

## BERG DRILL HOLE ASSAY SUMMARY

Drill Hole (BRG07-)	Mineral Horizon	From (m)	To (m)	Length (m)	Cu (%)	Cu-S (%)	Mo (%)	MoS <sub>2</sub> (%)	Ag (g/t)	CuEQ (%)	
138	Supergene	33.5	69.2	35.7	0.483	0.448	0.050	0.083	26.8	1.132	
	Hypogene	69.2	157.6	88.4	0.385	0.385	0.057	0.095	6.9	0.964	
	Combined	33.5	157.6	124.1	0.413	0.403	0.055	0.092	12.7	1.013	
				including	21.3	0.341	0.341	0.081	0.135	7.7	1.149
				including	24.3	0.584	0.584	0.046	0.077	11.0	1.088
139	Supergene	44.8	145.4	100.6	0.449	0.414	0.015	0.025	6.0	0.630	
				including	42.7	0.626	0.570	0.020	0.033	4.0	0.840
	Hypogene	145.4	163.7	18.3	0.282	0.282	0.007	0.012	3.0	0.368	
	Combined	44.8	163.7	118.9	0.423	0.393	0.014	0.023	5.5	0.591	
140	Supergene	54.1	133.2	79.1	0.539	0.499	0.010	0.017	4.2	0.661	
				including	51.6	0.665	0.614	0.008	0.013	4.7	0.772
141	Supergene	53.9	172.8	118.9	0.248	0.184	0.010	0.017	1.2	0.350	
142	Supergene	41.8	145.4	103.6	0.524	0.437	0.028	0.047	3.3	0.808	
				including	60.9	0.608	0.504	0.031	0.052	4.2	0.926
143	Hypogene	215.5	520.3	304.8	0.286	0.286	0.036	0.060	4.6	0.653	
				including	54.9	0.366	0.366	0.024	0.040	6.3	0.633
				including	79.2	0.329	0.329	0.054	0.090	5.7	0.872
144	Supergene	40.2	137.7	97.5	0.311	0.234	0.003	0.005	2.6	0.357	
	Hypogene	270.4	404.5	134.1	0.251	0.251	0.039	0.065	3.5	0.639	
				including	48.7	0.249	0.249	0.069	0.115	3.7	0.918
145	Supergene	38.7	148.4	109.7	0.454	0.406	0.049	0.082	4.2	0.940	
	Hypogene	148.4	236.8	88.4	0.321	0.321	0.058	0.097	4.0	0.890	
	Combined	38.7	236.8	198.1	0.394	0.363	0.053	0.088	4.1	0.917	
	*entire hole	Combined	7.2	389.2	382.0	0.241	0.223	0.047	0.078	2.9	0.699
146	Supergene	32.6	78.3	45.7	0.298	0.264	0.024	0.040	3.2	0.544	
	Hypogene	90.2	181.1	90.9	0.296	0.296	0.085	0.142	4.3	1.119	
	Hypogene	230.7	419.7	189.0	0.262	0.262	0.056	0.093	6.3	0.828	
	Hypogene	419.7	541.6	121.9	0.241	0.241	0.027	0.045	4.2	0.522	
	*entire hole	Combined	32.6	541.6	509.0	0.250	0.247	0.047	0.078	4.7	0.721
147	Supergene	44.8	105.8	61.0	0.282	0.260	0.007	0.012	1.7	0.359	
148	Supergene	25.0	81.4	56.4	0.344	0.290	0.017	0.028	2.4	0.519	
	Hypogene	94.6	397.8	303.2	0.108	0.108	0.016	0.027	6.3	0.300	
				including	53.8	0.137	0.137	0.023	0.038	1.9	0.365
149	Supergene	41.8	138.5	96.7	0.519	0.482	0.045	0.075	3.9	0.966	
	Hypogene	138.5	412.3	273.8	0.353	0.353	0.015	0.025	4.9	0.526	
				including	150.1	0.421	0.421	0.020	0.033	5.8	0.647
	Combined	41.8	412.3	370.5	0.396	0.386	0.023	0.038	4.7	0.643	
	*moly zone	Combined	26.5	157.6	131.1	0.446	0.418	0.048	0.080	4.3	0.923
150	Supergene	20.4	127.1	106.7	0.455	0.422	0.051	0.085	4.5	0.962	
	Hypogene	127.1	491.6	364.5	0.317	0.317	0.027	0.045	4.9	0.602	
	Combined	20.4	491.6	471.2	0.348	0.340	0.033	0.055	4.8	0.689	
	*moly zone	Combined	11.3	216.9	205.6	0.439	0.422	0.052	0.087	6.1	0.966

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151	Supergene	16.7	190.9	<b>174.2</b>	<b>0.566</b>	0.522	0.020	<b>0.033</b>	5.4	<b>0.789</b>
	Hypogene	190.9	320.8	<b>129.9</b>	<b>0.251</b>	0.251	0.014	<b>0.023</b>	3.6	<b>0.406</b>
	Combined	16.7	320.8	<b>304.1</b>	<b>0.432</b>	0.406	0.018	<b>0.030</b>	4.6	<b>0.631</b>
152	Supergene	18.9	185.0	<b>166.1</b>	<b>0.414</b>	0.395	0.019	<b>0.032</b>	5.0	<b>0.627</b>
	Combined	18.9	297.8	<b>278.9</b>	<b>0.356</b>	0.342	0.017	<b>0.028</b>	7.1	<b>0.563</b>
153	Supergene	29.1	118.6	<b>89.5</b>	<b>0.384</b>	0.355	0.008	<b>0.013</b>	7.6	<b>0.510</b>
	Hypogene	118.6	136.2	<b>17.6</b>	<b>0.307</b>	0.307	0.014	<b>0.023</b>	13.7	<b>0.531</b>
	Combined	29.1	136.2	<b>107.1</b>	<b>0.371</b>	0.346	0.009	<b>0.015</b>	8.6	<b>0.514</b>
154	Supergene	7.1	60.0	<b>52.9</b>	<b>0.313</b>	0.209	0.095	<b>0.159</b>	14.0	<b>1.297</b>
	Hypogene	101.9	325.2	<b>223.3</b>	<b>0.368</b>	0.368	0.030	<b>0.050</b>	6.5	<b>0.691</b>
155	Supergene	17.4	111.9	<b>94.5</b>	<b>0.386</b>	0.386	0.012	<b>0.020</b>	3.2	<b>0.518</b>
	Hypogene	218.5	340.5	<b>122.0</b>	<b>0.212</b>	0.212	0.076	<b>0.126</b>	5.6	<b>0.957</b>
156	LeachCap	6.7	72.2	<b>65.5</b>	<b>0.048</b>	0.041	0.083	<b>0.138</b>	4.9	<b>0.852</b>
	Supergene	72.2	152.1	<b>79.9</b>	<b>0.442</b>	0.404	0.058	<b>0.096</b>	6.2	<b>1.023</b>
	Hypogene	152.1	346.6	<b>194.5</b>	<b>0.313</b>	0.313	0.065	<b>0.109</b>	4.7	<b>0.954</b>
	*entire hole	Combined	6.7	including 428.9	<b>422.2</b>	<b>0.276</b>	0.268	0.063	<b>0.105</b>	4.7
157	Supergene	27.7	124.1	<b>96.4</b>	<b>0.254</b>	0.233	0.056	<b>0.093</b>	3.0	<b>0.796</b>
	Hypogene	182.0	407.5	<b>225.5</b>	<b>0.278</b>	0.278	0.049	<b>0.082</b>	3.3	<b>0.761</b>
			including	<b>60.9</b>	<b>0.261</b>	0.261	0.074	<b>0.123</b>	3.2	<b>0.969</b>
			including	<b>186.0</b>	<b>0.291</b>	0.291	0.044	<b>0.073</b>	3.4	<b>0.721</b>
158	Supergene	6.1	47.9	<b>41.8</b>	<b>0.347</b>	0.306	0.045	<b>0.074</b>	28.1	<b>0.953</b>
	Combined	6.1	407.5	<b>401.4</b>	<b>0.296</b>	0.285	0.058	<b>0.096</b>	7.2	<b>0.884</b>
			including	<b>76.2</b>	<b>0.390</b>	0.380	0.080	<b>0.134</b>	7.3	<b>1.190</b>
159	Hypogene	80.8	337.4	<b>256.6</b>	<b>0.266</b>	0.266	0.046	<b>0.076</b>	3.8	<b>0.720</b>
			including	<b>54.9</b>	<b>0.230</b>	0.230	0.073	<b>0.122</b>	2.8	<b>0.930</b>
160	Supergene	9.8	47.9	<b>38.1</b>	<b>0.419</b>	0.378	0.037	<b>0.061</b>	9.9	<b>0.829</b>
	Combined	9.8	322.2	<b>312.4</b>	<b>0.268</b>	0.263	0.044	<b>0.073</b>	5.1	<b>0.713</b>
			including	<b>173.7</b>	<b>0.282</b>	0.282	0.056	<b>0.094</b>	4.9	<b>0.842</b>
161	Supergene	23.6	84.4	<b>60.8</b>	<b>0.553</b>	0.497	0.065	<b>0.109</b>	3.1	<b>1.183</b>
	Hypogene	84.4	249.0	<b>164.6</b>	<b>0.321</b>	0.321	0.029	<b>0.048</b>	4.3	<b>0.620</b>
	Combined	23.6	249.0	<b>225.4</b>	<b>0.384</b>	0.368	0.039	<b>0.065</b>	3.9	<b>0.772</b>
162	Supergene	20.4	43.0	<b>22.6</b>	<b>0.318</b>	0.297	0.024	<b>0.039</b>	2.5	<b>0.560</b>
	Hypogene	43.0	377.0	<b>334.0</b>	<b>0.284</b>	0.284	0.059	<b>0.098</b>	26.0	<b>1.010</b>
	Combined	20.4	377.0	<b>356.6</b>	<b>0.286</b>	0.286	0.057	<b>0.095</b>	24.5	<b>0.983</b>
			including	<b>155.5</b>	<b>0.310</b>	0.310	0.076	<b>0.127</b>	26.2	<b>1.201</b>
163	Hypogene	130.1	410.6	<b>280.5</b>	<b>0.192</b>	0.192	0.043	<b>0.072</b>	3.8	<b>0.619</b>
			including	<b>137.2</b>	<b>0.283</b>	0.283	0.021	<b>0.035</b>	6.3	<b>0.520</b>
			including	<b>143.3</b>	<b>0.105</b>	0.105	0.064	<b>0.107</b>	1.4	<b>0.715</b>
164	Supergene	17.4	75.3	<b>57.9</b>	<b>0.304</b>	0.285	0.032	<b>0.054</b>	3.8	<b>0.632</b>
	Hypogene	75.3	157.0	<b>81.7</b>	<b>0.230</b>	0.230	0.053	<b>0.089</b>	2.3	<b>0.742</b>
	Combined	17.4	157.0	<b>139.6</b>	<b>0.261</b>	0.261	0.045	<b>0.074</b>	2.9	<b>0.697</b>

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Drill Hole (BRG07-)	Mineral Horizon	From (m)	To (m)	Length (m)	Cu (%)	Cu-S (%)	Mo (%)	MoS <sub>2</sub> (%)	Ag (g/t)	CuEQ (%)
165	Supergene	6.1	78.4	<b>72.3</b>	<b>0.237</b>	0.209	0.027	<b>0.045</b>	1.8	<b>0.500</b>
	Hypogene	78.4	500.2	<b>421.8</b>	<b>0.250</b>	0.250	0.047	<b>0.078</b>	3.8	<b>0.710</b>
			including	<b>103.7</b>	<b>0.234</b>	0.234	0.071	<b>0.118</b>	3.7	<b>0.921</b>
*entire hole	Combined	6.1	500.2	<b>494.1</b>	<b>0.248</b>	0.248	0.044	<b>0.073</b>	3.5	<b>0.683</b>
166	Supergene	125.8	137.7	<b>11.9</b>	<b>0.524</b>	0.501	0.033	<b>0.054</b>	5.2	<b>0.863</b>
	Hypogene	137.7	246.0	<b>108.3</b>	<b>0.300</b>	0.300	0.039	<b>0.065</b>	7.9	<b>0.720</b>
	Combined	125.8	246.0	<b>120.2</b>	<b>0.322</b>	0.322	0.039	<b>0.064</b>	7.6	<b>0.734</b>

\*Cu-S (copper sulphide) represents the percentage of copper attributed to sulphide mineralogy. Copper sulphide is calculated with the formula 'Cu-S' = 'Total Copper' - 'Non-sulphide Copper'. Non-sulphide copper analysis is completed on samples collected from the supergene horizon of the Berg deposit using ALS Chemex method code 'Cu-AA05'. MoS<sub>2</sub> values are provided for comparative purposes and are calculated by multiplying the Mo% assay by 1.6681. The copper equivalent (CuEQ%) calculation uses metal prices of US\$1.50/lb copper, US\$14/lb molybdenum, and US\$7/oz silver and is based on in-situ metal content.