

Appendix E: EL36/2003 Drill Core Assays

[illegible]

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H1000	Hole	From	To	Interval	Sample	Batch	Fe	S	Si	Al	Ca	Mg	Na	K	Ti	Mn	WO3	P	LOI	As	Co	Cr	Cu	Pb	U	V	Zn
H1001		m	m	m			%	%	%	%	%	%	%	%	%	%	%	%	%	ppm	ppm	ppm	ppm	ppm	ppm	ppm	ppm
D	DC002	175.3	177.5	2.2	DCD1045	AD10049743	6.92	0.763	16.15	4.67	4.03	14.7	0.562	0.581	0.588	0.1555	0.0038	0.0481	11.9	30	30	-7	-10	-10	-10	223	70
D	DC003	41.2	43	1.8	DCD1016	AD10040437	26.5	0.304	19.45	2.83	1.67	1.975	0.08	0.098	0.258	0.556	0.0126	0.0586	6.82	-10	340	-7	950	-10	-10	35	980
D	DC003	43	44.1	1.1	DCD1017	AD10040437	14.15	0.639	18.15	3.29	2.39	12.6	0.119	0.0245	0.416	0.1275	-0.0013	0.0561	7.58	-10	40	-7	170	-10	-10	130	90
D	DC003	44.1	45.5	1.4	DCD1018	AD10040437	9.73	0.814	19.4	3.7	0.917	15.35	0.1405	0.0161	0.373	0.0997	-0.0013	0.0525	7.62	-10	50	-7	140	-10	-10	108	50
D	DC003	45.5	48.4	2.9	DCD1019	AD10040437	10.05	0.816	19.25	4.57	1.335	14	0.637	0.077	0.353	0.0815	-0.0013	0.0434	6.75	-10	40	-7	120	-10	10	91	20
D	DC003	48.4	50.6	2.2	DCD1020	AD10040437	10.75	1.15	19.05	3.55	1.15	14.9	0.169	0.0182	0.378	0.0859	-0.0013	0.0828	6.68	-10	60	-7	240	-10	-10	106	60
D	DC003	50.6	52.3	1.7	DCD1021	AD10040437	10.35	1.045	18.75	3.98	2.38	13.6	0.519	0.0307	0.351	0.102	-0.0013	0.0472	7.44	-10	50	-7	250	-10	-10	113	10
D	DC003	52.3	55.8	3.5	DCD1022	AD10040437	8.38	1.275	17.45	4.72	1.56	15.65	0.222	0.0237	0.44	0.082	-0.0013	0.048	9.04	-10	50	-7	190	-10	10	122	-10
D	DC003	55.8	58	2.2	DCD1023	AD10040437	5.32	0.68	18.4	5.8	2.11	16	0.532	0.207	0.647	0.0879	-0.0013	0.0387	8.58	-10	30	-7	30	-10	10	157	10
D	DC003	58	60.6	2.6	DCD1024	AD10040437	4.02	0.956	24.6	8.1	1.025	6.45	4.27	0.868	0.548	0.0314	-0.0013	0.0567	4	-10	-10	-7	-10	-10	20	128	-10
D	DC003	60.6	62.6	2	DCD1025	AD10040437	5.78	2.51	24.1	7.32	0.813	4	4.21	0.719	0.523	0.0174	-0.0013	0.0479	4.75	-10	40	-7	30	-10	10	150	-10
D	DC003	62.6	64	1.4	DCD1026	AD10040437	3.31	0.732	21.3	7.57	0.64	12.35	2.42	0.167	0.513	0.0291	-0.0013	0.044	7.55	-10	30	-7	-10	-10	20	127	-10
D	DC003	64	65.5	1.5	DCD1027	AD10040437	3.22	0.927	15.7	5.51	2.78	18.5	0.153	0.0582	0.384	0.0761	-0.0013	0.035	13.3	-10	10	-7	-10	-10	20	83	-10
D	DC003	65.5	67.2	1.7	DCD1028	AD10040437	3.69	0.429	2.59	0.517	1.825	30.2	0.23	0.0091	0.066	0.041	-0.0013	0.0093	27.6	-10	-10	-7	60	-10	20	8	110
D	DC003	67.2	69.7	2.5	DCD1029	AD10040437	7.35	0.684	8.9	1.775	6.11	20.5	0.204	0.0164	0.12	0.0625	-0.0013	0.022	22.2	-10	-10	-7	160	-10	20	17	-10
D	DC003	69.7	71.5	1.8	DCD1030	AD10040437	8.03	1.345	13.1	3.06	5.4	16.45	0.0776	0.0577	0.314	0.155	-0.0013	0.0398	15.45	-10	10	-7	290	-10	10	95	30
D	DC003	71.5	73.8	2.3	DCD1031	AD10040437	4.95	0.963	23.7	7.66	1.2	6.73	3.76	0.878	0.553	0.0392	-0.0013	0.056	4.98	-10	30	-7	20	-10	-10	166	-10
D	RRDDH2	270	272	2	RRH1008	AD09133670	17.9	1.925	20.1	5.17	1.735	2.6	1.945	0.0355	1.065	0.189	-0.0013	0.175	4.67	10	80	-7	90	-10	-10	222	80
D	RRDDH2	272	274	2	RRH1009	AD09133670	16.7	0.788	21.4	5.95	1.675	2.88	2.76	0.0237	1.075	0.24	-0.0013	0.0875	3.85	-10	40	-7	80	-10	-10	365	80
D	RRDDH2	274	276	2	RRH1010	AD09133670	17.3	0.991	20.2	6.1	1.86	2.96	2.88	0.0202	1.065	0.295	-0.0013	0.0766	4.14	10	30	-7	60	-10	-10	359	80
D	RRDDH2	276	278	2	RRH1011	AD09133670	16.55	0.877	20.4	5.67	2.53	2.93	2.64	0.0367	1	0.415	-0.0013	0.0814	5.09	-10	30	-7	60	-10	-10	330	100
D	RRDDH2	278	280	2	RRH1012	AD09133670	18.3	2.03	19.15	5.62	1.83	1.9	3.11	0.0514	1.11	0.321	-0.0013	0.158	4.39	10	110	-7	70	10	-10	221	80
D	RRDDH2	280	282	2	RRH1013	AD09133670	16.4	2.33	20.4	5.1	2.63	1.28	2.96	0.0779	1.175	0.391	-0.0013	0.22	4.47	40	210	-7	70	-10	-10	112	70
D	RRDDH2	282	284	2	RRH1014	AD09133670	15.7	2.01	19.65	5.53	2.64	1.945	2.89	0.0803	1.145	0.652	-0.0013	0.237	5.75	20	100	-7	130	50	-10	139	160
D	RRDDH2	284	286	2	RRH1015	AD09133670	16.7	1.555	21.5	5.4	2.04	1.43	3.42	0.0787	1.11	0.379	-0.0013	0.191	3.14	10	80	-7	50	-10	-10	136	50
D	RRDDH2	314	316	2	RRH1001	AD09133670	18.65	0.44	21.2	5.4	1.485	3.01	1.205	0.567	1.125	0.425	-0.0013	0.115	4.4	30	30	-7	170	-10	-10	269	130
D	RRDDH2	316	318	2	RRH1002	AD09133670	24.6	1.06	20.3	3.76	1.215	2.09	0.604	0.149	0.919	0.296	0.0063	0.1015	3.07	10	30	-7	480	10	-10	207	110
D	RRDDH2	318	320	2	RRH1003	AD09133670	40.8	3.8	12.65	0.347	1.645	0.392	0.0547	0.0517	0.05	0.148	0.0303	0.0189	0.76	40	40	-7	1360	30	-10	20	60
D	RRDDH2	320	322	2	RRH1004	AD09133670	44.1	>6	10.9	1.295	1.13	1	0.129	0.0359	0.18	0.159	0.0214	0.0214	7.06	40	60	-7	730	30	-10	67	100
D	RRDDH2	322	324	2	RRH1005	AD09133670	22.9	2.39	19.65	4.37	0.706	2.18	0.294	0.213	0.695	0.245	0.0063	0.0684	3.99	20	30	-7	600	-10	-10	202	100
D	RRDDH2	324	326	2	RRH1006	AD09133670	19.8	1.2	21.8	5.23	0.855	2.24	1.07	0.1735	0.956	0.345	-0.0013	0.0775	3.14	20	20	-7	220	-10	-10	296	100
D	RRDDH2	326	328	2	RRH1007	AD09133670	16.3	1.45	24.2	4.94	0.754	1.785	1.985	0.0692	0.968	0.123	-0.0013	0.251	2.68	10	20	-7	270	-10	-10	98	40
D	RRDDH3	68	69	1	BASS147890	PH09127895	4.31	1.47	29.3	5.54	1.705	3.55	3.73	0.49	0.439	0.0639	-0.0013	0.0272	1.94	263	20	-7	50	440	-10	116	310
D	RRDDH3	69	70.8	1.8	BASS147891	PH09127895	8.1	0.669	19.6	5.98	2.44	11.35	1.515	0.449	0.848	0.113	-0.0013	0.0551	6.84	278	10	-7	-10	110	-10	337	250
D	RRDDH3	71	72	1	BASS147893	PH09127895	7.82	1.085	20	7.02	3.67	6.78	3.03	0.609	0.926	0.1195	-0.0013	0.0561	6.94	357	40	-7	240	-10	-10	383	40
D	RRDDH3	72	72.3	0.3	BASS147894	PH09127895	12.8	0.0908	16.85	1.115	10.45	9.37	0.0534	0.0175	0.158	0.224	-0.0013	0.0164	12.5	351	10	-7	160	-10	-10	79	-10
D	RRDDH3	72.3	73	0.7	BASS147895	PH09127895	9.17	0.789	21	6.34	4.13	5.97	3.14	0.723	0.945	0.173	-0.0013	0.0576	5.27	334	40	-7	140	20	-10	392	30
D	RRDDH3	73	73.8	0.8	BASS147896	PH09127895	13.5	2.3	17.95	4.38	3.78	8.47	1.45	0.235	0.655	0.178	-0.0013	0.0473	5.26	60	40	-7	80	-10	-10	282	240
D	RRDDH3	73.8	74	0.2	BASS147897	PH09127895	10.35	1.93	20.5	6.1	3.28	5.89	3	0.1775	0.933	0.1545	-0.0013	0.0593	4.38	30	50	-7	50	-10	-10	386	180
D	RRDDH3	74	75	1	BASS147898	PH09127895	9.14	2.01	20.4	5.98	4.48	5.81	3.21	0.199	0.899	0.153	-0.0013	0.0612	4.56	29	80	-7	10	-10	-10	388	20
D	RRDDH3	75	75.4	0.4	BASS147899	PH09127895	10.05	0.1945	22.5	6.55	3.15	5.67	3.84	0.1555	0.982	0.0912	-0.0013	0.0626	3.31	56	10	-7	-10	-10	-10	425	-10
D	RRDDH3	75.4	76	0.6	BASS147900	PH09127895	51.9	4.27	2.86	0.509	0.148	2.18	0.0504	-0.0008	0.06	0.0259	0.0441	0.017	3.28	377	340	-7	730	-10	10	2380	40
D	RRDDH3	76	76.8	0.8	BASS147901	PH09127895	54.2	>6	2.52	0.471	0.731	2.71	0.0393	-0.0008	0.09	0.05	0.0252	0.0436	4.93	461	420	102	990	-10	-10	2530	50
D	RRDDH3	76.8	77	0.2	BASS147902	PH09127895	55.3	3.11	2.32	0.451	0.294	2.51	0.018	-0.0008	0.101	0.0307	0.0479	0.0767	1.41	446	280	-7	580	-10	-10	2730	90
D	RRDDH3	77	78	1	BASS147903	PH09127895	58.5	5.65	2.07	0.445	0.478	2.42	0.0134	-0.0008	0.126	0.0326	0.0429	0.0567	3.77	548	410	-7	940	-10	-10	2820	90
D	RRDDH3	78	79	1	BASS147904	PH09127895	49.4	3.32	3.63	0.441	0.753	4.33	0.0077	-0.0008	0.281	0.0528	0.0353	0.0498	2.89	466	220	-7	560	-10	-10	2460	70
D	RRDDH3	79	79.8	0.8	BASS147905	PH09127895	15.8	3.17	14.45	0.247	0.488	16.3	0.0104	-0.0008	0.022	0.036	-0.0013	0.0162	10	313	110	-7	370	-10	-10	471	20
D	RRDDH3	80	81	1	BASS147907	PH09127895	31.9	>6	10.45	1.995	2.25	5.27	1.155	0.0415	0.282	0.0459											

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H1001		m	m	m			%	%	%	%	%	%	%	%	%	%	%	%	%	ppm	ppm	ppm	ppm	ppm	ppm	ppm	ppm
D	RRDDH3	245	246	1	BASS147914	PH09127895	5.62	0.0473	22.8	6.34	7.94	3.83	2.59	0.487	0.607	0.1255	-0.0013	0.0149	8.32	157	10	-7	-10	-10	-10	172	-10
D	RRDDH3	246	246.5	0.5	BASS147915	PH09127895	6.39	0.519	23.5	6.67	5.85	3.68	2.87	0.529	0.807	0.103	-0.0013	0.1315	6.27	236	90	7	-10	-10	-10	217	-10
D	RRDDH3	246.5	247	0.5	BASS147916	PH09127895	7.22	1.41	21	3.9	9.33	3.32	0.638	0.1945	0.736	0.14	-0.0013	0.274	12.2	195	110	-7	100	60	-10	273	-10
D	RRDDH3	247	247.5	0.5	BASS147917	PH09127895	7.02	0.0467	24.2	8.16	2.26	3.24	2.95	2.01	0.993	0.0605	-0.0013	0.0517	5.76	111	20	-7	-10	-10	-10	326	10
D	RRDDH3	247.5	247.6	0.1	BASS147918	PH09127895	11.35	0.254	20.5	4.86	7.72	2.64	2.29	0.288	0.468	0.118	-0.0013	0.0354	10.45	52	10	-7	-10	-10	-10	249	-10
D	RRDDH3	247.6	248	0.4	BASS147919	PH09127895	5.83	0.236	23	6.34	6.41	3.19	3.68	0.187	0.751	0.0871	-0.0013	0.0165	8.93	138	20	-7	-10	10	-10	189	-10
D	RRDDH3	248	249	1	BASS147920	PH09127895	7.59	0.1405	22.6	6.28	5.25	3.37	3.18	0.442	0.679	0.0821	-0.0013	0.0426	9.43	208	30	-7	-10	-10	-10	245	20
D	RRDDH3	249	250	1	BASS147921	PH09127895	5.02	0.1565	23	6.39	6.6	3.15	3.68	0.309	0.857	0.0697	-0.0013	0.0095	9.79	142	30	-7	-10	-10	-10	161	-10
D	RRDDH3	250	250.6	0.6	BASS147922	PH09127895	9.94	0.446	20.4	5.47	6.74	4.41	2.04	0.299	0.687	0.0999	-0.0013	0.0462	9.52	129	50	-7	10	30	-10	443	10
D	RRDDH3	250.6	251	0.4	BASS147923	PH09127895	12	0.47	20.5	5.81	4.31	4.47	1.965	0.1375	0.979	0.0856	-0.0013	0.0905	8.36	109	40	-7	20	-10	-10	500	10
D	RRDDH3	251	251.2	0.2	BASS147924	PH09127895	12.45	0.106	23.4	6.95	3.1	5.37	2.58	0.0552	0.925	0.0711	-0.0013	0.0544	-0.5	223	-10	-8	-10	-10	-10	463	-10
D	RRDDH3	251.2	251.6	0.4	BASS147925	PH09127895	35	>6	9.19	1.005	6.29	2.1	0.0408	0.0066	0.546	0.0613	-0.0013	0.162	4.92	411	390	-7	1010	10	-10	1400	-10
D	RRDDH3	251.7	252	0.3	BASS147927	PH09127895	16.35	2.61	17.2	4.92	4.11	5.23	1.215	0.0572	0.849	0.0741	-0.0013	0.0231	6.08	149	130	-7	380	10	-10	624	10
D	RRDDH3	252	252.7	0.7	BASS147928	PH09127895	14.5	1.58	18.8	5.44	3.16	5.06	1.565	0.0392	1.89	0.116	-0.0013	0.0804	6.19	209	70	-7	110	-10	-10	588	-10
D	RRDDH3	252.7	253	0.3	BASS147929	PH09127895	6.54	1.76	21.9	8.19	4.5	2.19	4.92	0.719	0.269	0.048	-0.0013	0.728	4.33	157	60	-7	40	-10	-10	213	-10
D	RRDDH3	253	253.6	0.6	BASS147930	PH09127895	14.5	1.455	15.5	4.26	8.98	4.02	1.25	0.0417	0.855	0.16	-0.0013	0.356	10.75	279	90	-7	150	40	-10	834	10
D	RRDDH3	253.6	253.7	0.1	BASS147931	PH09127895	28.5	1.41	12.7	3.72	4.68	3.36	0.818	0.139	0.861	0.107	-0.0013	0.277	5.4	390	50	-7	50	-10	-10	1930	-10
D	RRDDH3	253.7	254	0.3	BASS147932	PH09127895	18.7	3.93	8.51	1.23	16.55	2.27	0.012	-0.0008	0.611	0.203	-0.0013	0.65	12.9	289	230	-7	330	-10	-10	1290	-10
D	RRDDH3	254	255	1	BASS147933	PH09127895	12.25	0.48	21.2	5.97	4.4	4.47	2.18	0.0938	1.115	0.12	-0.0013	0.0984	5.6	242	50	-7	-10	-10	-10	553	10
EOF																											