9	0.6	7.0	2.9	3.7	11.0	7.1	34.8	6.4	31.6	5.5	94.8	10.4	1.4	95.1	13.6	28.5	40.8
NAA3	0.7	-0.1	2.4	1.0	0.3	1.7	1.4	9.2	1.4	13.1	1.4	13.1	1.6	1.5	13.8	2.2	3.0	1.0
NAA4	-0.1	-0.1	-0.1	0.8	0.2	2.1	-0.1	2.9	1.2	4.0	1.2	4.8	1.1	1.2	4.9	0.6	2.0	2.5
NAA5	0.7	-0.1	0.9	0.9	0.3	1.4	1.1	3.4	0.4	4.9	1.4	5.3	1.2	1.5	5.6	2.3	3.5	1.8
NAA6	0.8	-0.1	0.9	1.1	0.6	2.0	1.2	4.3	1.4	6.3	1.3	8.7	1.3	1.6	8.4	2.1	2.9	4.1
NAA7	0.7	-0.1	0.2	0.9	0.2	2.1	-0.1	2.6	1.1	3.7	1.3	4.7	1.1	1.3	4.8	1.7	2.2	2.7
NAA8	-0.1	-0.1	-0.1	0.8	0.2	1.1	-0.1	2.8	1.1	5.2	1.2	5.1	1.0	-0.1	5.1	-0.1	1.8	2.0
NAA9	0.8	-0.1	1.1	0.3	1.0	2.8	1.3	4.7	1.7	9.2	1.6	10.4	1.4	1.8	10.4	2.5	3.9	5.2
NB1	0.7	-0.1	0.2	1.1	0.7	2.6	-0.1	3.9	1.6	5.2	1.4	6.5	1.2	1.5	6.5	2.0	2.9	3.7
NB2	-0.1	-0.1	-0.1	0.7	0.1	0.6	-0.1	2.9	0.2	5.9	1.1	5.2	-0.1	-0.1	5.4	-0.1	1.7	2.0
NB3	-0.1	-0.1	-0.1	0.8	0.2	0.8	-0.1	2.2	0.4	3.0	1.1	3.0	-0.1	-0.1	3.2	-0.1	1.3	0.6
NB4	0.7	-0.1	1.0	0.9	0.3	1.7	-0.1	4.8	0.5	6.0	1.2	8.6	1.2	1.3	8.7	1.7	2.1	0.4
NBB1	0.7	-0.1	-0.1	0.3	0.7	2.4	-0.1	2.8	1.3	5.3	1.4	4.3	1.0	1.3	4.3	1.7	2.2	2.9
NBB10	0.8	-0.1	1.3	1.0	0.7	1.5	1.6	4.6	0.4	2.9	1.5	10.6	2.0	2.6	10.8	3.9	6.9	1.5
NBB11	1.0	-0.9	2.5	0.8	1.2	5.5	1.4	7.4	3.1	12.0	2.4	13.5	1.7	2.5	13.5	4.5	8.9	13.0
NBB2	0.8	0.9	1.9	1.1	0.8	1.8	1.8	7.8	0.5	5.4	1.6	14.3	2.2	1.8	14.4	2.7	4.2	5.7
NBB2-R	0.8	-0.1	1.6	0.3	0.7	1.5	1.6	8.1	0.4	4.7	1.4	10.7	1.9	1.6	11.0	2.3	3.5	4.7
NBB3	0.9	0.9	2.1	0.3	1.0	1.7	1.7	8.0	0.5	4.6	1.7	13.2	2.1	1.8	13.2	2.6	3.6	0.8
NBB4	0.7	-0.1	-0.1	0.9	0.2	0.5	-0.1	2.2	0.2	2.9	1.2	3.2	1.0	-0.1	3.4	-0.1	1.6	0.6
NBB5	-0.1	-0.1	-0.1	0.8	0.2	1.5	-0.1	2.7	1.3	3.9	1.2	4.7	1.0	-0.1	4.7	-0.1	1.8	0.4
NBB6	0.9	-0.2	2.0	0.4	1.0	4.2	1.6	7.5	2.2	7.3	1.9	13.7	1.9	1.7	14.0	2.9	4.9	6.6
NBB7	-0.1	-0.1	-0.1	-0.1	0.2	0.5	-0.1	1.7	0.3	2.6	-0.1	2.6	-0.1	-0.1	2.5	-0.1	-0.1	1.5
NBB8	-0.1	-0.1	0.9	0.3	0.5	1.8	1.1	6.2	0.9	4.7	1.3	10.1	1.2	1.2	4.6	1.7	2.4	3.1
NBB9	0.9	0.3	3.2	0.4	0.8	1.8	2.4	10.7	1.0	2.7	1.8	14.3	2.9	1.9	15.1	3.7	7.0	10.1
NC1	-0.1	-0.1	-0.1	0.7	0.2	0.6	-0.1	2.8	0.3	3.8	1.1	4.4	-0.1	-0.1	4.4	-0.1	1.7	2.0
NC2	-0.1	-0.1	-0.1	0.8	0.2	0.9	-0.1	2.9	0.5	5.2	1.2	5.2	1.0	1.1	5.4	1.5	1.9	2.3
NC3	-0.1	-0.1	0.7	0.7	0.2	-0.1	-0.1	3.0	0.3	4.7	1.1	5.7	1.0	-0.1	5.7	-0.1	1.4	1.7
NC4	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	1.8	0.3	3.0	-0.1	3.0	-0.1	-0.1	3.0	-0.1	-0.1	1.4
NC5	-0.1	-0.1	0.9	0.8	0.2	0.7	-0.1	3.5	0.4	5.2	1.2	6.3	1.1	-0.1	6.3	-0.1	1.7	2.0
NC6	0.7	-0.1	0.2	0.3	0.9	3.7	-0.1	3.5	2.0	2.5	1.5	4.9	1.1	1.4	4.9	1.8	2.4	3.0
NC7	-0.1	-0.1	-0.1	0.7	0.2	0.4	-0.1	0.3	0.5	2.9	1.4	3.6	1.0	-0.1	0.9	-0.1	0.8	2.2
NC8	-0.1	-0.1	0.6	0.8	0.2	0.6	-0.1	2.4	0.4	3.3	1.1	3.6	1.0	-0.1	3.7	-0.1	1.7	0.6
NCC1	0.8	-0.9	2.7	0.3	0.6	2.1	3.1	19.0	1.2	19.8	2.2	39.6	4.2	6.2	49.2	10.0	30.3	44.1
NCC1-R	0.8	0.9	2.6	1.2	1.0	1.9	3.2	18.2	1.2	19.1	2.2	45.6	4.4	6.7	46.2	13.3	29.8	43.8
NCC2	1.1	-1.1	4.0	1.1	2.0	7.7	3.1	0.9	4.6	13.9	2.4	28.7	4.0	8.5	29.2	17.5	36.3	55.2
NCC3	0.8	0.8	2.9	0.3	0.7	3.2	1.7	5.9	1.8	9.3	1.7	13.6	2.1	2.9	13.6	4.9	9.8	14.6
NCC4	1.1	1.1	3.4	0.4	1.4	2.0	3.4	12.2	1.3	18.8	2.3	30.6	4.5	6.2	30.6	12.1	26.5	40.2
NCC5	0.7	0.8	4.1	1.0	0.8	2.5	1.4	5.9	1.5	8.8	1.4	11.5	1.6	2.1	11.9	3.2	5.8	8.3
NCC6	0.7	-0.1	4.3	0.8	0.2	1.2	1.8	4.0	0.2	6.6	1.4	7.9	2.1	2.2	8.2	3.6	5.2	4.1
ND1	0.7	-0.1	-0.1	1.0	0.8	1.9	-0.1	2.5	1.1	3.5	1.3	3.6	1.0	1.6	3.8	2.1	2.9	3.8
ND10	-0.1	-0.1	1.0	0.8	0.2	0.5	-0.1	3.3	0.2	5.1	1.2	5.0	-0.1	-0.1	5.2	-0.1	1.3	0.4
ND2	0.7	-0.1	1.4	1.1	0.7	1.6	1.3	5.5	0.5	9.0	1.4	8.8	1.4	1.6	9.0	2.2	3.2	0.8
ND3	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.1	0.4	3.7	1.4	4.1	-0.1	-0.1	4.4	-0.1	-0.1	1.5
ND4	0.8	0.8	3.0	1.0	0.4	1.6	1.4	12.0	1.0	16.7	1.5	18.3	1.6	1.3	18.9	1.8	2.2	0.6
ND5	-0.1	-0.1	0.2	-0.1	0.7	-0.1	-0.1	3.6	1.1	5.3	1.2	4.7	-0.1	-0.1	4.8	-0.1	-0.1	0.3
ND6	-0.1	-0.1	0.8	0.8	0.2	0.9	-0.1	3.2	0.5	4.2	1.1	4.1	1.0	-0.1	4.3	-0.1	1.5	0.5
ND7	-0.1	-0.1	-0.1	0.8	0.2	1.4	-0.1	1.5	0.4	3.0	1.1	1.6	-0.1	-0.1	1.6	-0.1	1.7	2.0
ND8	-0.1	-0.1	0.7	0.8	0.2	0.7	-0.1	2.6	0.5	4.3	1.2	4.2	1.0	-0.1	4.1	-0.1	1.5	0.5
ND8-R	-0.1	-0.1	0.9	0.9	0.2	1.5	-0.1	3.4	0.5	5.0	1.1	5.3	1.0	-0.1	5.3	-0.1	1.6	0.6
ND9	0.7	-0.1	0.8	0.9	0.3	1.9	-0.1	3.8	1.1	5.7	1.2	5.4	1.0	-0.1	5.6	-0.1	1.7	2.0
NDD1	0.9	-0.9	2.3	1.2	1.8	7.1	1.8	8.9	3.7	11.8	1.9	10.3	2.0	2.4	10.7	3.2	4.8	6.4
NDD2	0.8	-0.1	1.1	0.3	1.1	2.3	1.2	4.1	1.4	6.2	1.5	8.3	1.4	1.9	8.6	2.9	4.3	1.1
NDD3	0.8	-0.1	0.8	0.4	0.8	4.3	-0.1	3.6	2.5	5.1	1.5	6.2	1.1	1.4	5.9	2.0	2.8	3.8
NDD4	0.7	-0.1	0.8	0.3	0.6	3.0	-0.1	3.1	2.0	4.9	1.5	4.6	1.1	1.4	4.6	1.9	2.6	0.8
NDD5	-0.1	-0.1	-0.1	0.8	0.2	1.4	-0.1	3.2	1.3	4.9	1.2	4.7	1.0	-0.1	5.0	-0.1	1.7	2.1
NE1	1.1	1.0	2.6	1.2	1.8	3.2	2.5	10.1	2.0	7.0	2.3	22.2	3.3	3.5	22.6	5.6	11.3	15.9
NE10	-0.1	-0.1	0.4	-0.1	0.2	0.5	-0.1	1.6	0.4	2.6	-0.1	2.5	-0.1	-0.1	2.7	-0.1	-0.1	1.2
NE11	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.1	0.3	3.1	1.1	3.5	-0.1	-0.1	3.6	-0.1	1.3	0.4
NE2	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.8	0.2	2.7	-0.1	2.7	-0.1	-0.1	2.7	-0.1	-0.1	1.3
NE3	-0.1	-0.1	-0.1	0.9	0.2	1.0	-0.1	2.3	1.0	3.7	1.2	4.5	1.0	-0.1	4.7	-0.1	1.8	2.2
NE4	-0.1	-0.1	0.4	0.3	0.3	1.6	-0.1	2.2	1.2	3.5	1.2	3.9	-0.1	-0.1	3.4	-0.1	1.7	2.0
NE4-R	-0.1	-0.1	-0.1	0.8	0.2	0.9	-0.1	1.6	0.6	2.4	1.1	2.4	-0.1	-0.1	2.4	-0.1	-0.1	1.5

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055-LPB	056-LBI	057-ALK	058-LPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NE5	0.8	-0.1	0.9	0.3	0.7	2.6	1.1	3.5	1.5	7.0	1.5	6.0	1.2	1.7	5.7	2.6	4.3	1.8
NE6	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.2	0.3	4.4	1.1	3.7	-0.1	-0.1	3.5	-0.1	-0.1	0.5
NE7	-0.1	-0.1	-0.1	0.9	0.2	-0.1	-0.1	1.6	0.2	2.3	1.1	2.2	1.0	-0.1	2.3	0.8	2.3	2.8
NE8	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.4	0.2	4.7	-0.1	4.1	-0.1	-0.1	4.1	-0.1	-0.1	1.2
NE9	0.7	-0.1	-0.1	0.8	0.3	1.9	-0.1	2.8	1.1	7.8	1.2	4.7	-0.1	1.1	4.7	-0.1	1.8	2.1
NEE1	0.7	-0.1	0.7	0.8	0.2	1.5	-0.1	3.2	0.5	6.6	1.2	5.5	1.1	1.3	5.8	1.6	1.8	0.7
NEE2	0.8	-0.1	3.6	0.4	0.8	1.7	1.4	5.0	1.0	8.9	1.7	12.8	1.7	1.5	13.2	2.0	2.8	3.5
NEE3	0.8	-0.1	1.1	1.1	0.7	2.0	1.3	5.5	1.3	9.1	1.4	11.0	1.4	1.4	11.0	1.8	2.5	3.2
NEE4	-0.1	-0.1	0.7	0.8	0.2	1.8	-0.1	2.8	0.2	3.6	1.2	5.0	1.1	1.2	5.1	-0.1	1.7	1.9
NEE5	-0.1	-0.1	-0.1	0.8	0.2	-0.1	-0.1	2.5	1.0	4.4	1.2	5.0	1.1	-0.1	5.0	-0.1	1.4	0.3
NEE6	0.8	-0.1	1.1	0.4	0.8	3.2	1.2	5.0	2.1	4.0	1.5	7.1	1.4	1.4	7.1	2.0	2.6	0.8
NEE7	0.9	-0.1	0.8	1.5	0.9	2.0	-0.1	3.2	3.1	5.0	1.7	5.2	1.0	1.6	5.3	1.9	1.9	1.6
NF1	0.8	-0.2	1.3	0.3	0.7	1.7	1.9	0.7	0.8	7.0	1.5	7.2	1.5	1.5	6.0	3.0	6.0	8.9
NF10	-0.1	-0.1	-0.1	0.8	0.2	0.7	-0.1	2.3	0.4	3.8	1.3	5.1	1.1	1.5	4.9	2.4	4.0	5.5
NF11	0.7	-0.1	0.4	0.3	0.9	4.3	1.4	5.6	1.9	1.4	1.3	12.9	0.4	2.8	13.2	6.5	7.0	8.9
NF2	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.3	0.2	3.6	-0.1	3.7	-0.1	-0.1	3.7	-0.1	-0.1	0.5
NF3	-0.1	-0.1	-0.1	0.7	0.2	0.6	-0.1	2.0	0.5	3.0	1.1	2.9	-0.1	-0.1	3.1	0.5	2.1	2.8
NF4	-0.1	-0.1	0.7	0.7	0.2	1.2	-0.1	2.9	0.4	4.2	1.1	5.1	1.0	-0.1	5.3	-0.1	1.8	2.4
NF5	-0.1	-0.1	0.4	0.7	0.2	-0.1	0.1	2.7	0.3	4.7	1.1	5.6	1.1	-0.1	5.6	0.5	2.0	2.5
NF6	-0.1	-0.1	1.2	0.8	0.2	-0.1	-0.1	4.5	0.2	5.6	1.1	8.5	0.3	0.4	8.7	0.9	2.7	0.5
NF7	1.2	1.0	0.5	1.3	1.9	5.9	1.6	10.1	3.6	8.5	2.8	18.9	1.9	4.5	19.4	7.6	15.7	22.1
NF8	-0.1	-0.1	-0.1	0.8	0.2	1.2	-0.1	2.3	1.3	3.6	1.2	3.8	-0.1	-0.1	3.8	1.8	2.4	3.2
NF8-R	0.7	-0.1	0.2	0.3	0.7	2.9	1.2	5.6	1.8	7.1	1.3	8.5	1.3	1.9	8.7	2.3	4.3	5.8
NF9	-0.1	-0.1	0.8	0.9	0.2	0.8	-0.1	3.2	1.1	4.4	1.2	5.9	1.1	1.4	6.1	1.9	2.6	1.2
NFF1	0.7	-0.1	-0.1	0.8	0.2	2.1	-0.1	3.3	1.2	7.2	1.2	7.0	1.1	1.3	6.8	1.9	2.7	1.3
NFF2	0.8	-0.1	1.5	1.3	0.8	2.2	1.6	6.5	2.1	10.1	1.6	11.7	1.8	2.4	12.1	3.3	4.7	3.0
NFF3	-0.1	-0.1	-0.1	0.7	0.2	-0.1	1.1	2.8	0.3	3.4	1.1	4.1	1.2	-0.1	4.3	0.4	1.8	0.8
NFF4	0.9	0.8	1.4	0.9	1.2	4.1	1.5	5.4	2.9	8.9	1.9	12.6	1.8	2.6	12.6	3.9	6.6	5.6
NFF5	0.8	-0.1	1.0	1.2	0.7	2.1	1.2	4.4	1.7	7.0	1.4	6.4	1.2	1.7	6.6	2.2	2.8	0.6
NFF6	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.2	0.3	3.5	1.1	4.2	1.0	-0.1	4.4	-0.1	1.4	0.5
NFF6-R	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	1.9	0.3	2.5	1.1	3.1	1.0	-0.1	3.1	-0.1	1.5	0.5
NFF7	0.7	-0.1	-0.1	0.9	0.6	0.7	-0.1	3.7	0.5	6.9	1.3	6.5	1.1	-0.1	6.8	1.7	2.1	0.5
NFF8	0.7	0.7	-0.1	0.9	0.5	1.6	-0.1	2.8	0.5	4.4	1.4	4.5	1.1	1.3	4.3	1.7	2.2	2.8
NG1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.6	-0.1	2.1	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	-0.1
NG10	-0.1	-0.1	0.0	0.8	0.2	-0.1	-0.1	2.0	0.2	4.6	1.2	4.7	1.1	-0.1	4.9	-0.1	1.4	0.4
NG11	-0.1	-0.1	1.0	0.7	0.1	-0.1	-0.1	3.5	0.2	4.2	-0.1	4.4	-0.1	-0.1	4.6	-0.1	1.2	1.4
NG2	-0.1	-0.1	0.8	-0.1	0.2	-0.1	-0.1	2.8	1.1	3.6	-0.1	3.5	-0.1	-0.1	3.7	-0.1	-0.1	1.2
NG2-R	-0.1	-0.1	0.7	-0.1	0.2	-0.1	-0.1	2.6	-0.1	3.4	-0.1	3.6	-0.1	-0.1	3.8	-0.1	-0.1	1.2
NG3	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	1.7	0.2	2.5	-0.1	2.5	-0.1	-0.1	2.6	-0.1	-0.1	1.3
NG4	-0.1	-0.1	-0.1	0.9	0.2	0.9	-0.1	2.1	0.6	3.1	1.2	2.8	-0.1	-0.1	2.9	-0.1	1.6	2.0
NG5	0.8	-0.1	1.6	0.3	0.7	2.6	1.3	6.1	1.6	8.6	1.6	10.7	1.5	1.5	10.4	2.3	3.5	4.7
NG6	-0.1	-0.1	-0.1	0.8	0.2	1.4	-0.1	2.1	0.3	3.1	1.1	3.2	-0.1	-0.1	3.3	-0.1	1.3	1.6
NG7	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.1	0.2	1.6	-0.1	1.2	-0.1	-0.1	1.2	-0.1	-0.1	1.2
NG8	-0.1	-0.1	-0.1	0.7	0.8	-0.1	-0.1	2.3	1.1	4.0	1.1	3.4	-0.1	-0.1	3.7	-0.1	1.3	0.4
NG9	-0.1	-0.1	0.1	0.9	0.3	1.6	-0.1	3.0	0.5	4.4	1.2	4.8	0.3	-0.1	4.6	0.6	1.9	2.3
NGG1	-0.1	-0.1	-0.1	0.7	0.1	0.5	-0.1	3.0	0.2	4.6	1.2	5.1	1.1	0.6	5.2	1.2	3.0	1.0
NGG10	0.7	0.7	0.1	0.4	0.2	1.2	1.5	5.5	0.3	3.6	1.3	11.1	1.9	1.5	11.3	2.3	3.8	5.3
NGG2	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.5	0.2	4.2	1.1	3.9	1.0	-0.1	4.0	0.6	1.9	0.7
NGG3	0.9	0.9	2.6	0.4	0.8	1.8	2.7	18.3	1.7	17.8	2.4	22.7	3.5	3.8	23.2	7.3	11.2	8.3
NGG4	0.7	-0.1	1.2	1.0	0.6	2.0	1.5	4.9	1.3	8.0	1.5	10.8	1.7	1.8	11.4	3.1	5.3	4.7
NGG5	0.7	-0.1	0.8	0.9	0.7	1.7	-0.1	3.3	1.1	5.1	1.1	6.1	1.3	1.9	6.3	2.7	4.4	1.6
NGG6	-0.1	-0.1	0.8	0.8	0.2	1.5	-0.1	3.4	0.5	6.3	1.3	6.3	1.2	1.5	6.1	2.1	3.0	1.4
NGG7	0.7	-0.1	0.8	0.9	0.2	1.5	1.2	4.3	0.5	7.3	1.3	7.9	1.3	1.9	8.2	2.6	4.0	1.8
NGG8	-0.1	-0.1	1.0	0.8	0.1	-0.1	1.3	4.4	0.2	6.3	1.2	8.2	1.5	-0.1	8.6	0.9	2.5	1.1
NGG9	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.2	0.2	3.1	1.1	3.4	-0.1	-0.1	3.5	-0.1	1.3	0.5
NH1	-0.1	-0.1	0.7	0.8	0.2	0.5	-0.1	3.0	0.3	5.3	1.2	4.8	-0.1	-0.1	4.8	-0.1	1.3	1.4
NH10	0.7	-0.1	-0.1	1.1	0.9	2.1	-0.1	2.2	1.7	3.1	1.2	2.8	-0.1	-0.1	3.0	1.4	1.5	1.6
NH10-R	0.8	-0.1	-0.1	1.5	1.3	4.3	-0.1	2.6	2.5	3.6	1.5	3.2	1.0	1.4	3.3	1.7	2.0	2.2
NH11	0.7	-0.1	0.8	1.0	0.6	2.2	-0.1	3.5	1.3	5.1	1.3	5.7	1.1	1.2	5.9	1.4	1.8	1.8
NH2	0.7	-0.1	0.8	1.1	0.7	1.4	-0.1	2.8	0.2	4.1	1.3	3.8	1.0	1.2	4.1	1.5	1.6	0.5
NH3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.6	0.2	2.3	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	1.0
NH4	0.7	-0.1	0.7	0.9	0.3	1.8	-0.1	3.1	1.2	5.0	1.2	5.3	1.0	-0.1	5.3	1.4	1.5	1.6
NH5	0.7	-0.1	-0.1	1.0	0.7	1.9	-0.1	2.5	1.3	4.1	1.2	3.2	-0.1	-0.1	3.4	-0.1	1.4	1.5
NH6	0.8	-0.1	-0.1	0.3	1.0	3.6	-0.1	2.8	2.1	4.1	1.4	3.5	1.0	1.4	3.6	1.7	1.9	2.0
NH7	1.0	0.7	0.2	1.1	1.5	7.1	1.1	6.2	4.7	9.6	1.9	10.5	0.3	1.9	10.5	2.3	2.8	3.1
NH8	0.7	-0.1	0.7	1.1	0.8	3.2	-0.1	2.9	1.8	4.2	1.4	4.0	1.0	1.3	4.2	1.5	1.6	1.7
NH9	0.8	-0.1	0.8	1.1	0.8	3.0	-0.1	0.4	1.7	4.1	1.3	4.1	1.0	1.3	4.2	1.5	1.7	1.8

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055-LPB	056-LBI	057-ALK	058-LPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NHH1	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.2	0.2	3.5	-0.1	4.1	1.1	-0.1	4.1	0.5	1.9	0.5
NHH10	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.3	0.2	4.1	1.1	4.2	1.0	-0.1	4.5	0.6	1.9	0.4
NHH11	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.7	1.2	3.8	1.1	4.1	1.1	0.5	4.4	1.1	2.8	1.3
NHH11-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.5	0.2	3.9	1.1	3.8	1.1	0.4	3.9	0.9	2.4	1.1
NHH12	-0.1	-0.1	1.4	0.7	0.2	0.5	-0.1	2.8	0.3	4.8	1.1	6.0	1.1	-0.1	6.0	0.9	2.6	3.5
NHH2	0.7	0.8	2.1	0.9	0.6	1.8	2.1	8.8	1.0	5.4	1.6	18.7	2.7	1.9	19.1	3.4	5.5	7.9
NHH3	-0.1	-0.1	0.7	0.7	0.2	0.6	-0.1	2.7	0.4	4.6	1.2	5.4	1.1	-0.1	5.4	0.8	2.5	1.1
NHH4	0.8	-0.1	1.4	1.0	0.8	1.2	1.7	4.7	0.2	7.1	1.5	10.6	2.0	3.4	10.9	5.2	9.2	8.4
NHH5	-0.1	-0.1	1.4	0.8	0.2	1.5	1.7	6.3	0.4	9.0	1.3	13.4	2.0	1.6	13.8	2.2	3.4	1.7
NHH6	-0.1	-0.1	0.8	0.8	0.2	0.7	-0.1	3.0	0.4	4.7	1.2	5.0	1.2	0.5	5.2	1.3	3.3	1.2
NHH7	0.7	-0.1	1.2	0.8	0.2	1.5	1.3	5.0	0.4	8.6	1.2	8.5	1.5	1.5	8.8	1.9	2.6	1.2
NHH8	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.8	0.3	4.6	1.1	5.7	1.1	-0.1	5.7	0.5	1.9	2.6
NHH9	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	3.0	0.3	4.9	1.2	6.2	1.2	1.1	6.5	0.5	1.9	0.8
NI1	1.3	1.0	3.1	1.3	2.1	3.3	2.3	17.3	2.4	10.0	2.8	30.3	2.8	3.5	30.9	4.7	6.2	7.3
NI10	0.7	-0.1	0.9	0.9	0.2	1.3	-0.1	4.4	0.2	6.5	1.2	6.2	1.1	-0.1	6.4	-0.1	1.6	1.8
NI12	-0.1	-0.1	-0.1	0.8	0.2	-0.1	-0.1	2.3	0.3	5.6	1.2	4.6	1.1	-0.1	3.7	-0.1	1.5	1.8
NI3	-0.1	-0.1	-0.1	0.8	0.2	0.5	-0.1	2.7	0.3	5.2	1.1	5.4	-0.1	-0.1	5.2	-0.1	1.5	0.4
NI4	0.9	-0.7	1.8	0.4	0.9	2.7	1.2	6.8	1.9	9.3	1.7	9.6	1.3	1.8	10.0	2.2	2.8	3.4
NI5	0.8	-0.1	0.9	1.2	0.8	2.3	-0.1	3.5	1.7	5.6	1.4	5.4	1.0	1.5	5.6	1.9	2.3	0.5
NI5-R	0.7	-0.1	0.8	1.1	0.7	2.1	-0.1	3.0	1.5	5.1	1.4	4.9	1.0	1.4	4.9	1.7	2.1	0.4
NI6	-0.1	-0.1	0.9	0.8	0.2	0.6	-0.1	2.5	0.3	4.9	1.1	4.7	1.0	-0.1	5.0	-0.1	1.4	1.5
NI7	-0.1	1.1	3.5	1.4	1.3	2.2	2.3	21.8	1.6	4.1	2.0	39.4	2.8	2.3	36.0	3.0	4.1	4.9
NI8	-0.1	-0.1	-0.1	-0.1	0.2	0.6	-0.1	2.4	0.2	4.2	1.1	3.5	-0.1	-0.1	3.5	-0.1	1.2	0.3
NI9	0.8	-0.1	0.8	1.1	0.9	2.6	-0.1	4.1	1.7	6.0	1.4	6.4	1.0	1.4	6.5	1.7	2.0	2.2
NII1	-0.1	-0.1	-0.1	-0.1	0.1	0.6	-0.1	1.4	0.2	2.1	1.1	1.7	-0.1	-0.1	1.7	-0.1	-0.1	1.2
NII10	0.7	-0.1	1.0	0.9	0.3	1.5	-0.1	5.1	0.4	3.8	1.2	6.9	1.2	1.3	7.1	1.7	2.1	0.3
NII11	0.7	-0.1	0.2	1.0	0.6	1.6	-0.1	2.8	0.4	3.6	1.3	4.1	1.0	1.3	4.3	1.7	2.2	2.7
NII11-R	0.7	-0.1	0.1	1.0	0.6	1.6	-0.1	3.4	0.5	4.8	1.3	5.0	1.1	1.5	5.7	1.9	2.7	3.4
NII12	-0.1	-0.1	-0.1	0.2	0.6	2.1	-0.1	2.5	1.1	3.5	1.2	3.7	1.0	1.2	3.7	0.6	2.2	2.7
NII13	1.0	0.9	1.8	1.0	1.5	3.0	1.8	11.5	1.7	10.3	1.8	18.1	0.6	2.5	18.3	3.6	5.4	6.8
NII14	-0.1	-0.1	-0.1	0.7	0.1	0.5	-0.1	1.9	0.3	2.5	1.1	2.5	-0.1	-0.1	2.6	-0.1	1.4	1.6
NII2	0.7	-0.1	0.9	0.3	0.6	2.3	-0.1	4.4	1.2	9.4	1.4	7.3	1.1	1.2	7.3	1.6	2.1	2.6
NII3	0.8	0.9	3.2	1.1	0.7	1.6	1.7	18.7	0.4	14.0	1.5	20.5	1.9	1.4	20.5	1.7	1.9	2.2
NII4	0.8	1.0	1.7	1.1	0.6	1.6	1.9	32.7	0.3	38.1	1.4	40.2	2.2	1.4	41.1	1.7	2.1	2.6
NII5	-0.1	-0.1	0.1	0.8	0.2	0.8	-0.1	3.9	0.4	5.8	1.1	5.9	1.1	-0.1	6.3	-0.1	1.6	1.8
NII6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.6	0.2	3.9	1.1	3.5	-0.1	-0.1	3.7	-0.1	1.3	1.5
NII7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.0	-0.1	3.5	-0.1	3.2	-0.1	-0.1	3.3	-0.1	-0.1	1.1
NII8	0.8	0.8	0.2	1.2	0.7	1.6	1.7	16.0	0.4	19.0	1.5	18.2	1.9	1.4	19.2	2.0	2.6	0.9
NII9	-0.1	-0.1	0.1	0.7	0.1	-0.1	-0.1	3.1	1.1	4.4	1.1	3.9	1.0	-0.1	4.0	-0.1	1.2	0.3
NJ1	0.8	-0.1	0.3	0.3	0.6	2.5	-0.1	9.1	1.6	7.0	1.3	6.8	1.1	1.3	6.6	1.5	2.3	2.8
NJ2	0.7	-0.1	1.0	1.1	0.7	1.9	-0.1	4.2	1.1	5.4	1.3	7.7	1.2	1.4	7.9	1.8	2.4	2.9
NJ3	0.7	-0.1	1.4	1.1	0.8	2.2	-0.1	4.7	1.3	6.4	1.3	7.8	1.2	1.3	8.2	1.8	2.3	2.6
NJ4	0.7	-0.1	0.8	0.3	0.5	2.0	-0.1	4.1	1.3	6.9	1.3	7.6	1.1	1.2	7.6	1.6	1.9	2.2
NJ5	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	2.3	0.4	3.0	1.1	3.0	1.0	-0.1	3.5	-0.1	1.4	1.5
NJ6	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	2.8	0.4	5.3	1.2	5.1	1.0	-0.1	4.9	-0.1	1.4	1.6
NJ7	0.8	-0.1	1.0	0.4	0.8	1.9	-0.1	3.6	1.6	5.4	1.3	5.5	1.1	1.4	5.4	1.9	2.3	2.8
NJJ1	0.7	-0.1	0.8	0.2	0.2	1.6	-0.1	2.7	0.4	4.0	1.2	4.1	-0.1	1.1	4.4	1.5	1.7	0.6
NJJ10	1.8	1.2	4.7	4.0	5.4	15.9	3.9	22.9	8.1	11.5	3.8	26.0	4.8	9.6	26.5	15.2	32.4	42.9
NJJ11	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.1	0.2	3.0	1.1	3.1	-0.1	-0.1	3.2	-0.1	1.2	1.4
NJJ12	0.7	-0.1	-0.1	1.0	0.7	2.7	-0.1	3.0	1.5	4.4	1.3	4.2	1.1	1.6	4.2	2.2	3.1	0.5
NJJ13	0.8	0.8	2.2	1.2	0.8	1.8	2.0	13.7	0.6	9.0	1.5	21.3	2.4	2.0	22.2	3.0	4.1	3.3
NJJ14	-0.1	-0.1	-0.1	-0.1	-0.1	0.5	-0.1	2.0	0.2	2.9	1.1	2.7	-0.1	-0.1	2.9	-0.1	1.3	0.4
NJJ2	0.7	-0.1	0.2	1.0	0.3	1.2	-0.1	4.1	1.4	6.1	1.2	5.8	1.1	1.2	5.9	1.5	1.6	2.1
NJJ3	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	-0.1	2.8	0.2	4.3	1.1	3.8	-0.1	-0.1	3.9	-0.1	1.3	0.4
NJJ4	0.7	0.9	2.5	0.9	0.2	1.3	2.1	15.6	1.4	16.0	1.2	26.6	2.5	1.2	27.4	1.6	2.0	2.4
NJJ5	0.7	-0.1	0.7	1.0	0.6	0.5	-0.1	2.5	0.2	3.8	1.2	3.8	1.0	-0.1	4.1	-0.1	1.7	2.0
NJJ6	0.7	-0.1	0.1	0.9	0.2	0.6	-0.1	3.7	0.2	5.6	1.2	5.7	1.1	-0.1	5.9	-0.1	1.6	1.9
NJJ7	0.7	-0.1	-0.1	0.8	0.2	0.5	-0.1	1.7	0.2	2.7	1.1	2.5	-0.1	-0.1	2.7	-0.1	1.4	1.6
NJJ8	-0.1	-0.1	0.7	0.8	0.2	-0.1	-0.1	2.5	0.2	3.7	1.1	3.6	1.0	-0.1	3.9	-0.1	1.3	1.5
NJJ8-R	0.7	-0.1	0.9	0.8	0.2	-0.1	-0.1	3.3	0.2	4.4	1.2	4.5	1.2	-0.1	4.6	-0.1	1.6	1.8
NJJ9	1.0	1.1	3.3	0.8	0.9	2.3	2.4	23.6	1.4	11.9	1.9	48.6	3.2	1.9	49.5	2.5	3.3	4.1
NK1	0.7	-0.1	0.2	0.2	0.5	1.9	-0.1	3.4	1.1	6.2	1.3	6.1	1.0	1.3	5.9	1.8	2.1	2.6
NK2	0.7	-0.1	-0.1	0.8	0.1	-0.1	-0.1	2.5	0.2	3.3	1.2	3.2	-0.1	-0.1	3.3	-0.1	1.5	0.4
NK3	0.7	-0.1	0.2	0.9	0.6	1.7	-0.1	3.0	1.0	5.5	1.2	5.3	1.0	1.2	5.4	1.7	1.8	2.1
NK4	0.7	-0.1	1.1	0.9	0.2	1.3	-0.1	3.7	0.6	4.7	1.3	5.1	1.0	1.2	5.2	1.4	1.3	0.2
NK4-R	0.7	-0.1	0.8	0.8	0.2	-0.1	-0.1	3.0	0.2	3.8	1.2	4.0	-0.1	-0.1	4.1	-0.1	1.2	0.3
NK5	-0.1	-0.1	1.0	1.0	0.7	2.4	-0.1	0.6	1.4	4.5	1.3	4.6	1.0	1.2	4.7	1.6	2.1	2.5

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055-LPB	056-LBI	057-ALK	058-LPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NK6	0.7	-0.1	0.7	0.9	0.2	1.7	-0.1	3.3	1.1	5.3	1.2	5.0	1.0	-0.1	5.1	-0.1	1.7	0.4
NK7	-0.1	-0.1	0.2	0.9	0.2	0.8	-0.1	2.9	0.4	3.9	1.2	4.5	1.0	-0.1	4.6	-0.1	1.7	1.9
NKK1	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.6	1.0	2.3	-0.1	2.3	1.0	-0.1	2.5	-0.1	1.4	1.8
NKK10	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.9	0.2	6.1	-0.1	5.6	1.0	-0.1	5.6	-0.1	1.5	0.5
NKK11	0.7	-0.1	1.0	1.0	0.6	1.6	1.2	6.5	0.4	4.6	1.6	5.1	1.3	1.6	5.3	2.2	3.2	1.5
NKK2	-0.1	-0.1	-0.1	0.8	0.2	-0.1	-0.1	2.5	0.3	3.6	1.2	3.7	1.1	-0.1	3.7	0.5	1.9	0.7
NKK3	0.7	-0.1	1.6	0.9	0.2	1.2	1.6	5.3	1.3	7.3	1.7	7.7	1.7	1.7	7.9	2.3	3.0	1.9
NKK4	0.9	1.5	5.6	1.1	0.7	1.6	6.4	36.6	1.6	41.1	1.8	66.3	8.3	3.9	67.5	5.9	10.5	9.5
NKK5	0.8	1.3	5.3	1.1	0.7	1.5	4.5	26.9	0.2	32.1	1.7	43.8	5.9	2.9	45.0	4.6	9.0	1.2
NKK6	0.7	-0.1	1.3	0.8	0.2	1.2	1.6	4.8	1.6	7.1	1.3	7.0	1.8	1.7	7.2	2.3	3.3	2.4
NKK7	-0.1	-0.1	1.0	0.7	-0.1	-0.1	1.4	3.2	1.1	4.7	1.2	5.6	1.5	0.7	5.7	1.3	3.3	1.2
NKK8	-0.1	-0.1	-0.1	0.8	0.2	1.6	-0.1	2.8	0.5	5.6	1.2	5.5	1.1	1.2	5.3	0.5	2.0	2.6
NKK9	-0.1	-0.1	-0.1	0.8	0.2	0.8	-0.1	4.0	0.5	7.5	1.2	6.9	1.1	0.5	7.1	0.9	2.5	1.1
NL1	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.9	0.2	3.3	-0.1	3.0	-0.1	-0.1	3.1	-0.1	1.7	0.6
NL2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.3	0.2	4.0	1.1	3.6	-0.1	-0.1	3.8	0.5	1.8	0.7
NL3	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.7	0.2	2.6	1.1	2.5	-0.1	-0.1	2.6	0.7	2.0	2.8
NL3-R	-0.1	-0.1	0.7	0.8	0.2	0.6	-0.1	2.8	0.4	4.8	1.2	4.7	1.0	1.5	4.8	2.0	2.7	4.0
NL4	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.1	0.2	3.4	-0.1	3.6	-0.1	-0.1	3.7	-0.1	1.4	1.8
NLL1	0.7	-0.1	0.4	0.9	0.2	-0.1	0.1	2.3	1.3	3.4	1.1	3.4	1.1	1.6	3.6	2.3	3.1	4.7
NLL10	-0.1	-0.1	-0.1	0.7	0.1	1.2	-0.1	3.1	0.2	4.2	1.1	4.7	1.1	-0.1	4.8	-0.1	1.6	2.1
NLL11	0.7	-0.1	0.7	0.9	0.2	1.6	-0.1	3.1	0.5	5.1	1.2	5.4	1.1	1.3	5.6	1.7	2.0	0.8
NLL12	-0.1	-0.1	-0.1	0.8	0.2	1.0	-0.1	2.1	1.3	1.3	1.2	3.9	1.0	-0.1	3.8	-0.1	1.8	0.7
NLL2	-0.1	-0.1	0.6	0.7	-0.1	-0.1	-0.1	3.1	1.1	4.6	1.1	4.7	1.1	-0.1	4.8	-0.1	1.4	1.8
NLL3	-0.1	-0.1	0.7	0.7	0.1	-0.1	-0.1	2.2	0.3	2.9	-0.1	2.9	-0.1	-0.1	2.8	-0.1	-0.1	1.7
NLL4	0.8	-0.1	2.0	1.2	0.7	1.5	1.4	6.5	0.5	8.6	1.6	9.7	-0.3	2.0	10.0	2.6	3.3	3.0
NLL5	0.9	1.1	3.8	1.0	0.2	1.3	3.5	24.0	1.4	13.7	1.4	37.8	4.4	2.0	38.7	2.6	3.5	2.4
NLL6	0.8	1.0	3.4	1.0	0.2	1.5	2.7	19.9	0.3	22.6	1.3	34.2	3.5	1.8	34.8	2.5	3.8	0.9
NLL7	1.2	1.7	5.2	1.2	1.6	2.6	6.5	41.4	1.6	42.6	2.6	78.0	8.7	4.4	79.5	8.2	15.1	22.4
NLL8	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	2.4	0.3	3.8	1.4	4.1	1.0	-0.1	4.1	-0.1	1.6	0.6
NLL9	0.9	0.1	2.2	0.3	0.8	1.7	2.3	8.6	0.5	13.2	1.9	12.4	3.2	0.3	11.9	4.2	8.4	13.1
NLL9-R	0.9	1.1	0.8	1.4	1.0	1.8	2.9	10.1	0.5	6.5	1.8	14.1	3.6	3.4	14.5	5.2	9.6	13.9
NM1	0.8	-0.1	1.2	1.1	0.8	1.9	1.3	4.5	1.1	7.0	1.4	6.1	1.4	1.6	6.3	2.1	2.9	0.4
NM2	-0.1	-0.1	-0.1	0.8	0.2	1.0	-0.1	2.5	0.5	4.8	1.1	4.7	1.0	-0.1	4.7	-0.1	1.8	0.3
NMM1	-0.1	-0.1	-0.1	0.9	0.2	1.0	-0.1	2.3	0.5	3.2	1.2	3.0	-0.1	-0.1	3.2	-0.1	1.3	1.4
NMM10	0.9	1.0	2.8	1.5	1.1	1.8	2.0	10.9	1.2	8.3	1.7	11.8	2.3	1.8	12.1	2.5	3.2	0.7
NMM2	0.8	-0.1	0.2	1.0	0.2	1.7	1.1	6.7	0.5	4.1	1.3	9.7	0.3	1.2	9.8	1.4	1.5	1.6
NMM3	-0.1	-0.1	0.8	0.9	0.2	0.6	-0.1	3.0	0.4	5.1	1.2	4.4	1.0	-0.1	4.6	-0.1	1.3	1.5
NMM3-R	-0.1	-0.1	0.6	0.8	0.2	0.6	-0.1	2.6	0.3	4.3	1.1	3.8	-0.1	-0.1	4.0	-0.1	1.3	1.4
NMM4	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	1.9	0.3	3.5	1.1	3.1	-0.1	-0.1	3.2	-0.1	-0.1	0.3
NMM5	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.0	0.2	3.0	1.1	2.9	-0.1	-0.1	3.0	-0.1	-0.1	1.3
NMM6	0.8	-0.1	0.2	0.3	0.8	1.7	1.3	6.7	0.5	5.5	1.5	10.1	1.4	1.5	10.4	2.0	2.7	3.5
NMM7	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.6	-0.1	2.3	-0.1	2.2	-0.1	-0.1	2.3	-0.1	-0.1	0.2
NMM8	-0.1	1.1	2.8	0.9	0.2	1.2	2.4	11.7	0.2	9.6	1.3	13.6	2.7	1.3	14.6	1.7	2.2	2.6
NMM9	0.8	0.7	-0.1	0.3	0.6	2.0	1.2	4.7	1.1	6.7	1.5	9.0	1.4	1.4	8.7	2.0	2.8	3.6
NN1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	1.4	0.2	2.1	-0.1	1.9	-0.1	-0.1	2.0	-0.1	-0.1	0.3
NNN1	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	3.2	1.1	6.6	-0.1	5.2	0.9	-0.1	5.4	-0.1	1.3	1.4
NNN10	0.7	-0.1	0.4	1.0	0.8	2.4	-0.1	2.4	1.4	0.5	1.1	2.6	1.0	-0.1	2.7	-0.1	1.7	1.9
NNN2	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	3.0	0.2	5.3	1.1	5.0	1.0	-0.1	5.3	-0.1	1.3	1.4
NNN3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.4	-0.1	1.9	-0.1	1.7	-0.1	-0.1	1.8	-0.1	-0.1	0.1
NNN4	0.8	-0.1	1.2	1.2	0.7	1.9	1.2	5.8	1.4	8.3	1.5	10.3	0.4	1.6	10.5	2.1	2.7	0.8
NNN5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.9	1.1	6.2	-0.1	4.7	-0.1	-0.1	4.7	-0.1	-0.1	0.3
NNN6	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.6	-0.1	2.1	-0.1	2.2	-0.1	-0.1	2.3	-0.1	-0.1	1.0
NNN7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.3	-0.1	1.5	-0.1	1.6	-0.1	-0.1	1.7	-0.1	-0.1	0.1
NNN8	0.9	0.8	0.3	1.4	1.1	2.6	1.3	5.9	1.7	4.1	1.6	12.1	0.4	1.6	12.3	2.0	2.7	3.1
NNN9	0.9	-0.1	1.3	1.2	0.8	1.6	1.2	5.3	0.4	6.8	1.3	6.8	1.4	1.3	6.9	1.7	2.1	2.4
NO1	-0.1	-0.1	0.7	0.8	0.1	-0.1	-0.1	2.8	0.2	3.9	1.1	4.4	1.2	0.6	4.5	2.0	2.9	1.0
NO2	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.1	0.2	1.3	-0.1	1.1	-0.1	-0.1	1.2	-0.1	-0.1	0.4
NOO1	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	2.3	1.1	3.1	-0.1	3.3	-0.1	-0.1	3.4	-0.1	-0.1	0.3
NOO2	-0.1	-0.1	0.9	0.8	0.2	-0.1	-0.1	3.5	0.2	5.0	1.2	5.0	1.1	-0.1	5.1	-0.1	1.3	0.5
NOO3	-0.1	-0.1	-0.1	0.7	0.1	0.7	-0.1	1.8	0.3	2.0	1.1	2.0	-0.1	-0.1	2.1	-0.1	1.3	0.4
NOO4	-0.1	-0.1	-0.1	0.9	0.2	1.3	-0.1	2.8	0.2	3.8	1.2	4.5	1.1	1.2	4.6	1.5	1.7	0.6
NOO5	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.0	1.0	2.7	-0.1	2.7	-0.1	-0.1	2.9	-0.1	-0.1	0.3
NOO6	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.6	-0.1	2.1	-0.1	2.3	-0.1	-0.1	2.3	-0.1	-0.1	0.3
NOO6-R	-0.1	-0.1	-0.1	0.7	0.7	-0.1	-0.1	1.6	1.0	2.2	-0.1	2.2	-0.1	-0.1	2.2	-0.1	-0.1	0.2
NOO7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.9	-0.1	2.9	-0.1	2.8	-0.1	-0.1	2.9	-0.1	-0.1	0.3
NOO8	0.8	0.9	2.4	1.1	0.8	1.5	1.9	11.7	0.2	17.5	1.5	23.6	2.3	1.8	24.1	2.4	3.4	4.2
NP1	-0.1	-0.1	-0.1	0.8	0.1	0.8	-0.1	2.7	0.2	4.7	1.1	3.9	1.0	-0.1	4.2	0.8	2.4	0.7

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

	055-LPB	056-LBI	057-ALK	058-LPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NP2	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.1	0.2	3.8	-0.1	3.3	-0.1	-0.1	3.3	0.5	1.7	0.7
NP3	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	2.0	0.2	3.3	-0.1	2.8	-0.1	-0.1	3.0	-0.1	1.6	0.6
NP4	1.1	0.2	2.2	0.4	1.1	2.7	2.2	9.0	1.8	8.2	3.0	24.2	3.0	3.8	23.7	10.5	25.6	39.6
NPP4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.6	-0.1	1.9	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	0.2
NPP1-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.5	-0.1	1.8	-0.1	2.0	-0.1	-0.1	2.1	-0.1	-0.1	0.2
NPP2	0.7	-0.1	1.2	1.0	0.3	1.6	1.3	6.5	0.5	11.1	1.4	10.3	1.4	1.5	9.8	2.2	2.9	2.1
NPP3	-0.1	-0.1	0.8	0.8	0.2	1.3	-0.1	4.6	1.2	8.2	1.2	9.4	1.4	1.2	9.0	0.6	2.1	2.9
NPP4	0.8	-0.1	0.8	1.2	-0.8	2.2	-0.1	4.8	1.6	8.6	1.4	8.4	1.3	1.8	8.4	2.4	3.4	1.5
NQ1	0.7	-0.1	-0.1	0.9	0.3	2.0	-0.1	1.8	1.2	2.8	1.3	2.6	1.0	1.6	2.6	2.1	2.9	4.3
NQ2	-0.1	-0.1	0.1	0.8	-0.2	-0.1	1.3	4.2	1.1	6.0	-0.1	5.6	-1.4	1.1	5.8	2.1	2.9	4.3
NQ2-R	0.7	-0.1	0.9	0.8	0.2	1.2	1.3	4.7	0.2	6.5	1.2	9.0	1.4	1.6	9.2	2.2	3.2	2.4
NQ3	-0.1	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	3.1	1.1	5.0	-0.1	4.8	1.0	-0.1	4.9	-0.1	1.3	0.2
NQ4	-0.1	-0.1	-0.1	0.8	0.2	0.7	-0.1	3.3	0.2	4.7	1.2	6.6	0.3	0.6	6.8	2.3	3.2	0.5
NQ5	0.7	-0.1	-0.1	1.0	-0.7	1.3	-0.1	2.3	1.2	3.4	1.3	3.7	1.1	1.8	3.8	2.5	3.9	0.8
NQ6	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	2.3	1.1	4.3	-0.1	3.7	-0.1	-0.1	3.8	-0.1	-0.1	0.4
NQ7	0.8	-0.1	0.1	1.1	-0.8	2.8	1.3	5.1	1.7	7.3	1.7	9.8	0.4	2.1	10.0	3.3	4.5	1.4
NQ8	-0.1	-0.1	-0.1	0.8	0.2	1.1	-0.1	2.8	1.0	4.5	1.1	4.6	1.1	-0.1	4.7	0.8	2.4	0.6
NQQ1	-0.1	-0.1	-0.1	0.8	-0.2	0.8	-0.1	2.4	0.5	4.5	1.1	3.8	-0.1	-0.1	3.9	-0.1	1.5	0.4
NR1	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.0	-0.1	2.7	-0.1	2.8	-0.1	-0.1	2.9	-0.1	-0.1	1.0
NR11	0.7	-0.1	1.1	1.0	-0.2	1.6	-0.1	3.8	0.4	5.0	1.2	5.2	1.1	1.1	5.3	1.3	1.3	1.6
NR12	0.7	-0.1	0.7	1.0	0.6	1.5	-0.1	3.1	0.4	4.9	1.2	5.4	1.0	1.1	5.6	-0.1	1.5	0.5
NR13	0.8	-0.1	1.9	0.3	0.7	1.9	1.5	9.3	0.5	4.6	1.3	10.1	1.8	1.4	16.7	2.0	2.8	3.4
NR14	-0.1	-0.1	0.2	0.7	0.1	-0.1	-0.1	3.0	1.1	4.4	1.1	4.1	1.0	-0.1	4.3	-0.1	-0.1	0.2
NR2	-0.1	-0.1	1.0	0.8	-0.2	-0.1	-0.1	3.4	0.2	5.4	1.2	5.0	1.0	-0.1	5.2	-0.1	1.3	0.3
NR3	0.8	-0.1	0.2	1.2	0.8	1.4	1.1	4.9	0.4	3.4	1.3	5.4	1.2	1.3	5.7	1.7	1.9	0.7
NR4	-0.1	-0.1	-0.1	-0.1	-0.1	0.6	-0.1	1.8	0.2	2.8	-0.1	2.5	-0.1	-0.1	2.6	-0.1	-0.1	0.3
NR5	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.8	1.1	2.7	-0.1	2.6	-0.1	-0.1	2.7	-0.1	1.2	1.3
NR6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.7	0.2	2.5	-0.1	2.4	-0.1	-0.1	2.5	-0.1	-0.1	0.3
NR7	0.8	-0.1	2.0	1.1	0.3	1.5	1.3	7.0	0.5	10.0	1.4	8.8	1.3	1.4	9.4	1.8	2.0	1.5
NR8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.1	1.1	3.0	-0.1	2.8	-0.1	-0.1	3.0	-0.1	-0.1	1.2
NS1	-0.1	-0.1	0.7	0.7	0.2	-0.1	-0.1	2.5	1.2	3.5	1.2	3.5	1.0	-0.1	0.8	0.5	1.7	1.0
NS10	0.7	-0.1	1.1	1.1	-0.8	2.7	1.3	5.1	1.6	7.3	1.4	12.9	1.6	1.7	13.1	3.9	6.0	9.4
NS11	0.7	-0.1	0.2	1.0	0.2	1.3	-0.1	3.9	1.5	5.9	1.3	5.5	1.0	1.3	5.7	1.6	1.9	2.5
NS12	0.7	-0.1	1.0	1.0	-0.7	2.1	1.2	4.0	1.2	5.9	1.4	8.5	1.3	1.5	8.5	2.1	3.3	4.5
NS13	0.8	-0.1	1.5	1.2	0.8	2.5	1.5	6.5	1.9	8.1	1.5	8.2	1.6	2.0	8.7	3.0	4.2	3.2
NS14	1.0	-0.9	2.3	1.2	1.1	2.8	1.7	6.2	3.6	11.5	2.4	13.8	2.1	3.4	14.2	5.9	9.3	7.2
NS2	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	3.0	1.1	5.0	1.1	6.2	1.2	0.4	6.4	0.8	2.4	1.1
NS3	0.7	-0.1	3.7	1.1	-0.8	1.4	1.3	3.8	0.3	6.2	1.5	7.9	1.6	2.3	7.9	9.3	8.6	6.2
NS4	-0.1	-0.1	1.6	0.7	0.2	1.2	1.0	5.3	0.3	7.2	1.2	8.0	1.0	1.1	8.3	1.5	1.8	2.5
NS4-R	-0.1	-0.1	1.2	0.7	-0.2	-0.1	-0.1	4.0	0.3	5.3	1.2	6.0	1.0	1.2	6.2	0.4	1.7	2.4
NS5	-0.1	-0.1	1.6	0.7	0.2	1.1	1.1	4.5	0.3	6.8	1.2	7.3	1.2	1.0	7.6	-0.1	1.4	1.8
NS6	-0.1	-0.1	0.4	0.7	-0.1	-0.1	-0.1	2.6	0.2	5.0	1.1	4.7	1.0	-0.1	4.8	0.5	2.0	2.7
NS7	-0.1	-0.1	1.8	0.8	0.2	1.4	1.5	7.3	0.3	9.5	1.2	10.1	1.7	1.4	10.4	1.0	2.9	1.2
NS8	0.8	-0.1	0.2	1.2	-0.8	2.1	1.0	5.3	1.5	8.9	1.6	9.2	1.8	2.4	9.5	3.7	6.2	4.9
NS9	-0.1	-0.1	0.7	-0.1	0.2	-0.1	-0.1	2.1	0.3	3.2	-0.1	3.2	-0.1	-0.1	3.3	-0.1	0.9	0.7
NT1	-0.1	-0.1	0.4	0.8	-0.2	0.8	-0.1	2.0	0.5	3.1	-0.1	3.0	-0.1	-0.1	3.1	0.4	1.9	2.3
NT10	-0.1	-0.1	-0.1	0.8	0.2	-0.1	-0.1	2.2	0.2	3.9	1.2	3.2	1.0	-0.1	3.4	0.8	2.2	3.2
NT11	-0.1	-0.1	0.7	-0.1	-0.2	-0.1	-0.1	2.0	0.2	2.4	1.4	2.7	-0.1	-0.1	2.8	-0.1	-0.1	1.8
NT12	0.7	-0.1	-0.1	0.9	0.2	1.4	-0.1	2.6	1.2	5.1	1.2	4.1	1.1	1.5	4.3	2.1	2.7	1.3
NT13	-0.1	-0.1	0.4	0.7	-0.1	-0.1	-0.1	2.8	0.2	3.5	1.1	3.6	1.0	-0.1	3.7	-0.1	1.7	0.6
NT14	-0.1	-0.1	-0.1	0.8	0.2	1.5	-0.1	2.4	0.6	4.6	1.2	4.7	1.0	1.1	4.6	-0.1	1.8	2.3
NT15	-0.1	-0.1	1.0	0.4	-0.2	1.6	1.2	4.1	0.4	5.9	1.4	7.7	1.3	1.6	8.0	2.6	3.6	1.7
NT16	0.8	-0.1	1.0	1.1	0.7	1.9	1.1	3.8	1.2	6.1	1.4	6.8	1.2	1.5	6.5	2.0	3.0	1.3
NT17	-0.1	-0.1	0.4	0.8	-0.2	0.6	-0.1	2.8	0.4	4.0	1.1	4.6	1.1	-0.1	4.6	-0.1	1.7	0.6
NT18	-0.1	-0.1	-0.1	-0.1	0.8	0.6	-0.1	1.5	0.2	1.9	-0.1	1.9	-0.1	-0.1	1.9	-0.1	-0.1	1.2
NT2	-0.1	-0.1	-0.1	0.8	0.2	1.1	-0.1	2.1	0.9	0.8	1.2	3.4	1.0	-0.1	3.5	1.0	3.0	3.9
NT3	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	1.9	0.2	3.1	-0.1	3.6	-0.1	-0.1	3.7	-0.1	-0.1	1.3
NT4	-0.1	-0.1	0.4	-0.1	-0.1	-0.1	-0.1	2.3	1.1	3.6	1.1	3.7	-0.1	-0.1	3.8	-0.1	-0.1	0.2
NT5	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	1.8	1.1	2.6	1.1	2.6	-0.1	-0.1	2.7	-0.1	1.4	1.6
NT6-R	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	-0.1	3.0	0.2	5.8	1.4	4.7	0.9	-0.1	5.0	-0.1	1.7	0.6
NT6	-0.1	-0.1	0.7	0.7	0.1	-0.1	-0.1	2.3	1.1	3.5	-0.1	3.6	-0.1	-0.1	3.7	-0.1	1.2	1.4
NT7	0.7	-0.1	1.4	0.9	-0.2	1.2	1.1	3.6	0.2	5.9	1.2	6.4	1.2	1.4	6.5	4.7	2.2	0.9
NT8	0.8	0.8	2.0	1.1	0.6	1.6	1.6	8.3	0.4	4.7	1.5	12.9	2.0	1.5	13.7	2.4	3.8	1.8
NT9	-0.1	-0.1	-0.1	0.8	-0.2	-0.1	-0.1	2.1	1.1	3.0	1.1	3.0	-0.1	-0.1	3.1	-0.1	1.4	1.7
NU1	-0.1	-0.1	0.7	0.8	0.2	-0.1	-0.1	2.3	1.2	2.9	1.1	3.1	-0.1	-0.1	3.3	-0.1	-0.1	0.2
NU10	0.7	-0.1	1.4	0.8	-0.2	1.5	1.4	5.6	0.4	4.1	1.3	12.6	0.5	2.1	12.6	3.5	5.9	8.0
NU11	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	2.2	0.3	0.5	1.1	2.6	1.0	-0.1	2.7	0.5	2.1	2.6

A14-06865

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Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055-LPB	056-LBI	057-ALK	058-LPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-LBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NU12	-0.1	-0.1	0.2	-0.1	0.2	-0.1	-0.1	3.0	0.2	5.2	-0.1	6.5	1.1	-0.1	6.2	-0.1	1.8	2.2
NU12-R	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.8	1.1	4.3	1.1	5.1	1.1	-0.1	5.2	-0.1	1.8	2.2
NU13	0.7	0.8	1.7	0.9	0.2	1.3	1.7	5.0	0.3	4.5	1.3	14.8	2.0	2.2	15.1	6.5	7.0	9.6
NU14	0.7	-0.1	0.8	0.8	0.2	1.7	-0.1	3.9	1.1	8.2	1.2	5.7	1.0	1.2	6.0	1.7	2.1	0.3
NU15	0.7	-0.1	0.9	0.9	0.2	1.5	1.2	6.7	0.3	5.2	1.2	7.1	1.3	1.4	7.5	2.1	3.2	4.3
NU16	0.8	0.8	0.3	0.3	0.8	1.9	1.8	11.3	1.2	7.4	1.7	21.4	2.2	2.1	21.8	3.2	5.3	7.4
NU17	0.7	-0.1	1.4	1.1	0.7	1.9	1.5	5.5	1.2	8.6	1.3	12.5	1.7	1.6	13.1	2.5	3.9	0.7
NU18	0.7	-0.1	0.8	0.8	0.2	1.1	-0.1	3.0	0.2	4.2	1.2	4.5	1.1	1.2	4.3	0.6	1.8	1.1
NU19	0.7	-0.1	0.8	0.3	0.3	2.2	-0.1	2.5	1.1	5.0	1.2	4.3	-0.1	1.1	4.4	0.6	2.3	3.0
NU2	0.9	0.2	2.7	0.4	0.7	1.6	3.0	15.7	0.6	19.6	1.7	20.2	4.1	1.7	20.9	3.5	6.3	9.2
NU3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.7	0.3	2.8	-0.1	2.9	-0.1	-0.1	3.0	-0.1	-0.1	1.8
NU4	0.9	1.0	3.8	1.2	0.7	1.7	2.7	19.9	0.5	2.8	1.9	27.3	3.5	2.7	28.1	5.3	10.7	16.0
NU5	-0.1	-0.1	0.7	0.1	-0.1	-0.1	-0.1	2.5	1.0	3.5	-0.1	3.7	-0.1	-0.1	3.8	-0.1	1.3	0.4
NU6	-0.1	-0.1	0.9	-0.1	0.2	-0.1	-0.1	3.0	0.2	3.9	-0.1	4.1	1.0	-0.1	4.3	-0.1	1.6	2.0
NU7	-0.1	-0.1	1.4	0.7	0.2	-0.1	-0.1	3.7	0.3	5.6	1.1	5.7	1.1	-0.1	5.9	-0.1	1.8	0.7
NU8	0.7	-0.1	1.3	0.9	0.3	1.7	1.2	5.5	1.0	8.3	1.3	10.1	1.4	1.4	10.1	2.0	2.8	1.0
NU9	0.7	-0.1	0.2	0.3	0.5	2.2	-0.1	3.8	1.3	2.0	1.8	6.2	1.2	1.7	6.2	2.6	4.8	6.5
NV1	0.8	-0.1	0.9	1.1	0.8	2.5	-0.1	4.4	1.5	8.1	1.4	5.8	1.2	1.3	5.6	1.6	1.9	2.1
NV10	-0.1	-0.1	-0.1	0.8	0.2	-0.1	0.1	2.6	0.2	4.0	-0.1	4.2	-0.1	-0.1	4.4	-0.1	1.4	0.3
NV11	-0.1	-0.1	-0.1	0.8	0.2	0.8	-0.1	2.1	0.5	2.9	1.1	3.0	-0.1	-0.1	3.1	-0.1	1.3	0.4
NV12	-0.1	-0.1	-0.1	0.8	0.2	1.0	-0.1	2.5	1.0	4.6	1.4	4.3	-0.1	-0.1	4.5	-0.1	1.4	1.6
NV13	0.7	-0.1	1.4	0.3	0.7	2.7	-0.1	4.2	1.6	7.5	1.3	8.6	1.2	1.3	8.6	1.6	2.0	2.5
NV14	0.7	-0.1	0.7	0.8	0.3	1.2	-0.1	2.9	1.5	5.5	1.2	5.2	-0.1	1.0	5.2	-0.1	0.1	1.9
NV15	-0.1	-0.1	-0.1	0.8	0.1	0.9	-0.1	2.4	0.4	3.7	1.1	3.8	-0.1	-0.1	4.0	-0.1	1.3	1.4
NV16	0.9	-0.1	-0.1	1.3	1.1	3.4	1.2	6.1	2.1	10.8	1.6	11.4	1.3	1.6	11.8	2.0	2.6	0.9
NV17	0.8	-0.1	0.7	0.3	0.7	2.2	-0.1	3.1	1.5	5.6	1.4	5.1	1.1	1.2	5.1	1.5	1.8	0.7
NV18	-0.1	-0.1	-0.1	0.9	0.3	0.7	-0.1	3.0	0.4	6.5	1.2	5.7	1.1	-0.1	5.8	-0.1	1.4	0.5
NV2	0.7	-0.1	0.8	0.9	0.7	1.4	-0.1	3.3	0.2	5.0	1.2	4.8	1.1	-0.1	4.9	1.6	1.9	0.6
NV3	-0.1	-0.1	-0.1	0.7	0.2	0.6	-0.1	2.3	0.3	3.8	1.4	4.1	-0.1	-0.1	4.0	-0.1	-0.1	1.4
NV4	0.7	-0.1	0.8	0.3	0.6	2.3	-0.1	3.8	1.3	8.9	1.3	6.9	1.1	1.2	6.9	1.6	2.0	2.5
NV5	0.7	-0.1	1.0	0.3	0.7	2.6	-0.1	5.8	1.8	12.5	1.3	10.9	1.2	1.5	10.9	1.9	2.6	0.9
NV5-R	0.8	-0.1	0.1	0.3	0.9	3.4	-0.1	4.7	2.2	7.8	1.4	9.2	1.2	1.5	9.2	1.8	2.3	2.8
NV6	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	1.9	0.3	2.4	1.4	2.5	-0.1	-0.1	2.6	0.4	1.8	2.2
NV7	-0.1	-0.1	-0.1	0.8	0.2	1.2	-0.1	0.3	-0.1	5.9	1.2	4.9	1.0	1.0	5.0	-0.1	-0.1	0.6
NV8	-0.1	-0.1	-0.1	0.8	0.2	0.5	-0.1	2.1	0.5	4.2	1.1	3.9	-0.1	-0.1	3.9	-0.1	1.5	1.8
NV9	-0.1	-0.1	0.7	0.8	0.2	0.7	-0.1	3.8	0.5	6.4	1.2	7.0	1.1	-0.1	7.1	-0.1	1.5	0.5
NW1	0.8	0.9	2.1	0.9	0.2	1.4	1.6	8.8	0.2	11.8	1.4	14.3	2.0	1.3	14.6	1.6	1.8	0.4
NW10	-0.1	-0.1	-0.1	0.7	0.2	0.5	-0.1	2.3	0.5	3.6	1.2	0.8	-0.1	1.0	3.4	-0.1	-0.1	0.4
NW11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.6	0.2	4.9	-0.1	4.3	-0.1	-0.1	4.4	-0.1	1.2	0.4
NW12	0.7	-0.1	-0.1	0.3	0.6	2.2	-0.1	0.4	1.4	4.4	1.3	4.8	1.1	1.3	4.8	1.6	2.0	0.7
NW13	0.8	-0.1	1.2	0.9	0.8	3.6	1.3	6.4	2.0	3.6	1.6	8.6	1.5	1.5	8.3	2.3	3.4	4.5
NW14	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.1	1.1	3.7	1.1	3.3	-0.1	-0.1	3.5	-0.1	1.2	1.4
NW15	0.8	0.8	1.4	0.4	1.0	3.2	1.3	5.0	1.9	7.2	1.6	7.0	1.5	1.6	7.3	2.1	2.9	3.7
NW16	-0.1	-0.1	-0.1	0.8	0.1	-0.1	-0.1	2.1	1.1	3.2	1.1	3.1	-0.1	-0.1	3.2	-0.1	1.6	0.6
NW17	0.7	-0.1	-0.1	0.7	0.3	0.6	-0.1	2.0	0.5	3.7	1.2	3.5	1.0	1.1	2.9	1.9	2.8	4.2
NW17-R	0.7	-0.1	-0.1	0.8	0.2	0.6	-0.1	2.0	0.5	3.3	1.2	3.3	1.0	1.1	3.2	1.9	2.7	4.1
NW18	-0.1	-0.1	-0.1	0.9	0.3	0.5	-0.1	2.6	0.3	0.5	1.2	3.5	1.1	-0.1	3.6	-0.1	1.5	0.4
NW2	0.7	-0.1	-0.1	0.8	0.3	0.5	-0.1	2.7	0.5	4.9	1.2	4.3	-0.1	-0.1	4.3	-0.1	1.4	1.7
NW2-R	-0.1	-0.1	-0.1	0.7	0.2	0.5	-0.1	2.5	0.5	4.1	1.1	3.3	-0.1	-0.1	3.4	-0.1	-0.1	1.6
NW3	0.6	-0.1	-0.1	0.7	0.2	0.5	-0.1	2.5	0.3	4.1	1.1	4.7	-0.1	-0.1	4.6	-0.1	-0.1	1.4
NW4	-0.1	-0.1	-0.1	0.7	0.2	0.5	-0.1	2.2	0.3	3.0	1.0	3.8	1.0	-0.1	3.7	-0.1	1.3	1.4
NW5	0.8	0.9	2.8	1.1	0.7	1.8	1.9	17.3	1.1	3.0	1.4	26.3	2.3	1.5	26.8	1.9	2.5	0.4
NW6	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	2.8	0.3	4.1	1.1	4.8	1.1	-0.1	5.0	-0.1	1.3	1.5
NW7	-0.1	-0.1	-0.1	0.7	0.2	0.4	-0.1	2.3	0.4	4.6	1.1	4.2	-0.1	-0.1	4.1	-0.1	-0.1	1.4
NW8	1.0	1.0	2.6	0.9	1.3	1.5	2.4	17.7	0.2	17.7	1.8	26.8	2.9	1.9	27.1	2.6	3.0	1.3
NW9	-0.1	-0.1	0.8	0.7	0.2	-0.1	-0.1	0.8	0.3	6.3	1.1	6.8	1.1	-0.1	6.5	-0.1	-0.1	1.4
NX1	-0.1	-0.1	-0.1	0.8	0.2	1.0	-0.1	1.8	1.0	2.8	1.1	2.5	-0.1	-0.1	2.7	-0.1	1.5	0.5
NX10	0.7	-0.1	0.8	0.8	0.2	1.6	1.1	3.0	0.6	5.2	1.3	5.0	1.2	1.4	5.0	2.1	3.4	4.7
NX11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.5	0.2	2.5	-0.1	2.3	-0.1	-0.1	2.3	-0.1	-0.1	1.5
NX12	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	1.7	0.3	2.7	-0.1	2.8	-0.1	-0.1	2.8	-0.1	-0.1	2.1
NX13	0.7	-0.1	0.9	1.0	0.6	2.1	-0.1	4.1	1.3	7.7	1.4	8.4	1.3	1.9	8.4	2.8	4.8	0.6
NX14	0.7	-0.1	0.1	0.9	0.2	1.5	-0.1	3.1	0.4	4.3	1.2	6.0	1.3	1.5	6.2	1.2	3.4	4.7
NX14-R	-0.1	-0.1	0.2	0.8	0.2	0.9	-0.1	2.8	0.5	4.3	1.2	5.0	1.1	1.3	5.9	0.9	2.9	4.0
NX15	-0.1	-0.1	0.7	0.7	0.2	1.4	-0.1	3.5	0.5	6.6	1.2	6.6	1.1	1.1	6.6	0.5	1.8	0.4
NX16	-0.1	-0.1	0.9	0.8	0.2	1.5	1.2	4.7	0.4	8.3	1.2	9.1	1.3	1.4	8.8	0.9	2.7	0.6
NX17	0.8	0.8	1.9	1.3	0.9	2.8	1.6	6.6	1.8	8.6	1.6	11.7	1.8	2.3	12.0	3.5	6.1	8.8
NX2	-0.1	-0.1	-0.1	0.7	0.2	0.6	-0.1	2.3	0.4	3.7	1.1	3.7	1.0	0.3	3.9	1.0	3.3	4.5

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	055-EPB	056-LBI	057-ALK	058-EPB	059-LPB	060-LPH	061-LBI	062-LBA	063-LPH	064-EBA	065-HPB	066-LBA	067-LBI	068-HPB	069-LA	070-HPB	071-HPB	072-HPB
NX3	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	2.2	1.2	4.1	1.1	3.7	1.0	-0.1	3.8	0.6	1.8	0.5
NX4	0.8	-0.1	1.6	1.1	0.6	1.7	1.7	7.1	1.4	8.5	1.6	8.7	1.9	2.4	9.0	3.8	5.5	4.0
NX5	-0.1	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	2.5	1.1	3.8	1.1	4.0	1.0	-0.1	4.2	-0.1	1.6	0.6
NX6	-0.1	-0.1	0.8	0.8	0.2	1.5	-0.1	3.5	0.4	4.8	1.2	6.7	1.2	-0.1	6.7	0.9	2.8	3.8
NX7	1.0	0.3	3.5	0.8	1.1	3.5	4.2	8.3	2.0	32.7	2.3	72.3	6.1	2.8	72.9	5.7	11.5	17.6
NX8	0.7	-0.1	0.1	1.0	0.5	2.2	1.2	4.2	1.4	6.6	1.3	7.4	1.3	1.4	7.5	2.2	3.5	4.8
NX9	0.8	-0.1	1.4	1.0	0.7	1.9	1.6	5.9	1.2	9.5	1.5	13.3	1.9	2.1	13.3	3.4	6.2	9.0
NY1	-0.1	-0.1	-0.1	0.7	0.2	0.4	-0.1	2.1	0.5	4.4	1.1	3.7	-0.1	-0.1	3.7	-0.1	-0.1	1.4
NY10	0.7	-0.1	1.2	0.3	0.9	2.8	-0.1	5.3	1.8	9.7	1.4	11.5	1.3	1.4	11.5	1.8	2.5	3.1
NY11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.0	0.2	4.0	-0.1	3.7	-0.1	-0.1	3.9	-0.1	1.1	0.3
NY12	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	2.0	0.2	2.7	1.1	2.7	1.0	-0.1	2.9	-0.1	1.5	0.4
NY13	-0.1	-0.1	-0.1	0.7	0.1	-0.1	-0.1	1.7	0.2	2.6	-0.1	2.4	-0.1	-0.1	2.5	-0.1	1.2	1.3
NY14	0.8	-0.1	1.0	0.7	0.9	2.1	1.2	4.2	1.4	6.2	1.5	5.8	1.2	1.8	6.0	2.4	3.4	4.3
NY15	0.8	-0.1	0.8	1.1	0.4	1.3	1.1	0.3	0.6	6.9	1.4	8.7	1.2	1.3	7.9	1.8	1.9	1.3
NY2	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	1.5	0.3	2.3	1.1	2.2	-0.1	-0.1	2.3	-0.1	1.2	1.3
NY3	-0.1	-0.1	-0.1	-0.1	0.7	-0.1	-0.1	1.8	-0.1	2.8	0.2	2.0	-0.1	-0.1	2.7	-0.1	-0.1	0.3
NY4	1.0	1.0	2.4	1.0	1.5	1.7	2.1	13.6	0.7	13.6	1.8	19.1	2.4	2.1	19.4	2.7	3.4	0.5
NY4-R	1.2	1.1	2.7	1.3	1.8	2.1	2.6	20.2	1.5	10.2	2.1	31.8	3.1	2.8	32.1	3.7	5.0	0.7
NY5	-0.1	-0.1	-0.1	0.7	0.2	-0.1	-0.1	1.9	0.4	2.8	1.1	3.2	-0.1	-0.1	3.4	-0.1	1.4	1.5
NY6	0.7	-0.1	0.8	0.3	0.5	1.7	-0.1	3.1	1.1	4.6	1.5	5.0	1.1	1.2	5.3	1.6	2.0	0.5
NY7	0.8	-0.1	1.1	0.3	0.7	2.2	1.2	5.0	1.6	8.9	1.5	8.2	1.2	1.6	7.8	2.1	2.9	0.6
NY8	-0.1	-0.1	-0.1	0.7	0.2	1.3	-0.1	2.3	0.4	3.7	1.1	4.1	-0.1	-0.1	4.1	-0.1	1.3	1.5
NY9	-0.1	-0.1	-0.1	0.9	0.6	1.8	-0.1	1.8	1.2	0.5	1.1	2.0	-0.1	-0.1	2.0	-0.1	1.5	1.6
NZ1	-0.1	-0.1	-0.1	0.8	0.2	0.5	-0.1	2.5	0.2	4.7	1.1	4.1	1.0	-0.1	4.1	-0.1	1.7	0.5
NZ10	1.1	0.8	2.2	1.4	2.2	8.2	1.2	6.4	4.3	2.8	2.3	11.4	1.4	2.1	12.0	2.9	4.0	4.6
NZ11	-0.1	-0.1	-0.1	0.8	0.2	0.6	-0.1	1.9	0.6	3.1	1.2	3.1	-0.1	-0.1	2.7	-0.1	-0.1	1.5
NZ12	-0.1	-0.1	1.0	0.3	0.6	1.8	-0.1	2.6	1.0	3.8	1.2	4.4	1.1	-0.1	4.4	-0.1	1.5	1.7
NZ13	-0.1	-0.1	-0.1	-0.1	0.2	-0.1	-0.1	1.7	0.3	2.7	1.2	2.0	-0.1	-0.1	2.8	-0.1	-0.1	1.3
NZ2	0.7	-0.1	0.9	0.8	0.2	-0.1	1.2	3.6	0.2	5.0	1.2	5.6	1.2	-0.1	5.8	-0.1	1.7	0.3
NZ2-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.6	1.2	2.3	1.1	2.4	-0.1	-0.1	2.5	-0.1	1.3	1.6
NZ3	-0.1	-0.1	0.7	0.8	0.2	1.3	-0.1	2.6	0.4	3.8	1.2	4.6	1.1	1.2	4.7	1.6	1.8	0.4
NZ4	-0.1	-0.1	-0.1	0.8	0.2	0.5	-0.1	2.6	0.2	3.8	1.2	3.9	1.1	-0.1	4.0	-0.1	1.7	0.5
NZ5	0.7	-0.1	-0.1	0.9	0.2	1.7	-0.1	2.7	0.5	4.3	1.2	3.9	1.0	1.3	4.0	1.6	1.8	0.6
NZ6	-0.1	-0.1	-0.1	0.8	0.2	1.0	-0.1	2.3	1.0	3.4	1.2	3.7	1.0	-0.1	3.7	-0.1	1.5	1.7
NZ7	0.7	-0.1	0.7	1.1	0.8	3.2	-0.1	3.8	1.9	8.8	1.3	8.3	1.1	1.4	8.3	1.8	2.3	2.8
NZ8	0.7	-0.1	-0.1	0.9	0.2	1.0	-0.1	2.7	1.1	4.0	1.2	4.2	-0.1	-0.1	4.4	1.4	1.5	0.5
NZ9	-0.1	-0.1	-0.1	0.8	0.2	0.8	-0.1	1.9	0.3	3.6	1.1	3.2	-0.1	-0.1	3.4	-0.1	1.2	1.2
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	4.6	-0.1	3.8	-0.1	-0.1	4.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	3.4	-0.1	3.4	-0.1	-0.1	3.4	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	2.1	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	1.6	1.0	2.2	-0.1	2.9	-0.1	-0.1	0.5	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	2.0	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	2.3	-0.1	2.3	-0.1	-0.1	2.4	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.3	-0.1	2.7	-0.1	2.4	-0.1	-0.1	2.4	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	2.2	-0.1	2.1	-0.1	-0.1	2.2	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	2.1	-0.1	2.0	-0.1	-0.1	2.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.3	-0.1	1.3	-0.1	1.3	-0.1	-0.1	1.3	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.3	-0.1	2.0	-0.1	1.9	-0.1	-0.1	2.0	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	0.3	-0.1	1.8	-0.1	1.7	-0.1	-0.1	1.7	-0.1	-0.1	-0.1

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	055 LPB	056 LBM	057 ALK	058 LPB	059 LPB	060 LPH	061 LBI	062 LBA	063 LPH	064 LBA	065 HPB	066 LBA	067 LBI	068 HPB	069 LA	070 HPB	071 HPB	072 HPB

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Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	079-HBA	074-HBA	075-HPB	076-LPH	077-MAR	078-ALK	079-LBL	080-LPH	081-MAR	082-LPH	083-HBA	084-HBA	085-LPH	086-LBL	087-MAR	088-HBA	089-THL	090-HPB
NA1	5.8	14.2	-0.1	1.1	2.6	1.9	0.6	2.5	2.4	2.8	25.0	1.3	24.6	3.1	4.4	30.9	-0.1	0.9
NA2	0.3	3.8	0.2	1.1	1.1	1.1	-0.1	1.1	1.1	1.1	7.7	0.5	7.3	2.1	0.6	9.2	-0.1	2.0
NAA1	3.0	9.4	3.0	1.7	1.6	2.7	-0.1	1.5	1.5	1.6	16.0	0.7	13.5	2.3	0.8	19.1	-0.1	0.5
NAA10	1.3	7.7	0.6	0.3	0.3	2.9	-0.1	1.4	1.5	1.5	11.3	2.7	1.8	2.2	2.8	11.3	-0.1	2.3
NAA11	6.8	18.8	10.5	4.5	4.8	11.7	2.3	7.1	2.5	8.5	32.7	7.4	28.4	3.1	4.5	33.0	-0.1	0.9
NAA2	68.4	45.0	33.9	10.6	8.7	14.5	8.1	17.4	6.1	20.9	83.7	20.4	172.0	12.4	10.8	172.0	2.6	4.1
NAA3	13.9	2.9	3.5	0.8	3.0	0.8	0.5	2.2	1.4	2.3	27.7	1.1	26.4	3.2	3.0	32.1	-0.1	2.5
NAA4	0.6	6.6	0.3	0.2	1.5	2.1	-0.1	1.3	1.3	1.5	2.5	0.7	11.5	2.3	2.5	6.5	-0.1	0.4
NAA5	3.2	9.3	4.5	0.4	2.2	3.2	-0.1	2.0	1.8	2.1	17.5	1.0	15.1	2.4	3.3	20.2	-0.1	0.5
NAA6	4.6	13.2	3.6	0.3	2.0	3.6	-0.1	1.8	1.6	1.9	24.2	0.9	20.1	2.6	3.1	15.0	-0.1	0.5
NAA7	0.6	6.8	0.3	0.3	1.5	2.1	-0.1	1.4	1.4	1.5	13.8	0.7	12.1	2.3	2.6	1.9	-0.1	2.3
NAA8	3.8	10.3	1.8	0.3	1.4	2.7	-0.1	1.3	1.2	1.3	13.2	0.7	2.0	2.2	2.2	1.9	-0.1	0.4
NAA9	7.5	14.0	4.5	0.6	2.3	4.7	-0.1	2.1	1.7	2.4	16.6	1.1	9.2	2.5	3.2	2.5	-0.1	0.5
NB1	0.8	6.5	3.2	0.2	1.6	0.6	-0.1	1.5	1.5	1.7	2.7	0.8	12.6	2.5	0.7	8.2	-0.1	0.5
NB2	4.7	10.0	1.8	1.2	1.1	2.7	-0.1	1.1	1.1	1.2	9.6	0.6	10.4	0.5	2.1	1.8	-0.1	1.9
NB3	0.4	5.8	1.7	1.1	1.2	1.6	-0.1	1.1	1.1	1.1	8.4	0.9	6.9	2.0	2.2	8.5	-0.1	2.0
NB4	0.8	9.4	0.4	1.5	1.5	0.7	-0.1	1.4	1.3	1.4	23.5	0.6	24.1	2.8	0.6	18.1	-0.1	2.2
NBB1	1.3	9.9	0.5	0.3	1.4	2.6	-0.1	1.3	1.3	1.4	12.4	2.3	1.9	0.5	2.5	2.0	-0.1	2.2
NBB10	8.7	4.2	0.7	1.6	4.0	1.2	1.1	3.7	2.5	4.4	29.1	1.6	27.6	3.4	1.5	20.0	-0.1	0.7
NBB11	8.6	17.9	11.3	1.9	3.4	6.5	1.0	3.2	2.7	3.6	29.1	1.0	12.1	3.1	4.9	16.6	-0.1	2.0
NBB2	5.2	14.9	4.8	1.7	4.0	6.5	0.9	3.4	1.7	3.9	6.6	1.6	25.4	3.0	3.2	15.4	-0.1	0.4
NBB2-R	4.3	13.3	0.1	1.4	3.3	1.4	0.7	2.6	1.6	3.2	24.9	1.2	22.6	2.7	2.9	14.3	-0.1	2.6
NBB3	11.1	11.6	0.8	1.7	3.5	1.3	0.7	3.0	1.7	3.3	4.9	1.3	21.4	2.7	0.9	23.6	-0.1	0.4
NBB4	0.6	4.5	1.8	0.2	1.5	1.6	-0.1	1.2	1.2	1.4	7.6	0.6	7.2	2.0	0.5	1.6	-0.1	2.1
NBB5	0.7	7.3	2.0	0.2	1.3	2.0	-0.1	1.3	1.2	1.3	12.7	0.6	10.9	2.1	2.3	1.9	-0.1	2.2
NBB6	6.1	16.1	5.0	1.9	0.5	6.6	1.7	3.0	1.9	3.6	28.7	5.4	12.4	3.0	3.6	4.3	-0.1	0.8
NBB7	0.4	4.2	1.4	0.3	1.0	1.1	-0.1	-0.1	1.0	1.1	6.8	1.9	0.6	1.9	2.0	0.9	-0.1	-0.1
NBB8	3.1	9.2	0.4	0.4	1.9	2.9	-0.1	1.7	1.3	1.9	16.9	3.1	13.7	2.2	2.6	19.5	-0.1	0.5
NBB9	15.4	18.3	-0.1	3.3	2.5	9.5	2.2	5.2	2.0	6.1	34.5	2.5	29.1	3.1	3.7	5.0	-0.1	1.2
NC1	0.6	9.7	1.8	0.2	1.1	2.1	-0.1	1.1	1.1	1.2	15.3	0.6	12.3	2.1	2.1	1.7	-0.1	1.9
NC2	3.5	11.1	2.1	1.3	1.2	2.6	-0.1	1.2	1.2	1.3	18.0	0.6	14.6	2.2	2.2	20.1	-0.1	2.1
NC3	1.3	9.2	1.5	0.2	1.2	2.2	-0.1	1.1	1.1	1.2	15.7	0.6	13.7	2.2	2.1	2.8	-0.1	2.0
NC4	3.6	4.5	0.2	0.2	1.1	1.3	-0.1	1.0	1.1	1.1	6.6	2.0	0.7	-0.1	2.0	7.1	-0.1	-0.1
NC5	1.3	8.0	0.2	-0.2	1.3	2.2	-0.1	1.2	1.2	1.3	14.5	2.2	12.9	2.2	0.5	2.1	-0.1	0.4
NC6	0.6	5.3	2.6	1.4	1.3	0.5	-0.1	1.3	1.4	1.4	8.1	0.6	9.2	2.3	0.7	1.5	-0.1	0.4
NC7	3.7	4.1	0.5	0.3	1.3	1.4	-0.1	1.2	1.1	1.3	9.4	2.3	8.3	2.2	2.2	1.6	-0.1	2.1
NC8	0.9	4.4	2.0	0.2	1.2	1.5	-0.1	1.2	1.1	1.2	7.8	0.6	7.0	2.0	2.2	1.1	-0.1	2.1
NCC1	38.1	36.3	36.6	3.5	8.3	2.5	4.4	7.6	4.2	8.7	87.6	8.1	158.0	13.3	7.6	40.8	3.2	4.5
NCC1-R	36.9	35.7	35.4	3.3	8.5	2.1	4.3	7.2	4.2	8.7	151.0	7.8	155.0	12.7	2.9	39.6	3.2	4.4
NCC2	9.5	29.9	46.3	5.0	9.9	15.1	4.0	10.1	6.8	12.8	73.2	5.5	64.8	5.6	4.3	52.6	-0.1	3.7
NCC3	7.6	21.3	1.1	1.9	3.8	7.4	1.0	3.9	2.2	4.7	37.8	1.7	13.7	3.1	4.0	25.0	-0.1	2.3
NCC4	13.2	33.3	33.8	5.6	7.7	16.5	4.0	11.8	4.8	13.7	72.5	11.9	12.1	5.9	8.2	44.7	-0.1	5.2
NCC5	5.5	15.1	7.1	1.4	3.1	1.6	0.6	2.8	1.8	3.3	27.9	1.2	23.7	2.9	3.3	29.6	-0.1	4.4
NCC6	5.0	13.9	6.7	1.9	5.1	1.7	-0.1	4.0	2.2	4.7	28.6	1.9	3.1	3.1	1.2	31.6	-0.1	1.3
ND1	0.6	6.2	3.3	0.3	1.5	2.0	-0.1	1.4	1.3	1.6	10.4	0.7	9.0	2.1	0.7	1.4	-0.1	2.3
ND10	4.3	7.6	1.4	0.2	1.2	1.9	-0.1	0.1	1.1	1.2	10.5	0.5	10.0	2.2	2.1	13.9	-0.1	1.9
ND2	6.9	19.0	0.6	1.0	2.6	5.4	-0.1	2.3	1.5	2.6	29.3	1.2	23.8	2.6	2.8	34.5	-0.1	2.9
ND3	4.7	6.5	4.5	0.3	1.1	1.6	-0.1	1.0	1.1	1.1	10.9	0.9	1.8	2.1	2.1	12.4	-0.1	1.9
ND4	23.1	21.0	2.8	0.4	0.4	7.4	1.6	2.3	1.6	2.6	39.6	1.3	39.9	4.2	3.0	49.2	-0.1	0.4
ND5	3.7	7.1	1.2	0.2	1.2	1.9	-0.1	1.4	1.0	1.1	9.3	2.0	3.8	2.1	2.1	10.3	-0.1	0.1
ND6	0.9	6.1	1.6	1.2	1.2	1.7	-0.1	1.1	1.2	1.2	11.0	0.5	10.0	2.2	0.5	12.8	-0.1	2.1
ND7	0.3	1.7	4.8	0.2	0.1	0.4	-0.1	0.1	-0.1	1.1	0.7	-0.1	3.1	-0.1	0.5	3.6	-0.1	0.1
ND8	1.0	7.1	1.6	1.2	1.2	2.0	-0.1	1.2	1.1	1.2	10.7	2.1	9.0	2.2	2.1	1.4	-0.1	2.1
ND8-R	6.6	7.9	1.8	0.2	1.3	2.3	-0.1	1.6	1.1	1.3	12.9	0.6	11.4	0.4	2.2	4.8	-0.1	2.1
ND9	0.9	9.6	1.8	0.2	1.3	0.9	-0.1	1.2	1.2	1.2	14.4	0.5	12.4	2.3	0.6	1.7	-0.1	2.2
NDD1	5.9	18.2	5.1	1.6	3.5	6.8	1.7	3.2	1.9	3.5	26.8	1.3	21.3	2.9	3.5	27.7	-0.1	0.5
NDD2	4.3	12.6	0.9	0.5	2.2	1.2	-0.1	2.0	1.5	2.2	21.5	1.0	17.8	2.4	2.8	23.9	-0.1	0.4
NDD3	1.1	10.1	0.3	0.3	1.6	2.6	-0.1	1.5	1.5	1.6	16.5	2.7	1.3	0.5	2.8	16.0	-0.1	2.4
NDD4	0.9	7.7	0.6	0.3	1.7	2.4	-0.1	1.5	1.5	1.7	13.1	2.7	11.2	2.3	2.9	13.7	-0.1	2.5
NDD5	3.1	11.0	4.9	1.4	1.5	2.6	-0.1	1.3	1.3	1.4	19.8	0.6	16.2	2.3	2.5	24.2	-0.1	2.1
NE1	17.1	19.4	13.1	2.8	6.1	9.2	2.8	5.6	2.6	6.8	7.8	3.1	37.8	4.0	4.7	21.0	-0.1	3.7
NE10	0.5	3.6	0.2	-0.1	0.1	0.9	-0.1	0.1	-0.1	0.1	5.0	-0.1	0.7	0.1	-0.1	6.6	-0.1	0.1
NE11	0.8	6.3	1.4	0.2	1.2	1.5	-0.1	1.1	1.1	1.2	11.2	0.6	10.3	2.1	0.5	15.8	-0.1	2.0
NE2	0.5	5.2	4.2	0.1	0.1	1.2	-0.1	0.1	-0.1	0.1	7.8	0.1	7.8	0.1	0.1	2.1	-0.1	0.1
NE3	0.7	8.3	0.2	0.2	1.2	1.9	-0.1	1.2	1.1	1.3	13.9	2.2	12.4	2.0	2.1	1.7	-0.1	2.0
NE4	0.8	7.2	0.2	0.3	1.1	1.6	-0.1	1.1	1.1	1.2	13.2	2.1	1.3	0.5	2.2	14.0	-0.1	2.0
NE4-R	0.4	4.1	1.4	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	6.3	-0.1	0.7	-0.1	2.0	6.3	-0.1	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBL	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBL	087-MAR	088-HBA	089-THL	090-HPB
NE5	1.5	10.6	5.6	0.3	2.0	4.0	-0.1	1.7	1.5	1.9	12.9	0.9	6.2	2.3	2.9	18.2	-0.1	0.4
NE6	0.9	7.6	1.6	0.2	1.3	2.2	-0.1	1.2	1.1	1.3	9.9	2.2	1.0	2.1	2.1	9.9	-0.1	2.1
NE7	0.3	3.3	2.5	1.2	1.1	0.4	-0.1	1.1	1.1	1.2	5.8	0.5	5.1	2.0	0.4	6.1	-0.1	1.9
NE8	0.8	8.5	1.1	-0.1	1.1	1.9	-0.1	-0.1	1.0	-0.1	9.4	-0.1	8.3	2.0	2.0	11.3	-0.1	-0.1
NE9	6.5	10.3	4.9	0.2	1.2	2.6	-0.1	1.1	1.2	1.2	4.7	2.1	1.6	2.1	2.3	10.7	-0.1	0.4
NEE1	3.7	12.1	2.1	0.2	1.6	2.9	-0.1	1.4	1.3	1.5	15.1	0.7	11.9	2.2	0.6	16.4	-0.1	2.3
NEE2	5.2	14.6	0.2	1.2	2.5	4.5	0.5	2.2	1.5	2.5	28.7	1.1	19.8	2.6	2.8	16.1	-0.1	0.4
NEE3	5.6	14.9	0.4	0.5	2.3	4.7	-0.1	1.9	1.5	2.1	25.7	0.9	23.0	2.7	2.8	16.4	-0.1	0.3
NEE4	0.6	5.0	0.1	0.2	1.4	1.8	-0.1	1.3	1.4	1.4	2.0	0.6	8.9	2.2	0.6	1.6	-0.1	0.5
NEE5	0.7	9.4	1.5	0.2	1.4	2.3	-0.1	1.3	1.3	1.4	14.4	0.6	11.8	2.1	2.2	1.7	-0.1	2.1
NEE6	3.8	10.1	0.8	0.4	2.1	3.2	-0.1	1.6	1.5	1.9	19.1	3.1	16.4	2.6	2.9	6.5	-0.1	0.6
NEE7	3.0	9.2	2.5	0.3	1.8	2.5	-0.1	1.4	1.6	1.6	14.6	0.7	13.2	2.2	3.0	16.4	-0.1	0.6
NF1	2.1	14.5	6.5	0.5	3.0	5.6	1.9	2.9	2.0	3.4	28.4	1.7	23.9	2.9	3.8	33.0	-0.1	0.8
NF10	0.7	6.3	4.6	1.4	1.3	2.2	-0.1	1.4	1.6	1.5	13.3	0.7	11.7	2.3	3.0	7.9	-0.1	2.4
NF11	8.3	2.1	7.6	0.6	1.9	0.6	0.7	1.9	1.7	2.1	35.7	1.0	5.9	4.8	1.0	34.8	-0.1	2.6
NF2	1.0	6.6	1.5	0.2	1.1	1.6	-0.1	1.1	1.0	1.2	11.5	2.0	10.0	2.1	2.1	13.8	-0.1	2.0
NF3	3.5	4.1	2.5	1.2	1.2	1.4	-0.1	1.1	1.1	1.2	7.4	2.1	6.9	2.1	2.2	8.4	-0.1	2.0
NF4	1.0	7.4	2.2	0.3	1.4	2.2	-0.1	1.3	1.3	1.4	16.4	2.4	15.3	2.5	2.4	12.4	-0.1	0.4
NF5	1.0	10.6	2.8	1.2	1.2	2.5	-0.1	1.2	1.2	1.3	16.3	0.6	14.6	2.4	0.6	2.3	-0.1	0.4
NF6	7.3	6.2	0.4	0.2	1.6	0.5	-0.1	1.5	1.3	1.6	18.6	0.7	18.9	2.9	0.7	8.4	-0.1	0.4
NF7	7.6	22.2	16.3	1.4	3.2	7.7	1.9	3.3	4.1	3.9	46.2	1.6	42.0	3.9	7.2	25.4	-0.1	3.2
NF8	0.6	9.1	2.8	1.1	1.1	2.0	-0.1	1.1	1.2	1.1	12.7	2.0	10.6	2.1	2.4	1.7	-0.1	2.1
NF8-R	1.2	8.9	4.6	0.2	1.7	0.7	-0.1	1.6	1.8	1.7	22.8	0.7	22.4	3.0	0.9	17.5	-0.1	0.5
NF9	0.8	7.7	3.1	0.2	1.5	2.5	-0.1	1.4	1.4	1.6	2.6	0.7	16.1	2.5	2.6	12.5	-0.1	0.5
NFF1	0.6	11.4	3.2	0.2	1.7	3.7	-0.1	1.5	1.4	1.7	13.6	2.7	7.9	2.4	2.7	2.1	-0.1	0.4
NFF2	6.0	16.1	6.3	1.3	3.8	2.0	0.7	2.8	2.0	3.2	27.6	1.2	26.3	3.0	1.1	32.7	-0.1	0.5
NFF3	4.7	1.1	2.2	0.4	2.0	0.5	-0.1	1.8	1.4	2.0	18.0	0.9	11.9	2.5	0.7	14.5	-0.1	0.4
NFF4	6.4	14.5	8.5	1.4	3.0	2.2	1.6	2.8	2.2	3.2	26.3	1.5	24.7	3.0	4.1	19.8	-0.1	0.7
NFF5	4.2	15.2	8.5	0.5	2.2	1.4	-0.1	1.8	1.6	2.0	20.0	0.8	16.7	2.4	0.6	23.4	-0.1	2.9
NFF6	0.8	8.2	1.6	0.2	1.2	1.9	-0.1	1.2	1.1	1.2	13.6	2.1	11.1	2.1	2.3	1.6	-0.1	2.1
NFF6-R	0.5	4.3	1.6	0.3	1.2	1.3	-0.1	1.2	1.1	1.2	7.8	2.1	6.8	2.0	2.2	1.9	-0.1	2.1
NFF7	3.9	12.9	2.3	0.3	1.5	1.1	-0.1	1.4	1.2	1.5	16.8	0.7	13.7	2.1	2.4	1.9	-0.1	2.3
NFF8	1.1	9.3	0.5	0.3	1.3	2.4	1.3	1.3	1.3	1.4	15.6	2.3	1.9	0.4	2.5	15.7	-0.1	2.2
NG1	0.6	2.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	5.3	-0.1	5.2	-0.1	-0.1	6.5	-0.1	-0.1
NG10	1.7	7.9	1.6	0.2	1.4	2.2	-0.1	1.2	1.2	1.3	18.0	0.6	11.4	2.2	0.5	14.4	-0.1	2.2
NG11	4.2	0.4	1.3	1.1	1.1	0.3	-0.1	-0.1	1.0	1.1	5.4	-0.1	5.2	2.0	-0.1	0.7	-0.1	-0.1
NG2	1.3	4.4	-0.1	1.1	1.2	1.4	-0.1	-0.1	1.1	1.1	7.8	2.0	7.8	2.1	0.5	9.7	-0.1	2.1
NG2-R	1.2	4.7	1.1	1.1	1.2	1.4	-0.1	-0.1	1.1	1.1	8.9	0.5	8.8	2.1	0.5	11.7	-0.1	-0.1
NG3	1.0	3.8	1.2	-0.1	1.0	1.1	-0.1	-0.1	1.0	-0.1	6.2	-0.1	5.6	-0.1	0.5	1.2	-0.1	0.1
NG4	0.7	4.1	1.8	1.1	1.1	1.2	-0.1	1.1	1.2	1.1	6.5	2.0	6.3	2.0	0.5	8.0	-0.1	2.1
NG5	1.9	13.2	0.5	0.4	2.5	4.5	-0.1	2.1	1.7	2.2	25.6	3.4	23.2	2.9	3.2	4.2	-0.1	1.5
NG6	1.9	4.5	1.5	1.3	1.2	1.1	-0.1	1.2	1.2	1.2	9.8	2.1	8.8	2.1	2.2	1.6	-0.1	2.3
NG7	0.3	1.4	1.1	1.1	0.1	0.1	-0.1	-0.1	-0.1	-0.1	3.0	-0.1	3.0	-0.1	-0.1	3.6	-0.1	-0.1
NG8	0.6	6.7	1.3	1.1	1.1	1.7	-0.1	-0.1	1.0	1.1	8.8	0.5	7.6	-0.1	0.5	1.9	-0.1	-0.1
NG9	0.6	7.4	2.0	0.2	1.3	0.7	-0.1	1.2	1.2	1.3	11.8	0.5	9.9	2.1	0.6	1.4	-0.1	2.3
NGG1	3.0	10.8	3.6	0.2	1.8	0.9	-0.1	1.6	1.4	1.4	19.2	0.7	16.0	2.3	0.8	23.2	-0.1	2.4
NGG10	9.5	11.0	0.6	1.6	0.7	5.2	0.8	3.2	1.5	3.7	3.8	1.5	22.0	3.0	2.9	14.7	-0.1	2.6
NGG2	0.6	9.1	2.2	0.2	1.3	0.9	-0.1	1.2	1.2	1.3	12.2	0.6	10.2	2.1	2.2	13.4	-0.1	2.1
NGG3	26.3	27.5	17.4	3.1	2.8	2.1	3.2	6.0	3.8	7.0	79.8	2.8	75.3	6.6	7.0	78.6	2.4	1.5
NGG4	5.6	15.9	6.7	1.4	3.2	5.9	0.6	2.6	1.7	3.1	30.0	1.5	24.7	2.8	3.2	18.9	-0.1	0.5
NGG5	3.1	7.2	1.2	0.5	2.0	1.0	-0.1	1.8	1.5	2.0	14.8	0.8	11.9	2.3	2.8	1.7	-0.1	0.4
NGG6	4.4	9.2	3.6	0.3	2.0	3.8	-0.1	1.7	1.5	1.9	15.0	0.7	7.1	2.3	2.9	2.2	-0.1	0.4
NGG7	5.2	18.6	4.9	2.1	2.0	1.7	-0.1	1.6	1.6	2.0	28.3	0.8	22.3	2.5	0.8	31.2	-0.1	0.4
NGG8	9.5	12.2	2.9	0.7	2.7	1.1	-0.1	2.0	1.4	2.2	25.2	0.9	22.3	2.7	2.6	17.5	-0.1	2.5
NGG9	0.6	5.6	1.7	0.2	1.5	1.9	-0.1	1.3	1.2	1.2	10.1	0.6	9.1	2.1	2.2	1.5	-0.1	2.1
NH1	1.3	9.4	1.3	1.1	1.2	2.3	-0.1	1.1	1.1	1.1	11.3	0.6	10.5	2.1	2.1	11.4	-0.1	2.1
NH10	0.4	5.7	1.5	-0.1	1.0	1.2	-0.1	-0.1	1.1	-0.1	8.7	-0.1	7.4	-0.1	2.1	9.9	-0.1	2.0
NH10-R	0.5	5.2	2.0	1.2	1.1	1.3	-0.1	1.1	1.1	1.1	8.3	0.5	7.2	-0.1	0.6	1.0	-0.1	2.2
NH11	0.8	8.9	1.7	0.2	1.3	2.1	-0.1	1.2	1.2	1.3	14.1	0.9	12.1	0.5	2.4	2.2	-0.1	2.4
NH2	0.7	6.7	1.7	1.2	1.2	1.8	-0.1	1.1	1.2	1.2	11.1	0.5	9.7	2.1	0.6	1.7	-0.1	2.2
NH3	0.3	3.8	0.2	-0.1	0.1	0.9	-0.1	-0.1	-0.1	-0.1	5.8	-0.1	5.2	-0.1	-0.1	1.3	-0.1	-0.1
NH4	0.7	7.0	1.4	1.1	1.2	1.7	-0.1	1.1	1.2	1.1	11.0	2.0	9.6	2.1	2.2	11.1	-0.1	2.1
NH5	0.5	4.5	1.5	1.1	1.1	1.1	-0.1	-0.1	1.2	1.1	7.9	1.9	7.3	0.1	2.2	1.7	-0.1	2.1
NH6	0.7	5.6	1.8	1.1	1.2	1.5	-0.1	1.1	1.3	1.1	8.6	2.0	7.5	2.1	0.7	1.5	-0.1	0.5
NH7	6.2	16.3	2.8	1.3	1.3	3.8	-0.1	1.3	1.7	1.3	26.3	2.2	21.2	2.7	3.2	3.6	-0.1	0.6
NH8	0.8	6.7	0.3	0.2	1.2	1.6	-0.1	1.1	1.3	1.1	10.9	2.0	9.8	2.2	2.5	12.7	-0.1	2.1
NH9	0.7	6.5	1.6	1.2	1.2	1.6	-0.1	1.1	1.3	1.2	11.9	0.9	9.9	2.2	2.5	1.7	-0.1	0.4

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Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

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NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBI	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBI	087-MAR	088-HBA	089-THI	090-HPB
NHH1	0.6	7.4	2.2	0.2	1.4	2.1	-0.1	1.3	1.2	1.4	13.1	0.6	11.8	2.1	0.6	15.5	-0.1	2.1
NHH10	3.1	9.8	2.3	0.2	1.3	2.3	-0.1	1.2	1.2	1.3	15.4	2.2	12.4	2.1	2.4	17.8	-0.1	0.4
NHH11	0.6	8.4	3.4	1.8	1.8	0.7	-0.1	1.5	1.4	1.6	16.7	0.7	14.8	2.3	2.6	20.8	-0.1	2.3
NHH11-R	0.6	7.7	2.8	0.2	1.6	2.3	-0.1	1.4	1.3	1.6	12.9	0.6	11.0	2.3	2.5	16.1	-0.1	2.2
NHH12	3.7	10.9	3.0	0.2	1.6	1.0	-0.1	1.5	1.2	1.7	16.9	0.8	14.2	2.1	2.4	18.9	-0.1	2.2
NHH2	16.7	9.6	0.3	2.1	5.8	6.6	2.4	4.3	2.0	5.0	49.6	2.1	45.9	4.5	3.7	35.0	-0.1	3.1
NHH3	3.4	10.5	3.0	0.3	1.6	2.6	-0.1	1.4	1.2	1.7	18.2	0.7	15.2	2.3	2.5	20.9	-0.1	2.4
NHH4	3.6	11.5	6.2	1.7	3.9	6.7	1.0	3.8	2.5	4.5	4.7	1.7	21.8	3.9	4.5	16.2	-0.1	2.2
NHH5	5.5	18.2	4.4	1.5	4.1	6.7	0.8	3.1	1.5	3.5	35.7	1.4	30.6	3.2	2.8	22.8	-0.1	2.9
NHH6	0.8	6.6	4.0	0.3	1.7	2.7	-0.1	1.5	1.2	1.7	11.3	0.8	9.5	2.2	0.7	1.3	-0.1	0.4
NHH7	5.8	14.5	3.2	0.7	2.7	1.8	-0.1	2.1	1.3	2.3	18.9	1.0	17.4	2.4	0.6	22.7	-0.1	2.6
NHH8	1.0	9.0	2.4	0.3	1.7	2.6	-0.1	1.5	1.2	1.6	17.0	0.8	14.3	2.3	2.3	2.3	-0.1	0.4
NHH9	1.2	11.4	2.4	0.3	1.9	3.2	-0.1	1.6	1.3	1.8	21.2	2.9	2.7	2.4	2.5	24.2	-0.1	0.4
NI1	23.5	10.3	-0.1	1.6	4.3	1.7	2.1	3.2	3.2	3.5	20.8	1.1	51.0	4.6	1.7	55.5	-0.1	3.6
NI10	4.2	12.2	1.7	1.4	1.4	2.8	-0.1	1.2	1.2	1.3	18.0	0.5	15.2	2.2	0.5	2.2	-0.1	2.2
NI2	1.1	8.0	1.7	0.2	1.2	2.6	-0.1	1.2	1.2	1.3	0.5	2.2	1.9	2.0	2.2	10.2	-0.1	0.4
NI3	0.9	9.8	1.6	1.1	1.1	2.4	-0.1	1.1	1.1	1.1	11.3	2.0	6.2	0.4	2.1	14.1	-0.1	2.1
NI4	12.7	13.8	0.4	0.4	2.4	4.5	1.2	1.8	1.9	1.9	24.8	0.9	24.0	3.2	1.1	28.3	-0.1	1.1
NI5	1.1	9.7	2.6	0.2	1.5	2.6	-0.1	1.3	1.5	1.4	14.1	0.6	12.2	2.3	0.8	15.5	-0.1	2.8
NI5-R	1.0	9.6	0.3	1.3	1.4	2.4	-0.1	1.2	1.4	1.3	13.0	2.2	11.0	2.3	2.7	14.1	-0.1	2.6
NI6	0.8	7.2	1.4	0.2	1.1	2.4	-0.1	1.1	1.1	1.2	7.6	0.5	1.6	2.0	0.5	9.4	-0.1	2.0
NI7	28.1	16.3	0.9	1.7	4.3	8.0	2.0	3.1	2.2	3.4	29.3	1.1	55.5	4.4	1.1	59.5	-0.1	4.0
NI8	0.5	6.9	1.2	1.1	1.1	1.7	-0.1	-0.1	1.1	1.1	7.5	0.5	6.7	-0.1	0.5	8.0	-0.1	-0.1
NI9	4.6	12.2	2.0	1.2	1.2	2.6	-0.1	1.1	1.3	1.2	17.8	0.5	15.2	2.3	0.7	2.9	-0.1	2.3
NII1	0.3	2.8	1.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	3.2	-0.1	3.3	-0.1	2.0	0.6	-0.1	-0.1
NII10	0.8	8.8	2.9	0.2	1.5	0.7	-0.1	1.3	1.3	1.5	2.8	0.6	14.2	2.4	0.6	10.1	-0.1	2.4
NII11	0.7	5.2	2.4	0.2	1.1	1.4	-0.1	1.1	1.2	1.2	9.0	2.1	7.7	2.1	0.6	1.4	-0.1	2.5
NII1-R	0.6	7.8	2.9	1.3	1.2	2.1	-0.1	1.2	1.3	1.3	12.0	0.6	10.1	2.2	0.7	2.5	-0.1	2.5
NII12	0.5	5.6	2.4	0.2	1.2	1.6	-0.1	1.2	1.2	1.2	9.8	0.6	8.4	2.2	2.3	10.1	-0.1	0.3
NII13	9.8	8.9	5.5	1.8	2.1	6.6	0.9	2.2	1.6	2.5	25.5	1.0	23.4	3.2	0.8	9.7	-0.1	3.0
NII14	0.4	4.2	1.5	-0.1	-0.1	1.1	-0.1	-0.1	-0.1	-0.1	7.2	0.6	6.4	-0.1	2.0	8.4	-0.1	2.1
NII2	1.7	13.3	2.5	0.3	1.3	3.2	-0.1	1.3	1.3	1.4	6.4	2.3	7.4	2.4	2.5	19.1	-0.1	2.2
NII3	27.2	29.3	2.0	0.7	2.8	8.0	1.6	2.2	1.5	2.3	41.4	1.1	34.8	3.4	2.8	25.7	-0.1	2.9
NII4	38.7	14.7	2.9	0.9	3.0	10.4	1.9	2.3	1.4	2.6	28.9	1.2	64.5	5.2	2.7	66.0	-0.1	2.9
NII5	0.7	8.5	1.7	1.3	1.3	2.7	-0.1	1.2	1.1	1.2	12.8	0.5	12.7	2.3	2.2	1.9	-0.1	2.1
NII6	0.6	7.4	1.3	1.1	1.2	1.8	-0.1	1.1	1.0	1.1	10.7	0.5	9.0	-0.1	2.1	1.9	-0.1	-0.1
NII7	0.5	7.4	1.1	-0.1	-0.1	1.6	-0.1	-0.1	-0.1	-0.1	10.1	-0.1	8.2	-0.1	-0.1	1.4	-0.1	-0.1
NII8	20.2	19.0	2.9	1.1	2.6	1.6	1.6	2.2	1.5	2.4	6.2	1.1	28.8	3.3	2.8	16.8	-0.1	3.0
NII9	0.8	7.2	1.3	1.1	1.1	1.9	-0.1	1.1	1.1	1.2	9.5	0.5	8.0	2.0	0.4	1.4	-0.1	-0.1
NJ1	4.6	9.8	2.5	0.3	1.4	3.8	-0.1	1.4	1.4	1.4	5.5	2.4	5.5	2.2	2.7	13.6	-0.1	0.5
NJ2	1.0	9.6	2.6	0.3	1.6	2.8	-0.1	1.4	1.5	1.5	3.7	0.7	17.7	2.5	0.8	3.2	-0.1	2.8
NJ3	4.3	11.8	2.4	0.2	1.4	2.9	-0.1	1.3	1.4	1.3	4.3	2.2	19.2	2.5	2.6	6.5	-0.1	2.5
NJ4	5.0	13.5	0.2	0.2	1.4	3.3	-0.1	1.3	1.3	1.3	17.9	0.6	18.3	2.4	2.5	22.9	-0.1	2.5
NJ5	0.5	4.5	0.2	0.2	1.1	1.8	-0.1	1.1	1.1	1.2	8.4	2.0	7.1	2.1	0.5	1.1	-0.1	2.1
NJ6	0.8	7.3	0.1	1.2	1.2	2.4	-0.1	1.1	1.1	1.2	9.7	2.0	9.4	2.0	0.6	10.5	-0.1	2.0
NJ7	6.2	7.7	0.2	0.3	0.2	2.2	-0.1	1.3	1.5	1.4	12.3	2.4	1.9	2.2	2.6	14.2	-0.1	0.6
NJJ1	1.2	6.2	1.8	1.3	1.3	1.7	-0.1	1.2	1.2	1.2	11.6	2.2	10.8	2.2	2.4	2.6	-0.1	2.3
NJJ10	36.3	18.9	36.7	13.0	11.4	17.9	4.7	9.9	7.0	12.7	109.0	5.8	93.9	7.2	4.4	56.4	2.7	2.9
NJJ11	0.4	5.3	1.3	-0.1	1.0	1.3	-0.1	-0.1	1.1	1.1	8.8	0.5	7.8	2.0	0.4	0.9	-0.1	-0.1
NJJ12	2.6	8.9	0.7	0.2	1.7	0.8	-0.1	1.5	1.4	1.7	15.2	0.8	13.9	2.2	2.7	17.8	-0.1	0.5
NJJ13	20.2	11.4	4.9	1.3	4.7	1.1	2.2	3.4	1.9	4.0	53.7	1.2	49.8	4.5	1.0	59.4	-0.1	3.1
NJJ14	3.8	5.7	4.4	0.2	1.1	1.4	-0.1	1.0	1.1	1.1	6.9	1.9	8.4	-0.1	2.1	1.6	-0.1	0.1
NJJ2	2.2	8.2	1.9	1.6	1.6	2.7	-0.1	1.4	1.3	1.5	11.5	0.7	10.3	2.2	0.5	12.5	-0.1	2.7
NJJ3	0.6	9.1	1.4	0.2	1.1	2.1	-0.1	1.1	1.1	1.1	8.7	2.0	8.4	-0.1	2.2	10.2	-0.1	0.1
NJJ4	17.5	6.8	2.1	3.5	3.3	1.2	2.0	2.7	1.5	3.1	20.2	1.2	38.1	3.9	0.5	39.3	-0.1	3.0
NJJ5	0.9	7.1	0.1	0.2	1.4	2.1	-0.1	1.3	1.1	1.4	11.4	0.6	10.0	2.1	0.4	1.1	-0.1	2.1
NJJ6	1.0	9.5	1.7	1.3	1.3	0.9	-0.1	1.2	1.2	1.3	13.3	0.5	10.9	2.2	0.5	1.5	-0.1	2.1
NJJ7	0.4	5.0	1.6	-0.1	1.0	1.3	-0.1	1.1	1.1	1.1	7.7	0.5	6.4	2.0	0.6	4.0	-0.1	2.0
NJJ8	0.8	6.7	1.4	0.2	1.4	1.9	-0.1	1.2	1.1	1.4	10.7	0.6	9.2	2.0	0.4	1.5	-0.1	2.1
NJJ8-R	3.8	7.0	4.5	0.2	1.6	2.8	-0.1	1.4	1.2	1.6	12.7	0.7	11.4	2.2	0.5	2.4	-0.1	2.2
NJJ9	26.7	13.4	3.4	1.4	3.8	7.9	2.4	3.0	2.0	3.5	31.2	1.3	62.7	5.2	1.0	52.2	-0.1	1.1
NK1	5.3	10.2	2.8	0.2	1.2	8.2	-0.1	1.2	1.3	1.2	14.5	2.1	2.1	2.1	2.4	14.3	-0.1	0.4
NK2	0.9	4.7	1.5	1.1	1.2	1.4	-0.1	1.1	1.2	1.1	8.2	0.5	7.6	2.1	0.6	9.1	-0.1	2.1
NK3	0.9	10.0	0.2	0.2	1.2	2.4	-0.1	1.3	1.3	1.3	13.4	2.0	12.7	2.2	2.4	15.7	-0.1	2.2
NK4	1.7	5.8	1.6	1.3	1.4	2.0	-0.1	1.2	1.5	1.3	10.8	2.2	10.7	2.3	0.6	12.4	-0.1	2.3
NK4-R	1.2	5.0	1.4	1.2	1.3	1.6	-0.1	1.1	1.3	1.2	9.3	0.5	9.1	2.2	0.6	10.7	-0.1	2.2
NK5	0.6	6.4	2.2	1.2	1.1	2.0	-0.1	1.2	1.3	1.2	10.6	0.6	1.6	0.5	2.4	9.3	-0.1	0.4

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBI	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBI	087-MAR	088-HBA	089-THI	090-HPB
NK6	3.6	11.2	1.8	1.2	1.1	2.4	-0.1	1.1	1.2	1.1	15.4	0.5	12.4	2.2	0.5	1.9	-0.1	2.2
NK7	0.8	6.3	0.2	0.2	1.2	1.7	-0.1	1.1	1.2	1.2	11.0	2.1	10.8	2.1	2.3	11.8	-0.1	0.4
NKK1	0.4	3.7	1.6	1.3	1.4	0.4	-0.1	1.2	1.1	1.3	7.5	0.6	7.4	2.1	0.5	1.4	-0.1	2.0
NKK10	4.6	9.8	1.7	1.3	1.3	3.4	-0.1	1.2	1.1	1.3	11.3	0.6	12.7	2.1	2.1	1.9	-0.1	0.4
NKK11	0.6	7.7	9.9	0.7	2.3	0.8	1.4	2.2	1.5	2.5	14.6	1.1	12.9	6.5	2.8	16.7	-0.1	2.6
NKK2	0.7	6.7	2.3	0.2	1.5	2.1	-0.1	1.4	1.3	1.5	12.8	0.7	11.1	2.2	2.4	14.4	-0.1	0.4
NKK3	4.8	12.3	3.9		3.8	0.6	3.2	1.7	3.7	27.3	1.4	25.0	3.1	0.9	36.0	-0.1	0.4	
NKK4	62.4	57.6	12.8	5.7	17.9	28.0	7.4	14.7	3.8	18.2	170.0	8.3	160.0	12.0	6.8	43.8	3.0	4.3
NKK5	43.8	39.9	4.5	4.7	8.3	3.6	4.6	10.1	2.9	12.3	95.6	10.8	92.7	7.3	5.0	23.6	-0.1	2.7
NKK6	4.4	12.9	4.1	1.3	3.7	1.4	0.6	3.0	1.7	3.5	22.1	1.4	19.5	2.7	0.9	26.2	-0.1	2.8
NKK7	1.0	8.5	3.9	1.2	3.4	1.0	-0.1	2.6	1.6	3.2	18.7	1.2	16.9	2.7	3.0	20.5	-0.1	2.6
NKK8	1.2	7.8	0.5	0.2	1.4	2.9	-0.1	1.4	1.3	1.5	-0.1	2.4	6.0	2.4	2.4	1.8	-0.1	0.5
NKK9	0.9	10.7	6.0	1.5	1.5	3.7	-0.1	1.4	1.2	1.5	13.2	0.6	14.1	2.3	2.4	1.6	-0.1	2.1
NL1	0.5	7.1	2.0	0.2	1.2	1.8	-0.1	1.2	1.2	1.2	9.1	0.5	7.9	2.0	2.2	1.3	-0.1	2.0
NL2	0.6	8.4	2.4	0.2	1.3	2.2	-0.1	1.2	1.2	1.3	10.3	0.5	9.0	2.1	0.7	2.0	-0.1	0.4
NL3	0.4	5.1	2.5	1.3	1.2	1.5	-0.1	1.2	1.3	1.3	8.4	0.6	7.7	2.0	2.5	1.6	-0.1	0.5
NL3-R	3.5	10.4	3.4	1.8	1.8	3.9	-0.1	1.6	1.5	1.8	14.6	0.8	12.8	2.2	2.9	2.7	-0.1	0.5
NL4	2.6	7.1	1.7	0.2	1.2	1.8	-0.1	1.2	1.1	1.3	11.9	2.1	10.3	2.0	2.2	14.4	-0.1	2.0
NLL1	0.6	5.6	3.9	1.8	2.0	0.5	-0.1	1.6	1.5	1.8	18.7	0.8	1.6	2.4	0.8	16.7	-0.1	0.3
NLL10	0.9	7.6	1.9	1.5	1.6	0.7	-0.1	1.4	1.2	1.5	15.3	0.7	13.7	2.3	2.3	18.2	-0.1	2.2
NLL11	0.8	8.5	2.3	0.2	1.4	0.9	-0.1	1.4	1.3	1.5	12.0	0.7	10.9	2.1	2.4	1.6	-0.1	2.3
NLL12	0.5	6.4	2.0	1.3	1.2	0.6	-0.1	1.2	1.1	1.3	7.4	0.5	7.9	0.4	2.2	1.1	-0.1	2.1
NLL2	0.9	7.4	1.6	0.2	1.4	2.3	-0.1	1.3	1.1	1.4	14.7	0.6	10.3	2.2	0.4	12.7	-0.1	2.1
NLL3	3.2	2.9	1.6	0.2	1.2	0.3	-0.1	1.1	1.0	1.2	5.5	0.6	5.1	2.0	0.4	6.0	-0.1	1.9
NLL4	7.4	10.5	4.3	1.5	4.8	1.9	3.3	1.8	3.9	22.0	1.4	21.5	3.1	3.3	25.0	-0.1	3.4	
NLL5	5.9	16.8	4.3	2.6	8.1	14.1	3.2	6.0	1.8	7.1	31.2	3.1	65.1	5.3	3.3	42.3	-0.1	3.4
NLL6	31.8	6.6	0.6	2.2	6.4	1.9	2.9	5.2	1.8	6.2	36.7	2.7	71.4	5.9	3.3	70.5	-0.1	3.2
NLL7	56.7	53.4	-0.1	6.4	11.0	25.7	6.6	13.1	3.5	16.5	68.1	14.2	128.0	9.5	2.0	118.0	2.5	4.2
NLL8	0.7	6.9	4.8	0.2	1.3	1.9	-0.1	1.2	1.1	1.3	11.6	0.6	9.7	2.2	2.2	1.6	-0.1	2.1
NLL9	7.9	22.0	11.8	4.1	3.0	12.3	2.7	6.7	2.6	8.6	38.1	4.5	3.4	0.6	4.8	35.7	-0.1	0.9
NLL9-R	17.1	18.6	11.6	3.0	7.3	2.2	3.7	7.1	2.6	8.7	44.1	3.8	34.6	3.6	1.5	44.1	-0.1	3.4
NM1	4.5	11.0	0.6	0.8	2.5	4.6	-0.1	2.2	1.6	2.5	18.0	1.1	14.2	2.3	2.9	18.3	-0.1	0.4
NM2	3.7	8.6	2.0	0.2	1.2	2.7	-0.1	1.2	1.2	1.3	10.8	0.6	2.0	0.5	2.2	1.7	-0.1	0.4
NMM1	0.5	5.3	1.3	-0.1	1.1	1.3	-0.1	-0.1	1.1	1.1	8.5	-0.1	7.5	-0.1	0.5	1.3	-0.1	-0.1
NMM10	7.3	17.6	0.6	1.5	3.8	1.7	1.6	3.1	1.7	3.5	20.9	1.3	25.9	3.9	0.9	32.7	-0.1	4.1
NMM2	8.3	4.9	0.2	1.4	1.5	0.6	-0.1	1.3	1.2	1.3	3.1	0.6	16.5	2.5	0.5	10.6	-0.1	2.1
NMM3	1.2	8.2	1.4	0.2	1.0	2.1	-0.1	1.0	1.2	1.1	8.9	2.0	8.6	2.1	0.6	10.8	-0.1	2.2
NMM3-R	0.8	7.3	1.3	1.0	1.1	1.7	-0.1	1.0	1.1	1.1	9.3	1.9	7.6	2.0	0.5	1.1	-0.1	2.1
NMM4	0.6	6.3	1.2	1.1	1.1	1.5	-0.1	-0.1	1.1	1.1	8.4	1.9	7.2	-0.1	0.5	0.3	-0.1	-0.1
NMM5	0.5	5.4	1.2	-0.1	1.0	1.3	-0.1	-0.1	-0.1	-0.1	8.8	-0.1	7.7	-0.1	0.5	1.4	-0.1	-0.1
NMM6	1.9	12.2	3.0	0.3	1.9	3.8	-0.1	1.7	2.5	1.9	22.1	0.9	19.3	2.8	1.5	12.2	-0.1	3.5
NMM7	0.4	4.2	1.1	-0.1	1.0	1.1	-0.1	-0.1	-0.1	1.1	6.6	-0.1	6.1	-0.1	0.4	1.0	-0.1	-0.1
NMM8	9.1	21.5	2.3	1.8	4.7	8.6	1.8	3.6	1.6	4.3	36.6	1.3	31.8	3.3	0.8	41.4	-0.1	4.7
NMM9	1.3	11.2	3.1	0.3	1.8	3.3	-0.1	1.7	1.5	1.8	20.2	0.9	17.5	2.6	2.8	3.0	-0.1	2.9
NN1	0.3	3.5	1.1	-0.1	0.1	0.9	-0.1	-0.1	-0.1	-0.1	5.6	-0.1	5.4	-0.1	-0.1	1.0	-0.1	-0.1
NNN1	4.7	9.5	1.3	1.2	1.2	2.9	-0.1	1.1	1.1	1.2	11.0	0.5	9.0	2.0	2.1	10.6	-0.1	2.0
NNN10	2.4	0.3	1.7	-0.1	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	4.0	-0.1	1.0	2.0	-0.5	0.8	-0.1	-0.1
NNN2	0.8	11.1	1.3	1.1	1.2	2.5	-0.1	1.1	1.1	1.1	12.1	0.5	11.7	2.1	0.5	1.8	-0.1	2.0
NNN3	0.4	2.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	5.3	-0.1	4.1	-0.1	-0.1	1.1	-0.1	-0.1
NNN4	1.5	12.9	3.0	0.2	1.7	1.1	-0.1	1.5	1.5	1.6	4.4	0.7	19.7	2.7	0.7	13.3	-0.1	2.9
NNN5	3.9	8.1	1.2	-0.1	1.1	2.3	-0.1	-0.1	-0.1	1.1	10.7	-0.1	8.3	-0.1	2.0	10.4	-0.1	-0.1
NNN6	0.4	3.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	6.2	-0.1	5.9	-0.1	-0.1	1.1	-0.1	-0.1
NNN7	0.4	2.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	4.5	-0.1	4.4	-0.1	-0.1	1.2	-0.1	-0.1
NNN8	8.2	9.3	2.7	0.6	1.8	0.8	-0.1	1.8	1.5	1.9	8.6	0.8	18.8	2.7	2.8	2.4	-0.1	3.0
NNN9	0.9	9.8	2.2	0.2	1.7	0.8	-0.1	1.6	1.4	1.7	17.0	0.7	14.9	2.3	0.7	2.0	-0.1	2.6
NO1	3.0	8.0	3.6	0.6	2.2	3.2	-0.1	1.9	1.4	2.2	15.2	0.9	13.6	2.3	2.7	2.0	-0.1	2.3
NO2	0.2	1.5	1.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.8	-0.1	2.5	-0.1	2.0	0.4	-0.1	-0.1
NOO1	0.6	4.1	1.1	-0.1	1.0	1.1	-0.1	-0.1	-0.1	1.1	1.1	-0.1	7.4	2.0	2.2	1.2	-0.1	-0.1
NOO2	1.0	7.7	1.6	0.2	1.5	0.7	-0.1	1.3	1.2	1.4	13.5	0.6	11.8	2.2	0.5	15.7	-0.1	2.4
NOO3	1.7	0.3	1.3	-0.1	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	0.6	-0.1	4.1	2.0	2.1	0.6	-0.1	2.0
NOO4	0.6	5.3	1.8	0.2	1.2	1.5	-0.1	1.2	1.2	1.2	9.9	0.6	8.9	2.2	0.5	1.4	-0.1	2.4
NOO5	0.5	4.6	1.1	1.1	1.1	1.1	-0.1	1.0	1.0	1.1	7.9	0.5	7.1	-0.1	0.4	1.4	-0.1	-0.1
NOO6	0.4	3.3	1.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	6.2	-0.1	5.9	-0.1	0.5	1.4	-0.1	-0.1
NOO6-R	0.4	3.3	1.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	6.0	-0.1	5.7	-0.1	0.4	1.2	-0.1	-0.1
NOO7	0.5	5.7	1.0	-0.1	-0.1	1.3	-0.1	-0.1	-0.1	-0.1	9.0	-0.1	7.6	-0.1	0.5	10.3	-0.1	-0.1
NOO8	11.0	20.7	3.6	1.2	3.1	6.2	1.9	2.8	1.7	3.2	35.7	1.1	34.2	3.4	0.8	4.9	-0.1	3.8
NP1	0.5	9.6	2.8	1.5	1.6	2.7	-0.1	1.3	1.2	1.5	11.4	0.6	9.8	2.1	2.4	2.1	-0.1	2.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBI	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBI	087-MAR	088-HBA	089-THI	090-HPB
NP2	0.5	7.7	2.0	1.3	1.3	2.1	-0.1	1.2	1.2	1.3	9.5	0.6	8.4	2.0	2.2	1.6	-0.1	2.0
NP3	0.5	6.1	1.8	1.2	1.2	1.8	-0.1	1.1	1.1	1.2	7.7	0.5	6.6	2.0	0.5	8.0	-0.1	2.0
NP4	10.5	21.8	35.1	4.0	2.9	12.4	3.3	7.6	5.6	9.6	8.0	4.8	45.9	4.7	9.9	23.6	-0.1	2.4
NPP4	0.4	2.6	1.1	0.2	1.1	0.1	-0.1	0.1	1.0	1.1	6.5	1.9	5.5	-0.1	2.1	6.9	-0.1	-0.1
NPP1-R	0.4	2.7	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	5.4	-0.1	5.4	-0.1	-0.1	7.1	-0.1	-0.1
NPP2	8.3	15.5	4.1	0.5	2.5	5.7	0.4	2.2	1.6	2.5	26.2	1.3	25.0	3.0	3.0	32.4	-0.1	0.5
NPP3	7.1	14.9	2.6	0.3	2.3	4.8	1.4	1.9	1.9	2.2	2.2	0.9	23.5	2.8	2.5	18.2	-0.1	2.3
NPP4	6.3	12.2	4.1	0.5	1.9	4.0	-0.1	1.8	1.5	2.0	16.6	0.8	15.8	2.4	2.7	10.3	-0.1	2.5
NQ1	0.5	4.4	3.7	1.5	1.4	1.5	-0.1	1.4	1.6	1.6	7.1	0.7	6.8	2.1	0.9	1.5	-0.1	2.5
NQ2	3.4	11.3	3.7	2.3	2.4	1.1	0.6	2.1	1.7	2.4	23.9	1.0	21.5	2.9	0.9	30.6	-0.1	0.4
NQ2-R	4.1	13.4	4.0	2.4	2.4	1.2	0.6	2.2	1.8	2.5	32.1	1.0	30.6	3.4	0.9	40.5	-0.1	0.4
NQ3	0.9	9.5	1.4	1.3	1.5	0.9	-0.1	1.3	1.2	1.4	15.0	0.6	12.9	2.2	0.5	18.2	-0.1	2.0
NQ4	2.8	10.0	0.7	2.1	2.1	1.0	0.4	1.9	1.4	2.2	22.1	1.0	20.9	2.6	2.8	26.9	-0.1	0.4
NQ5	2.4	7.9	4.8	2.0	1.8	2.5	-0.1	1.7	1.4	2.0	14.3	0.8	12.2	2.2	2.7	17.7	-0.1	0.4
NQ6	0.7	8.9	1.4	-0.1	1.0	1.7	-0.1	-0.1	-0.1	-0.1	10.3	1.9	8.8	-0.1	2.0	12.4	-0.1	-0.1
NQ7	4.2	10.3	5.3	1.0	2.5	1.5	0.6	2.4	1.7	2.8	4.1	1.2	20.1	2.9	3.2	14.1	-0.1	0.4
NQ8	0.6	7.7	2.9	1.5	1.4	2.7	-0.1	1.4	1.2	1.6	11.1	0.6	10.3	2.2	2.3	1.5	-0.1	2.1
NQQ1	0.6	6.9	1.6	1.2	1.2	2.2	-0.1	1.2	1.1	1.2	8.1	0.6	7.8	-0.1	2.1	1.4	-0.1	2.0
NR1	0.8	3.2	-0.1	1.1	1.1	1.0	-0.1	-0.1	-0.1	1.0	6.2	-0.1	6.2	-0.1	0.5	7.6	-0.1	-0.1
NR11	0.9	7.1	1.5	0.2	1.3	2.0	-0.1	1.2	1.2	1.2	10.7	0.8	9.7	2.2	0.4	11.7	-0.1	2.5
NR12	0.9	9.5	1.6	0.2	1.2	2.1	-0.1	1.2	1.1	1.2	14.4	2.1	11.7	0.4	2.2	2.8	-0.1	2.2
NR13	11.3	5.9	2.9	0.8	2.5	1.1	0.5	2.0	1.4	2.2	10.2	1.1	25.1	3.2	0.7	4.2	-0.1	3.8
NR14	0.9	6.8	1.2	1.2	1.2	1.8	-0.1	1.1	1.1	1.2	11.1	0.5	9.5	2.1	2.1	1.2	-0.1	-0.1
NR2	6.3	6.8	1.4	0.2	1.3	2.0	-0.1	1.2	1.2	1.2	9.7	0.6	8.6	2.2	0.4	10.5	-0.1	2.2
NR3	6.0	5.9	2.1	0.2	1.5	0.6	-0.1	1.4	1.2	1.5	10.1	0.6	9.4	2.3	0.6	11.1	-0.1	2.6
NR4	0.5	4.0	1.2	-0.1	-0.1	0.9	-0.1	-0.1	-0.1	-0.1	6.3	-0.1	5.8	-0.1	1.9	1.3	-0.1	-0.1
NR5	0.4	4.6	1.2	1.1	1.1	1.1	1.1	-0.1	1.0	1.1	7.9	1.9	7.4	-0.1	0.5	1.7	-0.1	-0.1
NR6	0.4	4.0	1.2	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	6.6	-0.1	6.2	-0.1	-0.1	1.4	-0.1	-0.1
NR7	3.4	13.3	2.5	0.5	2.1	4.4	1.2	1.7	1.6	1.8	21.7	0.8	19.9	2.8	0.7	26.0	-0.1	3.0
NR8	0.7	5.1	1.2	1.1	1.1	1.4	-0.1	-0.1	-0.1	1.1	7.8	-0.1	6.8	-0.1	0.4	9.0	-0.1	-0.1
NS1	4.3	4.1	2.2	0.3	1.5	1.9	-0.1	1.4	1.2	1.5	7.6	2.6	7.4	2.2	0.6	9.0	-0.1	0.4
NS10	5.1	15.4	6.0	0.9	3.0	5.2	0.8	2.7	2.0	3.2	33.9	1.5	5.5	3.5	1.1	36.0	-0.1	2.8
NS11	3.6	12.0	2.2	0.2	1.5	3.0	-0.1	1.4	1.3	1.4	18.4	0.6	15.2	2.3	2.6	21.6	-0.1	0.4
NS12	1.2	11.2	0.2	0.5	2.1	3.7	-0.1	2.0	1.6	2.2	4.4	1.0	18.2	2.6	2.9	12.6	-0.1	0.4
NS13	12.1	14.0	5.3	1.0	3.1	1.2	0.5	2.3	1.8	2.7	30.3	1.1	28.4	3.3	1.0	34.8	-0.1	1.2
NS14	8.1	20.7	12.5	1.7	4.1	2.2	1.8	3.5	3.0	4.1	36.9	1.7	34.2	3.5	5.5	21.6	-0.1	2.7
NS2	4.1	12.9	2.8	0.2	1.7	3.5	-0.1	1.6	1.2	1.8	17.7	0.8	16.8	2.3	2.4	22.4	-0.1	2.1
NS3	5.1	11.2	3.8	1.3	3.2	1.7	0.6	2.8	1.9	3.2	22.3	1.2	19.1	2.8	3.5	22.2	-0.1	0.5
NS4	9.6	9.2	2.3	0.3	0.3	3.7	1.2	1.6	1.4	1.7	16.8	2.9	16.9	2.8	2.5	22.5	-0.1	0.4
NS4-R	2.2	7.1	2.1	0.3	1.9	0.5	-0.1	1.4	1.2	1.6	13.9	2.0	14.1	2.6	2.4	18.6	-0.1	0.4
NS5	9.6	9.3	1.7	0.3	0.4	1.0	-0.1	1.8	1.3	2.0	17.7	1.0	18.3	3.0	2.5	23.5	-0.1	2.3
NS6	3.4	10.5	2.4	1.5	1.6	2.7	-0.1	1.4	1.2	1.5	14.6	0.6	12.3	2.2	0.6	16.9	-0.1	2.1
NS7	13.5	13.4	3.5	1.1	3.8	1.0	1.5	2.5	1.6	2.8	29.7	1.3	28.9	3.5	0.8	34.2	-0.1	0.3
NS8	2.1	15.5	6.1	1.2	4.1	1.6	1.9	3.1	2.1	3.6	28.0	1.3	26.4	3.3	3.9	32.4	-0.1	0.7
NS9	1.6	4.8	2.1	1.1	1.2	1.5	-0.1	1.1	1.1	1.2	8.0	2.1	7.8	2.1	2.2	10.4	-0.1	1.9
NT1	0.5	5.9	2.4	0.2	1.1	1.5	-0.1	1.1	1.1	1.1	7.3	2.0	7.1	2.8	0.5	6.7	-0.1	2.0
NT10	1.0	4.6	2.8	0.3	1.9	1.9	-0.1	1.5	1.3	1.7	8.9	0.8	8.3	2.1	0.7	10.9	-0.1	2.4
NT11	3.1	3.0	4.7	0.3	1.4	0.3	-0.1	1.3	1.2	1.4	6.2	2.3	6.0	2.1	0.6	7.4	-0.1	2.1
NT12	0.7	7.3	3.3	0.5	2.0	2.7	-0.1	1.7	1.4	2.0	9.5	0.9	8.5	2.1	0.7	10.5	-0.1	0.4
NT13	0.6	6.4	1.9	0.2	1.5	1.9	-0.1	1.3	1.2	1.4	10.3	0.6	9.4	2.1	2.2	1.9	-0.1	2.1
NT14	0.9	8.6	0.3	0.3	1.4	2.4	-0.1	1.3	1.2	1.5	14.2	2.4	1.7	2.2	2.3	1.7	-0.1	2.2
NT15	4.2	12.1	4.7	0.5	2.3	3.7	-0.1	2.0	1.4	2.2	23.4	1.0	17.5	2.9	2.6	13.9	-0.1	2.8
NT16	4.3	8.8	3.6	0.3	1.9	3.6	-0.1	1.7	1.4	1.9	12.1	0.8	12.5	2.4	0.7	1.9	-0.1	2.7
NT17	0.7	7.1	1.9	0.2	1.5	2.3	-0.1	1.4	1.1	1.5	12.3	0.7	10.5	2.2	2.2	4.5	-0.1	2.2
NT18	0.5	3.4	1.1	-0.1	-0.1	0.9	-0.1	-0.1	-0.1	-0.1	5.0	-0.1	4.5	-0.1	-0.1	1.5	-0.1	-0.1
NT2	0.5	4.9	3.5	0.2	1.3	1.5	-0.1	1.2	1.2	1.3	10.4	0.6	9.2	2.2	2.3	1.6	-0.1	0.4
NT3	0.8	7.6	1.2	-0.1	1.0	1.5	-0.1	-0.1	-0.1	-0.1	13.1	1.9	10.9	-0.1	0.5	16.8	-0.1	-0.1
NT4	2.7	7.2	1.2	1.1	1.1	1.6	-0.1	-0.1	-0.1	1.1	14.2	0.5	9.9	2.0	0.5	14.5	-0.1	-0.1
NT5	0.5	4.1	1.5	1.2	1.2	1.2	-0.1	1.1	1.1	1.2	7.1	0.5	6.6	-0.1	0.5	1.4	-0.1	-0.1
NT6-R	3.9	10.3	2.0	1.4	1.4	2.8	-0.1	1.2	1.2	1.3	11.8	0.6	9.8	2.1	0.6	2.2	-0.1	2.1
NT6	1.0	5.5	1.3	1.3	1.3	1.6	-0.1	1.2	1.1	1.2	8.5	0.6	8.0	2.0	0.4	1.9	-0.1	-0.1
NT7	2.4	6.2	2.6	0.3	1.9	0.6	-0.1	1.6	1.3	1.8	14.3	0.8	11.6	2.5	0.6	12.0	-0.1	2.3
NT8	3.0	20.7	4.9	1.6	4.0	7.9	1.9	3.2	1.8	3.7	43.5	1.5	39.3	0.6	1.0	53.4	-0.1	1.1
NT9	0.7	5.1	4.5	0.2	1.3	1.5	-0.1	1.2	1.1	1.3	8.4	0.5	7.8	2.0	0.5	1.4	-0.1	-0.1
NU1	0.9	4.1	1.4	0.2	1.5	1.6	-0.1	1.2	1.2	1.4	8.0	2.2	7.9	2.1	0.6	9.9	-0.1	2.1
NU10	10.7	5.5	5.6	0.6	2.1	0.9	1.7	2.0	1.7	2.2	38.1	1.0	39.0	4.9	1.0	40.2	-0.1	0.6
NU11	3.6	4.2	2.3	0.3	1.2	1.5	1.2	1.3	1.1	1.3	9.2	2.3	8.2	2.1	0.5	11.6	-0.1	2.2

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBI	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBL	087-MAR	088-HBA	089-THI	090-HPB
NU12	4.5	12.9	2.0	0.2	1.4	2.8	-0.1	1.3	1.1	1.4	22.0	0.6	16.2	2.3	2.2	26.3	-0.1	2.2
NU12-R	0.7	8.6	2.0	1.4	1.3	2.4	-0.1	1.3	1.1	1.4	15.5	0.6	13.7	2.2	0.6	1.8	-0.1	2.1
NU13	11.5	13.2	3.0	1.4	3.1	5.3	1.0	3.2	1.9	3.6	15.7	1.5	31.8	3.4	1.1	20.8	-0.1	0.5
NU14	5.4	9.6	2.4	1.5	1.3	3.2	-0.1	1.3	1.2	1.4	9.2	0.6	1.2	2.1	2.3	11.9	-0.1	2.3
NU15	0.9	10.7	6.7	2.1	1.9	1.0	-0.1	1.8	1.3	2.0	21.0	0.9	16.8	2.5	2.5	2.9	-0.1	2.4
NU16	7.0	17.4	0.4	1.4	3.4	1.6	2.0	3.1	1.8	3.5	18.7	1.3	42.9	4.0	3.4	7.1	-0.1	1.2
NU17	6.2	17.0	3.0	0.9	2.7	5.5	0.7	2.4	1.5	2.7	2.7	1.3	28.1	3.8	2.0	16.5	-0.1	3.2
NU18	1.1	6.0	2.3	0.3	1.7	2.4	-0.1	1.5	1.3	1.6	10.5	2.7	9.7	2.2	0.7	1.4	-0.1	0.4
NU19	1.1	7.3	2.7	0.3	1.2	2.2	-0.1	1.2	1.2	1.3	10.7	2.2	1.5	2.1	2.2	12.7	-0.1	0.4
NU2	27.2	7.1	0.7	4.6	3.2	2.8	3.3	6.4	2.1	7.8	64.8	4.0	58.8	5.1	3.9	78.0	-0.1	0.5
NU3	0.6	6.2	1.6	0.2	1.1	1.5	-0.1	1.1	1.0	1.1	9.8	0.5	8.9	1.9	2.1	11.0	-0.1	0.1
NU4	38.1	35.1	-0.1	3.1	7.6	2.7	3.3	6.3	3.0	7.5	85.5	7.2	82.8	6.9	5.5	102.0	2.4	1.7
NU5	1.3	4.5	1.4	1.2	1.2	1.5	-0.1	1.1	1.0	1.2	8.4	0.5	8.0	2.1	0.5	9.6	-0.1	0.1
NU6	1.9	5.0	0.3	0.2	1.4	1.9	-0.1	1.2	1.1	1.3	9.6	0.6	9.9	2.2	0.5	12.1	-0.1	2.1
NU7	2.6	9.1	2.0	0.2	1.7	8.0	-0.1	1.4	1.2	1.6	15.5	0.8	14.9	2.4	2.3	20.3	-0.1	2.3
NU8	1.8	12.1	3.3	0.3	2.0	1.3	-0.1	1.8	1.4	1.9	24.8	0.9	23.1	2.9	0.6	12.9	-0.1	2.7
NU9	0.8	8.2	6.5	0.3	1.4	2.5	-0.1	1.5	1.5	1.5	16.6	0.7	14.9	2.5	0.8	10.6	-0.1	0.5
NV1	4.0	11.8	1.8	1.5	1.5	2.9	-0.1	1.3	1.3	1.4	14.2	0.6	11.8	2.1	2.4	13.2	-0.1	0.4
NV10	3.0	9.7	1.6	0.2	1.9	1.9	-0.1	1.1	1.1	1.2	16.3	0.6	13.6	0.4	0.5	10.8	-0.1	2.0
NV11	0.4	4.4	1.5	1.2	1.1	1.0	-0.1	1.1	1.1	1.1	8.9	2.0	7.7	2.0	2.1	10.9	-0.1	-0.1
NV12	3.9	9.4	1.5	0.2	1.2	1.9	-0.1	1.1	1.1	1.2	13.6	0.5	11.2	2.0	2.1	15.1	-0.1	1.9
NV13	4.8	10.8	0.3	0.3	1.6	3.2	-0.1	1.6	1.2	1.7	17.5	0.7	13.3	2.2	2.3	2.4	-0.1	2.3
NV14	1.3	10.2	0.4	1.2	1.2	2.2	-0.1	1.2	1.1	1.2	16.5	2.2	2.1	2.2	2.2	16.8	-0.1	2.1
NV15	2.3	8.0	1.3	1.1	1.1	1.6	-0.1	1.1	1.0	1.1	13.3	2.0	10.8	2.0	0.4	15.5	-0.1	-0.1
NV16	7.7	18.8	2.9	0.3	1.8	4.4	-0.1	1.7	1.4	1.9	22.0	0.8	22.0	2.5	2.7	3.0	-0.1	2.7
NV17	1.2	10.7	2.1	0.3	1.4	2.4	-0.1	1.3	1.3	1.4	15.9	2.3	1.5	2.2	2.4	17.5	-0.1	0.4
NV18	3.9	8.2	1.6	0.3	1.3	2.4	-0.1	1.2	1.1	1.2	9.8	0.6	1.0	2.0	0.5	10.7	-0.1	2.1
NV2	3.0	9.3	2.1	1.5	1.5	2.4	-0.1	1.4	1.2	1.5	14.1	0.6	11.6	2.1	0.6	15.7	-0.1	2.1
NV3	0.8	7.6	4.3	1.1	1.1	1.7	-0.1	1.0	1.0	1.1	12.2	1.9	10.2	0.5	0.4	1.6	-0.1	0.1
NV4	1.5	8.6	0.3	0.3	1.3	2.9	-0.1	1.3	1.3	1.4	14.0	2.3	2.6	2.3	2.4	14.4	-0.1	0.4
NV5	7.0	13.6	2.0	0.2	1.6	4.2	-0.1	1.6	1.4	1.6	19.8	0.7	19.0	2.5	2.6	3.8	-0.1	2.5
NV5-R	5.1	14.5	0.2	0.2	1.4	3.1	-0.1	1.3	1.3	1.5	23.6	0.6	19.9	2.5	2.5	3.4	-0.1	2.4
NV6	0.4	3.1	1.9	0.2	1.1	0.8	-0.1	1.1	1.3	1.2	5.0	2.0	5.5	2.0	2.5	1.0	-0.1	2.0
NV7	1.5	11.7	1.8	0.2	1.2	2.4	-0.1	1.1	1.1	1.2	20.4	2.1	8.1	2.2	2.2	21.2	-0.1	0.4
NV8	0.9	6.8	0.3	0.2	1.1	1.6	-0.1	1.1	1.1	1.1	10.4	2.0	1.4	2.0	2.1	0.5	-0.1	2.0
NV9	1.1	10.1	1.7	0.3	1.4	2.6	-0.1	1.3	1.1	1.4	16.4	2.3	13.8	2.3	2.2	16.3	-0.1	2.1
NW1	5.9	17.5	4.8	1.2	2.8	1.6	-0.6	2.4	1.4	2.6	29.4	1.1	24.8	2.6	0.7	15.7	-0.1	2.7
NW10	0.7	4.6	1.5	0.3	1.2	1.3	-0.1	1.1	1.1	1.2	8.3	2.1	7.3	2.1	0.6	8.8	-0.1	2.0
NW11	3.2	10.8	1.3	1.1	1.1	2.1	-0.1	0.1	1.1	1.1	13.2	0.5	10.2	-0.1	0.5	14.2	-0.1	0.1
NW12	3.4	9.5	2.3	0.3	1.5	2.2	-0.1	1.5	1.2	1.6	15.6	2.5	12.7	2.2	2.4	19.0	-0.1	0.4
NW13	4.4	9.8	0.3	0.4	0.3	3.4	0.5	1.9	1.4	2.1	4.7	1.1	18.3	2.7	0.8	9.2	-0.1	0.5
NW14	0.5	7.1	1.3	1.2	1.2	1.7	-0.1	1.1	1.0	1.2	10.4	0.5	8.6	-0.1	0.4	11.6	-0.1	-0.1
NW15	4.5	13.1	3.2	0.6	2.3	1.2	-0.1	2.1	1.5	2.3	22.5	1.1	18.5	2.5	2.8	24.7	-0.1	0.4
NW16	0.5	6.7	1.8	1.2	1.2	1.6	-0.1	1.1	1.1	1.2	10.6	0.5	8.8	-0.1	0.5	12.3	-0.1	-0.1
NW17	0.8	7.1	4.1	0.4	0.3	2.0	-0.1	1.3	1.3	1.4	11.0	2.4	1.6	0.4	2.5	12.1	-0.1	2.2
NW17-R	0.7	6.0	3.8	0.3	0.2	1.7	-0.1	1.3	1.3	1.4	10.0	2.3	1.1	0.4	2.5	10.2	-0.1	2.1
NW18	0.5	3.6	1.7	0.3	1.2	0.4	-0.1	1.1	1.3	1.2	4.2	2.1	7.3	2.1	2.8	4.1	-0.1	2.5
NW2	1.0	9.3	1.6	0.3	1.2	2.0	-0.1	1.2	1.1	1.2	14.9	2.1	1.4	2.1	0.5	15.3	-0.1	0.4
NW2-R	1.1	8.5	1.5	0.2	1.2	1.8	-0.1	1.2	1.1	1.2	15.8	2.0	6.1	2.1	0.5	17.8	-0.1	2.0
NW3	0.9	8.2	0.2	0.2	1.2	1.9	-0.1	1.1	1.0	1.2	14.3	2.1	11.9	2.1	2.0	2.3	-0.1	-0.1
NW4	0.5	5.4	1.3	1.1	1.1	1.4	-0.1	1.1	1.0	1.1	10.0	1.9	8.6	2.0	0.5	1.4	-0.1	-0.1
NW5	3.5	12.0	2.8	1.3	0.8	1.5	1.9	2.8	1.5	3.1	23.7	1.3	53.4	4.4	0.7	59.7	-0.1	2.9
NW6	0.7	7.2	1.4	0.2	1.3	1.9	-0.1	1.2	1.1	1.3	11.5	0.6	9.8	2.1	0.5	1.7	-0.1	0.1
NW7	0.9	8.5	1.3	0.2	1.2	1.9	-0.1	1.1	1.0	1.2	9.6	2.1	1.3	2.0	2.0	11.1	-0.1	2.0
NW8	19.6	11.4	0.8	1.7	0.9	1.4	3.2	3.4	1.6	3.8	19.3	1.3	40.8	4.0	0.8	25.1	-0.1	3.2
NW9	1.7	12.2	1.3	0.3	0.2	2.7	-0.1	1.3	1.1	1.4	19.5	2.3	7.4	2.1	2.1	20.0	-0.1	-0.1
NX1	0.5	4.1	1.7	1.1	1.1	1.1	-0.1	0.1	1.1	1.1	6.8	1.9	0.8	-0.1	0.5	1.3	-0.1	2.0
NX10	3.9	10.6	4.2	0.3	1.9	3.3	-0.1	1.8	1.5	2.0	18.8	1.1	15.9	2.4	2.9	22.0	-0.1	0.6
NX11	0.5	4.1	1.4	-0.1	1.0	1.1	-0.1	0.1	0.1	1.1	6.4	0.9	0.6	-0.1	0.5	1.0	-0.1	0.1
NX12	0.5	4.9	0.2	0.2	1.1	1.3	-0.1	1.1	1.1	1.2	8.3	0.6	0.7	2.0	2.1	1.1	-0.1	1.9
NX13	5.6	11.0	0.8	0.3	1.8	4.4	-0.1	1.7	1.6	1.9	17.0	0.8	18.0	2.6	3.1	12.3	-0.1	0.5
NX14	0.8	8.1	0.3	0.3	1.8	0.9	-0.1	1.8	1.4	2.0	16.2	0.9	14.2	2.4	2.6	9.5	-0.1	0.4
NX14-R	3.4	10.4	0.4	0.3	1.5	2.6	-0.1	1.4	1.3	1.6	18.9	0.7	15.6	2.3	2.4	21.5	-0.1	0.4
NX15	7.3	12.8	2.4	0.8	1.6	3.6	-0.1	1.4	1.2	1.6	8.9	2.5	8.6	0.6	2.4	2.9	-0.1	0.4
NX16	6.5	18.9	0.7	0.9	2.1	4.8	-0.1	1.9	1.3	2.1	25.1	0.9	22.1	2.5	2.5	3.2	-0.1	2.5
NX17	1.5	14.2	0.3	1.5	3.4	1.4	1.7	3.1	1.8	3.5	28.4	1.3	25.1	3.0	3.4	15.9	-0.1	1.4
NX2	2.6	8.4	9.9	0.2	1.3	2.2	-0.1	1.4	1.3	1.5	15.2	0.7	13.0	2.2	2.6	20.4	-0.1	2.2

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-EPH	077-MAR	078-ALK	079-LBL	080-LPH	081-MAR	082-EPH	083-HBA	084-HBA	085-EPH	086-LBL	087-MAR	088-HBA	089-THL	090-HPB
NX3	0.5	8.2	2.2	1.5	1.4	2.2	-0.1	1.3	1.3	1.5	11.5	0.7	9.9	2.1	2.5	1.7	-0.1	2.1
NX4	12.3	2.7	6.9	1.2	3.2	0.7	1.0	2.8	2.4	3.3	36.0	1.3	35.1	3.7	4.4	41.1	-0.1	0.8
NX5	2.5	7.9	1.9	1.2	1.2	1.9	-0.1	1.2	1.2	1.2	14.0	0.6	12.3	0.5	2.2	17.3	-0.1	0.4
NX6	0.8	7.7	3.3	0.3	1.8	0.8	-0.1	1.6	1.4	1.8	19.2	0.8	18.3	2.8	0.7	14.3	-0.1	0.4
NX7	47.4	50.1	0.9	4.3	4.3	10.7	5.4	8.8	2.6	10.4	78.9	10.2	81.0	11.6	4.8	131.0	2.5	1.6
NX8	1.1	8.6	4.1	0.3	0.3	3.5	-0.1	1.8	1.5	2.0	18.6	1.0	15.9	2.6	2.6	10.1	-0.1	0.4
NX9	6.1	13.0	7.6	1.3	3.2	6.4	0.9	3.0	2.0	3.5	4.7	1.4	31.2	3.6	3.7	21.8	-0.1	0.5
NY1	1.0	6.0	1.3	0.3	1.2	2.0	-0.1	1.1	1.1	1.1	0.5	2.0	1.5	2.1	2.1	1.2	-0.1	2.0
NY10	7.8	14.9	2.8	0.3	1.7	3.8	-0.1	1.6	1.5	1.8	11.7	0.8	20.6	2.6	2.8	3.1	-0.1	0.4
NY11	3.1	10.3	1.2	1.2	1.0	2.0	-0.1	0.1	1.0	0.1	19.4	-0.1	1.3	-0.1	2.0	13.3	-0.1	0.1
NY12	0.4	4.4	1.6	1.2	1.2	1.2	-0.1	1.1	1.2	1.2	8.0	0.5	7.3	2.1	0.5	1.3	-0.1	2.2
NY13	0.4	5.0	1.3	-0.1	0.1	1.2	-0.1	-0.1	1.0	0.1	7.1	-0.1	6.0	-0.1	0.5	1.9	-0.1	0.1
NY14	3.9	11.5	3.8	0.3	1.8	3.3	-0.1	1.6	1.6	1.7	17.4	0.8	14.6	2.4	0.7	20.1	-0.1	3.1
NY15	5.7	15.8	2.5	0.3	1.6	3.4	-0.1	1.4	1.4	1.5	24.5	2.5	9.5	2.3	2.8	25.3	-0.1	0.5
NY2	0.4	3.8	1.3	-0.1	1.1	1.0	-0.1	-0.1	1.1	-0.1	5.6	1.9	0.6	-0.1	2.1	6.1	-0.1	0.4
NY3	0.4	5.0	1.1	-0.1	1.0	1.2	-0.1	-0.1	1.0	-0.1	5.9	-0.1	5.9	-0.1	0.5	1.7	-0.1	0.1
NY4	14.2	6.4	3.6	1.4	3.5	0.9	0.8	2.7	1.8	2.9	11.3	1.3	24.8	3.1	0.8	14.5	-0.1	3.6
NY4-R	22.9	12.6	0.5	1.6	4.6	7.7	2.3	3.5	2.1	3.9	22.5	1.4	44.7	4.2	1.1	30.3	-0.1	4.4
NY5	0.6	4.7	0.3	0.2	1.1	1.3	-0.1	1.1	1.1	1.1	8.0	2.0	6.9	2.0	2.2	1.1	-0.1	2.1
NY6	0.9	8.5	2.3	0.2	1.4	2.2	-0.1	1.2	1.4	1.3	14.2	2.2	12.5	2.3	0.7	1.9	-0.1	0.4
NY7	6.7	15.3	0.5	0.3	2.0	4.7	-0.1	1.6	1.8	1.7	20.6	0.8	20.1	2.6	1.0	26.4	-0.1	0.5
NY8	1.0	8.3	1.5	-0.3	1.2	1.8	-0.1	1.2	1.2	1.2	15.3	2.1	2.2	2.2	2.3	2.4	-0.1	0.4
NY9	1.6	0.2	1.5	-0.1	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	0.5	-0.1	3.3	-0.1	0.4	0.6	-0.1	1.9
NZ1	3.3	10.2	1.9	0.2	1.4	2.6	-0.1	1.3	1.2	1.3	12.6	0.6	1.2	2.1	2.2	13.7	-0.1	2.1
NZ10	4.8	14.3	4.0	0.3	1.8	3.5	-0.1	1.6	1.9	1.8	25.0	0.8	20.8	0.5	3.6	11.5	-0.1	2.2
NZ11	0.8	5.7	0.2	-0.1	0.1	1.2	-0.1	-0.1	1.1	-0.1	10.1	-0.1	1.4	0.5	2.1	9.2	-0.1	1.9
NZ12	0.6	5.5	1.5	0.2	0.2	1.6	-0.1	1.1	1.1	1.2	8.2	2.1	0.5	2.0	0.6	1.0	-0.1	2.1
NZ13	0.5	4.8	0.2	-0.1	1.0	1.2	-0.1	-0.1	1.1	-0.1	7.8	1.9	6.1	-0.1	2.1	0.9	-0.1	2.1
NZ2	1.0	7.8	2.0	0.6	3.0	0.7	-0.1	2.0	1.3	2.3	15.6	1.0	14.3	2.4	0.6	16.8	-0.1	2.4
NZ2-R	0.3	4.0	1.4	1.3	1.4	0.4	-0.1	1.2	1.1	1.3	7.7	0.6	7.7	2.0	0.5	1.5	-0.1	2.0
NZ3	0.5	7.4	2.3	0.3	1.7	2.2	-0.1	1.4	1.5	1.5	13.7	2.5	12.0	2.2	2.5	15.8	-0.1	0.5
NZ4	0.6	7.4	1.9	1.5	1.4	2.1	-0.1	1.3	1.2	1.4	12.4	0.6	10.6	2.1	0.5	13.4	-0.1	2.2
NZ5	0.7	8.7	2.0	1.4	1.4	2.2	-0.1	1.3	1.3	1.3	12.1	0.5	9.9	2.1	2.4	13.1	-0.1	2.2
NZ6	0.5	5.6	0.2	-0.2	1.2	1.7	-0.1	1.2	1.1	1.3	9.0	2.2	0.8	2.1	2.2	1.3	-0.1	2.1
NZ7	7.5	14.1	0.3	1.3	1.3	3.4	-0.1	1.3	1.3	1.3	13.7	0.6	8.0	0.4	2.5	2.6	-0.1	0.3
NZ8	0.6	7.4	4.5	1.2	1.3	1.7	-0.1	1.1	1.2	1.2	10.4	0.5	9.9	2.1	2.3	1.6	-0.1	0.3
NZ9	0.6	8.4	0.2	-0.1	-0.1	1.7	-0.1	-0.1	1.1	-0.1	10.6	-0.1	1.0	-0.1	2.1	1.6	-0.1	-0.1
LMB-QA	0.7	6.2	-0.1	-0.1	-0.1	2.4	-0.1	-0.1	-0.1	-0.1	6.1	-0.1	1.2	2.0	-0.1	8.5	-0.1	-0.1
LMB-QA	0.5	8.0	-0.1	-0.1	-0.1	1.6	-0.1	-0.1	-0.1	-0.1	6.9	-0.1	0.8	-0.1	-0.1	4.1	-0.1	-0.1
LMB-QA	0.5	3.2	-0.1	-0.1	-0.1	0.8	-0.1	-0.1	-0.1	-0.1	4.5	-0.1	0.6	-0.1	-0.1	4.4	-0.1	-0.1
LMB-QA	2.7	3.5	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	-0.1	-0.1	7.3	-0.1	1.2	-0.1	-0.1	7.4	-0.1	-0.1
LMB-QA	0.5	2.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	4.5	-0.1	0.9	-0.1	-0.1	4.7	-0.1	-0.1
LMB-QA	0.5	4.0	-0.1	-0.1	-0.1	0.9	-0.1	-0.1	-0.1	-0.1	5.6	-0.1	0.8	-0.1	-0.1	5.4	-0.1	-0.1
LMB-QA	0.4	3.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	3.2	-0.1	0.6	-0.1	-0.1	4.0	-0.1	-0.1
LMB-QA	0.5	2.9	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	-0.1	-0.1	8.9	-0.1	3.4	-0.1	-0.1	3.0	-0.1	-0.1
LMB-QA	0.4	3.2	-0.1	-0.1	-0.1	0.8	-0.1	-0.1	-0.1	-0.1	3.4	-0.1	0.7	-0.1	-0.1	3.5	-0.1	-0.1
LMB-QA	0.3	1.5	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	-0.1	-0.1	2.9	-0.1	2.8	-0.1	-0.1	3.0	-0.1	-0.1
LMB-QA	0.3	3.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	4.7	-0.1	0.6	-0.1	-0.1	4.5	-0.1	-0.1
LMB-QA	0.3	2.4	-0.1	-0.1	-0.1	0.1	-0.1	-0.1	-0.1	-0.1	4.0	-0.1	0.6	-0.1	-0.1	4.1	-0.1	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	073-HBA	074-HBA	075-HPB	076-LPH	077-MAR	078-ALK	079-LBI	080-LPH	081-MAR	082-LPH	083-HBA	084-HBA	085-LPH	086-LBI	087-MAR	088-HBA	089-THI	090-HPB

A14-06865

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
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	091-LBI	092-LPH	093-LA	094-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-LPH	101-MAR	102-MBI	103-LPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NA1	3.3	2.8	41.7	4.2	0.9	3.1	42.9	3.5	0.6	2.7	2.7	2.5	2.9	4.8	12.0	2.9	2.3	8.1
NA2	-0.1	-0.1	13.6	2.0	-0.1	2.1	14.1	-0.1	2.1	-0.1	2.1	1.2	2.1	2.6	4.9	1.7	2.0	7.4
NAA1	2.2	2.3	14.5	1.9	0.4	2.5	14.3	-0.1	0.5	2.3	2.2	1.5	0.8	3.0	5.4	-0.1	-0.1	7.4
NAA10	2.0	1.7	9.0	1.8	2.4	2.4	1.6	-0.1	2.3	2.2	2.1	1.3	2.3	2.7	3.8	-0.1	-0.1	-0.1
NAA11	4.4	1.6	25.2	4.6	2.5	2.4	25.7	2.0	2.2	2.2	2.6	3.2	2.3	3.9	16.7	1.8	2.0	7.5
NAA2	14.2	3.3	298.0	23.0	6.4	5.1	309.0	15.4	4.3	3.6	6.8	11.2	3.6	9.8	53.1	9.9	6.6	10.2
NAA3	3.3	1.5	28.3	3.5	0.8	2.3	30.3	-0.1	2.1	2.1	2.7	2.1	2.0	2.9	13.1	2.1	2.1	7.4
NAA4	2.2	1.6	22.4	2.4	2.5	2.2	22.9	1.0	2.1	2.1	2.1	1.4	2.3	2.7	5.8	1.7	2.0	-0.1
NAA5	2.5	1.6	18.5	2.6	3.2	2.3	18.2	0.5	2.2	2.2	2.3	1.8	2.2	3.4	8.1	1.7	2.1	7.6
NAA6	2.4	1.7	30.0	2.8	2.7	2.4	30.9	2.9	2.3	2.2	2.3	1.6	2.3	3.3	6.7	1.8	2.1	7.5
NAA7	2.1	1.6	19.5	2.2	0.5	2.3	19.5	-0.1	2.2	2.2	2.1	1.3	2.3	2.8	4.7	1.6	1.9	7.4
NAA8	2.0	1.6	16.7	2.0	2.2	2.1	18.7	-0.1	2.0	-0.1	2.0	1.2	2.1	2.4	4.1	1.5	-0.1	-0.1
NAA9	2.5	1.8	27.8	0.3	0.9	2.4	27.4	2.9	2.2	2.2	2.1	1.5	2.3	3.2	6.4	1.8	2.0	7.5
NB1	2.4	1.8	23.2	0.3	0.5	2.4	24.0	1.1	2.3	2.2	2.2	1.5	2.3	2.7	6.1	1.7	2.0	7.4
NB2	-0.1	-0.1	10.8	1.8	0.5	2.0	10.3	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	2.2	-0.1	-0.1	-0.1
NB3	-0.1	-0.1	7.5	1.7	0.4	2.0	7.1	-0.1	2.0	-0.1	1.1	1.1	2.1	2.2	2.6	-0.1	-0.1	-0.1
NB4	2.7	1.6	50.4	3.3	0.5	2.2	52.8	2.5	2.2	2.0	2.5	1.7	2.2	2.5	8.9	2.1	2.1	7.5
NBB1	-0.1	1.6	12.0	1.7	2.2	2.3	11.9	-0.1	2.2	2.1	2.0	1.1	2.2	2.4	2.6	-0.1	-0.1	-0.1
NBB10	4.5	1.6	57.0	6.0	1.1	2.4	60.3	4.5	2.3	2.2	2.8	3.4	2.3	5.3	16.6	4.0	2.5	8.1
NBB11	2.8	2.7	51.6	0.4	4.8	3.6	49.8	3.8	0.6	2.8	2.5	2.0	3.0	5.0	12.4	2.2	2.3	8.3
NBB2	3.6	1.6	42.3	4.3	1.4	2.3	41.1	3.5	2.1	2.1	2.4	2.4	2.2	3.2	11.8	2.3	2.1	7.4
NBB2-R	3.0	1.5	30.6	3.3	1.0	2.2	29.8	2.6	2.0	2.0	2.0	1.9	2.1	2.8	8.8	1.8	2.0	-0.1
NBB3	2.8	1.5	27.3	0.4	1.2	2.2	27.1	2.9	2.0	2.0	2.1	1.7	2.1	2.9	6.5	1.6	1.9	-0.1
NBB4	-0.1	-0.1	6.4	-0.1	0.4	-0.1	5.9	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NBB5	-0.1	1.7	16.0	1.8	2.2	2.2	14.9	-0.1	2.1	2.0	1.9	1.1	2.1	2.4	3.3	1.4	-0.1	-0.1
NBB6	2.8	2.0	39.3	3.2	4.0	2.9	39.6	3.3	2.6	2.4	2.4	1.7	2.3	3.2	1.7	1.8	2.2	7.6
NBB7	-0.1	-0.1	7.4	1.6	1.9	-0.1	7.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.0	-0.1	-0.1	-0.1	-0.1
NBB8	2.3	1.5	14.0	2.1	2.5	2.1	2.3	-0.1	2.1	2.1	2.2	1.5	2.1	2.5	5.1	-0.1	-0.1	-0.1
NBB9	3.5	1.6	34.8	3.5	2.2	2.3	33.3	2.7	2.2	2.2	2.3	2.4	2.2	3.1	10.5	1.6	2.0	7.4
NC1	-0.1	-0.1	15.4	1.8	0.3	2.0	15.3	-0.1	-0.1	-0.1	2.0	1.0	-0.1	2.1	2.6	1.3	-0.1	-0.1
NC2	-0.1	1.5	19.4	2.0	0.4	2.1	18.9	-0.1	2.0	2.0	2.1	1.2	2.1	2.3	3.6	1.4	-0.1	-0.1
NC3	-0.1	-0.1	20.1	2.0	2.1	2.0	20.0	-0.1	2.0	-0.1	2.0	1.2	-0.1	2.1	3.7	1.4	1.9	-0.1
NC4	-0.1	-0.1	5.7	1.6	2.0	1.9	5.6	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.0	-0.1	-0.1	-0.1
NC5	2.1	1.4	21.0	2.2	0.5	2.1	21.0	0.9	2.0	1.9	2.1	1.2	2.1	2.3	3.9	1.5	1.9	-0.1
NC6	-0.1	1.9	15.3	2.1	0.7	2.6	15.0	-0.1	2.3	2.2	2.0	1.1	2.2	2.4	3.7	1.3	-0.1	-0.1
NC7	2.0	1.4	10.5	1.9	2.2	2.0	11.0	-0.1	2.0	-0.1	2.1	1.3	2.0	2.2	3.7	-0.1	1.9	-0.1
NC8	-0.1	1.4	10.2	1.8	2.2	2.0	10.0	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.2	2.7	-0.1	-0.1	-0.1
NCC1	11.8	1.9	389.0	20.5	1.0	3.7	417.0	21.1	3.5	3.0	9.3	9.2	3.1	7.9	51.9	14.8	7.8	12.0
NCC1-R	12.2	1.9	357.0	20.8	1.0	3.8	390.0	19.7	3.1	2.9	9.0	8.4	2.9	8.2	49.8	14.6	7.1	11.8
NCC2	8.2	3.7	146.0	12.9	3.8	0.9	154.0	6.9	3.8	3.5	6.3	1.3	14.7	36.3	9.6	3.9	10.2	10.2
NCC3	3.4	2.0	47.7	4.1	4.6	2.7	46.5	3.2	2.6	2.4	2.6	2.3	2.4	3.7	12.6	2.3	2.1	7.8
NCC4	7.1	2.0	75.6	8.9	6.6	2.7	75.9	4.3	2.7	2.5	8.3	5.6	2.5	5.6	29.7	6.8	2.4	8.3
NCC5	3.0	1.9	46.2	3.4	5.6	0.4	41.1	3.3	2.4	2.3	2.4	2.0	2.3	3.1	10.7	1.9	2.0	7.5
NCC6	4.3	1.6	26.9	4.2	2.0	2.8	26.3	2.2	2.2	2.2	2.9	3.9	2.2	4.0	19.1	1.4	2.1	7.8
ND1	-0.1	1.5	9.1	1.7	0.6	2.1	8.2	-0.1	2.1	2.0	2.0	1.2	2.1	2.3	2.9	-0.1	-0.1	-0.1
ND10	-0.1	-0.1	8.6	-0.1	0.3	2.0	8.6	-0.1	-0.1	-0.1	1.9	1.1	-0.1	2.2	1.9	-0.1	-0.1	-0.1
ND2	2.7	1.6	23.7	2.5	1.1	2.2	22.5	2.1	2.1	2.1	2.3	1.9	2.2	3.0	7.7	1.3	2.0	7.4
ND3	-0.1	1.4	9.2	-0.1	2.0	2.1	1.7	-0.1	2.0	-0.1	2.0	1.0	-0.1	2.0	2.1	-0.1	-0.1	-0.1
ND4	3.3	1.7	42.6	3.5	3.7	2.6	28.4	2.6	2.5	2.4	3.1	2.4	2.3	2.8	13.1	1.8	2.3	7.7
ND5	-0.1	-0.1	7.6	1.6	0.4	-0.1	7.4	-0.1	-0.1	-0.1	-0.1	1.9	1.1	-0.1	2.1	2.2	-0.1	-0.1
ND6	-0.1	1.6	10.2	1.9	0.4	2.1	1.6	-0.1	2.1	2.0	2.0	1.2	2.1	2.2	3.4	1.4	-0.1	-0.1
ND7	-0.1	-0.1	8.4	1.9	-0.1	-0.1	8.4	-0.1	-0.1	-0.1	2.0	1.0	-0.1	2.1	2.0	-0.1	-0.1	-0.1
ND8	-0.1	1.5	12.1	1.9	2.2	2.1	11.9	-0.1	2.0	-0.1	1.9	1.2	2.1	2.2	3.3	1.4	-0.1	-0.1
ND8-R	2.1	1.5	15.4	0.2	0.6	2.1	15.2	-0.1	2.1	2.0	2.1	1.3	2.1	2.2	4.1	-0.1	-0.1	-0.1
ND9	2.1	1.6	12.8	2.0	0.5	2.1	12.5	-0.1	2.0	2.0	2.0	1.3	2.1	0.6	4.0	1.4	-0.1	-0.1
NDD1	3.3	3.0	22.9	3.5	1.1	3.0	21.3	0.4	0.5	2.7	2.6	2.2	1.0	3.3	10.8	2.0	2.0	7.8
NDD2	2.3	1.6	22.3	2.4	0.8	2.3	21.5	2.4	2.1	2.1	2.1	1.3	2.1	2.6	5.3	1.4	1.9	-0.1
NDD3	1.9	1.4	15.4	2.1	2.5	2.8	15.5	-0.1	0.6	2.3	2.2	1.3	2.5	2.2	4.4	1.5	2.0	7.4
NDD4	2.1	1.9	11.8	2.0	0.6	2.6	11.5	-0.1	2.4	2.2	2.1	1.5	2.4	3.2	5.0	1.4	1.9	-0.1
NDD5	2.2	1.7	16.9	2.1	0.4	2.8	17.0	-0.1	2.1	2.1	2.8	1.4	2.1	2.5	6.3	1.4	-0.1	-0.1
NE1	5.0	2.0	63.0	6.5	3.8	2.8	62.7	4.6	2.5	2.4	2.8	3.5	2.5	4.4	16.0	3.4	2.5	8.0
NE10	-0.1	-0.1	4.4	-0.1	1.9	1.9	4.0	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NE11	-0.1	1.4	11.2	-0.1	0.4	2.0	11.0	-0.1	-0.1	2.0	1.9	2.0	1.2	2.1	2.2	2.6	-0.1	-0.1
NE2	-0.1	-0.1	8.1	2.0	-0.1	-0.1	7.8	-0.1	-0.1	-0.1	-0.1	0.4	-0.1	-0.1	2.1	-0.1	-0.1	-0.1
NE3	-0.1	1.5	15.4	1.8	0.4	2.1	15.1	-0.1	2.0	-0.1	1.9	1.0	2.1	2.1	2.5	-0.1	-0.1	-0.1
NE4	-0.1	1.6	13.6	1.8	2.1	2.2	13.7	-0.1	2.2	2.1	2.0	1.1	2.2	2.1	2.4	-0.1	-0.1	-0.1
NE4-R	-0.1	-0.1	5.8	-0.1	1.9	1.9	5.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1

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Date: November 25, 2014
R=Replicate Sample

	D91-LBI	092-LPH	093-LA	D94-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-LPH	101-MAR	102-MBI	103-LPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NE5	2.2	1.6	11.7	2.0	3.0	2.4	1.6	2.0	2.2	2.1	2.1	1.4	2.2	2.6	4.2	-0.1	-0.1	-0.1
NE6	-0.1	1.4	7.5	1.6	2.2	2.1	1.2	-0.1	-0.1	-0.1	1.9	1.0	-0.1	2.0	2.3	-0.1	-0.1	-0.1
NE7	-0.1	-0.1	5.4	1.7	0.3	-0.1	5.3	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.1	1.3	-0.1	-0.1
NE8	-0.1	-0.1	7.7	-0.1	0.4	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NE9	-0.1	1.5	6.0	1.6	2.2	2.2	7.3	-0.1	2.0	2.0	1.9	1.0	2.1	2.3	-0.1	-0.1	-0.1	-0.1
NEE1	-0.1	1.6	13.1	1.7	0.5	2.1	10.7	-0.1	2.0	2.0	2.0	1.2	2.1	2.6	3.0	-0.1	-0.1	-0.1
NEE2	2.6	1.5	30.0	0.4	0.8	2.3	29.8	1.3	2.1	2.0	2.3	1.5	2.1	2.7	6.0	1.7	1.9	-0.1
NEE3	2.7	1.7	36.9	0.4	0.9	2.4	36.0	1.4	2.2	2.2	2.3	1.5	2.1	2.7	7.4	1.8	2.0	7.3
NEE4	2.1	1.4	16.2	2.0	0.4	2.4	13.1	-0.1	2.1	2.1	2.4	1.2	2.7	11.3	3.4	1.4	-0.1	7.6
NEE5	-0.1	-0.1	14.0	1.9	2.2	2.0	13.5	-0.1	-0.1	-0.1	1.9	1.1	2.0	2.3	2.9	1.3	-0.1	-0.1
NEE6	2.4	2.0	19.0	2.5	2.8	2.7	19.3	-0.1	2.4	2.3	2.2	1.5	2.4	2.9	5.9	1.8	2.0	7.4
NEE7	2.0	2.1	12.7	1.7	0.5	2.4	12.4	-0.1	2.3	2.3	2.1	1.4	0.7	2.9	3.9	-0.1	-0.1	7.1
NF1	3.4	1.7	27.5	3.4	4.1	2.5	15.2	2.1	2.4	2.3	2.1	3.1	2.3	3.7	17.0	1.5	2.1	7.8
NF10	2.3	1.6	27.3	2.8	0.4	2.2	28.2	1.4	2.1	2.0	2.1	1.4	2.2	3.7	6.4	2.2	2.1	7.6
NF11	4.2	2.1	116.0	0.5	0.4	8.1	123.0	3.6	0.6	2.6	8.5	2.6	2.7	4.1	14.3	5.3	3.1	8.9
NF2	-0.1	-0.1	10.3	1.7	0.4	-0.1	10.4	-0.1	-0.1	-0.1	2.0	1.1	-0.1	2.1	3.0	-0.1	-0.1	-0.1
NF3	-0.1	1.3	6.9	1.6	0.4	2.0	6.7	-0.1	2.0	-0.1	2.0	1.2	2.1	2.3	2.9	-0.1	-0.1	-0.1
NF4	2.4	1.5	23.7	2.6	0.6	2.2	25.9	-0.1	2.1	2.0	2.3	1.6	2.2	2.5	8.2	1.9	2.1	7.3
NF5	2.2	1.4	23.6	2.4	0.4	2.1	24.6	1.0	2.0	2.0	2.2	1.3	2.1	2.5	5.3	1.3	2.1	7.3
NF6	2.9	-0.1	51.0	0.3	0.5	2.2	54.3	2.6	2.1	2.1	2.6	1.9	2.1	2.8	9.3	3.0	2.3	7.4
NF7	4.0	3.0	86.2	5.2	5.5	3.7	86.7	6.2	0.6	3.0	3.0	2.9	1.1	7.4	18.2	3.6	0.4	9.0
NF8	-0.1	1.6	12.5	1.7	0.3	2.2	12.5	-0.1	2.1	2.0	2.0	1.1	2.2	2.4	2.5	-0.1	-0.1	-0.1
NF8-R	3.0	2.1	43.2	0.4	0.5	2.6	46.2	1.9	2.5	2.4	2.7	2.0	2.5	3.4	11.0	2.5	2.3	7.9
NF9	2.5	1.6	28.4	0.3	0.5	2.2	30.3	1.1	2.2	2.0	2.3	1.7	2.2	2.9	8.5	2.2	2.1	7.4
NFF1	2.3	1.6	22.3	2.4	0.6	2.2	22.6	-0.1	2.2	2.1	2.2	1.4	2.2	2.8	6.4	1.7	2.0	7.5
NFF2	3.5	1.9	33.3	4.0	1.3	2.5	33.0	2.8	2.3	2.3	2.7	2.7	2.4	3.8	13.7	2.3	2.1	7.7
NFF3	2.7	1.3	15.0	2.9	0.7	2.0	16.4	-0.1	2.0	2.0	2.3	1.8	2.1	2.9	8.8	1.9	2.1	7.4
NFF4	3.0	2.3	44.4	0.3	1.6	2.8	43.8	3.7	2.5	2.4	2.5	2.0	2.6	4.2	9.2	2.4	2.3	8.0
NFF5	2.4	1.9	16.4	2.2	1.0	2.4	15.8	0.5	2.2	2.1	2.2	1.6	2.3	3.0	5.9	1.6	-0.1	-0.1
NFF6	-0.1	-0.1	14.1	1.8	2.2	2.0	13.7	-0.1	1.9	-0.1	1.9	1.1	-0.1	2.2	2.6	1.3	-0.1	-0.1
NFF6-R	-0.1	-0.1	9.4	1.7	0.5	2.0	8.9	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.2	-0.1	-0.1	-0.1
NFF7	-0.1	1.6	14.2	1.9	0.5	2.1	13.5	2.1	2.0	2.0	2.0	1.1	2.1	2.3	3.2	1.3	-0.1	-0.1
NFF8	-0.1	1.5	14.4	1.6	2.2	2.2	1.9	-0.1	2.1	2.0	2.4	1.1	2.2	2.5	2.5	-0.1	-0.1	-0.1
NG1	-0.1	-0.1	4.8	-0.1	-0.1	-0.1	4.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NG10	2.0	1.4	11.3	1.7	0.6	2.1	11.0	1.8	2.0	2.0	2.0	1.2	2.1	2.4	3.1	-0.1	-0.1	-0.1
NG11	-0.1	-0.1	6.1	-0.1	0.4	-0.1	5.9	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NG2	-0.1	-0.1	7.3	-0.1	0.5	-0.1	7.3	-0.1	-0.1	-0.1	1.9	0.4	1.1	-0.1	2.2	2.1	-0.1	-0.1
NG2-R	-0.1	-0.1	8.7	-0.1	0.4	-0.1	8.6	-0.1	-0.1	-0.1	1.9	2.0	1.1	-0.1	2.1	2.1	-0.1	-0.1
NG3	-0.1	-0.1	5.6	-0.1	2.0	-0.1	5.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.0	-0.1	-0.1	-0.1	-0.1
NG4	-0.1	1.5	6.8	-0.1	0.5	2.1	5.8	-0.1	2.0	2.0	-0.1	1.1	2.1	2.4	-0.1	-0.1	-0.1	-0.1
NG5	2.6	1.9	43.2	3.2	5.4	2.8	42.9	0.4	0.4	2.4	2.3	1.7	2.4	3.1	8.1	1.9	2.2	7.6
NG6	-0.1	1.5	11.4	-0.1	2.4	2.0	1.5	2.0	2.0	2.0	-0.1	1.1	2.0	2.3	2.6	-0.1	-0.1	-0.1
NG7	-0.1	-0.1	3.0	-0.1	-0.1	-0.1	2.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NG8	-0.1	-0.1	7.2	-0.1	0.4	-0.1	6.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NG9	-0.1	1.5	16.1	2.0	0.7	2.4	14.9	2.1	2.0	1.9	1.0	1.1	-0.1	2.3	3.0	1.4	-0.1	-0.1
NGG1	2.3	1.5	16.4	2.2	0.6	2.1	16.4	2.1	2.1	2.2	2.2	1.6	2.2	2.7	5.7	1.5	-0.1	7.3
NGG10	3.3	1.5	30.6	3.8	1.1	2.1	30.3	2.8	2.1	2.1	2.4	2.2	2.1	2.7	10.0	2.0	2.0	7.4
NGG2	-0.1	1.4	10.3	1.7	0.5	2.0	9.1	-0.1	-0.1	1.9	1.9	1.2	-0.1	2.3	3.1	-0.1	-0.1	-0.1
NGG3	7.4	1.9	80.1	10.3	2.3	2.8	90.0	3.7	2.7	2.6	5.2	6.3	2.6	2.6	28.6	4.8	3.4	8.5
NGG4	2.9	1.7	33.3	3.2	1.3	2.4	33.3	2.8	2.2	2.1	2.3	1.8	2.1	2.8	8.7	1.8	2.0	-0.1
NGG5	2.3	1.6	17.8	2.3	0.7	2.3	16.7	-0.1	2.1	2.0	2.0	1.3	2.0	2.7	4.7	1.4	-0.1	-0.1
NGG6	2.4	1.5	20.9	2.4	3.2	2.2	20.7	-0.1	2.1	2.1	2.1	1.5	2.2	3.0	5.7	1.5	2.0	7.2
NGG7	2.4	1.6	21.6	2.4	0.6	2.2	21.4	2.2	2.1	2.1	2.2	1.6	2.2	3.0	5.9	1.6	1.9	7.4
NGG8	3.0	1.5	32.7	3.2	0.9	2.2	32.7	3.0	2.1	2.0	2.4	1.8	2.0	2.6	8.6	1.8	1.9	-0.1
NGG9	2.0	-0.1	10.1	1.9	2.2	2.0	9.9	-0.1	-0.1	-0.1	1.9	1.3	-0.1	2.4	3.8	-0.1	-0.1	-0.1
NH1	-0.1	-0.1	10.4	1.7	2.4	2.0	10.3	-0.1	1.9	-0.1	-0.1	1.1	-0.1	2.2	2.4	-0.1	-0.1	-0.1
NH10	-0.1	1.5	7.1	-0.1	0.4	2.0	6.9	-0.1	2.0	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NH10-R	-0.1	1.8	7.2	-0.1	0.5	2.2	6.1	-0.1	2.0	-0.1	-0.1	1.0	2.1	2.3	-0.1	-0.1	-0.1	-0.1
NH11	-0.1	1.7	15.3	1.9	0.7	2.2	14.6	2.3	2.1	2.1	2.0	1.1	2.2	2.3	2.8	1.3	-0.1	-0.1
NH2	-0.1	1.4	10.0	1.6	0.6	2.0	8.9	1.9	2.0	2.0	2.0	1.1	-0.1	2.3	2.1	-0.1	-0.1	-0.1
NH3	-0.1	-0.1	4.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NH4	-0.1	1.5	14.7	1.8	2.1	2.1	13.8	-0.1	-0.1	-0.1	-0.1	1.0	2.1	2.3	2.3	-0.1	-0.1	-0.1
NH5	-0.1	1.5	8.0	-0.1	2.1	2.1	6.9	-0.1	2.0	-0.1	-0.1	1.0	2.1	2.2	-0.1	-0.1	-0.1	-0.1
NH6	-0.1	1.9	7.0	1.6	0.3	2.3	6.5	-0.1	2.1	2.1	1.9	1.1	2.2	2.6	-0.1	-0.1	-0.1	-0.1
NH7	2.1	3.2	26.3	2.1	0.6	3.3	26.0	2.5	0.6	2.8	2.2	1.2	1.1	3.1	4.1	1.5	2.0	7.6
NH8	-0.1	1.8	8.9	1.6	0.4	2.4	8.6	-0.1	2.2	2.1	2.0	1.1	2.3	2.5	2.1	-0.1	-0.1	-0.1
NH9	-0.1	1.9	9.9	1.6	0.4	2.3	9.5	-0.1	0.4	2.2	2.0	1.1	2.3	2.5	2.3	-0.1	-0.1	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	091-LBI	092-LPH	093-LA	094-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-EPH	101-MAR	102-MBI	103-EPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NHH1	2.2	1.4	17.0	2.1	0.5	2.0	15.8	-0.1	-0.1	1.9	2.0	1.3	-0.1	2.4	4.3	1.5	-0.1	-0.1
NHH10	-0.1	1.4	11.8	1.7	0.4	2.0	7.6	-0.1	2.0	2.0	2.0	1.2	-2.1	2.4	3.2	-0.1	-0.1	-0.1
NHH11	2.5	1.4	15.4	2.4	0.5	2.0	15.4	-0.1	2.0	2.0	2.2	1.8	2.1	2.6	7.0	1.5	-0.1	-0.1
NHH11-R	2.2	1.4	11.4	2.0	0.5	2.0	1.7	-0.1	2.0	2.0	2.1	1.5	-0.1	2.5	4.7	1.4	-0.1	-0.1
NHH12	2.1	1.4	17.9	0.2	2.3	2.0	17.0	-0.1	-0.1	1.9	2.0	1.3	-0.1	2.3	4.0	1.4	-0.1	-0.1
NHH2	5.0	1.8	78.5	7.0	3.6	2.5	80.1	4.8	2.3	2.2	3.8	4.1	2.3	3.9	19.2	3.7	2.6	7.9
NHH3	2.1	1.5	20.3	2.1	2.6	2.1	19.9	-0.1	2.0	2.0	2.1	1.3	2.1	2.7	4.8	1.5	-0.1	-0.1
NHH4	3.7	1.6	34.2	4.4	1.5	2.4	34.8	3.3	2.1	2.1	2.5	2.3	2.1	4.4	12.4	2.5	2.1	7.8
NHH5	3.6	1.5	41.4	4.2	1.4	2.3	41.4	3.2	2.1	2.0	2.5	2.3	2.0	2.7	11.9	2.1	2.1	-0.1
NHH6	2.1	1.5	12.2	2.0	0.5	2.0	11.8	-0.1	-0.1	2.0	2.0	1.3	2.1	2.6	3.7	1.4	-0.1	-0.1
NHH7	2.4	1.5	16.8	2.2	0.9	2.1	16.1	2.2	2.0	2.0	2.1	1.5	2.1	2.5	4.8	1.5	-0.1	-0.1
NHH8	2.2	1.4	20.2	2.3	2.4	2.0	20.4	-0.1	2.0	2.0	2.1	1.4	2.1	2.4	4.9	1.5	1.9	-0.1
NHH9	2.3	1.4	19.2	2.3	2.4	2.2	19.9	-0.1	2.0	1.9	2.2	1.4	2.0	2.4	1.2	1.4	1.9	7.3
NI1	4.4	2.2	112.0	0.7	3.4	0.4	112.0	7.7	2.7	2.6	2.9	2.6	2.5	5.4	11.6	3.1	2.4	8.2
NI10	-0.1	1.4	13.6	1.8	0.6	2.0	13.4	0.5	2.0	2.0	2.1	1.2	-0.1	2.3	2.9	-0.1	-0.1	-0.1
NI2	-0.1	1.3	8.0	1.6	2.2	2.0	7.7	-0.1	2.0	2.0	2.0	1.0	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NI3	-0.1	1.4	12.5	1.7	2.1	2.0	12.0	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.2	2.4	-0.1	-0.1	-0.1
NI4	2.5	1.9	22.2	2.2	3.8	2.6	24.9	2.1	2.5	2.4	2.6	1.9	2.4	3.6	8.3	1.4	2.0	7.6
NI5	-0.1	1.8	13.4	1.7	3.0	2.4	12.3	2.1	2.2	2.2	2.0	1.3	2.3	3.0	3.8	-0.1	-0.1	-0.1
NI5-R	-0.1	1.7	11.1	1.6	2.7	2.3	10.3	2.0	2.1	2.1	2.0	1.2	2.2	2.8	0.8	-0.1	-0.1	-0.1
NI6	-0.1	1.3	11.4	1.7	0.5	2.0	10.5	-0.1	-0.1	-0.1	1.9	1.0	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NI7	4.1	1.7	104.0	10.4	2.6	2.6	107.0	7.0	2.5	2.3	2.7	2.3	2.2	3.8	10.4	2.6	2.3	7.5
NI8	-0.1	-0.1	5.8	-0.1	0.4	-0.1	5.3	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NI9	-0.1	1.6	15.8	1.8	0.5	2.3	15.4	-0.1	0.4	2.1	2.0	1.1	2.2	2.4	3.0	-0.1	-0.1	-0.1
NII1	-0.1	-0.1	3.5	-0.1	-0.1	2.0	3.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NII10	2.4	1.5	22.1	2.4	0.6	2.1	22.7	0.9	2.1	2.0	2.1	1.4	2.1	2.5	5.6	1.7	2.0	-0.1
NII11	-0.1	1.6	11.8	1.9	0.6	2.2	11.0	2.3	2.1	2.1	2.0	1.1	2.1	2.4	2.3	1.4	-0.1	-0.1
NII11-R	-0.1	1.6	18.2	2.2	0.6	2.3	17.9	2.7	2.1	2.0	2.0	1.1	2.0	2.5	3.8	1.6	1.9	7.4
NII12	-0.1	1.5	11.7	2.0	2.3	2.1	11.6	-0.1	2.0	2.0	2.0	1.2	2.1	2.4	3.6	1.5	1.9	-0.1
NII13	3.6	1.9	50.4	0.4	1.1	2.4	50.1	4.5	2.3	2.2	2.5	1.8	2.3	2.9	7.5	2.6	2.1	7.5
NII14	-0.1	-0.1	6.2	1.6	0.4	-0.1	5.9	-0.1	-0.1	-0.1	-0.1	0.9	-0.1	2.1	-0.1	-0.1	-0.1	-0.1
NII2	2.1	1.7	20.8	1.9	2.4	2.4	21.0	-0.1	2.3	2.1	2.1	1.2	2.3	2.4	3.4	-0.1	2.0	7.4
NII3	3.0	1.6	30.0	3.1	3.5	2.3	31.5	0.5	2.3	2.3	2.7	1.9	2.2	2.7	9.2	1.6	2.1	7.5
NII4	3.8	1.5	84.9	4.5	3.8	2.5	89.1	5.3	2.4	2.3	2.8	2.0	2.2	2.7	11.0	2.4	2.4	7.5
NII5	2.2	1.5	21.3	2.3	0.4	2.1	21.4	1.0	2.0	2.0	2.1	1.2	2.1	2.3	4.5	1.5	1.9	-0.1
NII6	-0.1	-0.1	6.3	1.6	0.4	-0.1	7.6	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.2	-0.1	-0.1	-0.1
NII7	-0.1	-0.1	7.5	-0.1	0.5	-0.1	0.9	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NII8	3.3	1.5	35.4	3.8	1.5	2.3	36.9	2.8	2.3	2.2	2.6	2.0	2.2	2.7	11.3	2.3	2.1	7.4
NII9	-0.1	-0.1	7.7	1.6	0.5	2.0	7.1	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.1	-0.1	-0.1	-0.1
NJ1	-0.1	1.6	17.9	2.0	2.6	2.3	17.9	-0.1	2.2	2.1	2.0	1.1	2.2	2.8	2.8	-0.1	1.9	7.3
NJ2	2.4	1.6	32.1	2.6	3.0	2.2	32.4	3.1	2.2	2.1	2.2	1.4	2.2	2.9	5.7	1.7	2.0	-0.1
NJ3	2.2	1.6	29.5	2.4	2.6	2.3	29.8	1.1	2.2	2.1	2.2	1.1	2.0	2.5	4.9	1.6	1.9	-0.1
NJ4	2.2	1.6	30.3	2.3	2.6	2.2	29.9	1.2	2.2	2.1	2.1	1.2	2.2	2.6	4.4	1.6	1.9	-0.1
NJ5	-0.1	-0.1	14.8	2.0	0.6	2.0	14.0	-0.1	2.0	-0.1	2.0	1.1	-0.1	2.2	2.6	1.4	-0.1	-0.1
NJ6	-0.1	-0.1	10.8	1.7	2.1	2.0	10.2	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NJ7	2.0	1.5	9.6	1.7	2.6	2.2	2.0	-0.1	2.2	2.1	2.1	1.2	2.2	2.8	3.0	-0.1	-0.1	-0.1
NJJ1	-0.1	1.5	12.6	1.6	0.7	2.2	7.1	2.0	2.1	2.1	2.1	1.2	2.2	2.4	3.1	-0.1	-0.1	7.3
NJJ10	14.3	5.4	106.0	19.4	1.7	5.0	119.0	3.0	0.6	4.1	7.6	12.9	1.7	15.3	54.3	10.5	1.1	10.9
NJJ11	-0.1	-0.1	8.2	1.6	0.4	2.0	7.8	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.1	-0.1	-0.1	-0.1
NJJ12	2.4	1.6	14.5	2.3	0.7	2.3	14.8	-0.1	2.0	2.1	2.1	2.2	1.7	2.3	2.9	7.1	1.5	7.1
NJJ13	5.6	1.7	60.9	6.6	1.3	2.4	62.7	3.6	2.3	2.3	3.5	4.2	2.3	3.2	18.3	3.1	2.3	7.7
NJJ14	-0.1	1.4	6.5	1.4	0.4	2.0	7.7	-0.1	-0.1	-0.1	1.9	1.1	-0.1	2.1	2.2	-0.1	-0.1	-0.1
NJJ2	2.1	1.5	10.4	1.8	1.0	2.1	9.8	2.1	2.1	2.0	2.1	1.3	2.1	2.4	3.5	-0.1	-0.1	-0.1
NJJ3	-0.1	-0.1	8.0	1.6	0.4	-0.1	7.4	-0.1	1.9	1.9	-0.1	1.1	-0.1	-0.1	2.0	-0.1	-0.1	-0.1
NJJ4	4.4	1.5	77.4	0.5	1.5	2.4	78.6	5.6	2.4	2.1	2.8	2.2	2.0	2.4	10.5	3.0	2.2	7.5
NJJ5	-0.1	1.3	10.1	1.7	0.5	2.0	9.5	-0.1	2.0	2.0	2.0	1.2	-0.1	2.2	2.7	-0.1	-0.1	-0.1
NJJ6	2.1	1.5	14.9	2.0	0.5	2.0	14.0	-0.1	2.0	1.9	2.0	1.2	2.1	2.3	3.7	1.5	-0.1	-0.1
NJJ7	-0.1	-0.1	8.6	-0.1	0.3	2.0	8.0	-0.1	2.0	-0.1	-0.1	1.0	2.1	2.1	-0.1	-0.1	-0.1	-0.1
NJJ8	-0.1	1.3	9.2	1.7	0.5	-0.1	1.2	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.1	2.6	-0.1	-0.1	-0.1
NJJ8-R	2.2	1.4	14.5	1.9	0.6	2.0	10.6	2.0	2.0	1.9	2.1	1.4	-0.1	2.1	4.6	-0.1	-0.1	-0.1
NJJ9	5.2	1.7	135.0	0.6	2.3	2.8	138.0	9.0	2.5	2.3	3.2	2.9	2.4	3.6	13.2	4.1	2.8	7.9
NK1	-0.1	1.5	15.7	1.8	2.2	2.1	15.0	-0.1	2.0	2.0	4.9	1.0	2.1	2.4	2.3	-0.1	-0.1	-0.1
NK2	-0.1	1.4	7.4	1.6	0.4	2.0	7.4	-0.1	-0.1	1.9	1.9	1.1	-0.1	2.4	2.4	-0.1	-0.1	-0.1
NK3	-0.1	1.5	16.5	2.0	2.2	2.1	16.3	-0.1	2.1	2.0	2.1	1.1	2.1	2.5	3.2	1.5	-0.1	-0.1
NK4	-0.1	1.6	9.8	1.5	0.6	2.2	10.6	2.0	2.1	2.1	2.0	1.2	2.2	2.9	3.1	-0.1	-0.1	-0.1
NK4-R	-0.1	1.5	8.8	-0.1	0.5	2.1	8.4	-0.1	2.0	2.0	2.0	1.2	2.1	2.6	2.4	-0.1	-0.1	-0.1
NK5	-0.1	1.6	13.6	1.8	2.2	2.2	12.6	-0.1	2.1	2.0	1.9	1.0	2.1	2.5	2.5	-0.1	-0.1	-0.1

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	091-LBI	092-LPH	093-LA	094-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-EPH	101-MAR	102-MBI	103-EPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NK6	-0.1	1.5	12.4	1.7	2.2	2.1	1.7	-0.1	2.1	2.0	2.0	1.1	0.6	2.4	2.7	1.3	-0.1	-0.1
NK7	-0.1	1.3	13.3	1.9	2.4	2.1	13.4	-0.1	2.0	1.9	2.0	1.1	2.1	2.4	3.1	-0.1	-0.1	-0.1
NKK1	1.9	1.3	9.0	1.7	0.4	2.0	7.8	-0.1	-0.1	-0.1	2.0	1.4	-0.1	2.2	4.0	-0.1	-0.1	-0.1
NKK10	-0.1	-0.1	13.5	1.9	0.5	2.0	13.1	-0.1	-0.1	-0.1	1.9	1.2	-0.1	2.2	3.1	1.4	-0.1	-0.1
NKK11	2.6	1.6	14.5	2.7	0.9	2.3	13.0	0.4	2.1	2.0	2.1	1.5	2.1	2.6	7.6	1.6	1.9	7.3
NKK2	2.3	1.4	12.0	2.1	2.4	2.1	12.1	-0.1	2.1	2.0	2.1	1.5	2.1	2.5	5.5	1.4	1.9	7.3
NKK3	4.6	1.5	26.4	4.8	0.9	2.2	27.8	2.1	2.2	2.1	8.0	3.9	2.2	3.2	20.6	1.9	2.2	7.7
NKK4	18.8	1.9	198.0	26.9	4.4	3.1	218.0	8.2	3.0	2.8	8.3	14.8	2.8	6.8	61.8	14.1	5.1	10.0
NKK5	10.5	1.8	97.5	14.3	2.6	2.8	107.0	4.3	2.5	2.5	5.0	7.5	2.3	4.7	39.7	5.4	3.3	8.3
NKK6	3.8	1.5	19.5	3.9	0.9	2.2	19.5	2.1	2.0	2.0	2.5	2.9	2.2	3.2	13.9	1.5	2.0	7.5
NKK7	3.5	1.4	17.5	3.3	0.8	2.1	18.2	0.5	2.1	2.0	2.5	3.0	2.1	3.0	14.3	1.4	2.0	7.4
NKK8	2.2	1.5	16.1	2.1	2.3	2.2	16.5	-0.1	2.2	2.1	2.1	1.4	2.2	2.4	4.8	1.5	2.0	7.3
NKK9	2.2	1.5	17.9	2.2	0.4	2.1	17.7	-0.1	2.0	-0.1	2.1	1.3	2.1	2.3	4.4	1.5	-0.1	-0.1
NL1	-0.1	-0.1	7.7	1.6	0.3	-0.1	7.3	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.2	2.8	-0.1	-0.1	-0.1
NL2	-0.1	-0.1	8.9	1.6	0.3	2.0	8.6	-0.1	-0.1	-0.1	2.0	1.3	-0.1	2.4	3.4	-0.1	-0.1	-0.1
NL3	-0.1	-0.1	8.2	1.6	0.4	2.0	8.2	-0.1	2.0	1.9	2.0	1.3	2.1	2.6	3.7	-0.1	-0.1	-0.1
NL3-R	2.2	1.6	12.9	1.9	0.5	2.2	12.9	-0.1	2.1	2.1	2.2	1.7	2.2	3.0	6.2	-0.1	-0.1	7.4
NL4	-0.1	-0.1	10.0	1.7	0.4	-0.1	9.8	-0.1	-0.1	-0.1	2.0	1.2	-0.1	2.2	3.0	-0.1	-0.1	-0.1
NLL1	2.6	1.5	13.9	2.2	0.6	2.1	15.4	1.9	2.1	2.1	2.4	2.1	3.0	10.7	4.3	2.0	7.3
NLL10	2.3	1.4	16.1	2.2	2.4	2.1	15.8	2.2	2.0	2.0	2.2	1.5	2.1	2.3	5.2	1.5	-0.1	-0.1
NLL11	2.1	1.5	14.6	2.0	2.5	2.4	13.8	-0.1	2.0	1.9	2.0	1.2	2.0	2.3	3.7	1.8	-0.1	-0.1
NLL12	-0.1	1.3	11.2	1.9	0.6	2.0	9.8	-0.1	-0.1	-0.1	1.9	1.1	2.1	2.2	2.7	-0.1	-0.1	-0.1
NLL2	2.2	1.4	10.6	2.0	0.5	2.0	10.8	-0.1	2.0	-0.1	2.0	1.4	-0.1	2.2	4.1	1.4	-0.1	-0.1
NLL3	-0.1	-0.1	5.5	1.6	0.4	-0.1	5.6	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.1	2.6	-0.1	-0.1	-0.1
NLL4	2.8	1.7	24.1	2.4	1.7	2.3	21.6	2.1	2.3	2.2	2.6	3.4	2.2	3.0	12.6	1.3	2.0	7.5
NLL5	6.6	1.6	75.0	7.5	1.9	2.4	77.4	4.5	2.3	2.3	3.4	4.2	2.2	2.8	17.7	2.9	2.3	7.5
NLL6	6.5	1.7	101.0	8.2	1.7	2.7	109.0	5.0	2.6	2.5	8.8	4.4	2.3	3.1	19.6	5.7	2.7	8.1
NLL7	12.7	2.0	181.0	17.3	4.9	3.2	187.0	10.0	2.9	2.7	5.1	8.2	2.6	4.6	35.7	6.2	3.5	8.3
NLL8	2.1	1.5	14.4	1.8	2.2	2.1	11.0	-0.1	2.1	2.0	2.0	1.2	2.1	2.3	3.2	1.4	-0.1	-0.1
NLL9	4.9	1.6	29.5	6.2	6.6	2.5	32.1	2.2	2.4	2.3	3.0	3.8	2.4	4.1	23.2	2.4	2.6	7.7
NLL9-R	5.6	1.6	34.2	6.6	2.8	2.5	34.5	0.6	2.4	2.4	2.9	3.9	2.4	3.9	18.3	2.9	2.2	8.0
NM1	2.6	1.7	13.6	2.5	0.8	2.2	13.0	0.5	2.2	2.2	2.2	1.7	2.2	2.7	6.5	1.4	-0.1	7.3
NM2	-0.1	1.5	14.0	1.8	0.4	2.0	10.6	-0.1	2.0	-0.1	2.0	1.1	2.1	2.2	2.6	1.8	-0.1	-0.1
NMM1	-0.1	1.4	7.4	1.6	0.3	2.0	6.8	-0.1	-0.1	-0.1	-0.1	1.0	2.1	2.1	-0.1	-0.1	-0.1	-0.1
NMM10	3.7	1.6	24.4	3.7	2.6	2.4	23.8	2.3	2.2	2.2	2.5	2.3	2.2	3.1	10.7	4.3	2.0	7.4
NMM2	2.4	1.5	22.6	2.4	0.5	2.1	23.1	1.0	2.1	2.0	2.2	1.4	2.1	2.3	5.1	1.6	1.9	-0.1
NMM3	-0.1	1.5	7.9	1.7	0.5	2.1	7.7	-0.1	2.0	-0.1	1.9	1.0	-0.1	2.2	2.2	-0.1	-0.1	-0.1
NMM3-R	-0.1	1.5	7.2	1.6	0.4	2.0	6.6	-0.1	2.0	-0.1	-0.1	1.0	-0.1	2.1	-0.1	-0.1	-0.1	-0.1
NMM4	-0.1	-0.1	6.8	-0.1	0.4	2.0	6.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.1	-0.1	-0.1	-0.1	-0.1
NMM5	-0.1	-0.1	7.7	-0.1	0.4	-0.1	7.5	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NMM6	2.7	1.6	24.7	2.9	1.1	2.3	25.3	2.7	2.3	2.2	2.3	1.5	2.2	3.3	6.6	1.8	2.0	7.5
NMM7	-0.1	-0.1	6.5	-0.1	0.4	-0.1	6.0	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NMM8	4.4	1.7	33.3	4.6	3.9	2.3	31.8	2.3	2.3	2.2	2.9	2.9	2.2	3.1	14.9	1.6	2.0	7.6
NMM9	2.4	1.6	32.4	2.7	3.5	2.4	32.1	3.2	2.2	2.2	2.2	1.4	2.3	3.0	6.3	1.8	2.1	7.6
NN1	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	4.8	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NNN1	-0.1	-0.1	8.4	1.6	0.4	-0.1	7.4	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.1	-0.1	-0.1	-0.1
NNN10	-0.1	1.5	9.0	1.7	-0.1	2.0	8.8	-0.1	2.0	-0.1	-0.1	1.0	2.1	2.1	-0.1	-0.1	-0.1	-0.1
NNN2	-0.1	-0.1	13.2	1.9	0.4	2.0	13.0	-0.1	1.9	-0.1	2.0	1.1	-0.1	2.2	2.9	1.3	-0.1	-0.1
NNN3	-0.1	-0.1	4.0	-0.1	-0.1	-0.1	3.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NNN4	2.6	1.7	32.7	3.0	1.1	2.4	31.8	3.2	2.2	2.2	2.3	1.5	2.2	2.8	6.5	1.8	2.0	7.5
NNN5	-0.1	-0.1	6.8	-0.1	0.4	-0.1	6.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NNN6	-0.1	-0.1	6.0	-0.1	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NNN7	-0.1	-0.1	4.4	-0.1	-0.1	-0.1	4.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NNN8	2.8	1.9	49.8	3.5	1.3	2.4	46.2	4.2	2.2	2.2	2.2	1.5	2.3	2.8	7.5	2.1	2.0	7.4
NNN9	2.2	1.5	9.9	2.1	0.9	2.1	14.1	0.4	2.1	2.0	2.0	1.3	2.1	2.3	3.7	-0.1	-0.1	-0.1
NO1	2.5	1.5	13.0	2.1	0.5	2.2	13.6	-0.1	2.0	2.0	2.2	1.7	2.0	2.4	6.7	-0.1	-0.1	-0.1
NO2	-0.1	-0.1	2.7	-0.1	-0.1	-0.1	2.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NOO1	-0.1	-0.1	10.1	1.7	0.5	2.0	9.9	-0.1	1.9	-0.1	-0.1	1.0	-0.1	2.2	-0.1	-0.1	-0.1	-0.1
NOO2	2.1	1.5	12.9	1.9	0.9	2.1	11.8	0.5	2.0	2.0	2.0	1.3	2.1	2.4	3.8	1.3	-0.1	-0.1
NOO3	-0.1	1.4	5.2	-0.1	0.4	2.0	5.0	-0.1	2.0	-0.1	-0.1	1.0	-0.1	2.0	-0.1	-0.1	-0.1	-0.1
NOO4	-0.1	1.5	14.4	1.9	0.8	2.1	13.9	2.5	2.1	2.0	2.0	1.2	2.1	2.3	2.6	1.4	-0.1	-0.1
NOO5	-0.1	-0.1	7.6	-0.1	0.5	-0.1	6.7	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.0	-0.1	-0.1	-0.1	-0.1
NOO6	-0.1	-0.1	5.8	-0.1	0.3	-0.1	5.2	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NOO6-R	-0.1	-0.1	5.9	-0.1	0.3	-0.1	5.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NOO7	-0.1	-0.1	7.2	-0.1	-0.1	-0.1	6.7	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NOO8	3.7	1.6	71.7	0.5	2.0	2.4	67.8	5.2	2.3	2.2	2.6	2.1	0.6	3.0	9.1	2.6	2.1	7.7
NP1	2.0	1.5	9.4	1.7	0.4	2.1	9.1	-0.1	2.0	2.0	2.0	1.9	2.1	2.4	4.3	-0.1	-0.1	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	091-LBI	092-LPH	093-LA	094-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-LPH	101-MAR	102-MBI	103-LPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NP2	-0.1	-0.1	7.9	1.7	0.3	2.0	7.6	-0.1	-0.1	-0.1	1.9	1.3	-0.1	2.3	3.2	-0.1	-0.1	-0.1
NP3	-0.1	-0.1	6.5	1.7	0.3	-0.1	6.1	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.1	2.8	-0.1	-0.1	-0.1
NP4	5.3	2.1	80.4	8.0	5.5	3.5	83.4	5.3	3.0	2.8	3.3	4.1	2.6	7.7	23.6	4.0	3.5	8.9
NPP4	-0.1	-0.1	5.4	-0.1	0.3	-0.1	5.3	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.2	1.9	-0.1	-0.1	-0.1
NPP1-R	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	4.9	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NPP2	3.2	1.6	26.3	3.7	0.9	2.4	27.5	0.4	2.3	2.2	2.7	2.3	2.3	2.9	12.6	2.4	2.2	7.6
NPP3	2.8	1.5	29.1	3.1	2.6	2.1	30.0	2.7	2.1	2.0	2.4	1.9	2.1	2.4	8.5	1.9	2.1	7.4
NPP4	2.3	1.7	17.6	2.3	0.6	2.3	17.3	-0.1	2.2	2.1	2.1	1.5	2.2	2.6	4.9	1.5	-0.1	7.1
NQ1	2.0	1.7	7.8	1.7	0.4	2.3	7.4	-0.1	2.3	2.2	2.1	1.5	2.3	3.8	4.7	-0.1	-0.1	7.8
NQ2	4.4	1.5	26.7	5.1	0.6	2.2	29.8	0.4	2.4	2.1	3.2	3.4	2.2	3.4	17.4	2.8	0.4	7.8
NQ2-R	4.1	1.5	46.2	5.1	0.6	2.2	50.1	1.6	2.2	2.2	3.2	3.1	2.2	3.4	14.8	3.1	2.4	7.8
NQ3	2.2	-0.1	12.6	2.1	0.4	2.1	12.8	-0.1	1.9	-0.1	2.1	1.4	1.9	2.4	5.7	1.3	-0.1	7.1
NQ4	3.1	1.5	33.9	0.3	0.5	2.2	34.8	1.3	2.1	2.0	2.5	2.2	2.2	2.9	10.3	2.3	2.1	7.3
NQ5	2.2	1.6	12.1	1.9	0.5	2.1	11.9	-0.1	2.1	2.0	2.0	1.6	2.1	2.6	5.1	-0.1	-0.1	7.1
NQ6	-0.1	-0.1	8.3	-0.1	-0.1	-0.1	0.9	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NQ7	3.2	1.9	44.4	3.9	1.0	2.5	43.2	3.9	2.3	2.2	2.4	2.1	2.4	3.2	4.3	2.4	2.1	7.5
NQ8	2.1	1.5	11.0	2.0	2.2	2.1	10.5	-0.1	2.0	-0.1	2.0	1.2	2.1	2.2	3.3	1.4	-0.1	-0.1
NQQ1	-0.1	1.5	7.0	1.8	0.4	2.0	6.3	-0.1	2.0	-0.1	-0.1	1.1	-0.1	2.1	2.0	-0.1	-0.1	-0.1
NR1	-0.1	-0.1	5.8	-0.1	2.0	-0.1	5.7	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NR11	-0.1	1.6	11.4	1.7	0.8	2.1	11.2	2.2	2.1	2.0	1.9	1.1	2.1	2.4	2.7	-0.1	-0.1	-0.1
NR12	-0.1	1.5	13.1	1.7	2.4	2.0	12.4	2.2	2.0	2.0	2.0	1.0	2.1	2.1	-0.1	-0.1	-0.1	-0.1
NR13	2.9	1.7	51.0	0.9	3.1	2.4	51.0	4.6	2.3	2.2	2.4	1.7	2.2	2.5	7.2	1.9	2.1	7.4
NR14	-0.1	1.4	9.2	1.7	0.5	-0.1	8.5	-0.1	-0.1	-0.1	1.9	1.0	-0.1	2.1	2.3	-0.1	-0.1	-0.1
NR2	-0.1	1.4	7.9	1.7	0.6	2.0	8.1	-0.1	2.0	1.9	2.0	1.1	-0.1	2.2	2.5	-0.1	-0.1	-0.1
NR3	2.1	1.5	10.3	1.8	0.8	2.1	2.1	0.4	2.0	2.0	2.0	1.2	-0.1	2.2	2.9	-0.1	-0.1	-0.1
NR4	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	4.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NR5	-0.1	-0.1	8.2	-0.1	0.5	-0.1	6.8	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NR6	-0.1	-0.1	5.8	-0.1	1.9	-0.1	5.3	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NR7	2.7	1.7	22.6	2.5	1.2	2.4	21.7	0.5	2.3	2.3	2.5	1.7	2.3	3.0	7.8	1.4	2.0	7.7
NR8	-0.1	-0.1	6.8	-0.1	0.4	-0.1	6.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NS1	-0.1	-0.1	7.2	1.6	0.5	2.1	8.0	-0.1	-0.1	-0.1	2.0	1.1	-0.1	2.2	3.0	-0.1	-0.1	-0.1
NS10	3.9	1.8	69.3	5.3	0.5	2.5	66.3	2.9	2.4	2.3	2.9	3.0	2.4	4.0	13.5	3.6	2.5	7.9
NS11	2.3	1.9	15.1	2.1	0.4	2.3	15.4	-0.1	0.4	2.2	2.2	1.5	2.3	2.6	5.9	1.4	-0.1	-0.1
NS12	2.7	1.7	38.4	0.3	3.0	2.3	39.7	3.5	2.2	2.1	2.2	1.8	2.3	3.6	7.9	2.2	2.1	7.4
NS13	3.6	2.0	30.9	4.0	1.4	2.6	33.0	0.4	2.5	2.4	3.0	2.8	2.4	3.6	16.4	2.2	2.2	7.7
NS14	3.9	2.7	46.2	4.5	6.1	3.6	45.0	2.7	3.2	3.0	3.0	3.0	3.1	5.7	17.0	2.6	2.5	8.0
NS2	2.4	1.4	21.3	2.3	0.5	2.0	22.0	-0.1	2.0	-0.1	2.1	1.5	2.1	2.4	5.7	1.6	-0.1	-0.1
NS3	3.0	1.5	24.5	3.9	1.0	2.2	24.7	2.6	2.1	2.1	2.3	2.3	2.2	3.1	11.0	1.9	2.1	7.7
NS4	2.1	1.4	16.2	1.7	2.4	2.2	10.8	-0.1	2.1	2.1	2.3	1.9	2.1	2.6	6.1	-0.1	-0.1	7.4
NS4-R	2.1	1.4	16.5	1.7	0.5	2.2	14.3	-0.1	2.0	2.0	2.2	1.5	2.0	2.4	5.3	-0.1	-0.1	-0.1
NS5	2.6	1.4	18.4	2.3	2.7	2.2	20.6	-0.1	2.1	2.1	2.5	2.5	2.1	2.6	10.2	1.4	2.0	7.4
NS6	2.1	1.4	12.1	1.8	0.4	2.0	11.5	-0.1	2.0	2.0	2.1	1.5	2.1	2.4	4.1	-0.1	-0.1	-0.1
NS7	3.8	1.5	1.6	3.9	0.6	2.2	30.3	-0.1	2.1	2.1	3.1	3.7	2.2	2.7	18.0	1.8	2.1	7.3
NS8	3.9	1.8	1.6	3.9	0.7	2.4	28.4	2.1	2.3	2.2	3.1	4.0	2.4	4.0	20.1	1.6	2.2	7.8
NS9	-0.1	-0.1	7.4	-0.1	0.5	-0.1	7.5	-0.1	-0.1	-0.1	1.9	1.2	-0.1	2.3	2.8	-0.1	-0.1	-0.1
NT1	-0.1	-0.1	8.9	1.7	0.4	2.0	8.7	-0.1	2.0	-0.1	-0.1	1.1	-0.1	2.1	2.4	1.3	-0.1	-0.1
NT10	1.9	1.4	9.0	1.6	0.6	2.0	8.3	-0.1	2.0	2.0	2.0	1.6	-0.1	2.4	4.2	-0.1	-0.1	-0.1
NT11	-0.1	-0.1	6.5	-0.1	2.2	2.0	6.8	-0.1	-0.1	1.9	1.9	1.3	-0.1	2.4	3.5	-0.1	-0.1	-0.1
NT12	2.0	1.5	8.9	1.7	0.5	2.2	8.1	-0.1	2.0	2.0	2.0	1.5	2.0	2.5	4.7	-0.1	-0.1	-0.1
NT13	-0.1	-0.1	9.6	-0.1	0.5	2.0	9.2	-0.1	-0.1	-0.1	2.0	1.3	-0.1	2.2	3.4	-0.3	-0.1	-0.1
NT14	2.1	1.5	16.3	2.0	2.3	2.2	16.0	-0.1	2.1	2.0	2.0	1.1	2.0	2.1	3.5	1.4	1.9	-0.1
NT15	2.1	1.5	24.9	2.4	3.0	2.1	21.4	2.4	2.0	2.0	2.4	1.5	2.1	2.9	5.1	-0.1	-0.1	-0.1
NT16	2.3	1.6	19.0	0.3	1.1	2.3	18.6	2.6	2.2	2.1	2.1	1.4	2.2	2.6	4.7	1.5	1.9	-0.1
NT17	2.2	1.5	15.9	0.3	2.4	2.1	15.5	-0.1	2.0	-0.1	2.0	1.4	2.1	2.2	4.3	-0.1	-0.1	-0.1
NT18	-0.1	-0.1	3.8	-0.1	-0.1	-0.1	3.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NT2	2.1	1.5	14.4	2.0	0.5	2.5	14.5	-0.1	2.2	2.0	2.0	1.1	2.0	2.3	4.4	1.6	1.9	-0.1
NT3	-0.1	-0.1	11.2	-0.1	1.9	-0.1	1.7	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	1.9	-0.1	-0.1	-0.1
NT4	-0.1	-0.1	9.4	-0.1	0.5	-0.1	9.2	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NT5	-0.1	-0.1	6.8	-0.1	0.3	-0.1	6.1	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.0	-0.1	-0.1	-0.1
NT6-R	-0.1	-0.1	9.8	1.6	0.4	2.1	9.2	-0.1	2.0	2.0	2.0	1.1	-0.1	2.2	3.2	-0.1	-0.1	-0.1
NT6	-0.1	-0.1	7.6	-0.1	0.4	-0.1	7.0	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.2	2.3	-0.1	-0.1	-0.1
NT7	2.2	1.3	10.9	1.9	1.0	2.1	12.2	-0.1	2.0	2.0	2.2	1.5	2.1	2.4	4.9	-0.1	-0.1	-0.1
NT8	3.8	1.6	42.0	4.8	2.5	2.6	45.0	0.5	2.5	2.4	3.1	3.1	2.3	3.2	16.8	2.5	2.4	7.8
NT9	-0.1	-0.1	7.8	-0.1	0.4	-0.1	7.5	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.1	2.6	-0.1	-0.1	-0.1
NU1	-0.1	1.4	8.0	-0.1	0.4	-0.1	7.8	-0.1	-0.1	-0.1	2.0	1.3	-0.1	2.3	3.0	-0.1	-0.1	-0.1
NU10	4.2	1.6	127.0	0.4	0.8	2.5	134.0	3.8	2.3	2.2	3.5	2.7	2.4	3.9	5.9	4.3	3.1	8.6
NU11	2.2	1.4	9.6	2.1	0.5	2.1	10.3	-0.1	2.0	2.0	2.2	1.3	2.0	2.4	5.0	1.5	1.9	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	D91-LBI	D92-LPH	D93-LA	D94-LBI	D95-MAR	D96-LPH	D97-HBA	D98-THI	D99-LPH	100-EPH	101-MAR	102-MBI	103-EPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH
NU12	2.2	1.4	25.3	2.2	0.6	2.0	25.9	-0.1	2.0	2.0	2.2	1.3	-0.1	2.3	4.7	1.5	1.9	-0.1
NU12-R	2.2	1.4	20.8	2.3	0.5	2.0	21.1	-0.1	-0.1	-0.1	2.0	1.3	-0.1	2.3	5.1	1.6	-0.1	-0.1
NU13	3.9	1.6	70.5	5.4	1.4	2.3	72.6	5.2	2.2	2.1	2.8	2.7	2.2	3.8	12.8	6.5	2.4	7.6
NU14	-0.1	1.6	10.9	1.7	2.4	2.1	0.9	-0.1	2.1	2.0	2.0	1.1	2.1	2.3	2.9	-0.1	-0.1	-0.1
NU15	2.5	1.5	32.4	0.3	0.8	2.2	32.4	3.0	2.1	2.1	2.5	1.6	2.2	2.5	7.2	1.6	1.9	7.3
NU16	4.4	1.9	96.3	0.6	1.9	2.6	96.3	6.8	2.4	2.3	2.8	2.9	2.3	3.2	13.2	2.8	2.3	7.7
NU17	3.3	1.7	49.6	0.4	1.7	2.3	49.9	3.6	2.3	2.2	2.2	2.2	2.2	2.6	10.7	2.0	2.0	7.3
NU18	2.2	1.5	11.3	2.1	1.0	2.1	11.5	0.4	2.0	2.0	2.1	1.6	2.1	2.6	5.5	1.4	1.9	-0.1
NU19	-0.1	1.6	16.0	1.9	2.2	2.2	14.4	-0.1	2.1	2.0	2.0	1.1	2.1	2.2	2.7	-0.1	-0.1	-0.1
NU2	8.0	1.6	57.9	10.7	2.2	2.5	35.7	0.5	2.4	2.4	4.5	6.2	2.3	3.2	34.8	3.0	2.7	7.8
NU3	-0.1	-0.1	3.6	1.6	-0.5	-0.1	0.6	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.1	-2.0	-0.1	-0.1	-0.1
NU4	9.3	1.9	93.0	13.0	3.8	2.9	110.0	0.5	2.9	2.8	6.9	8.6	2.6	5.6	47.1	5.9	4.2	9.1
NU5	-0.1	-0.1	7.7	1.6	0.4	-0.1	8.0	-0.1	-0.1	-0.1	-0.1	1.2	-0.1	2.1	2.9	-0.1	-0.1	-0.1
NU6	2.0	-0.1	9.8	1.7	0.5	2.0	10.5	-0.1	2.0	2.0	2.1	1.4	-0.1	2.2	4.1	-0.1	-0.1	-0.1
NU7	2.3	1.4	15.1	2.0	0.7	2.2	15.7	-0.1	2.0	2.0	2.2	1.4	2.0	2.5	6.2	1.2	-0.1	-0.1
NU8	2.8	1.7	39.6	0.4	1.1	2.4	40.8	3.3	2.2	2.1	2.4	1.9	2.2	2.6	9.5	2.0	2.1	-0.1
NU9	2.4	1.7	30.9	2.7	0.8	2.5	31.2	3.2	2.4	2.2	2.8	1.5	2.3	3.3	6.7	1.9	2.1	7.7
NV1	2.1	1.6	10.3	1.8	0.6	2.2	9.4	-0.1	2.0	-0.1	2.0	1.2	2.2	2.3	3.3	-0.1	-0.1	-0.1
NV10	-0.1	-0.1	12.6	1.6	0.3	-0.1	12.2	-0.1	-0.1	-0.1	4.9	1.1	-0.1	2.1	-2.4	-0.1	-0.1	-0.1
NV11	-0.1	-0.1	7.9	-0.1	0.4	2.0	7.5	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NV12	-0.1	1.5	10.7	-0.1	0.3	2.0	1.1	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	-0.1	1.9	-0.1	-0.1
NV13	-0.1	1.6	19.2	2.0	0.8	2.2	18.0	-0.1	2.1	2.0	1.9	1.1	2.1	2.2	3.1	-0.1	-0.1	-0.1
NV14	-0.1	1.5	17.9	1.7	2.3	2.3	18.1	-0.1	2.1	-2.0	2.0	1.0	2.1	2.2	-2.4	-0.1	1.9	-0.1
NV15	-0.1	-0.1	10.2	-0.1	0.4	-0.1	9.5	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NV16	2.2	1.9	28.2	2.1	1.0	2.4	27.0	2.5	2.3	2.2	2.2	1.2	2.3	2.5	3.7	1.4	-0.1	-0.1
NV17	-0.1	1.7	11.8	1.7	0.6	2.4	1.7	-0.1	2.1	2.1	2.0	1.0	2.1	2.2	2.4	-0.1	-0.1	-0.1
NV18	-0.1	1.4	11.6	1.7	-0.5	2.0	10.9	-0.1	1.9	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NV2	-0.1	1.4	10.9	1.7	0.4	2.0	10.2	-0.1	2.0	2.0	2.0	1.2	2.1	2.2	2.8	-0.1	-0.1	-0.1
NV3	-0.1	-0.1	14.1	1.7	0.3	2.1	10.9	-0.1	-0.1	-0.1	-0.1	0.9	-0.1	1.9	2.1	-0.1	-0.1	-0.1
NV4	2.1	1.7	20.2	2.1	0.5	2.3	19.8	0.8	2.2	2.1	2.0	1.2	2.2	2.3	3.4	1.4	1.9	-0.1
NV5	2.2	1.7	24.4	2.2	0.7	2.3	23.9	-0.1	2.3	-2.1	2.1	1.3	2.3	2.4	4.7	1.5	2.1	-0.1
NV5-R	2.2	2.1	27.2	2.2	0.6	2.5	27.4	0.9	0.5	2.2	2.1	1.3	2.4	2.5	4.3	1.5	1.9	7.2
NV6	-0.1	-0.1	6.8	1.7	0.3	2.4	6.5	-0.1	2.0	1.9	1.0	1.1	2.1	2.3	-0.1	-0.1	-0.1	-0.1
NV7	-0.1	1.4	17.9	1.8	2.2	2.1	18.4	-0.1	2.0	2.0	2.1	1.1	2.1	2.2	2.8	-0.1	-0.1	-0.1
NV8	-0.1	-0.1	10.2	1.7	2.0	2.0	10.0	-0.1	2.0	-0.1	-0.1	1.0	2.0	-0.1	-0.1	-0.1	-0.1	-0.1
NV9	2.2	1.5	22.3	2.3	2.2	2.1	21.8	1.0	2.0	-0.1	2.0	1.2	2.1	2.1	3.6	1.4	-0.1	-0.1
NW1	2.8	1.5	20.5	2.9	1.0	2.4	27.5	2.7	2.0	2.0	2.2	1.6	2.1	2.5	6.5	1.7	1.9	-0.1
NW10	-0.1	1.4	7.0	1.7	2.0	2.1	7.1	-0.1	2.0	1.9	1.9	1.0	2.0	2.1	2.3	-0.1	-0.1	-0.1
NW11	-0.1	-0.1	9.3	-0.1	0.4	-0.1	8.6	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NW12	2.0	1.6	13.9	1.6	0.6	2.3	2.7	-0.1	2.2	2.0	2.0	1.2	2.1	2.2	2.8	-0.1	-0.1	-0.1
NW13	2.7	1.8	18.3	2.9	0.9	2.5	18.8	2.2	2.4	2.1	2.2	1.7	2.3	2.5	7.0	1.8	2.1	-0.1
NW14	-0.1	-0.1	8.1	1.6	0.3	-0.1	7.3	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.0	-0.1	-0.1	-0.1	-0.1
NW15	2.5	2.0	20.0	2.5	0.9	2.5	2.3	0.5	2.3	2.2	2.2	1.7	2.4	2.7	1.4	1.4	1.9	7.4
NW16	-0.1	-0.1	8.4	1.6	0.3	-0.1	7.8	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	2.1	-0.1	-0.1	-0.1
NW17	2.0	1.4	8.9	1.7	2.1	2.4	9.0	-0.1	2.1	2.0	2.0	1.2	2.1	2.3	2.8	-0.1	-0.1	-0.1
NW17-R	-0.1	1.4	8.0	1.7	2.1	2.1	7.9	-0.1	2.0	2.0	2.0	1.1	2.1	2.3	2.5	-0.1	-0.1	-0.1
NW18	-0.1	1.4	10.8	2.0	0.7	2.2	11.5	-0.1	2.1	-2.0	2.1	1.2	2.1	2.9	3.6	1.4	-0.1	-0.1
NW2	-0.1	1.4	10.7	1.6	0.5	2.0	1.9	-0.1	2.0	-0.1	2.0	1.0	2.0	2.1	2.1	-0.1	-0.1	-0.1
NW2-R	-0.1	1.4	12.2	1.8	2.1	2.2	2.2	-0.1	2.0	2.0	2.0	0.9	1.9	2.0	2.2	-0.1	-0.1	-0.1
NW3	-0.1	-0.1	17.4	1.9	0.4	2.0	17.1	-0.1	-0.1	-0.1	1.9	1.0	-0.1	-0.1	2.6	-0.1	-0.1	-0.1
NW4	-0.1	-0.1	11.8	1.7	0.4	2.0	11.4	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NW5	4.1	1.7	87.0	0.6	1.4	2.5	89.7	5.8	2.4	2.3	3.0	2.8	2.3	2.6	12.9	2.4	2.2	7.8
NW6	-0.1	1.4	15.4	1.9	0.5	2.0	14.7	-0.1	-0.1	-0.1	-0.1	1.1	-0.1	2.1	-2.7	1.3	-0.1	-0.1
NW7	-0.1	-0.1	10.9	1.7	2.2	2.0	10.2	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NW8	4.1	1.6	60.3	4.6	1.6	2.3	60.9	4.7	2.2	-2.1	2.6	2.4	2.2	2.6	10.8	2.0	2.1	7.4
NW9	2.0	1.4	17.7	1.9	2.3	2.0	17.8	-0.1	-0.1	-0.1	2.0	1.1	-0.1	2.1	2.7	-0.1	-0.1	-0.1
NX1	-0.1	1.5	5.9	-0.1	-0.1	2.1	5.6	-0.1	2.0	-0.1	-0.1	1.0	-0.1	2.1	2.1	-0.1	-0.1	-0.1
NX10	2.6	1.6	16.9	2.6	2.6	2.3	8.0	-0.1	2.2	2.2	2.4	1.9	2.2	2.9	8.6	1.4	2.0	7.7
NX11	-0.1	-0.1	5.4	1.6	0.5	-0.1	0.7	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NX12	-0.1	-0.1	8.3	1.7	0.4	2.0	7.9	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	2.0	2.6	-0.1	-0.1	-0.1
NX13	2.6	1.7	27.1	2.9	0.7	2.3	27.4	-0.1	2.3	-2.2	2.2	1.6	2.2	2.8	7.2	1.4	2.0	7.3
NX14	2.5	1.5	22.5	2.7	0.8	2.1	22.6	0.9	2.1	2.0	2.2	1.6	2.1	2.4	6.8	1.7	1.9	-0.1
NX14-R	2.1	1.5	20.7	2.2	0.6	2.1	20.8	-0.1	2.1	-2.0	2.1	1.3	2.1	2.3	4.4	1.4	-0.1	-0.1
NX15	2.1	1.6	17.2	2.0	2.4	2.1	17.2	-0.1	2.1	2.1	2.1	1.4	2.2	2.3	4.1	1.4	-0.1	-0.1
NX16	2.6	1.5	25.5	2.8	2.8	2.1	25.4	2.4	2.1	2.0	2.2	1.7	2.1	2.3	6.4	1.5	-0.1	-0.1
NX17	3.2	2.1	32.7	3.5	4.1	2.7	32.7	2.9	2.5	2.3	2.4	2.3	2.5	3.0	11.0	1.8	2.1	7.7
NX2	2.4	1.5	14.5	2.4	0.3	2.1	15.7	-0.1	2.1	2.0	2.5	1.9	2.2	2.9	8.1	1.8	2.0	7.4

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	091-LB1	092-LPH	093-LA	094-LB1	095-MAR	096-LPH	097-HBA	098-TH1	099-LPH	100-EPH	101-MAR	102-NB1	103-EPH	104-MAR	105-ALK	106-MB1	107-MB1	108-LPH
NX3	2.1	1.4	10.6	1.9	0.3	2.0	10.4	-0.1	2.0	2.0	2.2	1.7	2.1	2.8	1.3	1.3	-0.1	7.5
NX4	5.4	1.9	43.5	7.4	0.7	2.5	50.7	2.3	2.4	2.4	4.0	4.0	2.5	4.9	23.7	3.3	3.0	8.1
NX5	2.1	1.4	12.7	2.0	0.3	2.0	12.9	-0.1	2.0	-0.1	2.1	1.4	2.1	2.4	4.7	1.4	-0.1	-0.1
NX6	2.9	1.5	40.2	0.4	0.6	2.2	42.0	2.0	2.2	2.1	2.5	2.0	2.2	2.7	9.4	2.1	2.1	7.4
NX7	12.1	2.0	266.0	19.6	3.0	3.2	277.0	14.5	2.8	2.5	6.0	9.7	2.6	4.2	41.4	8.3	4.9	8.3
NX8	2.6	1.8	27.1	0.3	0.7	2.4	26.9	1.1	2.3	2.2	2.2	1.7	2.3	2.7	8.1	1.8	2.0	7.5
NX9	4.2	1.7	57.3	5.6	1.2	2.4	58.5	4.4	2.3	2.4	2.8	3.0	2.3	3.5	14.2	3.2	2.3	7.8
NY1	-0.1	-0.1	8.4	1.7	2.0	2.0	8.2	-0.1	1.9	-0.1	-0.1	1.1	-0.1	2.2	2.2	-0.1	-0.1	-0.1
NY10	2.4	1.7	43.2	3.0	2.8	2.4	42.3	2.0	2.2	2.1	2.2	1.5	2.2	3.0	7.4	2.0	2.0	7.4
NY11	-0.1	-0.1	9.0	-0.1	1.9	-0.1	7.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NY12	-0.1	1.4	8.2	1.6	0.4	2.0	7.4	-0.1	2.0	2.0	2.0	1.2	-0.1	2.5	2.6	-0.1	-0.1	7.2
NY13	-0.1	-0.1	5.7	-0.1	0.4	-0.1	5.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NY14	2.4	1.8	16.8	2.2	1.2	2.4	15.2	2.1	2.3	2.2	2.2	1.5	2.3	3.0	5.6	-0.1	-0.1	7.4
NY15	2.1	1.5	22.1	2.0	2.9	2.2	22.3	2.1	2.1	2.0	2.1	1.2	2.1	2.7	3.5	-0.1	1.9	-0.1
NY2	-0.1	-0.1	5.1	-0.1	0.4	-0.1	4.8	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NY3	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	5.0	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NY4	3.3	1.6	30.9	3.5	2.0	2.2	30.6	3.1	2.1	2.1	2.3	1.9	2.1	2.9	7.5	1.7	1.9	-0.1
NY4-R	4.8	1.7	67.8	5.9	2.7	2.4	69.3	4.9	2.3	2.2	3.0	3.0	2.2	3.6	12.4	2.9	2.2	7.6
NY5	-0.1	-0.1	9.6	1.8	2.1	2.0	9.5	-0.1	2.0	-0.1	2.0	1.1	-0.1	2.2	2.7	1.4	-0.1	-0.1
NY6	2.1	1.6	14.2	2.0	2.5	2.2	14.4	-0.1	2.1	2.1	2.1	1.3	2.2	2.8	4.7	1.5	-0.1	-0.1
NY7	2.7	1.8	20.2	2.6	1.2	2.4	12.3	2.0	2.3	2.2	2.5	2.0	2.3	3.5	10.2	1.5	2.0	7.5
NY8	-0.1	1.4	15.6	1.8	2.3	2.1	16.1	-0.1	2.1	2.0	2.3	1.1	2.1	2.4	2.8	-0.1	-0.1	-0.1
NY9	-0.1	-0.1	4.9	-0.1	0.4	2.0	4.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NZ1	-0.1	1.4	9.3	1.6	0.5	2.0	8.7	-0.1	1.9	-0.1	1.9	1.2	2.1	2.3	2.6	-0.1	-0.1	-0.1
NZ10	2.3	2.7	37.2	2.5	1.6	3.2	34.2	3.0	0.5	2.7	2.3	1.3	2.8	3.4	5.2	1.6	2.0	7.7
NZ11	-0.1	-0.1	7.5	-0.1	1.9	2.0	7.2	-0.1	1.9	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NZ12	-0.1	-0.1	11.9	1.8	2.2	2.0	10.9	-0.1	1.9	-0.1	-0.1	1.0	2.0	2.2	2.4	-0.1	-0.1	-0.1
NZ13	-0.1	-0.1	6.3	1.6	0.4	-0.1	6.2	-0.1	-0.1	-0.1	-0.1	1.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
NZ2	2.5	1.4	13.1	2.1	0.6	2.0	13.4	2.0	2.0	2.0	2.3	2.2	2.1	2.5	7.5	-0.1	-0.1	-0.1
NZ2-R	-0.1	-0.1	8.4	1.6	0.6	-0.1	7.7	-0.1	-0.1	-0.1	1.9	1.3	-0.1	2.3	3.2	-0.1	-0.1	-0.1
NZ3	2.1	1.4	12.2	1.8	2.4	2.1	12.4	-0.1	2.0	2.0	2.1	1.4	2.1	2.6	4.1	-0.1	-0.1	-0.1
NZ4	2.1	1.4	10.6	1.9	0.5	2.1	10.1	-0.1	2.0	1.9	2.0	1.2	-0.1	2.3	3.6	1.4	-0.1	-0.1
NZ5	2.0	1.6	9.8	1.7	0.4	2.1	1.4	-0.1	2.1	2.0	2.0	1.2	2.1	2.5	2.9	-0.1	-0.1	-0.1
NZ6	-0.1	1.4	12.6	1.9	2.3	2.1	12.0	-0.1	2.0	-0.1	1.9	1.1	2.1	2.3	2.8	-0.1	-0.1	-0.1
NZ7	-0.1	1.8	17.0	1.9	2.3	2.3	16.4	-0.1	2.2	2.1	2.1	1.1	2.3	2.5	2.8	1.3	-0.1	-0.1
NZ8	-0.1	1.7	9.3	-0.1	0.4	2.4	8.9	-0.1	2.1	-0.1	2.0	1.1	2.2	2.3	0.6	-0.1	-0.1	-0.1
NZ9	-0.1	-0.1	7.4	-0.1	-0.1	1.9	1.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	6.4	-0.1	-0.1	-0.1	5.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	6.3	-0.1	-0.1	-0.1	6.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.6	-0.1	-0.1	-0.1	3.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	6.4	-0.1	-0.1	-0.1	5.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	0.5	-0.1	-0.1	-0.1	4.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	0.4	-0.1	-0.1	-0.1	4.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.5	-0.1	-0.1	-0.1	3.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.4	-0.1	-0.1	-0.1	3.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.3	-0.1	-0.1	-0.1	3.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	2.8	-0.1	-0.1	-0.1	2.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.9	-0.1	-0.1	-0.1	3.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	-0.1	3.8	-0.1	-0.1	-0.1	3.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	091-LBI	092-LPH	093-LA	094-LBI	095-MAR	096-LPH	097-HBA	098-THI	099-LPH	100-LPH	101-MAR	102-MBI	103-LPH	104-MAR	105-ALK	106-MBI	107-MBI	108-LPH

A14-06865

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109 MAR	110 HBA	111 MAR	112 MBI	113 HBA	114 MBI	115 MBI	116 MAR	117 HA	118 MRH	119 HBA	120 THI	121 MRH	122 MPH	123 MPH	124 MBI	125 HAR	126 MPH
NA1	22.3	70.5	7.1	12.1	73.5	6.1	16.9	5.8	125.0	7.0	80.4	-0.1	8.4	9.2	8.0	9.8	6.7	7.4
NA2	10.5	35.7	-0.1	9.7	41.7	1.6	11.4	4.5	133.0	-0.1	78.3	-3.9	-0.1	7.1	7.0	8.4	4.5	6.8
NAA1	10.4	38.1	-0.1	9.0	38.1	10.6	10.8	3.9	57.3	6.7	35.4	-0.1	7.5	8.0	7.1	8.2	5.1	-0.1
NAA10	7.9	24.0	-0.1	8.6	24.7	1.7	8.6	3.4	35.4	-0.1	4.5	-0.1	-0.1	6.0	-0.1	7.7	4.3	-0.1
NAA11	18.3	58.5	7.2	10.8	54.0	18.0	20.3	4.9	48.0	-0.1	39.9	-0.1	-0.1	8.3	-0.1	9.6	6.4	-0.1
NAA2	78.9	295.0	12.0	31.3	312.0	17.7	35.1	15.6	333.0	8.1	225.0	5.8	9.3	20.3	8.9	17.1	12.6	8.1
NAA3	17.1	69.0	7.8	11.5	68.4	2.5	10.2	5.0	72.9	-0.1	57.6	-0.1	-0.1	7.9	-0.1	8.9	5.1	-0.1
NAA4	9.8	37.2	-0.1	9.4	40.8	1.6	11.3	4.4	84.8	-0.1	55.5	-0.1	-0.1	6.5	-0.1	8.2	4.8	-0.1
NAA5	12.6	43.2	7.0	10.0	42.9	2.1	9.4	4.0	66.6	6.4	37.8	-0.1	7.3	7.9	6.8	8.2	5.6	6.7
NAA6	10.2	46.8	-0.1	9.8	49.2	1.8	10.9	4.1	72.9	-0.1	47.1	-0.1	-0.1	7.0	-6.7	8.3	5.1	-6.7
NAA7	8.5	33.0	-0.1	9.0	35.4	1.5	10.8	3.7	63.6	6.4	38.1	-0.1	-0.1	6.7	6.9	8.1	4.5	-0.1
NAA8	7.4	33.0	-0.1	8.8	34.8	1.5	9.9	3.8	71.1	-0.1	41.1	-0.1	-0.1	5.9	-0.1	7.6	4.3	-0.1
NAA9	9.3	38.4	-0.1	9.4	39.9	1.4	10.7	3.9	76.8	-0.1	44.7	-0.1	-0.1	6.8	6.8	8.0	5.1	-0.1
NB1	9.4	33.6	-0.1	9.2	35.1	11.7	11.5	3.6	44.7	-0.1	35.0	-0.1	-0.1	6.6	-0.1	8.0	4.9	-0.1
NB2	5.7	20.1	-0.1	-0.1	20.6	-0.1	7.7	-0.1	33.6	-0.1	22.4	-0.1	-0.1	5.6	-0.1	-0.1	3.9	-0.1
NB3	6.0	18.4	-0.1	-0.1	18.7	8.8	8.0	-0.1	23.4	-0.1	17.2	-0.1	-0.1	5.8	-0.1	3.9		-0.1
NB4	12.1	68.1	7.4	11.5	74.4	2.1	12.6	5.3	126.0	-0.1	88.5	4.2	-0.1	7.0	-0.1	9.1	4.7	-0.1
NBB1	6.0	24.4	-0.1	-0.1	25.1	8.7	8.3	-0.1	36.6	-0.1	25.9	-0.1	-0.1	5.6	-0.1	7.4	4.1	-0.1
NBB10	31.2	81.0	7.5	13.7	85.2	6.3	18.2	7.3	103.0	6.8	86.7	-0.1	7.6	10.6	6.8	10.0	7.5	6.9
NBB11	14.9	65.1	7.1	11.2	69.6	22.9	27.8	5.0	174.0	7.4	83.7	4.0	8.6	8.9	8.0	11.0	6.8	7.6
NBB2	14.5	49.8	7.0	10.5	51.0	2.0	11.1	4.6	91.2	-0.1	44.4	-0.1	-0.1	7.7	-0.1	8.1	5.8	-0.1
NBB2-R	10.6	42.9	-0.1	9.4	42.6	1.8	9.4	4.0	57.9	-0.1	39.3	-0.1	-0.1	6.8	-0.1	7.8	5.2	-0.1
NBB3	8.3	32.1	-0.1	8.7	32.1	1.7	8.4	3.7	30.9	-0.1	23.0	-0.1	-0.1	6.0	-0.1	7.3	4.8	-0.1
NBB4	5.6	14.2	-0.1	-0.1	14.4	-0.1	-0.1	-0.1	11.6	-0.1	10.8	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NBB5	6.4	25.9	-0.1	-0.1	26.8	10.0	10.4	-0.1	47.7	-0.1	24.1	-0.1	-0.1	5.8	-0.1	7.5	4.3	-0.1
NBB6	11.7	47.1	6.9	9.9	48.0	2.0	11.5	3.9	63.3	6.5	37.2	-0.1	7.5	7.0	7.0	8.5	5.2	6.8
NBB7	5.3	14.2	-0.1	-0.1	15.2	-0.1	-0.1	-0.1	20.7	-0.1	13.7	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
NBB8	9.0	35.4	-0.1	9.0	34.5	1.6	9.6	3.6	36.0	-0.1	27.9	-0.1	-0.1	6.3	-0.1	7.9	4.5	-0.1
NBB9	11.3	45.0	-0.1	9.2	43.2	1.8	9.5	3.6	48.6	-0.1	28.2	-0.1	-0.1	7.0	-0.1	7.7	5.4	-0.1
NC1	6.1	26.5	-0.1	-0.1	26.6	8.5	9.2	-0.1	65.7	-0.1	24.3	-0.1	-0.1	5.4	-0.1	7.2	3.8	-0.1
NC2	6.9	36.0	-0.1	8.9	36.9	9.6	8.8	3.4	54.9	-0.1	34.2	-0.1	-0.1	6.1	-0.1	7.7	4.1	-0.1
NC3	6.8	31.8	-0.1	8.7	32.7	1.8	8.1	3.6	44.1	-0.1	28.6	-0.1	-0.1	5.9	-0.1	7.4	4.2	-0.1
NC4	5.6	16.0	-0.1	-0.1	16.5	-0.1	-0.1	-0.1	21.4	-0.1	15.2	-0.1	-0.1	5.7	-0.1	-0.1	4.0	-0.1
NC5	6.8	28.9	-0.1	8.7	30.3	1.5	8.5	3.6	44.7	-0.1	27.3	-0.1	-0.1	5.8	-0.1	7.4	4.2	-0.1
NC6	7.2	23.4	-0.1	8.2	26.5	10.1	10.2	3.6	39.3	-0.1	24.2	-0.1	-0.1	6.4	6.8	7.7	4.4	-0.1
NC7	7.2	22.6	-0.1	8.6	23.8	8.6	8.0	3.6	22.3	-0.1	17.6	-0.1	-0.1	5.7	-0.1	7.8	4.2	-0.1
NC8	6.2	17.5	-0.1	-0.1	17.9	-0.1	7.9	-0.1	24.8	-0.1	14.6	-0.1	-0.1	5.3	-0.1	-0.1	3.8	-0.1
NCC1	96.3	549.0	18.6	55.2	672.0	33.6	73.8	30.3	1500.0	9.1	864.0	56.9	11.6	22.3	10.8	29.3	10.3	8.8
NCC1-R	92.7	558.0	15.7	51.3	696.0	12.5	70.2	31.8	1380.0	9.4	894.0	58.8	11.3	23.5	10.8	29.2	10.7	8.5
NCC2	79.2	201.0	6.5	23.0	220.0	12.6	36.3	16.3	504.0	8.8	329.0	25.0	10.3	16.7	10.6	15.2	13.1	9.0
NCC3	15.4	68.1	7.0	10.9	69.3	6.6	16.9	4.7	120.0	6.5	60.3	-0.1	7.6	8.6	7.1	9.2	6.3	6.8
NCC4	33.3	99.0	7.7	13.5	93.9	2.6	13.5	5.9	86.4	6.9	69.0	-0.1	7.8	12.0	7.0	9.4	10.3	6.9
NCC5	12.4	51.3	-0.1	10.1	52.5	15.4	17.5	4.4	98.1	-0.1	43.2	-0.1	-0.1	7.4	-0.1	9.0	5.6	-0.1
NCC6	25.7	70.2	7.7	12.1	63.6	1.7	13.7	6.1	79.8	6.6	51.6	-0.1	-0.1	10.6	6.8	9.6	7.0	6.7
ND1	6.3	20.1	-0.1	-0.1	20.1	-0.1	10.5	-0.1	22.0	-0.1	16.1	-0.1	-0.1	5.8	-0.1	7.4	4.2	-0.1
ND10	5.4	20.7	-0.1	-0.1	20.8	-0.1	7.6	-0.1	20.6	-0.1	16.8	-0.1	-0.1	5.7	-0.1	7.3	4.0	-0.1
ND2	11.1	55.2	7.2	9.6	52.2	12.6	12.9	3.8	68.1	-0.1	42.3	-0.1	-0.1	7.0	6.5	8.6	5.1	-0.1
ND3	5.7	22.6	-0.1	2.2	23.4	-0.1	7.7	-0.1	29.8	-0.1	19.8	-0.1	-0.1	6.1	-0.1	7.5	3.8	-0.1
ND4	14.5	82.2	9.1	12.1	87.6	11.9	11.0	4.9	84.9	-0.1	58.2	-0.1	-0.1	7.7	6.7	9.5	5.5	6.8
ND5	5.6	18.5	-0.1	-0.1	18.7	-0.1	-0.1	-0.1	20.4	-0.1	16.9	-0.1	-0.1	5.7	-0.1	-0.1	3.9	-0.1
ND6	7.3	26.1	-0.1	8.6	26.0	9.5	8.5	3.2	35.4	-0.1	21.7	-0.1	-0.1	6.3	-0.1	7.5	4.1	-0.1
ND7	4.9	10.3	-0.1	-0.1	10.5	-0.1	-0.1	-0.1	19.0	-0.1	9.0	-0.1	-0.1	5.7	-0.1	-0.1	-0.1	-0.1
ND8	6.5	22.9	-0.1	-0.1	23.7	9.3	9.1	-0.1	37.5	-0.1	22.6	-0.1	-0.1	6.0	-0.1	7.4	4.3	-0.1
ND8-R	7.3	28.4	-0.1	8.6	29.1	9.5	9.4	3.6	42.9	-0.1	26.3	-0.1	-0.1	6.9	-0.1	7.7	4.4	-0.1
ND9	7.4	29.4	-0.1	8.8	30.3	9.5	9.2	3.7	50.1	-0.1	28.0	-0.1	-0.1	6.8	6.4	7.6	4.7	-0.1
NDD1	14.9	53.1	7.2	10.9	51.9	5.9	15.3	4.7	110.0	6.7	48.3	-0.1	7.9	7.8	7.4	9.3	5.3	7.0
NDD2	8.0	36.0	-0.1	8.4	38.7	10.0	9.9	3.7	53.7	-0.1	30.0	-0.1	-0.1	6.4	-0.1	7.7	4.7	-0.1
NDD3	7.7	31.6	-0.1	8.9	32.7	1.5	11.0	3.8	48.0	6.7	30.9	-0.1	7.5	6.4	7.0	8.2	4.6	-0.1
NDD4	8.9	29.0	-0.1	8.9	28.4	9.8	9.8	3.7	33.9	-0.1	25.1	-0.1	-0.1	6.4	-0.1	7.8	4.8	-0.1
NDD5	11.9	48.3	7.0	9.7	46.8	1.8	9.2	3.9	58.5	-0.1	43.4	-0.1	-0.1	7.1	-0.1	8.8	4.6	-0.1
NE1	22.1	70.5	7.4	12.4	71.7	4.9	21.3	5.6	82.8	7.0	58.5	-0.1	7.8	9.2	7.3	9.8	7.2	6.9
NE10	5.1	10.9	-0.1	-0.1	11.1	-0.1	-0.1	-0.1	12.6	-0.1	10.1	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	-0.1
NE11	6.1	26.5	-0.1	8.4	26.8	8.6	8.4	-0.1	40.8	-0.1	22.6	-0.1	-0.1	6.9	-0.1	7.7	4.4	-0.1
NE2	5.0	17.3	-0.1	-0.1	17.5	-0.1	-0.1	-0.1	18.8	-0.1	15.0	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NE3	5.4	28.0	-0.1	-0.1	27.1	-0.1	8.3	-0.1	33.3	-0.1	22.5	-0.1	-0.1	5.6	-0.1	-0.1	4.0	-0.1
NE4	5.6	22.6	-0.1	-0.1	23.2	-0.1	8.9	-0.1	29.1	-0.1	20.0	-0.1	-0.1	5.6	-0.1	7.4	3.9	-0.1
NE4-R	4.9	12.7	-0.1	-0.1	13.1	-0.1	-0.1	-0.1	16.9	-0.1	12.7	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1

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Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109-MAR	110-HBA	111-MAR	112-MBI	113-HBA	114-MBI	115-MBI	116-MAR	117-HA	118-MPH	119-HBA	120-THI	121-MPH	122-MPH	123-MPH	124-MBI	125-HAR	126-MPH
NE5	7.2	27.0	-0.1	8.5	26.5	9.5	9.9	3.5	38.1	-0.1	24.1	-0.1	-0.1	6.4	-0.1	7.6	4.8	-0.1
NE6	5.4	18.2	-0.1	-0.1	19.5	-0.1	-0.1	-0.1	24.6	-0.1	3.2	-0.1	-0.1	5.7	-0.1	-0.1	4.2	-0.1
NE7	5.8	14.4	-0.1	-0.1	14.9	-0.1	8.0	-0.1	15.2	-0.1	12.9	-0.1	-0.1	5.7	-0.1	-0.1	3.7	-0.1
NE8	5.0	17.5	-0.1	-0.1	17.4	-0.1	-0.1	-0.1	23.7	-0.1	15.9	-0.1	-0.1	5.7	-0.1	-0.1	-0.1	-0.1
NE9	5.5	18.0	-0.1	-0.1	18.4	-0.1	8.9	-0.1	34.2	-0.1	3.6	-0.1	-0.1	5.4	-0.1	7.4	3.8	-0.1
NEE1	6.3	25.8	-0.1	8.3	25.7	-0.1	9.4	-0.1	50.1	-0.1	22.9	-0.1	-0.1	6.5	-0.1	7.6	4.4	-0.1
NEE2	8.2	33.9	-0.1	8.9	33.9	1.7	9.4	3.6	69.6	-0.1	28.5	-0.1	-0.1	6.8	-0.1	7.4	4.6	-0.1
NEE3	10.6	46.8	-0.1	9.7	48.6	1.6	10.8	4.3	96.3	-0.1	44.4	-0.1	-0.1	6.8	6.8	8.0	5.0	-0.1
NEE4	6.7	22.9	-0.1	8.8	23.8	1.7	8.3	3.4	31.2	-0.1	19.7	-0.1	-0.1	6.1	-0.1	7.6	4.5	-0.1
NEE5	6.1	25.2	-0.1	-0.1	25.4	1.6	7.7	-0.1	33.6	-0.1	21.8	-0.1	-0.1	5.5	-0.1	-0.1	4.1	-0.1
NEE6	8.8	37.2	-0.1	9.4	38.1	1.8	10.5	4.0	48.6	6.7	38.0	-0.1	7.4	6.4	-0.1	6.2	4.7	-0.1
NEE7	7.9	31.5	-0.1	8.6	30.6	10.1	10.3	3.5	29.6	-0.1	23.3	-0.1	-0.1	6.4	6.8	7.8	4.6	-0.1
NF1	25.4	73.6	7.8	12.0	69.6	1.9	12.9	5.3	78.0	6.7	59.4	-0.1	7.5	10.3	7.0	9.6	6.8	6.8
NF10	12.7	42.3	-0.1	10.1	47.4	2.2	11.0	4.8	133.0	6.8	79.2	5.4	7.7	7.3	7.1	8.5	5.5	6.9
NF11	33.0	154.0	8.6	49.9	190.0	16.3	32.4	10.8	661.0	7.9	372.0	37.8	10.0	11.6	9.6	14.7	6.4	8.0
NF2	6.4	27.5	-0.1	8.4	27.9	1.7	7.7	-0.1	35.7	-0.1	24.3	-0.1	-0.1	5.9	-0.1	7.3	4.1	-0.1
NF3	7.2	20.7	-0.1	-0.1	21.2	8.8	8.3	-0.1	24.1	-0.1	18.4	-0.1	-0.1	6.0	-0.1	7.4	4.1	-0.1
NF4	12.4	52.5	7.4	10.6	55.2	2.2	10.2	4.8	85.8	-0.1	52.5	-0.1	-0.1	7.3	-0.1	8.6	4.9	-0.1
NF5	8.6	41.7	5.6	9.4	45.6	1.8	10.1	4.0	61.3	-0.1	48.5	-0.1	-0.1	6.6	6.8	6.3	4.5	-0.1
NF6	16.5	72.0	7.5	12.8	83.7	3.1	14.1	6.6	156.0	-0.1	100.0	6.4	-0.1	8.0	-0.1	9.3	5.4	-0.1
NF7	21.9	109.0	7.7	13.9	114.0	5.9	22.4	6.2	197.0	8.1	103.0	-0.1	9.1	13.4	8.4	11.0	9.2	7.9
NF8	5.9	27.6	-0.1	-0.1	29.1	9.2	9.1	-0.1	45.0	-0.1	31.8	-0.1	-0.1	6.9	-0.1	7.7	4.7	-0.1
NF8-R	15.8	75.0	8.0	12.4	79.8	2.2	15.5	5.6	144.0	7.0	85.2	-0.1	8.0	10.2	7.3	9.8	6.4	7.1
NF9	14.0	51.3	7.1	10.6	54.0	2.1	10.7	4.9	74.7	-0.1	55.2	-0.1	-0.1	7.3	-0.1	8.5	5.3	-0.1
NFF1	10.0	41.7	-0.1	9.5	44.4	1.6	11.1	3.9	39.4	-0.1	46.8	-0.1	-0.1	7.2	7.0	8.2	4.8	-0.1
NFF2	19.3	67.8	7.4	11.7	66.6	2.0	13.9	5.6	104.0	6.7	66.3	-0.1	7.5	9.0	7.0	9.2	6.2	6.9
NFF3	13.0	40.2	-0.1	10.4	39.9	10.0	8.9	4.7	40.2	-0.1	37.5	-0.1	-0.1	7.0	6.7	8.0	5.0	-0.1
NFF4	14.9	60.6	7.2	11.2	65.4	1.9	16.0	5.2	138.0	7.0	79.8	4.1	8.0	7.8	7.3	9.3	5.8	7.0
NFF5	9.4	39.3	-0.1	9.2	37.8	10.4	10.1	3.9	59.7	-0.1	34.2	-0.1	-0.1	7.1	-0.1	8.0	5.0	-0.1
NFF6	5.7	24.8	-0.1	-0.1	25.4	8.2	7.8	-0.1	35.1	-0.1	22.8	-0.1	-0.1	5.5	-0.1	-0.1	4.1	-0.1
NFF6-R	5.9	16.8	-0.1	-0.1	17.5	-0.1	-0.1	-0.1	23.8	-0.1	16.3	-0.1	-0.1	5.4	-0.1	-0.1	3.9	-0.1
NFF7	6.4	26.8	-0.1	-0.1	28.5	8.6	8.1	-0.1	35.1	-0.1	21.6	-0.1	-0.1	6.0	-0.1	-0.1	4.2	-0.1
NFF8	5.6	26.9	-0.1	8.4	26.3	8.6	8.2	-0.1	31.2	-0.1	23.0	-0.1	-0.1	6.0	-0.1	7.6	4.3	-0.1
NG1	4.8	12.7	-0.1	-0.1	13.5	-0.1	-0.1	-0.1	12.8	-0.1	11.2	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NG10	6.5	28.6	-0.1	8.7	29.0	1.5	8.4	3.6	45.6	-0.1	26.7	-0.1	-0.1	6.2	-0.1	7.6	4.2	-0.1
NG11	4.9	12.1	-0.1	-0.1	12.8	-0.1	7.9	-0.1	14.8	-0.1	11.7	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NG2	5.6	18.4	-0.1	-0.1	19.6	-0.1	7.7	-0.1	29.0	-0.1	13.0	-0.1	-0.1	5.7	-0.1	-0.1	4.0	-0.1
NG2-R	5.8	4.2	-0.1	-0.1	21.9	-0.1	7.8	-0.1	22.2	-0.1	16.2	-0.1	-0.1	5.6	-0.1	-0.1	3.7	-0.1
NG3	4.9	14.6	-0.1	-0.1	15.3	-0.1	7.5	-0.1	16.7	-0.1	13.5	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
NG4	5.5	16.0	-0.1	-0.1	16.3	-0.1	9.9	-0.1	22.1	-0.1	13.8	-0.1	-0.1	5.7	-0.1	8.4	4.0	-0.1
NG5	10.2	47.1	-0.1	10.1	51.3	1.9	15.5	4.3	88.5	6.8	41.4	-0.1	7.6	8.1	7.0	9.8	5.3	6.7
NG6	6.3	23.1	-0.1	-0.1	23.8	-0.1	10.3	-0.1	43.5	-0.1	20.4	-0.1	-0.1	7.5	-0.1	7.7	4.4	-0.1
NG7	-0.1	8.7	-0.1	-0.1	9.3	-0.1	-0.1	-0.1	2.1	-0.1	8.7	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NG8	5.3	16.7	-0.1	-0.1	16.6	-0.1	-0.1	-0.1	20.8	-0.1	14.0	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NG9	6.2	23.1	-0.1	-0.1	24.8	9.8	10.1	3.5	46.2	-0.1	25.4	-0.1	-0.1	6.0	-0.1	7.4	4.1	-0.1
NGG1	9.7	40.2	-0.1	9.2	39.0	9.9	9.3	3.6	38.7	-0.1	31.8	-0.1	-0.1	6.8	-0.1	7.9	4.8	-0.1
NGG10	12.1	40.2	-0.1	9.8	39.9	1.8	9.5	4.3	63.7	-0.1	28.8	-0.1	-0.1	6.9	-0.1	7.8	5.5	-0.1
NGG2	6.1	24.5	-0.1	-0.1	24.6	-0.1	9.1	-0.1	42.9	-0.1	22.6	-0.1	-0.1	6.4	-0.1	7.5	4.2	-0.1
NGG3	42.9	179.0	10.5	21.8	179.0	4.7	20.0	10.8	127.0	7.0	123.0	-0.1	7.8	13.8	7.0	12.8	8.0	7.1
NGG4	10.8	47.4	-0.1	9.5	46.2	1.7	11.1	3.8	56.1	-0.1	35.7	-0.1	-0.1	6.7	-0.1	8.0	5.0	-0.1
NGG5	7.1	25.3	-0.1	8.2	27.8	10.1	10.7	3.8	48.0	-0.1	24.7	-0.1	-0.1	6.3	-0.1	7.5	4.8	-0.1
NGG6	8.5	34.8	-0.1	9.1	34.5	10.7	11.0	3.8	56.1	-0.1	33.0	-0.1	-0.1	6.8	6.8	8.0	4.9	-0.1
NGG7	8.6	45.9	-0.1	9.2	44.4	10.1	10.0	3.8	52.8	-0.1	35.7	-0.1	-0.1	6.6	6.7	8.0	4.7	-0.1
NGG8	11.3	45.6	-0.1	9.6	44.7	1.7	9.6	4.0	53.1	-0.1	35.1	-0.1	-0.1	6.8	-0.1	7.7	4.8	-0.1
NGG9	7.1	24.0	-0.1	-0.1	24.1	8.4	8.3	3.5	32.7	-0.1	20.7	-0.1	-0.1	5.9	-0.1	-0.1	4.3	-0.1
NH1	5.5	23.9	-0.1	-0.1	24.5	-0.1	8.3	-0.1	37.2	-0.1	21.4	-0.1	-0.1	5.5	-0.1	7.3	3.9	-0.1
NH10	5.3	15.7	-0.1	-0.1	15.6	-0.1	8.3	-0.1	17.0	-0.1	12.0	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	-0.1
NH10-R	5.1	16.4	-0.1	-0.1	16.3	-0.1	9.9	-0.1	22.4	-0.1	14.1	-0.1	-0.1	5.4	-0.1	7.2	-0.1	-0.1
NH11	5.9	26.3	-0.1	-0.1	26.8	9.7	9.7	-0.1	36.9	-0.1	23.7	-0.1	-0.1	6.0	-0.1	7.6	4.1	-0.1
NH2	5.6	21.6	-0.1	-0.1	21.4	-0.1	8.1	-0.1	27.5	-0.1	16.6	-0.1	-0.1	5.6	-0.1	-0.1	3.7	-0.1
NH3	-0.1	12.4	-0.1	-0.1	12.7	-0.1	-0.1	-0.1	14.8	-0.1	11.4	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
NH4	5.5	20.3	-0.1	-0.1	21.4	-0.1	9.0	-0.1	40.2	-0.1	19.4	-0.1	-0.1	5.8	-0.1	7.2	3.9	-0.1
NH5	5.4	17.4	-0.1	-0.1	18.3	-0.1	8.0	-0.1	23.9	-0.1	14.5	-0.1	-0.1	6.0	-0.1	-0.1	3.7	-0.1
NH6	5.6	19.3	-0.1	-0.1	19.7	-0.1	10.3	-0.1	25.7	-0.1	17.2	-0.1	-0.1	5.8	-0.1	7.6	4.0	-0.1
NH7	6.7	45.3	7.0	9.2	46.8	14.4	10.2	3.6	76.2	7.0	43.5	-0.1	8.0	6.6	-0.1	8.9	4.5	7.2
NH8	5.9	22.4	-0.1	-0.1	23.1	-0.1	9.8	-0.1	24.0	-0.1	18.6	-0.1	-0.1	5.6	-0.1	7.6	3.8	-0.1
NH9	5.9	24.6	-0.1	8.3	25.2	-0.1	9.8	-0.1	32.1	-0.1	21.2	-0.1	-0.1	6.5	-0.1	7.6	4.2	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109-MAR	110-HBA	111-MAR	112-MBI	113-HBA	114-MBI	115-MBI	116-MAR	117-HA	118-MPH	119-HBA	120-THI	121-MPH	122-MPH	123-MPH	124-MBI	125-HAR	126-MPH
NHH1	7.4	30.6	-0.1	8.8	31.2	1.4	9.5	3.7	52.5	-0.1	28.6	-0.1	-0.1	6.0	-0.1	7.6	4.3	-0.1
NHH10	7.1	28.4	-0.1	8.5	27.7	8.7	7.8	-0.1	39.6	-0.1	23.4	-0.1	-0.1	6.0	-0.1	7.4	4.1	-0.1
NHH11	11.2	39.9	-0.1	9.1	37.8	1.8	8.5	3.6	33.0	-0.1	27.7	-0.1	-0.1	6.9	-0.1	7.7	4.6	-0.1
NHH11-R	8.6	28.6	-0.1	8.7	27.7	1.5	8.6	3.7	30.3	-0.1	22.4	-0.1	-0.1	6.4	-0.1	7.4	4.5	-0.1
NHH12	6.5	29.8	-0.1	-0.1	29.4	9.0	8.7	-0.1	39.6	-0.1	24.6	-0.1	-0.1	5.9	-0.1	7.3	4.3	-0.1
NHH2	28.2	108.0	8.0	15.1	113.0	3.4	17.5	7.1	153.0	6.6	103.0	-4.1	7.5	19.6	6.8	10.2	6.9	-0.1
NHH3	7.7	37.8	-0.1	8.9	38.7	10.5	11.0	3.7	59.1	-0.1	36.6	-0.1	-0.1	6.5	-0.1	7.9	4.6	-0.1
NHH4	17.9	51.0	6.7	10.5	54.6	2.1	10.6	5.1	63.4	-0.1	44.1	-0.1	-0.1	7.4	8.1	6.8	8.3	6.5
NHH5	14.0	56.7	6.5	9.9	59.4	2.0	10.4	5.0	66.9	-0.1	44.4	-0.1	-0.1	7.8	-0.1	8.0	6.0	-0.1
NHH6	7.1	21.5	-0.1	-0.1	21.7	9.4	8.8	-0.1	27.0	-0.1	18.3	-0.1	-0.1	5.8	-0.1	7.2	4.1	-0.1
NHH7	7.6	33.3	-0.1	8.7	33.0	9.2	8.9	3.7	43.5	-0.1	28.1	-0.1	-0.1	6.1	-0.1	7.5	4.4	-0.1
NHH8	8.4	32.7	-0.1	8.9	34.2	1.8	8.7	3.6	48.9	-0.1	29.3	-0.1	-0.1	5.9	-0.1	7.6	4.3	-0.1
NHH9	8.6	41.1	-0.1	8.7	42.9	1.8	8.3	3.8	47.7	-0.1	31.5	-0.1	-0.1	6.4	-0.1	7.7	4.7	-0.1
NI1	15.8	85.8	7.9	13.3	93.6	2.8	15.5	5.6	136.0	7.0	73.5	-0.1	7.8	8.6	7.3	9.2	6.4	7.0
NI10	6.0	28.7	-0.1	8.4	28.7	8.6	7.7	-0.1	30.9	-0.1	20.8	-0.1	-0.1	5.7	-0.1	7.3	4.0	-0.1
NI2	5.2	16.9	-0.1	-0.1	17.3	-0.1	7.1	-0.1	29.0	-0.1	17.1	-0.1	-0.1	5.6	-0.1	7.2	3.9	-0.1
NI3	5.6	22.2	-0.1	-0.1	23.1	-0.1	8.2	-0.1	36.3	-0.1	21.0	-0.1	-0.1	5.8	-0.1	7.2	3.9	-0.1
NI4	11.9	55.8	7.8	10.1	60.0	11.3	11.2	4.1	47.7	-0.1	38.1	-0.1	-0.1	6.8	-0.1	8.6	4.8	6.8
NI5	6.9	30.9	-0.1	8.6	31.2	11.1	11.8	3.5	51.6	-0.1	26.0	-0.1	-0.1	6.6	-0.1	8.0	4.5	-0.1
NI5-R	6.4	25.5	-0.1	8.4	25.7	9.5	9.6	-0.1	33.9	-0.1	20.9	-0.1	-0.1	6.2	-0.1	7.6	4.3	-0.1
NI6	5.3	17.6	-0.1	-0.1	19.0	-0.1	8.7	-0.1	36.6	-0.1	19.2	-0.1	-0.1	6.3	-0.1	7.2	3.8	-0.1
NI7	12.2	75.0	7.3	11.3	80.4	2.7	11.8	4.5	79.5	-0.1	53.7	-0.1	-0.1	7.0	-0.1	8.3	5.3	-0.1
NI8	5.2	16.1	-0.1	-0.1	16.0	-0.1	7.7	-0.1	23.6	-0.1	14.9	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NI9	6.2	29.2	-0.1	-0.1	29.4	9.6	9.7	-0.1	36.9	-0.1	24.1	-0.1	-0.1	5.7	-0.1	7.5	4.0	-0.1
NII1	4.9	11.3	-0.1	-0.1	11.4	-0.1	-0.1	-0.1	17.9	-0.1	2.5	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NII10	9.6	36.6	-0.1	9.1	39.0	1.7	10.0	4.1	53.8	-0.1	41.4	-0.1	-0.1	6.6	-0.1	7.8	4.6	-0.1
NII11	6.2	19.3	-0.1	-0.1	20.0	10.0	9.6	-0.1	28.8	-0.1	17.8	-0.1	-0.1	5.8	6.6	7.5	3.9	-0.1
NII11-R	7.6	27.2	-0.1	8.9	29.1	10.0	9.9	3.8	53.7	-0.1	35.4	-0.1	7.4	6.2	6.6	7.8	4.3	-0.1
NII12	7.4	24.3	-0.1	8.7	26.1	1.5	10.1	3.8	39.0	-0.1	27.2	-0.1	-0.1	6.1	-0.1	7.7	4.3	-0.1
NII13	13.6	51.6	6.9	10.8	54.3	5.4	15.1	5.2	80.7	-0.1	54.9	-0.1	-0.1	7.2	-0.1	8.6	5.0	-0.1
NII14	5.8	16.5	-0.1	-0.1	17.0	-0.1	7.7	-0.1	19.4	-0.1	15.5	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NII2	6.6	30.9	-0.1	8.8	33.9	1.6	10.8	3.6	66.9	-0.1	34.8	-0.1	-0.1	6.8	6.6	8.9	4.4	6.6
NII3	11.8	64.2	7.9	10.5	66.6	2.0	10.0	3.9	84.9	-0.1	46.8	-0.1	-0.1	7.3	-0.1	8.7	5.1	-0.1
NII4	12.7	87.6	9.0	12.3	95.7	12.3	11.0	5.0	66.7	-0.1	58.5	-0.1	-0.1	7.4	-0.1	8.0	5.0	-0.1
NII5	8.7	34.5	-0.1	8.9	37.8	1.6	10.5	3.7	77.7	-0.1	44.7	-0.1	-0.1	6.2	-0.1	7.9	4.1	-0.1
NII6	5.7	20.6	-0.1	-0.1	20.7	-0.1	-0.1	-0.1	27.0	-0.1	18.6	-0.1	-0.1	5.6	-0.1	-0.1	3.9	-0.1
NII7	5.2	18.2	-0.1	-0.1	18.3	-0.1	-0.1	-0.1	26.3	-0.1	17.8	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NII8	14.7	60.9	7.6	11.1	62.1	2.5	10.5	4.7	70.5	-0.1	46.2	-0.1	-0.1	7.5	-0.1	8.8	4.9	-0.1
NII9	5.7	18.8	-0.1	-0.1	18.9	-0.1	7.7	-0.1	26.0	-0.1	17.6	-0.1	-0.1	5.7	-0.1	-0.1	3.9	-0.1
NJ1	6.5	22.9	-0.1	8.5	24.3	10.6	10.9	-0.1	41.4	-0.1	4.8	-0.1	-0.1	5.9	-0.1	7.8	4.2	-0.1
NJ2	8.8	39.6	-0.1	9.3	41.1	1.6	10.9	3.9	48.9	-0.1	33.0	-0.1	-0.1	5.9	-0.1	7.8	4.3	-0.1
NJ3	8.1	42.3	-0.1	9.2	43.5	1.5	9.9	3.6	55.8	-0.1	34.2	-0.1	-0.1	5.7	-0.1	7.3	3.9	-0.1
NJ4	7.4	38.1	-0.1	8.9	39.9	1.6	10.3	3.4	64.2	-0.1	33.0	-0.1	-0.1	5.9	-0.1	7.8	4.1	-0.1
NJ5	6.2	18.8	-0.1	-0.1	20.1	1.6	8.6	-0.1	32.7	-0.1	17.2	-0.1	-0.1	5.7	-0.1	7.4	3.8	-0.1
NJ6	5.1	18.0	-0.1	-0.1	18.9	-0.1	8.6	-0.1	28.3	-0.1	17.9	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NJ7	6.2	22.6	-0.1	-0.1	23.7	8.8	8.4	-0.1	27.0	-0.1	19.0	-0.1	-0.1	5.7	-0.1	7.5	4.3	-0.1
NJJ1	7.2	28.6	-0.1	8.6	29.8	10.1	9.7	-0.1	51.9	-0.1	24.5	-0.1	-0.1	7.5	-0.1	7.9	4.3	-0.1
NJJ10	133.0	366.0	18.7	38.4	348.0	20.2	34.5	26.2	366.0	9.2	351.0	-7.7	10.6	28.3	9.9	23.2	13.7	9.5
NJJ11	6.0	21.0	-0.1	-0.1	21.5	-0.1	7.7	-0.1	30.9	-0.1	21.9	-0.1	-0.1	5.9	-0.1	-0.1	4.0	-0.1
NJJ12	11.5	42.9	-0.1	9.5	40.5	10.3	10.0	4.2	54.9	-0.1	38.7	-0.1	-0.1	7.4	-0.1	6.3	5.1	-0.1
NJJ13	26.0	106.0	8.5	14.4	104.0	2.9	12.7	6.6	96.3	-0.1	73.2	-0.1	-0.1	10.3	6.8	9.6	6.0	-0.1
NJJ14	6.2	21.3	-0.1	-0.1	21.1	-0.1	7.4	-0.1	26.2	-0.1	18.5	-0.1	-0.1	5.8	-0.1	-0.1	3.8	-0.1
NJJ2	7.1	23.6	-0.1	8.4	23.6	8.8	8.5	-0.1	30.9	-0.1	3.6	-0.1	-0.1	6.0	-0.1	7.4	4.3	-0.1
NJJ3	5.6	18.4	-0.1	-0.1	18.6	-0.1	-0.1	-0.1	28.1	-0.1	8.2	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NJJ4	16.6	75.0	7.5	12.5	78.3	2.9	12.0	5.6	136.0	4.1	71.1	-0.1	-0.1	8.4	-0.1	8.5	5.0	-0.1
NJJ5	6.2	22.9	-0.1	-0.1	23.0	-0.1	7.8	-0.1	28.3	-0.1	18.6	-0.1	-0.1	5.8	-0.1	-0.1	4.8	-0.1
NJJ6	6.7	27.0	-0.1	8.7	28.9	9.3	9.1	3.7	57.0	-0.1	28.9	-0.1	-0.1	6.2	-0.1	7.5	4.2	-0.1
NJJ7	5.8	17.7	-0.1	-0.1	19.0	-0.1	8.0	-0.1	32.4	-0.1	18.3	-0.1	-0.1	5.6	-0.1	7.4	3.7	-0.1
NJJ8	6.5	20.9	-0.1	-0.1	20.5	-0.1	-0.1	-0.1	22.9	-0.1	16.3	-0.1	-0.1	5.6	-0.1	-0.1	3.9	-0.1
NJJ8-R	8.9	30.0	-0.1	8.8	29.3	1.5	8.0	3.8	45.3	-0.1	23.7	-0.1	-0.1	6.3	-0.1	7.6	4.4	-0.1
NJJ9	20.6	103.0	7.9	15.4	115.0	3.8	16.1	7.2	193.0	-0.1	106.0	6.3	7.7	8.9	7.2	10.0	5.9	6.8
NK1	5.4	23.8	-0.1	-0.1	24.5	9.5	9.7	-0.1	42.9	-0.1	26.9	-0.1	-0.1	5.7	-0.1	7.5	4.0	-0.1
NK2	6.2	20.8	-0.1	-0.1	21.1	-0.1	7.8	-0.1	26.3	-0.1	18.1	-0.1	-0.1	5.8	-0.1	-0.1	4.0	-0.1
NK3	6.7	28.0	-0.1	-0.1	28.9	9.7	9.0	-0.1	39.8	-0.1	26.6	-0.1	-0.1	5.6	-0.1	7.4	3.9	-0.1
NK4	6.5	26.9	6.9	8.3	28.3	9.5	9.5	-0.1	29.3	-0.1	21.2	-0.1	-0.1	6.0	-0.1	7.8	4.3	-0.1
NK4-R	5.9	4.6	-0.1	8.4	22.6	8.9	8.6	-0.1	22.6	-0.1	17.5	-0.1	-0.1	5.9	-0.1	7.5	4.2	-0.1
NK5	5.6	19.6	-0.1	-0.1	21.2	10.9	11.1	-0.1	35.4	-0.1	19.4	-0.1	-0.1	5.9	-0.1	7.5	3.8	-0.1

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109-MAR	110-HBA	111-MAR	112-MBI	113-HBA	114-MBI	115-MBI	116-MAR	117-HA	118-MPH	119-HBA	120-THI	121-MPH	122-MPH	123-MPH	124-MBI	125-HAR	126-MPH
NK6	6.0	27.2	-0.1	8.4	26.9	9.9	9.9	-0.1	42.3	-0.1	23.6	-0.1	-0.1	6.2	-0.1	7.6	4.2	-0.1
NK7	6.3	26.4	-0.1	-0.1	27.5	1.5	8.8	3.5	35.1	-0.1	22.5	-0.1	-0.1	5.8	-0.1	7.4	4.1	-0.1
NKK1	8.3	22.5	-0.1	-0.1	22.1	8.5	7.5	-0.1	21.8	-0.1	17.1	-0.1	-0.1	6.3	-0.1	-0.1	4.2	-0.1
NKK10	6.3	25.4	-0.1	-0.1	26.0	8.6	8.2	3.5	39.6	-0.1	24.9	-0.1	-0.1	5.8	-0.1	7.3	4.1	-0.1
NKK11	10.0	33.0	-0.1	8.7	34.8	10.0	10.0	4.3	51.9	-0.1	27.9	-0.1	-0.1	7.4	-0.1	7.7	5.4	-0.1
NKK2	8.9	34.2	-0.1	8.9	33.9	1.6	8.7	3.8	55.5	-0.1	31.8	-0.1	-0.1	7.4	-0.1	7.9	5.0	-0.1
NKK3	28.4	78.9	7.9	13.1	72.6	2.8	10.7	5.6	79.2	-0.1	60.0	-0.1	-0.1	10.8	-0.1	10.8	6.6	-0.1
NKK4	124.0	396.0	14.6	41.7	396.0	10.9	31.5	24.9	315.0	7.6	306.0	9.9	8.5	26.7	7.6	19.8	13.1	7.5
NKK5	57.6	194.0	10.0	21.4	201.0	2.6	17.0	12.1	129.0	6.8	133.0	-0.1	7.5	15.7	7.0	12.7	9.0	6.8
NKK6	18.8	51.0	6.9	10.3	47.4	10.1	9.1	5.0	54.0	-0.1	40.2	-0.1	-0.1	8.5	-0.1	8.3	6.2	-0.1
NKK7	22.4	55.5	7.5	10.9	51.0	2.3	9.3	5.0	45.6	-0.1	42.6	-0.1	-0.1	9.2	-0.1	6.6	5.9	-0.1
NKK8	8.5	30.9	-0.1	9.1	33.0	1.7	10.1	3.9	69.3	-0.1	36.6	-0.1	-0.1	7.3	-0.1	8.0	4.6	-0.1
NKK9	7.9	30.9	-0.1	-0.1	32.1	9.9	9.0	6.7	53.1	-0.1	33.6	-0.1	-0.1	6.2	-0.1	7.6	4.4	-0.1
NL1	6.9	20.2	-0.1	-0.1	20.4	-0.1	7.8	-0.1	26.3	-0.1	18.2	-0.1	-0.1	6.0	-0.1	7.2	4.0	-0.1
NL2	7.5	23.9	-0.1	8.4	23.8	8.2	7.7	-0.1	28.3	-0.1	20.7	-0.1	-0.1	6.2	-0.1	7.3	4.4	-0.1
NL3	8.4	24.7	-0.1	8.2	24.6	-0.1	8.5	3.4	29.8	-0.1	22.1	-0.1	-0.1	6.4	-0.1	7.5	4.6	-0.1
NL3-R	11.9	36.0	-0.1	9.1	35.1	9.9	9.8	4.0	43.5	-0.1	30.9	-0.1	-0.1	7.2	-0.1	8.0	5.1	-0.1
NL4	6.7	25.5	-0.1	-0.1	25.5	8.5	7.8	-0.1	26.7	-0.1	21.2	-0.1	-0.1	6.0	-0.1	7.3	4.2	-0.1
NLL1	17.9	54.0	7.7	10.7	50.4	10.8	9.7	5.0	49.2	-0.1	44.3	-0.1	-0.1	8.2	-0.1	6.8	5.0	-0.1
NLL10	8.3	34.5	-0.1	8.9	33.9	1.8	8.5	3.8	45.0	-0.1	27.4	-0.1	-0.1	6.6	-0.1	7.5	4.5	-0.1
NLL11	6.8	24.8	-0.1	-0.1	25.0	9.2	8.6	-0.1	33.2	-0.1	22.2	-0.1	-0.1	6.1	-0.1	7.8	4.1	-0.1
NLL12	5.7	19.6	-0.1	-0.1	20.3	-0.1	8.3	-0.1	31.2	-0.1	18.9	-0.1	-0.1	6.1	-0.1	-0.1	4.1	-0.1
NLL2	7.3	24.6	-0.1	8.5	24.4	1.7	7.6	3.6	25.5	-0.1	18.3	-0.1	-0.1	5.9	-0.1	-0.1	4.3	-0.1
NLL3	6.3	17.3	-0.1	-0.1	18.5	-0.1	7.6	-0.1	24.7	-0.1	16.0	-0.1	-0.1	6.3	-0.1	-0.1	4.1	-0.1
NLL4	18.1	45.6	7.7	9.8	45.0	10.0	9.4	4.2	59.7	-0.1	32.4	-0.1	-0.1	9.0	-0.1	8.2	6.3	-0.1
NLL5	19.5	48.3	8.4	12.8	88.5	12.1	10.4	5.6	66.0	-0.1	50.7	-0.1	-0.1	8.6	-0.1	8.6	6.1	-0.1
NLL6	26.3	122.0	9.8	15.7	129.0	15.4	13.4	5.9	186.0	6.5	80.3	-0.1	7.6	10.7	6.9	10.2	7.4	6.8
NLL7	46.8	174.0	9.5	19.7	168.0	5.1	17.1	9.0	121.0	6.7	96.0	-0.1	7.6	12.9	7.3	11.2	8.7	6.8
NLL8	6.6	24.7	-0.1	8.6	25.6	8.8	8.5	3.5	48.3	-0.1	23.6	-0.1	-0.1	6.3	-0.1	7.6	4.2	-0.1
NLL9	23.4	70.2	7.4	12.3	63.6	2.9	11.4	5.6	60.3	6.8	48.0	-0.1	7.6	9.7	7.0	9.2	7.3	6.8
NLL9-R	24.3	69.3	7.3	11.6	64.8	2.1	11.2	5.3	60.3	6.3	48.5	-0.1	7.6	9.2	6.9	8.8	7.0	6.8
NM1	10.3	33.6	-0.1	9.1	32.4	9.8	9.5	3.9	43.8	-0.1	28.3	-0.1	-0.1	6.8	-0.1	7.8	5.0	-0.1
NM2	5.9	22.1	-0.1	-0.1	23.2	9.2	8.8	-0.1	36.6	-0.1	22.8	-0.1	-0.1	6.7	-0.1	7.8	4.3	-0.1
NMM1	5.5	17.2	-0.1	-0.1	17.4	-0.1	7.6	-0.1	19.1	-0.1	14.7	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NMM10	13.6	55.8	7.4	10.1	54.0	1.8	9.8	4.6	77.4	-0.1	42.9	-0.1	-0.1	7.7	-0.1	6.7	6.4	-0.1
NMM2	7.7	34.8	-0.1	8.9	36.3	1.7	9.0	3.8	45.3	-0.1	31.8	-0.1	-0.1	6.7	-0.1	7.6	4.5	-0.1
NMM3	5.8	4.1	-0.1	-0.1	20.6	-0.1	7.9	-0.1	26.6	-0.1	17.6	-0.1	-0.1	5.6	-0.1	7.4	3.8	-0.1
NMM3-R	5.2	16.7	-0.1	-0.1	17.0	-0.1	7.9	-0.1	24.4	-0.1	16.2	-0.1	-0.1	5.7	-0.1	7.2	-0.1	-0.1
NMM4	5.4	16.7	-0.1	-0.1	17.3	-0.1	7.9	-0.1	29.2	-0.1	17.0	-0.1	-0.1	6.2	-0.1	-0.1	3.9	-0.1
NMM5	5.6	19.0	-0.1	-0.1	19.7	-0.1	7.8	-0.1	27.1	-0.1	17.2	-0.1	-0.1	6.3	-0.1	-0.1	3.7	-0.1
NMM6	8.9	39.0	7.0	9.7	40.5	1.9	10.2	4.1	50.4	-0.1	32.4	-0.1	-0.1	6.5	-0.1	6.6	5.3	6.8
NMM7	5.2	16.7	-0.1	-0.1	17.5	-0.1	-0.1	-0.1	23.1	-0.1	15.3	-0.1	-0.1	5.6	-0.1	-0.1	3.9	-0.1
NMM8	19.3	72.9	7.5	11.5	69.6	2.7	9.5	5.0	112.0	-0.1	54.6	-0.1	-0.1	8.6	-0.1	8.8	5.6	-0.1
NMM9	8.9	41.7	-0.1	9.6	46.8	1.6	12.6	4.1	103.0	6.7	46.5	-0.1	7.6	7.8	7.1	8.4	4.9	6.8
NN1	4.8	14.7	-0.1	-0.1	14.8	-0.1	-0.1	-0.1	15.7	-0.1	13.6	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NNN1	5.8	18.8	-0.1	-0.1	18.9	-0.1	7.5	-0.1	30.3	-0.1	17.8	-0.1	-0.1	5.7	-0.1	-0.1	3.9	-0.1
NNN10	5.0	14.7	-0.1	-0.1	16.6	-0.1	9.1	-0.1	31.2	-0.1	24.9	-0.1	-0.1	6.7	-0.1	-0.1	-0.1	-0.1
NNN2	6.5	27.9	-0.1	8.3	29.3	9.1	8.5	3.4	50.1	-0.1	27.5	-0.1	-0.1	5.8	-0.1	7.4	3.8	-0.1
NNN3	-0.1	10.6	-0.1	-0.1	11.0	-0.1	-0.1	-0.1	11.6	-0.1	9.9	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NNN4	9.6	43.5	-0.1	9.8	46.2	1.6	13.6	4.5	94.8	-0.1	44.7	-0.1	7.4	6.8	6.9	8.5	4.7	6.7
NNN5	4.7	15.5	-0.1	-0.1	16.8	-0.1	-0.1	-0.1	24.3	-0.1	15.8	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NNN6	5.0	14.2	-0.1	-0.1	14.7	-0.1	7.6	-0.1	14.2	-0.1	11.8	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
NNN7	-0.1	11.2	-0.1	-0.1	11.7	-0.1	-0.1	-0.1	1.9	-0.1	10.0	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NNN8	10.0	44.1	-0.1	9.9	46.8	4.9	13.8	4.1	122.0	-0.1	48.3	-0.1	-0.1	7.2	-0.1	6.6	8.3	5.0
NNN9	6.8	28.9	-0.1	8.2	28.7	8.8	8.4	3.5	27.3	-0.1	20.9	-0.1	-0.1	5.9	-0.1	7.4	4.1	-0.1
NO1	9.9	34.5	-0.1	8.4	34.8	8.5	8.3	4.1	32.4	-0.1	25.0	-0.1	-0.1	7.2	-0.1	7.6	5.2	-0.1
NO2	5.0	9.5	-0.1	-0.1	10.0	-0.1	-0.1	-0.1	11.5	-0.1	9.5	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NOO1	5.2	17.7	-0.1	-0.1	19.3	-0.1	8.0	-0.1	34.5	-0.1	20.5	-0.1	-0.1	5.7	-0.1	-0.1	-0.1	-0.1
NOO2	7.2	30.6	-0.1	8.7	30.6	9.4	9.2	3.7	54.6	-0.1	27.2	-0.1	-0.1	6.5	-0.1	7.7	4.3	-0.1
NOO3	5.1	10.1	-0.1	-0.1	10.7	-0.1	9.0	-0.1	11.1	-0.1	10.1	-0.1	-0.1	5.5	-0.1	7.2	-0.1	-0.1
NOO4	6.0	21.2	-0.1	-0.1	22.2	9.7	9.6	3.5	35.1	-0.1	19.9	-0.1	-0.1	5.9	-0.1	7.5	4.1	-0.1
NOO5	5.4	17.4	-0.1	-0.1	17.3	-0.1	-0.1	-0.1	23.9	-0.1	15.0	-0.1	-0.1	5.6	-0.1	-0.1	3.6	-0.1
NOO6	4.8	14.6	-0.1	-0.1	14.9	-0.1	-0.1	-0.1	16.5	-0.1	12.4	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
NOO6-R	5.0	14.7	-0.1	-0.1	15.3	-0.1	7.6	-0.1	17.5	-0.1	13.4	-0.1	-0.1	5.8	-0.1	-0.1	-0.1	-0.1
NOO7	5.0	17.5	-0.1	-0.1	18.2	-0.1	-0.1	-0.1	26.0	-0.1	17.1	-0.1	-0.1	6.8	-0.1	-0.1	-0.1	-0.1
NOO8	13.9	61.8	7.1	11.0	65.4	2.0	12.8	4.7	178.0	-0.1	69.6	-0.1	7.7	8.6	6.9	8.6	5.7	6.7
NP1	8.8	25.4	-0.1	-0.1	25.1	9.2	8.5	3.4	31.2	-0.1	21.5	-0.1	-0.1	6.5	-0.1	7.4	4.4	-0.1

A14-06865

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Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109 MAR	110 HBA	111 MAR	112 MBI	113 HBA	114 MBI	115 MBI	116 MAR	117 HA	118 MPH	119 HBA	120 THI	121 MPH	122 MPH	123 MPH	124 MBI	125 HAR	126 MPH
NP2	7.3	23.1	-0.1	-0.1	22.8	-0.1	8.0	3.5	29.5	-0.1	20.7	-0.1	-0.1	6.4	-0.1	7.3	4.4	-0.1
NP3	7.1	18.8	-0.1	-0.1	18.6	-0.1	7.5	-0.1	23.4	-0.1	16.4	-0.1	-0.1	5.9	-0.1	-0.1	3.9	-0.1
NP4	32.4	89.4	7.1	13.8	99.3	3.6	19.8	6.7	120.0	7.3	76.2	-0.1	8.3	12.8	7.7	10.9	9.8	7.4
NPP4	5.8	14.5	-0.1	-0.1	15.3	-0.1	-0.1	-0.1	1.2	-0.1	13.2	-0.1	-0.1	6.0	-0.1	-0.1	4.0	-0.1
NPP1-R	5.1	13.2	-0.1	-0.1	13.5	-0.1	-0.1	-0.1	12.4	-0.1	11.6	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NPP2	16.3	65.1	7.6	11.9	64.5	2.4	12.8	6.5	82.8	6.6	56.4	-0.1	-0.1	8.7	6.8	9.4	5.6	6.8
NPP3	12.2	51.0	7.1	10.2	51.9	2.1	10.0	4.6	75.3	-0.1	45.3	-0.1	-0.1	7.4	-0.1	8.2	5.0	-0.1
NPP4	8.6	31.8	-0.1	8.8	32.4	10.0	9.9	3.7	46.8	-0.1	28.8	-0.1	-0.1	5.9	-0.1	7.6	4.2	-0.1
NQ1	11.5	24.7	-0.1	8.8	24.7	10.6	11.0	3.8	39.0	6.8	27.7	-0.1	7.6	8.2	6.8	8.3	5.8	7.0
NQ2	30.6	96.9	8.5	14.9	90.9	14.5	11.5	7.5	86.4	6.4	83.7	-0.1	7.4	10.6	6.8	10.3	5.9	-0.1
NQ2-R	26.2	102.0	7.9	14.3	104.0	2.8	14.5	7.2	113.0	-0.1	97.5	-0.1	-0.1	10.3	6.6	10.2	6.1	-0.1
NQ3	9.9	35.7	-0.1	8.7	37.8	1.9	8.3	4.0	37.5	-0.1	29.9	-0.1	-0.1	6.8	-0.1	7.7	4.7	-0.1
NQ4	16.7	62.1	7.0	10.8	60.3	2.1	11.8	4.9	84.3	-0.1	57.3	-0.1	-0.1	7.7	-0.1	8.6	5.3	-0.1
NQ5	8.7	32.1	-0.1	8.6	1.4	9.2	9.0	3.6	31.2	-0.1	24.2	-0.1	-0.1	6.4	-0.1	7.7	4.7	-0.1
NQ6	5.1	19.0	-0.1	-0.1	19.5	-0.1	7.5	-0.1	26.8	-0.1	18.0	-0.1	-0.1	5.8	-0.1	-0.1	-0.1	-0.1
NQ7	13.7	51.0	-0.1	10.5	54.0	5.3	15.1	4.8	122.0	6.6	62.1	3.4	-0.1	8.2	6.9	8.6	5.8	-0.1
NQ8	6.8	22.1	-0.1	-0.1	22.0	8.9	8.7	-0.1	28.4	-0.1	19.0	-0.1	-0.1	5.6	-0.1	7.2	4.0	-0.1
NQQ1	5.8	16.9	-0.1	-0.1	16.9	-0.1	8.0	-0.1	24.2	-0.1	15.7	-0.1	-0.1	5.6	-0.1	-0.1	3.8	-0.1
NR1	5.3	14.6	-0.1	-0.1	15.6	-0.1	-0.1	-0.1	16.8	-0.1	12.0	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NR11	5.7	22.4	-0.1	-0.1	22.4	9.3	9.2	-0.1	29.5	-0.1	18.0	-0.1	-0.1	6.2	-0.1	7.4	4.2	-0.1
NR12	5.4	23.4	-0.1	-0.1	24.0	-0.1	8.8	-0.1	31.5	-0.1	20.3	-0.1	-0.1	5.9	-0.1	7.4	3.9	-0.1
NR13	9.6	41.4	7.0	9.9	43.8	2.2	10.6	4.1	56.4	-0.1	30.9	-0.1	-0.1	6.6	-0.1	8.0	4.6	-0.1
NR14	5.6	23.1	-0.1	-0.1	25.3	8.1	7.7	-0.1	33.3	-0.1	20.9	-0.1	-0.1	6.2	-0.1	7.4	4.2	-0.1
NR2	5.9	21.0	-0.1	8.2	21.9	9.0	8.5	-0.1	26.0	-0.1	18.9	-0.1	-0.1	5.9	-0.1	7.4	4.1	-0.1
NR3	6.3	21.3	-0.1	-0.1	22.2	-0.1	9.6	-0.1	25.0	-0.1	15.6	-0.1	-0.1	5.7	-0.1	7.4	3.8	-0.1
NR4	4.7	13.6	-0.1	-0.1	14.4	-0.1	8.6	-0.1	15.9	-0.1	12.8	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NR5	5.5	18.0	-0.1	-0.1	17.9	-0.1	-0.1	-0.1	22.1	-0.1	14.8	-0.1	-0.1	5.6	-0.1	-0.1	3.8	-0.1
NR6	4.8	14.3	-0.1	-0.1	14.3	-0.1	-0.1	-0.1	15.9	-0.1	12.4	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
NR7	11.8	55.2	7.5	10.5	55.5	1.6	11.5	4.3	93.3	6.5	46.8	-0.1	7.6	7.5	7.0	8.9	4.8	6.8
NR8	5.4	17.6	-0.1	-0.1	18.4	-0.1	7.3	-0.1	24.7	-0.1	16.5	-0.1	-0.1	5.8	-0.1	-0.1	-0.1	-0.1
NS1	6.8	19.8	-0.1	-0.1	20.9	-0.1	7.7	-0.1	20.7	-0.1	15.9	-0.1	-0.1	6.1	-0.1	7.4	4.5	-0.1
NS10	26.2	87.9	7.4	13.7	93.9	3.2	15.0	6.7	179.0	6.5	110.0	6.9	7.7	9.8	7.2	10.0	6.4	7.0
NS11	10.9	42.3	-0.1	9.2	41.7	1.4	9.7	4.1	52.5	-0.1	39.3	-0.1	-0.1	7.0	-0.1	8.3	4.8	-0.1
NS12	13.4	49.8	-0.1	10.1	53.1	2.0	12.2	4.6	119.0	-0.1	72.0	-0.1	4.1	7.3	-6.7	6.4	5.3	-0.1
NS13	23.0	89.7	8.3	13.1	87.0	2.4	14.7	6.1	99.3	6.5	78.6	-0.1	7.5	9.1	7.0	10.4	5.7	6.9
NS14	24.6	95.7	6.2	13.4	92.7	5.2	25.2	6.3	133.0	7.5	84.6	-0.1	8.7	10.5	8.5	12.6	7.4	7.8
NS2	8.7	39.9	-0.1	9.1	40.8	1.6	8.8	3.8	49.8	-0.1	36.6	-0.1	-0.1	6.3	-0.1	7.7	4.7	-0.1
NS3	14.5	43.5	-0.1	9.7	43.2	1.6	11.6	4.2	54.6	6.4	36.3	-0.1	7.4	7.7	6.7	8.3	6.6	-0.1
NS4	9.5	39.6	7.4	9.0	39.9	9.5	8.8	3.4	37.5	-0.1	28.3	-0.1	-0.1	7.4	-0.1	8.2	5.0	-0.1
NS4-R	9.6	36.0	7.0	8.7	38.7	9.2	9.5	3.6	42.0	-0.1	27.6	-0.1	-0.1	7.0	-0.1	8.2	5.1	-0.1
NS5	14.9	49.5	7.9	10.2	50.1	1.7	9.4	4.3	56.1	-0.1	36.9	-0.1	-0.1	8.0	-0.1	8.5	5.4	6.6
NS6	8.0	29.3	-0.1	8.7	28.5	8.9	9.5	3.6	67.5	-0.1	24.3	-0.1	-0.1	6.4	-0.1	7.5	4.5	-0.1
NS7	22.6	77.4	8.5	12.3	74.1	2.0	10.8	5.6	69.6	-0.1	57.9	-0.1	-0.1	9.5	-0.1	9.3	5.7	-0.1
NS8	28.6	78.3	6.7	12.5	74.7	1.9	12.2	6.0	79.2	6.8	67.2	-0.1	7.6	11.0	6.8	10.0	6.9	6.8
NS9	7.1	21.4	-0.1	8.3	21.6	-0.1	7.5	-0.1	21.0	-0.1	18.6	-0.1	-0.1	6.0	-0.1	7.3	4.2	-0.1
NT1	6.0	18.0	-0.1	-0.1	19.0	9.6	8.8	-0.1	24.6	-0.1	18.2	-0.1	-0.1	5.5	-0.1	7.5	4.7	-0.1
NT10	8.5	21.9	-0.1	-0.1	1.1	-0.1	7.8	-0.1	24.3	-0.1	17.8	-0.1	-0.1	6.3	-0.1	-0.1	4.6	-0.1
NT11	7.6	18.4	-0.1	-0.1	18.9	-0.1	7.7	-0.1	18.9	-0.1	15.6	-0.1	-0.1	6.1	-0.1	-0.1	4.4	-0.1
NT12	9.2	23.8	-0.1	8.1	24.7	9.2	9.1	3.6	33.6	-0.1	3.8	-0.1	-0.1	6.7	-0.1	7.5	4.7	-0.1
NT13	7.3	23.3	-0.1	-0.1	23.0	-0.1	9.2	-0.1	25.5	-0.1	18.4	-0.1	-0.1	6.2	-0.1	-0.1	4.4	-0.1
NT14	6.8	25.7	-0.1	8.6	27.1	9.9	10.0	3.6	57.3	-0.1	29.3	-0.1	-0.1	5.9	-0.1	7.6	4.3	-0.1
NT15	7.6	33.3	-0.1	8.7	32.4	1.1	8.3	3.9	32.4	-0.1	23.9	-0.1	-0.1	6.5	-0.1	7.4	4.4	-0.1
NT16	7.5	27.0	-0.1	8.8	27.4	10.9	11.3	3.7	47.1	-0.1	25.2	-0.1	-0.1	6.3	6.5	7.8	4.6	-0.1
NT17	7.7	25.1	-0.1	-0.1	25.8	9.8	9.4	3.4	41.7	-0.1	24.5	-0.1	-0.1	5.8	-0.1	7.4	4.3	-0.1
NT18	-0.1	10.1	-0.1	-0.1	10.0	-0.1	8.1	-0.1	9.2	-0.1	9.2	-0.1	-0.1	5.0	-0.1	-0.1	-0.1	-0.1
NT2	7.6	27.8	-0.1	9.1	29.8	10.8	11.2	3.9	39.5	-0.1	33.6	-0.1	-0.1	6.0	-0.1	3.8	5.0	-6.7
NT3	5.1	25.0	-0.1	-0.1	25.0	-0.1	-0.1	-0.1	24.6	-0.1	19.5	-0.1	-0.1	5.8	-0.1	-0.1	-0.1	-0.1
NT4	5.3	21.2	-0.1	-0.1	21.3	-0.1	-0.1	-0.1	23.2	-0.1	17.5	-0.1	-0.1	5.7	-0.1	-0.1	-0.1	-0.1
NT5	5.8	17.1	-0.1	-0.1	17.5	-0.1	7.7	-0.1	20.8	-0.1	14.8	-0.1	-0.1	5.9	-0.1	-0.1	4.0	-0.1
NT6-R	7.6	25.6	-0.1	-0.1	26.8	9.5	9.1	-0.1	51.0	-0.1	26.9	-0.1	-0.1	6.9	-0.1	7.7	4.1	-0.1
NT6	5.9	17.9	-0.1	-0.1	18.1	-0.1	7.4	-0.1	17.9	-0.1	14.4	-0.1	-0.1	5.7	-0.1	-0.1	4.0	-0.1
NT7	8.0	29.8	7.4	8.9	30.6	8.7	8.5	3.9	35.4	-0.1	28.7	-0.1	-0.1	6.7	-0.1	7.7	4.8	-0.1
NT8	20.5	92.1	7.9	12.9	90.0	2.8	12.6	5.5	99.0	6.7	65.7	-0.1	7.7	9.4	7.2	9.8	6.0	6.8
NT9	6.4	19.5	-0.1	-0.1	19.7	-0.1	7.6	-0.1	21.8	-0.1	16.6	-0.1	-0.1	6.0	-0.1	-0.1	4.1	-0.1
NU1	6.8	19.4	-0.1	-0.1	19.9	-0.1	7.6	-0.1	19.6	-0.1	14.7	-0.1	-0.1	6.3	-0.1	-0.1	4.1	-0.1
NU10	39.3	158.0	9.4	19.6	188.0	10.1	30.9	10.4	480.0	7.2	324.0	25.1	8.6	9.9	8.1	16.3	5.8	7.2
NU11	8.6	32.1	-0.1	9.1	32.1	1.7	9.2	3.9	36.3	-0.1	29.4	-0.1	-0.1	6.2	-0.1	8.0	4.5	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

	109 MAR	110 HBA	111 MAR	112 MBI	113 HBA	114 MBI	115 MBI	116 MAR	117 HA	118 MPH	119 HBA	120 THI	121 MPH	122 MPH	123 MPH	124 MBI	125 HAR	126 MPH
NU12	7.9	42.6	-0.1	9.0	42.3	9.5	9.2	3.7	51.6	-0.1	35.4	-0.1	-0.1	6.2	-0.1	7.6	4.5	-0.1
NU12-R	8.9	36.9	-0.1	9.2	37.5	1.5	9.3	3.9	48.6	-0.1	37.8	-0.1	-0.1	6.2	-0.1	7.7	4.5	-0.1
NU13	23.6	27.1	7.6	13.0	80.7	6.3	17.5	5.6	96.6	-0.1	74.4	-0.1	-0.1	9.8	6.8	9.5	5.9	-0.1
NU14	6.0	22.0	-0.1	-0.1	22.1	-0.1	10.8	-0.1	63.9	-0.1	3.5	-0.1	-0.1	7.0	-0.1	7.6	4.1	-0.1
NU15	10.1	42.9	-0.1	9.4	42.3	1.7	9.8	3.7	44.2	-0.1	33.6	-0.1	-0.1	6.7	-0.1	7.6	4.6	-0.1
NU16	17.1	71.1	7.6	12.1	72.9	2.6	12.7	5.3	108.0	6.7	56.7	-0.1	7.5	8.6	6.8	8.7	6.2	6.7
NU17	13.0	50.1	-0.1	9.8	48.6	1.7	10.7	4.0	65.4	-0.1	36.5	-0.1	-0.1	7.1	-0.1	7.9	5.4	-0.1
NU18	9.3	26.7	-0.1	8.7	26.3	1.5	8.5	3.7	24.7	-0.1	20.0	-0.1	-0.1	6.5	-0.1	7.5	4.8	-0.1
NU19	6.2	22.1	-0.1	-0.1	22.3	1.6	8.9	-0.1	33.9	-0.1	18.7	-0.1	-0.1	5.7	-0.1	7.4	3.8	-0.1
NU2	42.9	146.0	9.5	19.4	134.0	16.7	13.3	10.0	129.0	6.7	96.6	-0.1	7.4	12.9	6.6	11.3	7.5	6.8
NU3	5.3	21.4	-0.1	-0.1	21.6	-0.1	-0.1	-0.1	24.4	-0.1	17.9	-0.1	-0.1	5.7	-0.1	-0.1	-0.1	-0.1
NU4	87.3	321.0	14.7	32.1	309.0	7.6	25.9	18.5	295.0	7.4	287.0	7.1	8.3	22.6	7.5	20.5	9.9	7.5
NU5	6.8	4.7	-0.1	-0.1	23.6	-0.1	7.6	-0.1	25.1	-0.1	19.8	-0.1	-0.1	6.0	-0.1	7.6	4.1	-0.1
NU6	8.7	27.1	-0.1	8.5	27.5	8.8	7.9	-0.1	27.2	-0.1	21.1	-0.1	-0.1	6.3	-0.1	7.5	4.4	-0.1
NU7	9.4	39.6	5.7	8.6	41.7	9.0	9.8	4.0	49.8	-0.1	30.6	-0.1	-0.1	6.9	-0.1	7.8	5.0	-0.1
NU8	11.2	51.3	7.1	10.4	53.4	2.1	12.1	4.4	85.5	-0.1	44.4	-0.1	-0.1	7.4	6.8	8.4	5.4	-0.1
NU9	10.7	41.7	-0.1	9.9	44.7	1.7	13.9	4.0	79.5	6.6	41.7	-0.1	-0.1	7.6	8.7	7.0	9.4	5.7
NV1	6.8	24.9	-0.1	-0.1	24.3	8.5	8.1	3.5	40.5	-0.1	21.9	-0.1	-0.1	6.1	-0.1	7.3	4.2	-0.1
NV10	5.5	27.2	-0.1	-0.1	27.0	8.8	9.5	-0.1	28.7	-0.1	24.8	-0.1	-0.1	5.7	-0.1	7.3	3.9	-0.1
NV11	5.1	17.7	-0.1	-0.1	18.4	-0.1	8.1	-0.1	0.8	-0.1	15.8	-0.1	-0.1	6.6	-0.1	-0.1	4.0	-0.1
NV12	5.6	22.6	-0.1	-0.1	22.5	-0.1	8.1	-0.1	25.9	-0.1	18.0	-0.1	-0.1	6.1	-0.1	-0.1	3.8	-0.1
NV13	5.8	5.2	-0.1	-0.1	24.8	9.0	8.9	-0.1	40.5	-0.1	21.4	-0.1	-0.1	5.5	-0.1	7.2	4.1	-0.1
NV14	5.2	26.7	-0.1	-0.1	27.1	-0.1	-0.1	-0.1	64.8	-0.1	24.3	-0.1	-0.1	5.6	-0.1	7.5	4.0	-0.1
NV15	4.8	20.8	-0.1	-0.1	20.3	-0.1	-0.1	-0.1	22.8	-0.1	16.8	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NV16	6.2	39.9	-0.1	8.7	40.2	10.6	11.1	-0.1	59.1	-0.1	35.9	-0.1	-0.1	6.3	-0.1	7.9	4.4	-0.1
NV17	5.8	26.4	-0.1	-0.1	27.8	9.5	9.5	-0.1	38.4	-0.1	22.1	-0.1	-0.1	5.9	-0.1	7.7	4.1	-0.1
NV18	5.3	17.9	-0.1	-0.1	18.4	-0.1	-0.1	-0.1	25.9	-0.1	16.9	-0.1	-0.1	5.9	-0.1	-0.1	-0.1	-0.1
NV2	6.1	24.3	-0.1	-0.1	24.2	-0.1	7.7	-0.1	32.4	-0.1	20.4	-0.1	-0.1	6.0	-0.1	-0.1	4.3	-0.1
NV3	5.5	22.5	-0.1	-0.1	22.9	-0.1	7.5	-0.1	29.3	-0.1	19.0	-0.1	-0.1	5.6	-0.1	-0.1	3.7	-0.1
NV4	6.8	24.9	-0.1	8.6	26.4	1.4	9.4	3.6	52.5	-0.1	27.3	-0.1	-0.1	6.0	6.7	7.7	4.3	-0.1
NV5	7.3	36.0	-0.1	8.9	36.6	10.6	11.0	3.6	73.8	-0.1	33.9	-0.1	-0.1	6.7	-0.1	7.9	4.5	-0.1
NV5-R	7.1	40.2	-0.1	9.0	40.8	11.0	11.3	3.5	65.4	-0.1	39.0	-0.1	-0.1	6.3	-0.1	8.1	4.4	-0.1
NV6	5.6	14.6	-0.1	-0.1	15.4	-0.1	8.2	-0.1	17.8	-0.1	13.8	-0.1	-0.1	10.2	-0.1	7.8	4.7	-0.1
NV7	6.2	32.7	-0.1	8.6	32.7	9.6	8.9	-0.1	42.3	-0.1	27.9	-0.1	-0.1	5.9	-0.1	7.7	3.9	-0.1
NV8	5.3	16.7	-0.1	-0.1	17.5	-0.1	-0.1	-0.1	26.3	-0.1	16.4	-0.1	-0.1	5.4	-0.1	-0.1	3.9	-0.1
NV9	7.0	26.6	-0.1	-0.1	27.3	1.7	8.2	3.3	39.9	-0.1	22.3	-0.1	-0.1	5.7	-0.1	7.2	3.8	-0.1
NW1	8.4	8.6	-0.1	9.2	41.4	1.7	8.6	3.9	76.8	-0.1	30.6	-0.1	-0.1	6.5	-0.1	7.6	4.7	-0.1
NW10	6.0	4.1	-0.1	-0.1	22.2	9.0	8.6	-0.1	29.1	-0.1	18.6	-0.1	-0.1	6.3	-0.1	7.4	3.9	-0.1
NW11	5.0	20.3	-0.1	-0.1	20.4	-0.1	-0.1	-0.1	28.1	-0.1	18.0	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NW12	6.1	28.8	-0.1	8.2	28.7	-0.1	9.5	-0.1	38.7	-0.1	23.5	-0.1	-0.1	6.4	-0.1	7.7	4.3	-0.1
NW13	9.6	37.8	7.0	9.6	37.8	1.9	10.6	4.2	47.7	-0.1	29.2	-0.1	-0.1	6.8	-0.1	8.1	4.7	-0.1
NW14	5.5	18.1	-0.1	-0.1	17.9	-0.1	-0.1	-0.1	23.2	-0.1	15.1	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NW15	9.9	43.5	-0.1	9.5	41.7	11.1	11.4	4.1	77.4	6.6	34.5	-0.1	-0.1	7.0	-0.1	8.3	5.0	-0.1
NW16	5.7	19.1	-0.1	-0.1	18.9	-0.1	-0.1	-0.1	21.8	-0.1	16.4	-0.1	-0.1	5.6	-0.1	-0.1	3.9	-0.1
NW17	6.5	22.1	-0.1	-0.1	22.0	1.8	7.6	-0.1	27.9	-0.1	19.8	-0.1	-0.1	5.7	-0.1	7.4	4.0	-0.1
NW17-R	6.4	19.4	-0.1	-0.1	20.0	1.7	7.7	-0.1	23.9	-0.1	17.6	-0.1	-0.1	5.8	-0.1	7.4	3.9	-0.1
NW18	6.9	23.6	-0.1	8.7	25.0	1.5	8.3	3.6	31.6	-0.1	24.4	-0.1	-0.1	5.8	6.8	7.6	4.6	6.6
NW2	5.8	23.7	-0.1	-0.1	24.2	-0.1	7.7	-0.1	29.0	-0.1	20.0	-0.1	-0.1	5.3	-0.1	7.3	-0.1	-0.1
NW2-R	5.4	27.1	-0.1	-0.1	26.9	-0.1	7.8	-0.1	29.1	-0.1	21.4	-0.1	-0.1	5.7	-0.1	7.3	3.8	-0.1
NW3	5.4	26.4	-0.1	-0.1	27.1	-0.1	8.5	-0.1	43.8	-0.1	24.4	-0.1	-0.1	5.6	-0.1	7.3	3.9	-0.1
NW4	5.5	19.4	-0.1	-0.1	20.6	-0.1	8.2	-0.1	28.6	-0.1	17.8	-0.1	-0.1	5.9	-0.1	-0.1	-0.1	-0.1
NW5	14.6	83.7	8.1	12.2	85.2	2.3	12.0	4.7	85.5	6.4	57.9	-0.1	7.4	7.9	6.9	8.9	5.3	6.8
NW6	5.7	20.5	-0.1	-0.1	21.4	8.9	8.4	-0.1	35.7	-0.1	19.4	-0.1	-0.1	5.6	-0.1	7.3	4.0	-0.1
NW7	5.2	17.9	-0.1	-0.1	18.5	-0.1	8.0	-0.1	31.5	-0.1	16.9	-0.1	-0.1	5.4	-0.1	-0.1	3.9	-0.1
NW8	11.2	11.7	7.0	10.3	54.3	2.1	10.1	4.3	73.5	-0.1	35.4	-0.1	-0.1	7.1	-0.1	8.0	5.0	-0.1
NW9	5.6	27.4	-0.1	-0.1	27.6	1.7	-0.1	-0.1	36.9	-0.1	22.0	-0.1	-0.1	5.5	-0.1	-0.1	4.0	-0.1
NX1	5.7	16.1	-0.1	-0.1	17.1	-0.1	7.6	-0.1	20.4	-0.1	15.6	-0.1	-0.1	6.5	-0.1	-0.1	3.8	-0.1
NX10	13.0	46.8	7.2	10.3	44.7	1.9	10.5	4.6	57.0	-0.1	39.9	-0.1	-0.1	8.3	6.5	8.7	5.4	6.8
NX11	5.9	15.7	-0.1	-0.1	15.9	-0.1	-0.1	-0.1	18.1	-0.1	14.2	-0.1	-0.1	5.8	-0.1	-0.1	3.9	-0.1
NX12	6.3	18.6	-0.1	-0.1	19.3	-0.1	8.0	-0.1	27.2	-0.1	18.1	-0.1	-0.1	6.3	-0.1	-0.1	4.1	-0.1
NX13	10.4	37.5	-0.1	8.8	37.8	1.7	10.6	3.9	52.5	-0.1	31.5	-0.1	-0.1	6.6	-0.1	7.9	5.0	-0.1
NX14	9.8	32.1	-0.1	9.1	31.8	1.6	9.2	3.9	38.4	-0.1	25.5	-0.1	-0.1	6.3	-0.1	7.4	4.7	-0.1
NX14-R	7.1	33.9	-0.1	8.4	33.6	9.4	9.0	3.4	38.6	-0.1	26.5	-0.1	-0.1	5.9	-0.1	7.4	4.4	-0.1
NX15	7.3	33.9	-0.1	8.7	33.6	9.4	9.3	3.6	59.1	-0.1	33.6	-0.1	-0.1	6.2	-0.1	7.7	4.5	-0.1
NX16	8.9	42.3	-0.1	8.9	40.8	1.5	9.1	3.5	50.7	-0.1	30.3	-0.1	-0.1	6.4	-0.1	7.6	4.7	-0.1
NX17	12.1	46.8	-0.1	9.6	46.8	13.7	14.4	3.9	65.7	6.7	36.0	-0.1	-0.1	7.5	7.9	6.7	8.5	5.8
NX2	14.2	49.8	7.1	10.2	48.6	2.2	9.5	5.1	66.3	-0.1	60.5	-0.1	-0.1	7.5	-0.1	8.8	5.1	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109 MAR	110 HBA	111 MAR	112 MBI	113 HBA	114 NBI	115 MBI	116 MAR	117 HA	118 MPH	119 HBA	120 THI	121 MPH	122 MPH	123 MPH	124 MBI	125 HAB	126 MPH
NX3	13.9	36.6	7.0	9.4	35.7	9.3	8.7	4.4	53.4	-0.1	39.6	-0.1	-0.1	8.0	-0.1	8.2	5.3	-0.1
NX4	43.2	166.0	9.5	19.9	160.0	4.5	18.5	10.6	184.0	6.8	155.0	-0.1	7.6	12.4	7.3	19.0	6.9	6.9
NX5	8.9	36.3	-0.1	8.9	36.3	1.5	8.5	4.0	40.8	-0.1	33.3	-0.1	-0.1	6.6	-0.1	7.8	4.5	-0.1
NX6	16.5	57.6	7.0	11.1	62.1	2.4	12.9	4.9	101.0	-0.1	60.6	-0.1	-0.1	7.9	-6.7	8.7	4.9	-0.1
NX7	60.6	220.0	10.0	26.3	228.0	6.8	23.9	13.7	250.0	6.8	160.0	7.5	7.7	16.1	6.8	13.2	9.1	6.8
NX8	10.9	38.4	-0.1	9.7	41.1	1.6	12.5	3.8	78.0	6.4	38.4	-0.1	-0.1	7.9	7.0	8.2	5.0	-0.1
NX9	22.4	71.7	7.4	12.6	73.2	2.6	13.9	6.1	105.0	6.7	63.6	-0.1	7.6	9.7	6.8	9.1	6.7	6.8
NY1	6.0	18.0	-0.1	-0.1	19.5	-0.1	8.0	3.5	33.0	-0.1	20.9	-0.1	-0.1	5.6	-0.1	7.3	4.0	-0.1
NY10	10.6	48.0	-0.1	9.8	52.8	1.4	14.6	4.5	138.0	-0.1	68.7	3.8	-0.1	6.8	-0.1	8.5	4.9	-0.1
NY11	4.5	20.0	-0.1	-0.1	19.4	-0.1	-0.1	-0.1	27.8	-0.1	19.2	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NY12	7.0	20.8	-0.1	-0.1	21.3	-0.1	8.0	-0.1	41.4	-0.1	19.6	-0.1	-0.1	6.5	-0.1	7.5	4.1	-0.1
NY13	4.9	14.6	-0.1	-0.1	15.6	-0.1	-0.1	-0.1	26.0	-0.1	13.7	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NY14	9.6	36.9	-0.1	9.2	36.3	12.2	12.9	4.0	67.8	-0.1	32.1	-0.1	7.5	7.5	6.7	8.3	5.0	6.7
NY15	6.4	36.6	-0.1	8.0	36.0	1.7	8.5	-0.1	44.1	-0.1	29.5	-0.1	-0.1	5.9	-0.1	7.6	4.2	-0.1
NY2	6.0	14.3	-0.1	-0.1	14.9	-0.1	-0.1	-0.1	21.4	-0.1	14.6	-0.1	-0.1	5.7	-0.1	-0.1	4.1	-0.1
NY3	4.5	14.6	-0.1	-0.1	15.1	-0.1	-0.1	-0.1	22.8	-0.1	15.5	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1
NY4	9.8	6.7	-0.1	9.4	35.4	1.6	10.3	3.5	26.0	-0.1	21.7	-0.1	-0.1	6.3	-0.1	7.6	4.6	-0.1
NY4-R	18.6	77.1	7.5	12.7	77.7	5.0	11.6	5.9	69.6	-0.1	54.6	-0.1	-0.1	8.2	-0.1	8.7	5.5	-0.1
NY5	6.9	19.8	-0.1	-0.1	21.5	1.8	8.0	3.4	32.1	-0.1	21.3	-0.1	-0.1	5.6	-0.1	7.4	3.8	-0.1
NY6	8.9	34.8	-0.1	9.2	34.8	1.7	9.9	3.7	45.0	-0.1	30.9	-0.1	-0.1	6.2	-0.1	8.0	4.3	-0.1
NY7	15.8	64.8	7.5	11.2	63.0	2.0	12.5	5.5	85.2	-0.1	63.9	-0.1	-0.1	8.1	6.6	9.5	5.3	6.8
NY8	6.0	29.0	-0.1	8.5	29.6	9.5	9.5	-0.1	41.7	-0.1	25.9	-0.1	-0.1	6.8	-0.1	7.7	4.2	-0.1
NY9	-0.1	9.6	-0.1	-0.1	10.2	6.4	14.1	-0.1	11.7	-0.1	10.1	-0.1	-0.1	5.4	-0.1	7.4	-0.1	-0.1
NZ1	6.4	22.3	-0.1	-0.1	22.6	-0.1	-0.1	-0.1	39.0	-0.1	21.4	-0.1	-0.1	6.4	-0.1	7.6	4.0	-0.1
NZ10	8.0	42.9	-0.1	9.2	44.4	15.4	17.1	3.6	103.0	6.8	38.7	-0.1	7.9	6.9	7.4	8.9	4.6	7.0
NZ11	4.8	15.3	-0.1	-0.1	15.4	-0.1	-0.1	-0.1	18.4	-0.1	15.8	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NZ12	5.7	17.8	-0.1	-0.1	18.5	-0.1	8.7	-0.1	44.1	-0.1	19.5	-0.1	-0.1	5.9	-0.1	-0.1	4.1	-0.1
NZ13	4.5	15.4	-0.1	-0.1	16.2	-0.1	-0.1	-0.1	24.3	-0.1	13.8	-0.1	-0.1	5.5	-0.1	-0.1	-0.1	-0.1
NZ2	12.5	34.8	6.8	8.9	33.0	9.1	8.3	3.7	33.6	-0.1	23.2	-0.1	-0.1	6.8	-0.1	7.6	4.7	-0.1
NZ2-R	7.2	21.9	-0.1	-0.1	22.0	-0.1	7.9	-0.1	60.9	-0.1	18.5	-0.1	-0.1	6.8	-0.1	7.3	4.3	-0.1
NZ3	7.5	30.6	-0.1	8.8	30.0	9.0	8.6	3.7	31.5	-0.1	24.8	-0.1	-0.1	6.2	-0.1	7.7	4.6	-0.1
NZ4	7.4	26.0	-0.1	8.5	25.7	8.9	8.6	3.5	39.6	-0.1	20.6	-0.1	-0.1	5.8	-0.1	7.4	4.0	-0.1
NZ5	6.5	23.5	-0.1	8.6	23.4	9.3	9.1	3.5	35.7	-0.1	21.4	-0.1	-0.1	6.1	-0.1	7.5	4.3	-0.1
NZ6	6.0	20.6	-0.1	-0.1	22.1	9.6	9.8	3.5	61.6	-0.1	24.9	-0.1	-0.1	7.9	-0.1	7.5	4.4	-0.1
NZ7	6.0	28.7	-0.1	8.4	29.3	9.8	10.1	-0.1	66.6	-0.1	35.4	-0.1	-0.1	6.1	-0.1	7.8	4.3	-0.1
NZ8	5.9	20.9	-0.1	-0.1	20.6	-0.1	8.1	-0.1	26.4	-0.1	15.9	-0.1	-0.1	5.9	-0.1	-0.1	4.1	-0.1
NZ9	5.0	16.2	-0.1	-0.1	16.2	-0.1	-0.1	-0.1	19.1	-0.1	2.8	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	14.7	-0.1	-0.1	1.6	-0.1	-0.1	-0.1	28.6	-0.1	19.1	-0.1	-0.1	5.7	-0.1	-0.1	4.1	-0.1
LMB-QA	-0.1	14.0	-0.1	-0.1	13.8	-0.1	-0.1	-0.1	17.0	-0.1	13.0	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	9.4	-0.1	-0.1	9.5	-0.1	-0.1	-0.1	11.4	-0.1	10.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	13.7	-0.1	-0.1	13.4	-0.1	-0.1	-0.1	15.7	-0.1	18.0	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	11.5	-0.1	-0.1	11.8	-0.1	-0.1	-0.1	14.6	-0.1	12.2	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	10.9	-0.1	-0.1	11.1	-0.1	-0.1	-0.1	13.5	-0.1	11.0	-0.1	-0.1	5.2	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	8.9	-0.1	-0.1	9.1	-0.1	-0.1	-0.1	12.2	-0.1	10.3	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	9.2	-0.1	-0.1	9.1	-0.1	-0.1	-0.1	12.1	-0.1	10.3	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	9.7	-0.1	-0.1	9.8	-0.1	-0.1	-0.1	12.9	-0.1	10.6	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	8.1	-0.1	-0.1	8.1	-0.1	-0.1	-0.1	8.6	-0.1	8.4	-0.1	-0.1	5.4	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	9.6	-0.1	-0.1	9.8	-0.1	-0.1	-0.1	11.5	-0.1	9.8	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1
LMB-QA	-0.1	9.1	-0.1	-0.1	9.2	-0.1	-0.1	-0.1	10.3	-0.1	9.3	-0.1	-0.1	5.3	-0.1	-0.1	-0.1	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	109 MAR	110 HBA	111 MAR	112 MBI	113 HBA	114 MBI	115 MBI	116 MAR	117 HA	118 MPH	119 HBA	120 THU	121 MPH	122 MPH	123 MPH	124 MBI	125 HAR	126 MPH

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
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R=Replicate Sample

	127 MRH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MRH	137 HBI	138 HBI	139 MRH	140 HPH	141 HBI	142 MRH	143 HA	144 HBI
NA1	8.0	7.9	8.0	8.7	7.5	272.0	61.8	124.0	56.1	51.3	68.1	65.7	51.0	51.3	51.3	-0.1	224.0	47.4
NA2	7.3	7.0	6.7	7.2	6.8	232.0	63.9	107.0	51.6	49.5	68.1	68.4	-0.1	-0.1	56.4	-0.1	489.0	52.5
NAA1	7.1	6.5	5.9	7.4	6.6	94.2	-0.1	75.3	49.5	-0.1	51.6	52.2	-0.1	-0.1	-0.1	-0.1	103.0	-0.1
NAA10	7.0	-0.1	6.2	8.6	-0.1	78.3	-0.1	60.6	49.5	-0.1	49.8	50.4	-0.1	-0.1	-0.1	-0.1	96.6	-0.1
NAA11	-0.1	6.5	6.5	8.1	-0.1	101.0	-0.1	74.1	-0.1	-0.1	51.6	50.4	-0.1	-0.1	-0.1	-0.1	88.8	-0.1
NAA2	8.5	8.9	11.3	14.8	7.7	555.0	75.9	248.0	59.4	55.2	86.7	83.1	51.3	52.2	54.0	50.1	372.0	49.8
NAA3	-0.1	6.6	6.3	8.0	-0.1	159.0	-0.1	108.0	-0.1	-0.1	59.1	59.4	-0.1	-0.1	-0.1	-0.1	143.0	-0.1
NAA4	7.3	6.8	6.4	7.0	-0.1	149.0	-0.1	82.8	49.8	-0.1	57.9	56.4	-0.1	-0.1	-0.1	-0.1	244.0	-0.1
NAA5	6.9	6.8	6.3	7.9	6.6	106.0	-0.1	83.1	49.8	-0.1	52.8	53.4	-0.1	-0.1	-0.1	-0.1	109.0	-0.1
NAA6	7.0	6.5	6.0	7.8	6.8	115.0	-0.1	75.0	51.6	-0.1	54.3	53.1	-0.1	-0.1	48.6	-0.1	152.0	-0.1
NAA7	7.2	6.6	6.2	7.1	6.7	108.0	-0.1	69.3	49.2	-0.1	55.2	56.1	-0.1	-0.1	-0.1	-0.1	174.0	44.7
NAA8	-0.1	-0.1	-0.1	7.1	-0.1	109.0	-0.1	70.2	49.5	-0.1	52.5	51.3	-0.1	-0.1	-0.1	-0.1	170.0	-0.1
NAA9	6.9	6.5	6.0	7.5	6.9	125.0	-0.1	66.3	53.4	-0.1	52.2	51.3	-0.1	-0.1	-0.1	-0.1	159.0	-0.1
NB1	7.0	-0.1	6.9	6.8	-0.1	76.8	-0.1	62.7	-0.1	-0.1	51.0	50.1	-0.1	-0.1	-0.1	-0.1	119.0	-0.1
NB2	-0.1	-0.1	-0.1	-0.1	-0.1	62.1	-0.1	54.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	94.8	-0.1
NB3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.6	-0.1
NB4	-0.1	6.6	6.0	7.4	-0.1	171.0	-0.1	96.0	-0.1	-0.1	59.1	59.7	-0.1	-0.1	51.0	-0.1	279.0	47.4
NBB1	-0.1	-0.1	-0.1	-0.1	-0.1	64.8	-0.1	55.8	49.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	100.0	-0.1
NBB10	7.1	7.6	8.3	10.5	6.8	268.0	65.1	146.0	50.7	-0.1	64.5	64.5	-0.1	-0.1	49.2	-0.1	170.0	-0.1
NBB11	7.9	7.3	7.2	8.6	7.7	215.0	57.9	91.5	58.5	54.6	66.5	64.2	50.1	50.1	55.2	49.5	274.0	49.2
NBB2	-0.1	-0.1	6.1	7.7	-0.1	101.0	-0.1	75.0	-0.1	-0.1	51.3	50.1	-0.1	-0.1	-0.1	-0.1	110.0	-0.1
NBB2-R	-0.1	-0.1	5.6	7.1	-0.1	72.6	-0.1	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.1	-0.1
NBB3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	85.2	-0.1
NBB4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	41.7	-0.1
NBB5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	96.3	-0.1
NBB6	7.0	6.5	6.2	7.8	6.7	95.7	-0.1	68.7	51.0	-0.1	50.7	51.3	-0.1	-0.1	-0.1	-0.1	20.3	-0.1
NBB7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.8	-0.1
NBB8	-0.1	-0.1	5.7	7.3	-0.1	-0.1	-0.1	65.1	-0.1	-0.1	48.9	48.5	-0.1	-0.1	-0.1	-0.1	93.9	-0.1
NBB9	-0.1	-0.1	6.3	6.7	-0.1	64.2	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.4	-0.1
NC1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	93.9	-0.1
NC2	-0.1	-0.1	6.0	-0.1	-0.1	86.1	-0.1	54.9	50.4	-0.1	48.3	48.9	-0.1	-0.1	-0.1	-0.1	136.0	-0.1
NC3	-0.1	-0.1	-0.1	-0.1	-0.1	60.6	-0.1	56.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	101.0	-0.1
NC4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	14.1	-0.1
NC5	-0.1	-0.1	-0.1	-0.1	-0.1	50.6	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	115.0	-0.1
NC6	-0.1	-0.1	5.6	-0.1	-0.1	54.6	-0.1	60.3	-0.1	-0.1	49.8	48.9	-0.1	-0.1	-0.1	-0.1	100.0	-0.1
NC7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	42.3	-0.1
NC8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NCC1	8.4	9.9	12.0	15.4	8.6	1076.0	215.0	666.0	65.7	74.4	221.0	210.0	64.2	59.4	105.0	54.8	2580.0	96.6
NCC1-R	8.3	10.0	12.0	15.5	8.5	1200.0	227.0	678.0	64.5	71.4	223.0	214.0	57.3	55.5	102.0	53.1	2390.0	91.5
NCC2	10.1	14.1	14.3	16.5	9.3	719.0	131.0	346.0	57.8	63.9	130.0	124.0	58.2	58.8	64.2	55.6	825.0	59.7
NCC3	7.1	6.8	6.5	8.0	6.9	140.0	-0.1	81.3	52.8	50.1	53.4	53.4	-0.1	-0.1	48.9	-0.1	24.0	-0.1
NCC4	7.3	7.5	8.9	9.2	6.9	166.0	-0.1	80.1	52.2	50.1	54.3	53.4	49.8	50.7	-0.1	-0.1	141.0	-0.1
NCC5	-0.1	-0.1	6.1	7.5	-0.1	94.2	-0.1	69.0	-0.1	-0.1	49.5	50.1	-0.1	-0.1	-0.1	-0.1	119.0	-0.1
NCC6	-0.1	7.0	7.2	8.9	-0.1	137.0	-0.1	97.8	49.5	-0.1	55.2	53.7	-0.1	-0.1	-0.1	-0.1	91.5	-0.1
ND1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1
ND10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.1	-0.1
ND2	-0.1	-0.1	5.9	7.4	-0.1	95.1	-0.1	67.2	50.4	-0.1	50.7	49.8	-0.1	-0.1	-0.1	-0.1	111.0	-0.1
ND3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.5	-0.1
ND4	-0.1	6.6	6.2	7.7	-0.1	118.0	53.4	83.1	-0.1	-0.1	55.8	54.6	-0.1	-0.1	48.9	-0.1	159.0	-0.1
ND5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.4	-0.1
ND6	-0.1	-0.1	-0.1	-0.1	-0.1	57.9	-0.1	56.1	-0.1	-0.1	47.4	-0.1	-0.1	-0.1	-0.1	-0.1	82.2	-0.1
ND7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	60.6	-0.1
ND8	-0.1	-0.1	-0.1	-0.1	-0.1	54.0	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	14.2	-0.1
ND8-R	-0.1	-0.1	-0.1	-0.1	-0.1	64.8	-0.1	59.7	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	72.6	-0.1
ND9	-0.1	-0.1	-0.1	-0.1	-0.1	67.5	-0.1	62.1	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	93.0	-0.1
NDD1	7.3	6.8	6.4	8.0	7.0	148.0	-0.1	93.9	53.1	-0.1	57.0	56.7	-0.1	-0.1	49.8	-0.1	139.0	-0.1
NDD2	-0.1	-0.1	5.6	7.1	-0.1	64.8	-0.1	59.7	-0.1	-0.1	49.2	48.3	-0.1	-0.1	-0.1	-0.1	99.0	-0.1
NDD3	7.3	-0.1	5.7	7.2	6.5	82.2	-0.1	49.8	-0.1	-0.1	52.8	51.6	-0.1	-0.1	-0.1	-0.1	114.0	-0.1
NDD4	6.8	6.5	5.8	7.3	6.5	73.8	-0.1	70.2	-0.1	-0.1	51.0	50.1	-0.1	-0.1	-0.1	-0.1	88.2	-0.1
NDD5	-0.1	6.7	6.5	7.2	-0.1	145.0	-0.1	94.2	49.5	-0.1	57.6	57.9	-0.1	-0.1	-0.1	-0.1	187.0	-0.1
NE1	7.1	6.9	6.9	8.7	6.8	136.0	-0.1	85.8	51.3	-0.1	54.9	53.4	-0.1	-0.1	-0.1	-0.1	138.0	-0.1
NE10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	61.5	-0.1
NE11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.3	-0.1
NE2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	44.9	-0.1
NE3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.1	-0.1
NE4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	87.3	-0.1
NE4-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NE5	-0.1	-0.1	-0.1	-0.1	-0.1	64.5	-0.1	56.1	56.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	94.2	-0.1
NE6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.6	-0.1
NE7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.1	-0.1
NE8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NE9	-0.1	-0.1	-0.1	-0.1	-0.1	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	95.7	-0.1
NEE1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	81.9	-0.1
NEE2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	90.3	-0.1
NEE3	-0.1	-0.1	5.8	7.3	-0.1	104.0	-0.1	71.1	-0.1	-0.1	50.4	50.7	-0.1	-0.1	-0.1	-0.1	136.0	-0.1
NEE4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.1	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	92.4	-0.1
NEE5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	86.1	-0.1
NEE6	6.9	-0.1	5.6	7.2	6.5	86.7	-0.1	67.6	-0.1	-0.1	51.6	50.4	-0.1	-0.1	-0.1	-0.1	109.0	-0.1
NEE7	-0.1	-0.1	5.8	7.1	-0.1	56.7	-0.1	65.1	-0.1	-0.1	50.4	49.5	-0.1	-0.1	-0.1	-0.1	83.4	-0.1
NF1	7.1	7.0	7.4	9.3	6.7	149.0	-0.1	100.0	51.0	48.6	55.6	54.0	-0.1	-0.1	-0.1	-0.1	140.0	-0.1
NF10	7.3	7.1	6.5	8.4	6.7	215.0	64.2	111.0	51.9	50.4	65.4	63.6	-0.1	-0.1	51.6	-0.1	318.0	48.0
NF11	8.5	8.5	8.6	9.9	8.1	636.0	143.0	306.0	60.9	65.4	142.0	131.0	59.7	56.7	102.0	57.0	2680.0	95.1
NF2	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	56.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.4	-0.1
NF3	-0.1	-0.1	5.9	6.6	-0.1	-0.1	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	44.4	-0.1
NF4	-0.1	6.5	6.2	7.7	-0.1	126.0	-0.1	86.4	49.2	-0.1	53.1	53.1	-0.1	-0.1	-0.1	-0.1	128.0	-0.1
NF5	-0.1	-0.1	6.2	6.6	-0.1	118.0	-0.1	67.6	60.4	-0.1	52.3	51.9	-0.1	-0.1	-0.1	-0.1	77.7	-0.1
NF6	-0.1	6.6	6.4	8.6	-0.1	212.0	-0.1	118.0	-0.1	-0.1	62.4	62.4	-0.1	-0.1	49.5	-0.1	234.0	-0.1
NF7	8.1	7.6	8.1	10.0	7.6	229.0	62.1	102.0	58.8	53.1	61.2	60.1	51.3	52.5	57.3	-0.1	245.0	49.8
NF8	-0.1	-0.1	-0.1	-0.1	-0.1	69.9	-0.1	57.0	50.1	-0.1	49.2	48.6	-0.1	-0.1	-0.1	-0.1	16.9	-0.1
NF8-R	7.4	6.9	6.7	8.3	6.9	175.0	58.5	103.0	61.6	50.4	58.2	58.5	-0.1	-0.1	51.6	-0.1	257.0	48.0
NF9	-0.1	6.6	6.3	8.0	-0.1	144.0	-0.1	93.0	-0.1	-0.1	55.2	55.2	-0.1	-0.1	-0.1	-0.1	132.0	-0.1
NFF1	-0.1	6.6	6.5	7.0	6.9	148.0	-0.1	69.5	52.8	48.0	52.8	53.1	-0.1	50.1	-0.1	-0.1	170.0	-0.1
NFF2	7.0	6.9	6.8	8.7	6.8	192.0	-0.1	109.0	52.2	-0.1	59.4	58.2	-0.1	-0.1	-0.1	-0.1	141.0	-0.1
NFF3	-0.1	6.5	6.4	7.5	-0.1	104.0	-0.1	84.6	-0.1	-0.1	56.5	54.6	-0.1	-0.1	-0.1	-0.1	114.0	-0.1
NFF4	7.3	7.1	6.5	8.4	7.2	205.0	63.0	103.0	54.9	51.0	61.2	60.0	-0.1	49.8	50.1	-0.1	211.0	46.5
NFF5	-0.1	-0.1	6.0	7.4	-0.1	93.0	-0.1	70.2	49.5	-0.1	49.8	50.7	-0.1	-0.1	-0.1	-0.1	101.0	-0.1
NFF6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	96.0	-0.1
NFF6-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.9	-0.1
NFF7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.6	-0.1
NFF8	-0.1	-0.1	6.6	7.0	-0.1	54.9	-0.1	54.3	-0.1	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	106.0	-0.1
NG1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.3	-0.1
NG10	-0.1	-0.1	-0.1	-0.1	-0.1	64.2	-0.1	59.1	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	98.4	-0.1
NG11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	65.4	-0.1
NG2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.3	-0.1
NG2-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	11.3	-0.1
NG3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.9	-0.1
NG4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	11.6	-0.1
NG5	7.2	6.5	5.9	7.4	6.3	86.8	-0.1	67.2	-0.1	-0.1	52.5	51.3	-0.1	-0.1	48.3	-0.1	14.6	-0.1
NG6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.8	-0.1
NG7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.7	-0.1
NG8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.1	-0.1
NG9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	42.5	-0.1
NGG1	-0.1	-0.1	5.8	7.4	-0.1	69.9	-0.1	63.0	-0.1	-0.1	49.8	48.9	-0.1	-0.1	-0.1	-0.1	91.2	-0.1
NGG10	-0.1	-0.1	5.0	7.2	-0.1	53.6	-0.1	63.3	-0.1	-0.1	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	91.8	-0.1
NGG2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.2	-0.1
NGG3	7.2	7.4	8.4	10.3	6.8	257.0	58.5	158.0	51.0	-0.1	64.2	63.6	-0.1	-0.1	49.5	-0.1	138.0	46.2
NGG4	-0.1	-0.1	5.8	7.2	-0.1	77.7	-0.1	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	106.0	-0.1
NGG5	-0.1	-0.1	5.6	-0.1	-0.1	59.4	-0.1	57.0	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	104.0	-0.1
NGG6	-0.1	-0.1	5.8	7.3	6.4	81.6	-0.1	61.5	49.8	-0.1	50.7	49.5	-0.1	-0.1	-0.1	-0.1	19.4	-0.1
NGG7	-0.1	-0.1	5.8	7.2	-0.1	80.7	-0.1	60.6	51.0	-0.1	49.8	48.9	-0.1	-0.1	-0.1	-0.1	111.0	-0.1
NGG8	-0.1	-0.1	5.9	7.2	-0.1	72.3	-0.1	63.3	-0.1	-0.1	47.7	48.3	-0.1	-0.1	-0.1	-0.1	104.0	-0.1
NGG9	-0.1	-0.1	-0.1	-0.1	-0.1	78.3	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.3	-0.1
NH1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	17.4	-0.1
NH10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.3	-0.1
NH10-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.4	-0.1
NH11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	93.0	-0.1
NH2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.2	-0.1
NH3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.4	-0.1
NH4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.1	-0.1
NH5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1
NH6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	83.7	-0.1
NH7	7.5	-0.1	5.6	7.1	7.0	90.6	-0.1	62.1	52.2	49.5	52.5	51.3	-0.1	-0.1	49.2	-0.1	140.0	-0.1
NH8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	17.2	-0.1
NH9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	15.3	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NHH1	-0.1	-0.1	-0.1	-0.1	-0.1	60.9	-0.1	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	91.2	-0.1
NHH10	-0.1	-0.1	-0.1	-0.1	-0.1	52.8	-0.1	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.9	-0.1
NHH11	-0.1	-0.1	6.2	6.6	-0.1	66.3	-0.1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.8	-0.1
NHH11-R	-0.1	-0.1	5.6	7.0	-0.1	55.2	-0.1	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.2	-0.1
NHH12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	16.2	-0.1
NHH2	7.1	7.0	8.0	8.6	6.6	235.0	-0.1	117.0	59.7	-0.1	61.8	59.4	-0.1	-0.1	48.6	-0.1	183.0	-0.1
NHH3	-0.1	-0.1	5.6	7.1	-0.1	75.9	-0.1	62.1	-0.1	-0.1	49.5	48.6	-0.1	-0.1	-0.1	-0.1	13.4	-0.1
NHH4	-0.1	6.9	6.8	8.6	6.6	106.0	-0.1	81.3	-0.1	-0.1	52.8	51.6	-0.1	-0.1	-0.1	-0.1	92.1	-0.1
NHH5	-0.1	-0.1	6.1	7.6	-0.1	90.6	-0.1	69.6	-0.1	-0.1	49.5	48.6	-0.1	-0.1	-0.1	-0.1	14.6	-0.1
NHH6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.3	-0.1
NHH7	-0.1	-0.1	-0.1	-0.1	-0.1	65.7	-0.1	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	89.7	-0.1
NHH8	-0.1	-0.1	5.6	7.0	-0.1	58.7	-0.1	62.4	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	99.9	-0.1
NHH9	-0.1	-0.1	5.7	7.0	-0.1	68.4	-0.1	60.0	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	98.1	-0.1
NI1	7.1	6.8	6.3	7.9	6.7	138.0	-0.1	90.0	51.0	-0.1	55.8	54.6	-0.1	-0.1	-0.1	-0.1	247.0	46.5
NI10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	80.4	-0.1
NI2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	83.1	-0.1
NI3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	86.7	-0.1
NI4	7.0	6.6	6.0	7.8	-0.1	91.8	-0.1	78.3	-0.1	-0.1	54.3	52.8	-0.1	-0.1	-0.1	-0.1	12.7	-0.1
NI5	-0.1	-0.1	-0.1	7.0	-0.1	59.4	-0.1	59.1	-0.1	-0.1	49.5	-0.1	-0.1	-0.1	-0.1	-0.1	91.5	-0.1
NI5-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	79.3	-0.1
NI6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	47.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	101.0	-0.1
NI7	-0.1	-0.1	6.2	6.8	-0.1	93.9	-0.1	65.4	-0.1	-0.1	50.7	49.5	-0.1	-0.1	-0.1	-0.1	161.0	-0.1
NI8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	14.5	-0.1
NI9	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.1	-0.1
NII1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.0	-0.1
NII10	-0.1	-0.1	6.2	6.7	-0.1	97.5	-0.1	67.2	-0.1	-0.1	51.0	49.8	-0.1	-0.1	-0.1	-0.1	113.0	-0.1
NII11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.0	-0.1
NII1-R	-0.1	-0.1	5.7	7.2	-0.1	82.5	-0.1	66.3	-0.1	-0.1	50.7	51.0	-0.1	-0.1	-0.1	-0.1	115.0	-0.1
NII12	-0.1	-0.1	5.7	7.1	-0.1	73.5	-0.1	67.2	-0.1	-0.1	51.0	50.1	-0.1	-0.1	-0.1	-0.1	91.5	-0.1
NII13	-0.1	6.6	6.1	7.7	-0.1	112.0	-0.1	83.7	-0.1	-0.1	52.8	52.8	-0.1	-0.1	-0.1	-0.1	118.0	-0.1
NII14	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.9	-0.1
NII2	6.9	-0.1	5.6	7.1	6.8	97.5	-0.1	59.7	51.3	48.9	50.1	50.7	-0.1	-0.1	48.6	-0.1	144.0	-0.1
NII3	-0.1	6.5	6.5	6.9	-0.1	116.0	-0.1	71.1	49.5	-0.1	51.6	52.2	-0.1	-0.1	-0.1	-0.1	23.2	-0.1
NII4	-0.1	-0.1	6.0	7.5	-0.1	116.0	-0.1	78.6	-0.1	-0.1	58.1	51.9	-0.1	-0.1	-0.1	-0.1	143.0	-0.1
NII5	-0.1	-0.1	6.1	6.6	-0.1	122.0	-0.1	70.8	48.6	-0.1	52.5	52.5	-0.1	-0.1	-0.1	-0.1	167.0	-0.1
NII6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	60.7	-0.1
NII7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	81.6	-0.1
NII8	-0.1	6.5	6.1	7.8	-0.1	113.0	-0.1	84.0	-0.1	-0.1	52.5	52.8	-0.1	-0.1	-0.1	-0.1	15.7	-0.1
NII9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.8	-0.1
NJ1	-0.1	-0.1	-0.1	7.0	-0.1	66.0	-0.1	54.6	50.1	-0.1	49.5	-0.1	-0.1	-0.1	48.3	-0.1	115.0	-0.1
NJ2	-0.1	-0.1	5.6	7.0	-0.1	69.0	-0.1	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	96.6	-0.1
NJ3	-0.1	-0.1	-0.1	-0.1	-0.1	75.3	-0.1	62.4	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	13.7	-0.1
NJ4	-0.1	-0.1	-0.1	-0.1	-0.1	77.4	-0.1	60.0	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	17.6	-0.1
NJ5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	67.0	-0.1
NJ6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	14.6	-0.1
NJ7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	82.8	-0.1
NJJ1	-0.1	-0.1	5.7	7.1	-0.1	53.7	-0.1	60.3	-0.1	-0.1	48.3	48.9	-0.1	-0.1	-0.1	-0.1	13.2	-0.1
NJJ10	9.8	12.5	16.1	20.1	9.0	350.0	93.0	676.0	55.4	56.7	188.0	177.0	55.8	57.0	65.4	53.4	600.0	58.5
NJJ11	-0.1	-0.1	-0.1	-0.1	-0.1	52.2	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	95.1	-0.1
NJJ12	-0.1	6.6	6.1	7.6	-0.1	106.0	-0.1	79.2	49.2	-0.1	54.0	52.8	-0.1	-0.1	-0.1	-0.1	108.0	-0.1
NJJ13	-0.1	6.8	6.9	8.4	-0.1	150.0	-0.1	101.0	-0.1	-0.1	55.2	55.2	-0.1	-0.1	-0.1	-0.1	121.0	-0.1
NJJ14	-0.1	-0.1	-0.1	-0.1	-0.1	77.7	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	77.7	-0.1
NJJ2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.3	-0.1
NJJ3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NJJ4	-0.1	6.5	6.5	7.0	-0.1	145.0	-0.1	83.4	-0.1	-0.1	53.4	53.1	-0.1	-0.1	-0.1	-0.1	137.0	-0.1
NJJ5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.9	-0.1
NJJ6	-0.1	-0.1	-0.1	-0.1	-0.1	65.1	-0.1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	95.1	-0.1
NJJ7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	89.1	-0.1
NJJ8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.0	-0.1
NJJ8-R	-0.1	-0.1	5.6	-0.1	-0.1	63.6	-0.1	66.9	-0.1	-0.1	48.0	48.6	-0.1	-0.1	-0.1	-0.1	76.3	-0.1
NJJ9	7.2	6.8	6.6	8.1	-0.1	198.0	-0.1	111.0	-0.1	-0.1	59.7	58.2	-0.1	-0.1	-0.1	-0.1	180.0	-0.1
NK1	-0.1	-0.1	-0.1	-0.1	-0.1	59.6	-0.1	-0.1	49.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	139.0	-0.1
NK2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	13.3	-0.1
NK3	-0.1	-0.1	-0.1	-0.1	-0.1	63.9	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	15.4	-0.1
NK4	-0.1	-0.1	-0.1	-0.1	-0.1	48.3	-0.1	58.8	-0.1	-0.1	49.8	48.9	-0.1	-0.1	-0.1	-0.1	90.9	-0.1
NK4-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.6	-0.1
NK5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	96.3	-0.1

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	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NK6	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	90.0	-0.1
NK7	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	14.5	-0.1
NKK1	-0.1	-0.1	5.6	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	122	-0.1
NKK10	-0.1	-0.1	-0.1	-0.1	-0.1	65.1	-0.1	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	96.9	-0.1
NKK11	-0.1	-0.1	5.9	7.3	-0.1	70.8	-0.1	66.6	-0.1	-0.1	49.6	48.9	-0.1	-0.1	-0.1	-0.1	62.5	-0.1
NKK2	-0.1	-0.1	5.8	7.4	-0.1	78.3	-0.1	68.4	-0.1	-0.1	51.3	50.4	-0.1	-0.1	-0.1	-0.1	96.9	-0.1
NKK3	-0.1	7.0	7.6	9.1	-0.1	172.0	-0.1	117.0	-0.1	-0.1	58.2	56.7	-0.1	-0.1	-0.1	-0.1	100.0	-0.1
NKK4	7.6	9.6	13.6	17.4	7.2	738.0	100.0	444.0	54.6	51.0	118.0	114.0	50.7	52.2	55.8	-0.1	342.0	51.3
NKK5	7.4	7.5	9.3	10.5	6.3	360.0	56.7	172.0	-0.1	-0.1	69.8	66.3	-0.1	-0.1	-0.1	-0.1	146.0	-0.1
NKK6	-0.1	6.7	6.7	8.1	-0.1	110.0	-0.1	82.5	-0.1	-0.1	52.2	51.0	-0.1	-0.1	-0.1	-0.1	89.7	-0.1
NKK7	-0.1	6.8	5.6	8.5	-0.1	136.0	-0.1	102.0	-0.1	-0.1	58.7	54.0	-0.1	-0.1	-0.1	-0.1	92.1	-0.1
NKK8	6.9	6.5	5.8	7.3	6.7	95.7	-0.1	68.4	51.0	-0.1	52.8	51.6	-0.1	-0.1	-0.1	-0.1	131.0	-0.1
NKK9	-0.1	-0.1	5.5	-0.1	-0.1	82.8	-0.1	59.7	49.2	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	145.0	-0.1
NL1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NL2	-0.1	-0.1	5.6	-0.1	-0.1	58.7	-0.1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.6	-0.1
NL3	-0.1	-0.1	6.1	6.6	-0.1	61.8	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NL3-R	-0.1	6.5	6.1	7.6	-0.1	87.9	-0.1	72.9	48.9	-0.1	50.4	50.7	-0.1	-0.1	-0.1	-0.1	84.9	-0.1
NL4	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	52.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.0	-0.1
NLL1	-0.1	6.6	5.8	7.9	-0.1	127.0	-0.1	102.0	-0.1	-0.1	56.5	55.5	-0.1	-0.1	-0.1	-0.1	99.3	-0.1
NLL10	-0.1	-0.1	5.5	-0.1	-0.1	57.3	-0.1	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	86.7	-0.1
NLL11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	81.2	-0.1
NLL12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	85.5	-0.1
NLL2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	70.2	-0.1
NLL3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NLL4	-0.1	6.6	6.5	7.9	-0.1	78.3	-0.1	76.3	-0.1	-0.1	51.9	50.4	-0.1	-0.1	-0.1	-0.1	83.1	-0.1
NLL5	-0.1	9.8	6.1	7.5	-0.1	89.4	-0.1	69.9	-0.1	-0.1	50.1	49.2	-0.1	-0.1	-0.1	-0.1	115.0	-0.1
NLL6	-0.1	6.9	5.8	8.5	6.6	148.0	-0.1	96.3	-0.1	-0.1	58.7	54.0	-0.1	-0.1	-0.1	-0.1	158.0	-0.1
NLL7	7.4	7.5	8.4	9.0	6.5	190.0	52.5	98.7	49.2	-0.1	56.7	54.9	-0.1	-0.1	47.1	-0.1	144.0	-0.1
NLL8	-0.1	-0.1	-0.1	6.4	-0.1	62.7	-0.1	54.3	-0.1	-0.1	49.6	48.9	-0.1	-0.1	-0.1	-0.1	87.5	-0.1
NLL9	7.2	6.9	7.2	8.9	6.5	127.0	-0.1	81.6	51.0	-0.1	53.4	52.2	-0.1	-0.1	-0.1	-0.1	102.0	-0.1
NLL9-R	7.1	6.9	7.0	8.6	6.7	104.0	-0.1	75.0	50.4	-0.1	54.9	50.7	-0.1	-0.1	-0.1	-0.1	96.3	-0.1
NM1	-0.1	-0.1	5.9	7.2	-0.1	77.1	-0.1	63.3	49.2	-0.1	49.5	48.9	-0.1	-0.1	-0.1	-0.1	89.4	-0.1
NM2	-0.1	-0.1	-0.1	-0.1	-0.1	62.4	-0.1	50.7	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	83.0	-0.1
NMM1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	70.8	-0.1
NMM10	-0.1	6.5	5.4	6.8	-0.1	104.0	-0.1	69.6	-0.1	-0.1	54.3	50.4	-0.1	-0.1	-0.1	-0.1	103.0	-0.1
NMM2	-0.1	-0.1	-0.1	-0.1	-0.1	65.7	-0.1	66.0	-0.1	-0.1	49.5	48.6	-0.1	-0.1	-0.1	-0.1	93.6	-0.1
NMM3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.3	-0.1
NMM3-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.6	-0.1
NMM4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.5	-0.1
NMM5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NMM6	-0.1	-0.1	6.4	7.2	-0.1	72.9	-0.1	58.2	-0.1	-0.1	50.4	48.9	-0.1	-0.1	-0.1	-0.1	93.6	-0.1
NMM7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	11.9	-0.1
NMM8	-0.1	6.6	6.8	7.4	-0.1	144.0	-0.1	85.2	49.2	-0.1	52.2	52.8	-0.1	-0.1	-0.1	-0.1	111.0	-0.1
NMM9	7.0	6.5	6.1	6.8	6.7	104.0	-0.1	63.3	51.6	50.4	53.4	52.5	-0.1	48.6	48.6	-0.1	158.0	-0.1
NN1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.4	-0.1
NNN1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.5	-0.1
NNN10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	106.0	-0.1
NNN2	-0.1	-0.1	-0.1	-0.1	-0.1	69.3	-0.1	54.3	-0.1	-0.1	49.5	48.6	-0.1	-0.1	-0.1	-0.1	103.0	-0.1
NNN3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	60.3	-0.1
NNN4	-0.1	-0.1	5.8	7.2	6.4	95.4	-0.1	69.9	50.7	-0.1	51.6	50.7	-0.1	-0.1	-0.1	-0.1	121.0	-0.1
NNN5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NNN6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	64.2	-0.1
NNN7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	60.3	-0.1
NNN8	-0.1	-0.1	6.2	7.2	-0.1	89.7	-0.1	69.0	-0.1	-0.1	51.0	49.8	-0.1	-0.1	-0.1	-0.1	139.0	-0.1
NNN9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.3	-0.1
NO1	-0.1	-0.1	5.6	7.2	-0.1	55.2	-0.1	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	74.7	-0.1
NO2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	62.4	-0.1
NOO1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	94.5	-0.1
NOO2	-0.1	-0.1	-0.1	-0.1	-0.1	50.0	-0.1	60.6	-0.1	-0.1	48.9	48.3	-0.1	-0.1	-0.1	-0.1	89.4	-0.1
NOO3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	65.1	-0.1
NOO4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.1	-0.1
NOO5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.0	-0.1
NOO6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	65.4	-0.1
NOO6-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	64.2	-0.1
NOO7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.0	-0.1
NOO8	-0.1	-0.1	6.6	7.0	6.9	148.0	-0.1	69.9	52.8	-0.1	53.1	51.6	-0.1	-0.1	-0.1	-0.1	22.6	-0.1
NP1	-0.1	-0.1	6.1	6.5	-0.1	63.3	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NP2	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	77.4	-0.1
NP3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1
NP4	7.6	7.7	8.7	10.8	7.3	210.0	-0.1	103.0	55.2	51.9	58.8	57.0	48.6	-0.1	50.1	-0.1	21.1	46.2
NPP4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	63.9	-0.1
NPP1-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	61.8	-0.1
NPP2	7.0	6.8	6.4	8.3	6.8	148.0	-0.1	94.8	51.6	-0.1	56.7	55.2	-0.1	-0.1	48.9	-0.1	129.0	-0.1
NPP3	-0.1	-0.1	6.0	7.4	6.8	129.0	-0.1	73.2	51.3	-0.1	51.0	51.3	-0.1	-0.1	-0.1	-0.1	136.0	-0.1
NPP4	-0.1	-0.1	5.6	-0.1	-0.1	79.2	-0.1	62.1	49.2	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	98.1	-0.1
NQ1	7.4	7.3	6.6	8.4	6.9	111.0	52.2	93.0	51.9	49.8	56.7	57.0	-0.1	-0.1	49.5	-0.1	102.0	45.9
NQ2	-0.1	7.1	7.4	9.2	6.5	233.0	56.7	147.0	-0.1	-0.1	63.6	63.3	-0.1	-0.1	-0.1	-0.1	131.0	-0.1
NQ2-R	-0.1	7.1	7.6	8.2	-0.1	240.0	56.7	124.0	-0.1	-0.1	64.8	62.7	-0.1	-0.1	-0.1	-0.1	165.0	-0.1
NQ3	-0.1	-0.1	5.8	7.2	-0.1	75.3	-0.1	68.1	-0.1	-0.1	49.8	48.9	-0.1	-0.1	-0.1	-0.1	87.3	-0.1
NQ4	-0.1	6.8	6.9	7.3	-0.1	140.0	-0.1	79.5	-0.1	-0.1	54.3	53.1	-0.1	-0.1	-0.1	-0.1	138.0	-0.1
NQ5	-0.1	-0.1	5.7	7.0	-0.1	55.5	-0.1	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	77.7	-0.1
NQ6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	74.1	-0.1
NQ7	-0.1	6.6	6.3	7.8	6.8	124.0	-0.1	76.8	51.0	-0.1	53.7	52.9	-0.1	-0.1	50.1	-0.1	196.0	-0.1
NQ8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	77.7	-0.1
NQQ1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.6	-0.1
NR1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1
NR11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	82.8	-0.1
NR12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	92.7	-0.1
NR13	-0.1	-0.1	5.7	7.1	-0.1	60.9	-0.1	63.0	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	91.2	-0.1
NR14	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	16.9	-0.1
NR2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	13.3	-0.1
NR3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.0	-0.1
NR4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	63.6	-0.1
NR5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.3	-0.1
NR6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.6	-0.1
NR7	6.9	6.6	6.0	7.4	6.6	106.0	-0.1	78.0	50.1	-0.1	52.5	52.5	-0.1	-0.1	-0.1	-0.1	20.4	-0.1
NR8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.6	-0.1
NS1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NS10	7.2	7.5	7.6	9.6	6.9	291.0	67.8	152.0	51.9	49.2	69.6	69.3	-0.1	-0.1	50.1	-0.1	245.0	46.2
NS11	7.2	6.7	6.5	7.0	6.5	128.0	-0.1	82.2	49.5	-0.1	56.7	55.5	-0.1	-0.1	-0.1	-0.1	125.0	-0.1
NS12	-0.1	6.8	6.9	8.0	6.6	176.0	-0.1	97.5	49.8	-0.1	56.4	56.7	-0.1	-0.1	-0.1	-0.1	198.0	-0.1
NS13	7.1	7.2	7.0	8.7	6.8	233.0	56.1	143.0	50.7	-0.1	68.7	68.1	-0.1	-0.1	49.8	-0.1	182.0	46.2
NS14	8.6	7.9	7.6	9.8	7.2	247.0	57.0	137.0	55.5	54.0	71.7	69.0	50.7	50.4	50.7	47.7	200.0	48.6
NS2	-0.1	-0.1	5.6	-0.1	-0.1	78.9	-0.1	61.2	48.6	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	105.0	-0.1
NS3	-0.1	6.5	6.7	7.2	-0.1	93.9	-0.1	61.2	49.8	-0.1	50.4	49.2	-0.1	-0.1	-0.1	-0.1	105.0	-0.1
NS4	-0.1	6.5	6.3	6.8	-0.1	63.9	-0.1	61.2	-0.1	-0.1	51.0	50.1	-0.1	-0.1	-0.1	-0.1	100.0	-0.1
NS4-R	-0.1	-0.1	5.6	7.2	-0.1	66.8	-0.1	67.8	-0.1	-0.1	53.9	51.0	-0.1	-0.1	-0.1	-0.1	112.0	-0.1
NS5	-0.1	6.5	6.2	7.7	-0.1	92.7	-0.1	78.9	-0.1	-0.1	52.8	51.6	-0.1	-0.1	-0.1	-0.1	110.0	-0.1
NS6	-0.1	-0.1	5.7	7.1	-0.1	57.9	-0.1	60.3	-0.1	-0.1	48.3	49.9	-0.1	-0.1	-0.1	-0.1	82.8	-0.1
NS7	-0.1	6.7	6.7	8.2	-0.1	158.0	-0.1	114.0	-0.1	-0.1	60.0	58.2	-0.1	-0.1	-0.1	-0.1	18.8	-0.1
NS8	7.1	7.3	7.6	9.6	6.7	219.0	54.0	145.0	50.1	-0.1	69.0	66.3	-0.1	-0.1	50.1	-0.1	155.0	46.5
NS9	-0.1	-0.1	-0.1	-0.1	-0.1	50.4	-0.1	62.4	-0.1	-0.1	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	78.3	-0.1
NT1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.5	-0.1
NT10	-0.1	-0.1	5.7	7.1	-0.1	48.3	-0.1	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	67.8	-0.1
NT11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	42.1	-0.1
NT12	-0.1	-0.1	5.6	7.1	-0.1	61.5	-0.1	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NT13	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.7	-0.1
NT14	-0.1	-0.1	-0.1	7.0	-0.1	69.6	-0.1	58.5	48.9	-0.1	48.0	48.3	-0.1	-0.1	-0.1	-0.1	115.0	-0.1
NT15	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	63.1	-0.1
NT16	-0.1	-0.1	5.7	7.0	-0.1	59.7	-0.1	58.2	-0.1	-0.1	48.0	49.2	-0.1	-0.1	-0.1	-0.1	102.0	-0.1
NT17	-0.1	-0.1	-0.1	-0.1	-0.1	58.7	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.9	-0.1
NT18	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	60.9	-0.1
NT2	-0.1	-0.1	6.2	6.5	-0.1	79.8	-0.1	60.6	-0.1	-0.1	51.0	50.1	-0.1	-0.1	-0.1	-0.1	106.0	-0.1
NT3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.3	-0.1
NT4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.0	-0.1
NT5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.0	-0.1
NT6-R	-0.1	-0.1	-0.1	6.5	-0.1	82.2	-0.1	60.9	48.9	-0.1	49.2	49.5	-0.1	-0.1	-0.1	-0.1	93.9	-0.1
NT6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.9	-0.1
NT7	-0.1	-0.1	5.7	7.1	-0.1	58.2	-0.1	65.4	-0.1	-0.1	50.3	49.5	-0.1	-0.1	-0.1	-0.1	90.3	-0.1
NT8	7.2	6.7	7.0	7.8	6.5	158.0	-0.1	91.2	-0.1	-0.1	57.6	57.6	-0.1	-0.1	-0.1	-0.1	131.0	-0.1
NT9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.4	-0.1
NU1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.9	-0.1
NU10	7.5	7.6	7.7	9.7	7.0	432.0	102.0	253.0	64.6	54.3	107.0	103.0	49.5	50.7	63.9	-0.1	906.0	59.1
NU11	-0.1	-0.1	-0.1	6.5	-0.1	73.2	-0.1	63.0	-0.1	-0.1	49.8	48.6	-0.1	-0.1	-0.1	-0.1	80.7	-0.1

A14-06865

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Alex Pleson
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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NU12	-0.1	-0.1	5.6	7.0	-0.1	75.6	-0.1	60.9	-0.1	-0.1	48.6	-0.1	-0.1	-0.1	-0.1	-0.1	106.0	-0.1
NU12-R	-0.1	-0.1	5.8	7.1	-0.1	87.9	-0.1	69.3	-0.1	-0.1	49.2	49.8	-0.1	-0.1	-0.1	-0.1	108.0	-0.1
NU13	6.9	6.9	7.1	9.1	6.5	196.0	-0.1	111.0	48.9	-0.1	56.7	56.4	-0.1	-0.1	-0.1	-0.1	137.0	-0.1
NU14	-0.1	-0.1	-0.1	-0.1	-0.1	60.0	-0.1	-0.1	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.2	-0.1
NU15	-0.1	-0.1	6.3	6.8	-0.1	76.2	-0.1	56.4	-0.1	-0.1	48.0	48.0	-0.1	-0.1	-0.1	-0.1	120.0	-0.1
NU16	7.0	6.7	6.6	8.0	6.6	108.0	-0.1	75.0	-0.1	49.5	52.8	51.6	-0.1	-0.1	-0.1	-0.1	182.0	-0.1
NU17	-0.1	-0.1	5.6	-0.1	-0.1	78.6	-0.1	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	132.0	-0.1
NU18	-0.1	-0.1	5.8	7.1	-0.1	50.1	-0.1	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.8	-0.1
NU19	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.1	-0.1
NU2	-0.1	7.0	7.9	9.6	-0.1	221.0	-0.1	137.0	-0.1	-0.1	60.9	59.1	-0.1	-0.1	-0.1	-0.1	113.0	-0.1
NU3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	75.3	-0.1
NU4	7.8	8.5	12.7	14.1	7.1	846.0	66.0	402.0	53.4	50.1	123.0	116.0	-0.1	51.0	54.6	-0.1	315.0	51.0
NU5	-0.1	-0.1	-0.1	-0.1	-0.1	48.8	-0.1	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	40.8	-0.1
NU6	-0.1	-0.1	5.4	6.4	-0.1	60.0	-0.1	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	79.8	-0.1
NU7	-0.1	-0.1	5.6	7.1	-0.1	65.1	-0.1	64.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.4	-0.1
NU8	-0.1	-0.1	6.0	8.0	-0.1	92.4	-0.1	65.7	-0.1	-0.1	49.8	48.6	-0.1	-0.1	-0.1	-0.1	127.0	-0.1
NU9	7.0	-0.1	6.3	7.6	6.6	93.6	-0.1	71.1	50.1	49.2	51.8	51.3	-0.1	-0.1	-0.1	-0.1	17.5	-0.1
NV1	-0.1	-0.1	-0.1	-0.1	-0.1	59.7	-0.1	54.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	85.5	-0.1
NV10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.6	-0.1
NV11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	66.3	-0.1
NV12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	70.8	-0.1
NV13	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	89.7	-0.1
NV14	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	103.0	-0.1
NV15	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.8	-0.1
NV16	-0.1	-0.1	-0.1	-0.1	-0.1	76.5	-0.1	56.8	51.0	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1
NV17	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	54.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	90.6	-0.1
NV18	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	79.8	-0.1
NV2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	80.7	-0.1
NV3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.6	-0.1
NV4	-0.1	-0.1	-0.1	6.9	-0.1	61.5	-0.1	58.2	-0.1	-0.1	48.0	-0.1	-0.1	-0.1	-0.1	-0.1	110.0	-0.1
NV5	-0.1	-0.1	-0.1	7.0	-0.1	72.9	-0.1	58.6	60.1	-0.1	48.0	48.3	-0.1	-0.1	-0.1	-0.1	15.8	-0.1
NV5-R	-0.1	-0.1	-0.1	7.0	-0.1	81.9	-0.1	62.7	49.5	-0.1	49.2	49.5	-0.1	-0.1	-0.1	-0.1	17.9	-0.1
NV6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	70.8	-0.1
NV7	-0.1	-0.1	-0.1	-0.1	-0.1	62.4	-0.1	54.9	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	110.0	-0.1
NV8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	79.8	-0.1
NV9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	95.4	-0.1
NW1	-0.1	-0.1	6.0	6.4	-0.1	71.4	-0.1	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	46.5	-0.1
NW10	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	86.1	-0.1
NW11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.2	-0.1
NW12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	82.2	-0.1
NW13	-0.1	-0.1	5.8	7.3	-0.1	68.4	-0.1	66.0	-0.1	-0.1	48.6	49.2	-0.1	-0.1	-0.1	-0.1	88.2	-0.1
NW14	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.0	-0.1
NW15	6.9	6.5	5.8	8.7	-0.1	83.7	-0.1	64.2	-0.1	-0.1	51.6	50.7	-0.1	-0.1	-0.1	-0.1	99.6	-0.1
NW16	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.0	-0.1
NW17	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	85.5	-0.1
NW17-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.6	-0.1
NW18	-0.1	-0.1	-0.1	-0.1	-0.1	57.9	-0.1	63.3	-0.1	-0.1	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	87.6	-0.1
NW2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	91.2	-0.1
NW2-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.8	-0.1
NW3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	52.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	118.0	-0.1
NW4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.9	-0.1
NW5	7.0	6.6	6.2	7.6	6.6	106.0	-0.1	78.3	-0.1	-0.1	52.8	53.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1
NW6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	83.7	-0.1
NW7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	80.4	-0.1
NW8	-0.1	-0.1	6.1	8.5	-0.1	63.3	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	87.3	-0.1
NW9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	89.7	-0.1
NX1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	51.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.6	-0.1
NX10	7.0	6.7	6.4	7.9	6.6	100.0	-0.1	77.7	50.4	-0.1	51.6	52.2	-0.1	-0.1	-0.1	-0.1	97.2	-0.1
NX11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	65.7	-0.1
NX12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.0	-0.1
NX13	-0.1	-0.1	5.8	7.4	-0.1	78.0	-0.1	62.1	49.5	-0.1	48.0	-0.1	-0.1	-0.1	-0.1	-0.1	98.1	-0.1
NX14	-0.1	-0.1	5.7	7.1	-0.1	55.8	-0.1	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	81.9	-0.1
NX14-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	49.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	65.5	-0.1
NX15	-0.1	-0.1	5.6	-0.1	-0.1	91.8	-0.1	57.9	51.0	-0.1	49.2	48.3	-0.1	-0.1	-0.1	-0.1	117.0	-0.1
NX16	-0.1	-0.1	6.0	6.9	-0.1	69.0	-0.1	51.9	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	97.5	-0.1
NX17	7.2	6.5	6.6	7.0	-0.1	78.9	-0.1	57.6	50.4	-0.1	50.4	49.5	-0.1	-0.1	-0.1	-0.1	111.0	-0.1
NX2	7.0	7.0	6.5	8.2	6.7	206.0	53.7	127.0	50.4	-0.1	68.7	66.9	-0.1	-0.1	-0.1	50.7	203.0	46.8

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI
NX3	-0.1	7.0	6.5	8.3	6.6	149.0	52.2	108.0	50.1	-0.1	61.5	60.0	-0.1	-0.1	-0.1	-0.1	124.0	-0.1
NX4	7.6	8.4	8.7	9.8	6.7	516.0	62.1	238.0	50.7	50.4	100.0	98.3	-0.1	-0.1	53.7	-0.1	348.0	49.2
NX5	-0.1	-0.1	5.8	7.2	-0.1	83.4	-0.1	70.8	-0.1	-0.1	49.8	50.4	-0.1	-0.1	-0.1	-0.1	93.3	-0.1
NX6	-0.1	6.6	6.9	7.3	-0.1	156.0	-0.1	87.3	-0.1	-0.1	55.2	55.2	-0.1	-0.1	-0.1	-0.1	192.0	-0.1
NX7	-0.1	7.4	8.8	11.0	6.6	342.0	-0.1	180.0	49.5	48.6	68.4	65.7	-0.1	-0.1	-0.1	-0.1	182.0	-0.1
NX8	7.1	6.5	6.5	7.0	-0.1	99.0	-0.1	64.5	50.4	-0.1	50.4	50.7	-0.1	-0.1	-0.1	-0.1	145.0	-0.1
NX9	7.1	6.9	7.1	8.7	6.7	147.0	-0.1	91.5	51.0	-0.1	54.6	53.1	-0.1	-0.1	-0.1	-0.1	16.5	-0.1
NY1	-0.1	-0.1	-0.1	-0.1	-0.1	59.7	-0.1	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	93.6	-0.1
NY10	6.9	-0.1	5.9	7.5	6.8	153.0	-0.1	80.1	51.6	-0.1	54.9	53.7	-0.1	-0.1	-0.1	-0.1	192.0	-0.1
NY11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.4	-0.1
NY12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	61.5	-0.1	-0.1	48.3	48.9	-0.1	-0.1	-0.1	-0.1	81.3	-0.1
NY13	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	67.8	-0.1
NY14	-0.1	-0.1	5.8	7.2	-0.1	77.4	-0.1	66.6	49.8	-0.1	49.5	50.1	-0.1	-0.1	-0.1	-0.1	93.9	-0.1
NY15	-0.1	-0.1	-0.1	-0.1	-0.1	66.3	-0.1	54.3	49.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	114.0	-0.1
NY2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	76.2	-0.1
NY3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	50.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	79.8	-0.1
NY4	-0.1	-0.1	-0.1	6.4	-0.1	-0.1	-0.1	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.5	-0.1
NY4-R	-0.1	6.5	6.2	7.7	-0.1	119.0	-0.1	89.7	-0.1	-0.1	51.3	51.6	-0.1	-0.1	-0.1	-0.1	100.0	-0.1
NY5	-0.1	-0.1	-0.1	-0.1	-0.1	58.5	-0.1	61.5	-0.1	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	-0.1	88.5	-0.1
NY6	-0.1	-0.1	5.7	7.2	-0.1	87.5	-0.1	73.8	-0.1	-0.1	51.9	52.2	-0.1	-0.1	-0.1	-0.1	105.0	-0.1
NY7	7.0	6.9	6.4	8.1	6.7	206.0	53.1	126.0	50.1	-0.1	65.1	63.6	-0.1	-0.1	-0.1	-0.1	155.0	-0.1
NY8	-0.1	-0.1	-0.1	-0.1	-0.1	55.5	-0.1	55.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	99.9	-0.1
NY9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	72.0	-0.1
NZ1	-0.1	-0.1	-0.1	-0.1	-0.1	57.0	-0.1	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	67.0	-0.1
NZ10	7.5	-0.1	6.1	6.6	6.8	89.7	-0.1	56.4	52.2	48.6	50.4	50.4	-0.1	-0.1	48.6	-0.1	16.9	-0.1
NZ11	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	73.5	-0.1
NZ12	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	53.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	87.6	-0.1
NZ13	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	71.7	-0.1
NZ2	-0.1	-0.1	6.1	6.7	-0.1	63.9	-0.1	64.2	-0.1	-0.1	49.2	48.6	-0.1	-0.1	-0.1	-0.1	75.9	-0.1
NZ2-R	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	74.7	-0.1
NZ3	-0.1	-0.1	6.3	6.7	-0.1	61.5	-0.1	56.4	-0.1	-0.1	50.1	49.5	-0.1	-0.1	-0.1	-0.1	15.6	-0.1
NZ4	-0.1	-0.1	-0.1	-0.1	-0.1	52.2	-0.1	56.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	78.6	-0.1
NZ5	-0.1	-0.1	-0.1	-0.1	-0.1	54.6	-0.1	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	84.9	-0.1
NZ6	-0.1	-0.1	-0.1	-0.1	-0.1	56.7	-0.1	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	110.0	-0.1
NZ7	-0.1	-0.1	-0.1	-0.1	7.0	105.0	-0.1	56.4	53.7	-0.1	48.3	48.9	-0.1	-0.1	-0.1	-0.1	145.0	-0.1
NZ8	-0.1	-0.1	-0.1	-0.1	-0.1	70.2	-0.1	56.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	70.2	-0.1
NZ9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	69.9	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	48.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	88.5	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.7	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	61.5	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	58.4	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	68.1	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	63.3	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	63.0	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	55.4	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	67.2	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	56.4	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	61.5	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	59.1	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	127 MPH	128 MPH	129 HAR	130 HAR	131 MPH	132 ALK	133 HAR	134 HAR	135 MPH	136 MPH	137 HBI	138 HBI	139 HPH	140 HPH	141 HBI	142 HPH	143 HA	144 HBI

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	145-HBA	146-HPH	147-HBI	148-HRH	149-HBI	150-HPH	151-NBI	152-HPH	153-HPH	154-HRH	155-HPH	156-HBI	157-HAR	158-HBA	159-HBA	160-NBI	161-HA	162-HPH
NA1	119.0	-0.1	46.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	228.0	-0.1	-0.1	279.0	-0.1
NA2	225.0	-0.1	48.3	-0.1	46.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	390.0	-0.1	-0.1	474.0	-0.1
NAA1	75.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	244.0	-0.1
NAA10	71.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	296.0	-0.1
NAA11	67.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	231.0	-0.1
NAA2	159.0	46.5	48.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	269.0	-0.1	-0.1	336.0	-0.1
NAA3	98.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	255.0	-0.1
NAA4	111.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	224.0	-0.1	-0.1	278.0	-0.1
NAA5	77.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	241.0	-0.1
NAA6	91.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	261.0	-0.1
NAA7	94.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	230.0	-0.1	-0.1	287.0	-0.1
NAA8	95.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	263.0	-0.1
NAA9	85.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	218.0	-0.1	-0.1	271.0	-0.1
NB1	77.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	255.0	-0.1
NB2	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	227.0	-0.1
NB3	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	226.0	-0.1
NB4	144.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	263.0	-0.1	-0.1	321.0	-0.1
NBB1	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	237.0	-0.1
NBB10	90.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	213.0	-0.1	-0.1	258.0	-0.1
NBB11	124.0	45.9	47.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	243.0	-0.1	-0.1	303.0	-0.1
NBB2	71.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	239.0	-0.1
NBB2-R	69.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	239.0	-0.1
NBB3	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	231.0	-0.1
NBB4	54.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	243.0	-0.1
NBB5	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	236.0	-0.1
NBB6	78.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	214.0	-0.1	-0.1	259.0	-0.1
NBB7	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	226.0	-0.1
NBB8	69.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	257.0	-0.1
NBB9	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	234.0	-0.1
NC1	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	169.0	-0.1	-0.1	230.0	-0.1
NC2	79.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	214.0	-0.1	-0.1	267.0	-0.1
NC3	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	251.0	-0.1
NC4	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NC5	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	247.0	-0.1
NC6	71.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	255.0	-0.1
NC7	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	221.0	-0.1
NC8	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	219.0	-0.1
NCC1	1040.0	50.7	78.6	46.8	64.2	-0.1	66.9	-0.1	174.0	-0.1	-0.1	171.0	189.0	1030.0	191.0	176.0	1220.0	-0.1
NCC1-R	969.0	49.2	74.7	48.0	60.3	-0.1	60.9	-0.1	173.0	-0.1	-0.1	168.0	189.0	867.0	180.0	173.0	1100.0	-0.1
NCC2	303.0	50.4	52.2	49.8	48.9	47.4	48.9	46.3	176.0	193.0	163.0	-0.1	170.0	387.0	170.0	173.0	474.0	-0.1
NCC3	87.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	227.0	-0.1	-0.1	274.0	-0.1
NCC4	81.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	214.0	-0.1	-0.1	267.0	-0.1
NCC5	75.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	255.0	-0.1
NCC6	69.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	287.0	-0.1
ND1	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	220.0	-0.1
ND10	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	228.0	-0.1
ND2	75.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	243.0	-0.1
ND3	10.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	249.0	-0.1
ND4	18.8	-0.1	46.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	233.0	-0.1	-0.1	285.0	-0.1
ND5	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	222.0	-0.1
ND6	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
ND7	65.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	162.0	-0.1	244.0	-0.1
ND8	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	229.0	-0.1
ND8-R	59.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	239.0	-0.1
ND9	69.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	233.0	-0.1
NDD1	89.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	212.0	-0.1	-0.1	269.0	-0.1
NDD2	69.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	247.0	-0.1
NDD3	80.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	249.0	-0.1
NDD4	70.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	230.0	-0.1
NDD5	84.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	261.0	-0.1
NE1	81.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	213.0	-0.1	-0.1	266.0	-0.1
NE10	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	216.0	-0.1
NE11	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	246.0	-0.1
NE2	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	231.0	-0.1
NE3	66.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	235.0	-0.1
NE4	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	228.0	-0.1
NE4-R	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	219.0	-0.1

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SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 HBA	146 HPH	147 HBI	148 HPH	149 HBI	150 HPH	151 HBI	152 HPH	153 HPH	154 HPH	155 HPH	156 HBI	157 HAR	158 HBA	159 HBA	160 HBI	161 HA	162 HPH
NE5	67.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	255.0	-0.1
NE6	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	229.0	-0.1
NE7	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	161.0	-0.1	-0.1	227.0	-0.1
NE8	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	224.0	-0.1
NE9	67.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	196.0	-0.1	-0.1	269.0	-0.1
NEE1	65.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	231.0	-0.1
NEE2	65.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	166.0	-0.1	-0.1	233.0	-0.1
NEE3	78.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	207.0	-0.1	-0.1	251.0	-0.1
NEE4	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	248.0	-0.1
NEE5	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	231.0	-0.1
NEE6	78.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	196.0	-0.1	-0.1	248.0	-0.1
NEE7	67.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	238.0	-0.1
NF1	79.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	201.0	-0.1	-0.1	246.0	-0.1
NF10	159.0	-0.1	47.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	264.0	-0.1	-0.1	330.0	-0.1
NF11	934.0	54.6	75.6	48.9	64.5	48.9	81.6	53.4	190.0	196.0	75.0	204.0	236.0	2370.0	228.0	214.0	2780.0	158.0
NF2	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	233.0	-0.1
NF3	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	226.0	-0.1
NF4	85.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	248.0	-0.1
NF5	86.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	215.0	-0.1	-0.1	268.0	-0.1
NF6	123.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	256.0	-0.1
NF7	131.0	46.2	46.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	277.0	-0.1	-0.1	345.0	-0.1
NF8	82.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	215.0	-0.1	-0.1	268.0	-0.1
NF8-R	10.0	45.0	47.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	275.0	-0.1	-0.1	333.0	-0.1
NF9	83.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	240.0	-0.1
NFF1	88.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	218.0	-0.1	-0.1	270.0	-0.1
NFF2	88.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	253.0	-0.1
NFF3	88.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	257.0	-0.1
NFF4	109.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	227.0	-0.1	-0.1	276.0	-0.1
NFF5	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	288.0	-0.1
NFF6	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	228.0	-0.1
NFF6-R	59.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	221.0	-0.1
NFF7	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	236.0	-0.1
NFF8	80.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	249.0	-0.1	-0.1	342.0	-0.1
NG1	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	172.0	-0.1	-0.1	222.0	-0.1
NG10	72.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	245.0	-0.1
NG11	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	172.0	-0.1	-0.1	217.0	-0.1
NG2	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	249.0	-0.1
NG2-R	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	227.0	-0.1
NG3	63.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	238.0	-0.1
NG4	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	218.0	-0.1
NG5	79.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	212.0	-0.1	-0.1	264.0	-0.1
NG6	66.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	229.0	-0.1
NG7	51.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	168.0	-0.1	-0.1	-0.1	-0.1
NG8	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	228.0	-0.1
NG9	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	236.0	-0.1
NGG1	68.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	234.0	-0.1
NGG10	56.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	234.0	-0.1
NGG2	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	232.0	-0.1
NGG3	4.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	226.0	-0.1	-0.1	278.0	-0.1
NGG4	70.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	242.0	-0.1
NGG5	68.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	246.0	-0.1
NGG6	78.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	221.0	-0.1	-0.1	275.0	-0.1
NGG7	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	243.0	-0.1
NGG8	67.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	238.0	-0.1
NGG9	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	221.0	-0.1
NH1	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	241.0	-0.1
NH10	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NH10-R	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	230.0	-0.1
NH11	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	241.0	-0.1
NH2	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	230.0	-0.1
NH3	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	172.0	-0.1	-0.1	218.0	-0.1
NH4	63.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	237.0	-0.1
NH5	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	227.0	-0.1
NH6	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	247.0	-0.1
NH7	93.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	230.0	-0.1	-0.1	281.0	-0.1
NH8	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	207.0	-0.1	-0.1	259.0	-0.1
NH9	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	245.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 HBA	146 HPH	147 HBI	148 HPH	149 HBI	150 HPH	151 HBI	152 HPH	153 HPH	154 HPH	155 HPH	156 HBI	157 HAR	158 HBA	159 HBA	160 HBI	161 HA	162 HPH
NHH1	66.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	236.0	-0.1
NHH10	63.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	235.0	-0.1
NHH11	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	221.0	-0.1
NHH11-R	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	216.0	-0.1
NHH12	64.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	236.0	-0.1
NHH2	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	264.0	-0.1
NHH3	74.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	237.0	-0.1
NHH4	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	235.0	-0.1
NHH5	68.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	241.0	-0.1
NHH6	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	230.0	-0.1
NHH7	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	231.0	-0.1
NHH8	67.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	226.0	-0.1
NHH9	69.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	233.0	-0.1
NI1	11.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	248.0	-0.1	-0.1	309.0	-0.1
NI10	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	225.0	-0.1
NI2	63.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	257.0	-0.1
NI3	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	225.0	-0.1
NI4	11.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	230.0	-0.1	-0.1	280.0	-0.1
NI5	69.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	236.0	-0.1
NI5-R	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	228.0	-0.1
NI6	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	254.0	-0.1
NI7	84.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	261.0	-0.1
NI8	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	229.0	-0.1
NI9	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	196.0	-0.1	-0.1	251.0	-0.1
NII1	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	223.0	-0.1
NII10	74.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	165.0	-0.1	-0.1	232.0	-0.1
NII11	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	239.0	-0.1
NII1-R	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	245.0	-0.1
NII12	69.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	225.0	-0.1
NII13	74.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	235.0	-0.1
NII14	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	221.0	-0.1
NII2	12.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	226.0	-0.1	-0.1	266.0	-0.1
NII3	8.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	223.0	-0.1	-0.1	271.0	-0.1
NII4	10.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	205.0	-0.1	-0.1	250.0	-0.1
NII5	90.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	207.0	-0.1	-0.1	251.0	-0.1
NII6	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	260.0	-0.1
NII7	62.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	219.0	-0.1
NII8	77.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	240.0	-0.1
NII9	60.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	224.0	-0.1
NJ1	73.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	250.0	-0.1
NJ2	67.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	235.0	-0.1
NJ3	72.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	240.0	-0.1
NJ4	75.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	250.0	-0.1
NJ5	60.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	221.0	-0.1
NJ6	63.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	235.0	-0.1
NJ7	66.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	166.0	-0.1	-0.1	234.0	-0.1
NJJ1	72.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	256.0	-0.1
NJJ10	361.0	48.0	54.9	36.5	49.2	-0.1	47.7	46.8	175.0	-0.1	-0.1	-0.1	176.0	321.0	163.0	-0.1	414.0	-0.1
NJJ11	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	240.0	-0.1
NJJ12	78.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	242.0	-0.1
NJJ13	81.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	241.0	-0.1
NJJ14	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
NJJ2	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NJJ3	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	220.0	-0.1
NJJ4	80.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	246.0	-0.1
NJJ5	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	226.0	-0.1
NJJ6	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	227.0	-0.1
NJJ7	63.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	234.0	-0.1
NJJ8	56.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	214.0	-0.1
NJJ8-R	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NJJ9	95.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	255.0	-0.1
NK1	72.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	234.0	-0.1
NK2	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	226.0	-0.1
NK3	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	240.0	-0.1
NK4	72.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	247.0	-0.1
NK4-R	63.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	233.0	-0.1
NK5	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	236.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145-HBA	146-HPH	147-HBI	148-HPH	149-HBI	150-HPH	151-HBI	152-HPH	153-HPH	154-HPH	155-HPH	156-HBI	157-HAR	158-HBA	159-HBA	160-HBI	161-HA	162-HPH
NK6	66.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	254.0	-0.1
NK7	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	240.0	-0.1
NKK1	56.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	217.0	-0.1
NKK10	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	233.0	-0.1
NKK11	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	222.0	-0.1
NKK2	75.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	244.0	-0.1
NKK3	74.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NKK4	190.0	45.6	49.5	-0.1	45.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	256.0	-0.1	-0.1	312.0	-0.1
NKK5	10.1	-0.1	49.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	263.0	-0.1
NKK6	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	226.0	-0.1
NKK7	59.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	165.0	-0.1	-0.1	226.0	-0.1
NKK8	86.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	216.0	-0.1	-0.1	263.0	-0.1
NKK9	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	286.0	-0.1
NL1	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	220.0	-0.1
NL2	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	160.0	-0.1	-0.1	220.0	-0.1
NL3	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	219.0	-0.1
NL3-R	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
NL4	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	221.0	-0.1
NLL1	72.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	168.0	-0.1	-0.1	229.0	-0.1
NLL10	66.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	231.0	-0.1
NLL11	64.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	282.0	-0.1
NLL12	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	227.0	-0.1
NLL2	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	216.0	-0.1
NLL3	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	235.0	-0.1
NLL4	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NLL5	74.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	238.0	-0.1
NLL6	8.7	-0.1	45.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	224.0	-0.1	-0.1	273.0	-0.1
NLL7	78.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	258.0	-0.1
NLL8	72.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	196.0	-0.1	-0.1	248.0	-0.1
NLL9	73.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	251.0	-0.1
NLL9-R	72.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	248.0	-0.1
NM1	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NM2	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	286.0	-0.1
NMM1	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	223.0	-0.1
NMM10	72.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	243.0	-0.1
NMM2	71.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	230.0	-0.1
NMM3	60.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	284.0	-0.1
NMM3-R	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
NMM4	61.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NMM5	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	219.0	-0.1
NMM6	67.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	229.0	-0.1
NMM7	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	222.0	-0.1
NMM8	77.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	241.0	-0.1
NMM9	90.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	229.0	-0.1	-0.1	286.0	-0.1
NN1	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	219.0	-0.1
NNN1	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NNN10	71.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	250.0	-0.1
NNN2	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	239.0	-0.1
NNN3	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	167.0	-0.1	-0.1	209.0	-0.1
NNN4	75.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	241.0	-0.1
NNN5	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	172.0	-0.1	-0.1	223.0	-0.1
NNN6	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	214.0	-0.1
NNN7	55.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	212.0	-0.1
NNN8	75.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	251.0	-0.1
NNN9	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	221.0	-0.1
NO1	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	218.0	-0.1
NO2	54.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	221.0	-0.1
NOO1	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	230.0	-0.1
NOO2	70.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	255.0	-0.1
NOO3	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	220.0	-0.1
NOO4	65.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	238.0	-0.1
NOO5	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	219.0	-0.1
NOO6	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	218.0	-0.1
NOO6-R	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	171.0	-0.1	-0.1	215.0	-0.1
NOO7	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	228.0	-0.1
NOO8	86.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	263.0	-0.1
NP1	60.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	227.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 HBA	146 HPH	147 HBI	148 HPH	149 HBI	150 HPH	151 HBI	152 HPH	153 HPH	154 HPH	155 HPH	156 HBI	157 HAR	158 HBA	159 HBA	160 HBI	161 HA	162 HPH
NP2	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	231.0	-0.1
NP3	55.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	172.0	-0.1	-0.1	246.0	-0.1
NP4	9.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	222.0	-0.1	-0.1	284.0	-0.1
NPP4	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	213.0	-0.1
NPP1-R	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	169.0	-0.1	-0.1	214.0	-0.1
NPP2	87.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	206.0	-0.1	-0.1	269.0	-0.1
NPP3	79.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	211.0	-0.1	-0.1	257.0	-0.1
NPP4	57.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	226.0	-0.1
NQ1	78.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	201.0	-0.1	-0.1	246.0	-0.1
NQ2	88.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	232.0	-0.1
NQ2-R	103.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	244.0	-0.1
NQ3	55.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	230.0	-0.1
NQ4	82.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	246.0	-0.1
NQ5	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	220.0	-0.1
NQ6	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	225.0	-0.1
NQ7	86.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	259.0	-0.1
NQ8	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	223.0	-0.1
NQQ1	57.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	212.0	-0.1
NR1	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	220.0	-0.1
NR11	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	234.0	-0.1
NR12	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	240.0	-0.1
NR13	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	249.0	-0.1
NR14	68.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	244.0	-0.1
NR2	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	233.0	-0.1
NR3	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	221.0	-0.1
NR4	56.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	170.0	-0.1	-0.1	215.0	-0.1
NR5	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	170.0	-0.1	-0.1	213.0	-0.1
NR6	56.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	168.0	-0.1	-0.1	213.0	-0.1
NR7	80.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	207.0	-0.1	-0.1	253.0	-0.1
NR8	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	225.0	-0.1
NS1	60.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NS10	124.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	226.0	-0.1	-0.1	267.0	-0.1
NS11	88.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	242.0	-0.1
NS12	101.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	253.0	-0.1
NS13	7.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	208.0	-0.1	-0.1	255.0	-0.1
NS14	122.0	-0.1	47.7	-0.1	-0.1	-0.1	-0.1	56.7	177.0	-0.1	-0.1	-0.1	-0.1	241.0	-0.1	-0.1	300.0	-0.1
NS2	69.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	237.0	-0.1
NS3	70.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	209.0	-0.1	-0.1	280.0	-0.1
NS4	7.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	204.0	-0.1	-0.1	258.0	-0.1
NS4-R	66.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	209.0	-0.1	-0.1	265.0	-0.1
NS5	7.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	251.0	-0.1
NS6	64.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	232.0	-0.1
NS7	95.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	251.0	-0.1
NS8	106.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	211.0	-0.1	-0.1	266.0	-0.1
NS9	66.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NT1	59.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	218.0	-0.1
NT10	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	218.0	-0.1
NT11	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	224.0	-0.1
NT12	60.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	229.0	-0.1
NT13	59.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	238.0	-0.1
NT14	72.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	245.0	-0.1
NT15	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	249.0	-0.1
NT16	68.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	201.0	-0.1	-0.1	245.0	-0.1
NT17	53.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	164.0	-0.1	-0.1	224.0	-0.1
NT18	53.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	169.0	-0.1	-0.1	212.0	-0.1
NT2	71.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	157.0	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	228.0	-0.1
NT3	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	229.0	-0.1
NT4	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	224.0	-0.1
NT5	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	215.0	-0.1
NT6-R	68.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	283.0	-0.1
NT6	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	170.0	-0.1	-0.1	215.0	-0.1
NT7	73.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	235.0	-0.1
NT8	4.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	213.0	-0.1	-0.1	266.0	-0.1
NT9	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	175.0	-0.1	-0.1	220.0	-0.1
NU1	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	221.0	-0.1
NU10	303.0	46.5	53.4	-0.1	47.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	296.0	-0.1	-0.1	356.0	-0.1
NU11	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	219.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 -HBA	146 -HPH	147 -HBI	148 -HPH	149 -HBI	150 -HPH	151 -HBI	152 -HPH	153 -HPH	154 -HPH	155 -HPH	156 -HBI	157 -HAR	158 -HBA	159 -HBA	160 -HBI	161 -HA	162 -HPH
NU12	70.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	259.0	-0.1
NU12-R	70.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	236.0	-0.1
NU13	79.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	237.0	-0.1
NU14	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	226.0	-0.1
NU15	72.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	207.0	-0.1	-0.1	258.0	-0.1
NU16	16.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	232.0	-0.1	-0.1	289.0	-0.1
NU17	88.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	242.0	-0.1
NU18	60.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	229.0	-0.1
NU19	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	229.0	-0.1
NU2	12.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	238.0	-0.1
NU3	58.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	223.0	-0.1
NU4	184.0	45.9	49.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	227.0	-0.1	-0.1	287.0	-0.1
NU5	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
NU6	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	226.0	-0.1
NU7	58.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	164.0	-0.1	-0.1	238.0	-0.1
NU8	78.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	251.0	-0.1
NU9	73.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	205.0	-0.1	-0.1	248.0	-0.1
NV1	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	237.0	-0.1
NV10	56.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	165.0	-0.1	-0.1	233.0	-0.1
NV11	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	220.0	-0.1
NV12	60.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	223.0	-0.1
NV13	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	235.0	-0.1
NV14	59.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	240.0	-0.1
NV15	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	231.0	-0.1
NV16	77.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	219.0	-0.1	-0.1	266.0	-0.1
NV17	68.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	242.0	-0.1
NV18	59.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	230.0	-0.1
NV2	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	229.0	-0.1
NV3	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	228.0	-0.1
NV4	69.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	243.0	-0.1
NV5	73.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	209.0	-0.1	-0.1	254.0	-0.1
NV5-R	80.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	220.0	-0.1	-0.1	266.0	-0.1
NV6	69.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	227.0	-0.1
NV7	72.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	201.0	-0.1	-0.1	244.0	-0.1
NV8	59.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	226.0	-0.1
NV9	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NW1	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	237.0	-0.1
NW10	67.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	202.0	-0.1	-0.1	254.0	-0.1
NW11	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	223.0	-0.1
NW12	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	233.0	-0.1
NW13	70.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	242.0	-0.1
NW14	57.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	224.0	-0.1
NW15	73.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	244.0	-0.1
NW16	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	219.0	-0.1
NW17	64.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NW17-R	62.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NW18	59.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	232.0	-0.1
NW2	66.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	235.0	-0.1
NW2-R	66.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	238.0	-0.1
NW3	69.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	197.0	-0.1	-0.1	253.0	-0.1
NW4	61.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	190.0	-0.1	-0.1	237.0	-0.1
NW5	11.3	-0.1	46.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	251.0	-0.1	-0.1	306.0	-0.1
NW6	62.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	228.0	-0.1
NW7	61.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	227.0	-0.1
NW8	64.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	227.0	-0.1
NW9	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	235.0	-0.1
NX1	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	237.0	-0.1
NX10	72.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	248.0	-0.1
NX11	57.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	174.0	-0.1	-0.1	212.0	-0.1
NX12	58.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	173.0	-0.1	-0.1	218.0	-0.1
NX13	55.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	238.0	-0.1
NX14	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	228.0	-0.1
NX14-R	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	226.0	-0.1
NX15	73.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	189.0	-0.1	-0.1	238.0	-0.1
NX16	58.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	238.0	-0.1
NX17	75.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	256.0	-0.1
NX2	138.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	234.0	-0.1	-0.1	289.0	-0.1

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 HBA	146 HPH	147 HBI	148 HPH	149 HBI	150 HPH	151 HBI	152 HPH	153 HPH	154 HPH	155 HPH	156 HBI	157 HAR	158 HBA	159 HBA	160 HBI	161 HA	162 HPH
NX3	88.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	198.0	-0.1	-0.1	242.0	-0.1
NX4	217.0	-0.1	48.0	-0.1	45.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	253.0	-0.1	-0.1	321.0	-0.1
NX5	68.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NX6	76.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	239.0	-0.1
NX7	9.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	244.0	-0.1
NX8	79.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	214.0	-0.1	-0.1	259.0	-0.1
NX9	75.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	199.0	-0.1	-0.1	248.0	-0.1
NY1	67.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	184.0	-0.1	-0.1	230.0	-0.1
NY10	93.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	210.0	-0.1	-0.1	263.0	-0.1
NY11	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	224.0	-0.1
NY12	65.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	191.0	-0.1	-0.1	233.0	-0.1
NY13	57.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	171.0	-0.1	-0.1	216.0	-0.1
NY14	67.8	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	192.0	-0.1	-0.1	235.0	-0.1
NY15	75.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	201.0	-0.1	-0.1	253.0	-0.1
NY2	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	228.0	-0.1
NY3	63.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	225.0	-0.1
NY4	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	183.0	-0.1	-0.1	223.0	-0.1
NY4-R	71.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	168.0	-0.1	-0.1	229.0	-0.1
NY5	65.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	225.0	-0.1
NY6	77.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	248.0	-0.1
NY7	104.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	203.0	-0.1	-0.1	256.0	-0.1
NY8	58.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	195.0	-0.1	-0.1	238.0	-0.1
NY9	56.7	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	225.0	-0.1
NZ1	64.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	188.0	-0.1	-0.1	235.0	-0.1
NZ10	85.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	237.0	-0.1	-0.1	294.0	-0.1
NZ11	60.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	162.0	-0.1	-0.1	229.0	-0.1
NZ12	63.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	187.0	-0.1	-0.1	233.0	-0.1
NZ13	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	178.0	-0.1	-0.1	217.0	-0.1
NZ2	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	185.0	-0.1	-0.1	231.0	-0.1
NZ2-R	61.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	160.0	-0.1	-0.1	226.0	-0.1
NZ3	75.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	206.0	-0.1	-0.1	259.0	-0.1
NZ4	63.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	176.0	-0.1	-0.1	223.0	-0.1
NZ5	65.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	233.0	-0.1
NZ6	71.4	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	193.0	-0.1	-0.1	238.0	-0.1
NZ7	78.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	218.0	-0.1	-0.1	263.0	-0.1
NZ8	69.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	222.0	-0.1
NZ9	58.2	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	182.0	-0.1	-0.1	221.0	-0.1
LMB-QA	63.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	200.0	-0.1	-0.1	244.0	-0.1
LMB-QA	58.5	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	186.0	-0.1	-0.1	227.0	-0.1
LMB-QA	54.6	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	169.0	-0.1	-0.1	213.0	-0.1
LMB-QA	59.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	165.0	-0.1	-0.1	233.0	-0.1
LMB-QA	60.0	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	194.0	-0.1	-0.1	238.0	-0.1
LMB-QA	64.3	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	179.0	-0.1	-0.1	249.0	-0.1
LMB-QA	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	177.0	-0.1	-0.1	216.0	-0.1
LMB-QA	54.9	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	181.0	-0.1	-0.1	228.0	-0.1
LMB-QA	56.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	180.0	-0.1	-0.1	226.0	-0.1
LMB-QA	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	169.0	-0.1	-0.1	206.0	-0.1
LMB-QA	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	171.0	-0.1	-0.1	214.0	-0.1
LMB-QA	53.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	-0.1	168.0	-0.1	-0.1	212.0	-0.1

A14-06865

Pleson Geoscience for Canstar Resources Inc.
Alex Pleson
-1=Reporting Limit of 1pg/g (ppt=parts per trillion)

SOIL GAS HYDROCARBONS
(SGH) by GC/MS
NORTH SURVEY AREA
KENORA PROJECT

Activation Laboratories Ltd.
Date: November 25, 2014
R=Replicate Sample

	145 HBA	146 HPM	147 HBI	148 HPH	149 HBI	150 HPH	151 HBI	152 HPM	153 HPH	154 HPH	155 HPM	156 HBI	157 HAR	158 HBA	159 HBA	160 HBI	161 HA	162 HPH