



# ASX ANNOUNCEMENT

16 January 2023



## REE Discovery at Hines Hill REE Project

### HIGHLIGHTS

- ❖ Maiden aircore drilling program of 49 drill holes has intersected zones up to 1,602ppm TREO (from 8metres in drillhole HHAC38), with mineralisation open in all directions
- ❖ Significant results to date include:
  - **25m @ 837ppm TREO** (173ppm MREO) from 6m, including **3m @1602ppm TREO** from 6m (HHC038)
  - **36m @ 639ppm TREO** (131ppm MREO) from surface, including **3m @1126ppm TREO** from 12m (HHC037)
  - **16m @ 693ppm TREO** (inc 144ppm MREO) from 24m (HHC008)
  - **12m @ 637ppm TREO** (inc 148ppm MREO) from 42m (HHC012)
  - **13m @ 604ppm TREO** (inc 111ppm MREO) from 36m, including **4m @ 1182ppm TREO** from 45m (HHC029)
- ❖ Addition of contiguous tenement E70/6136 grows the Hines Hill project area to ~576km<sup>2</sup>
- ❖ Geochemical roadside sampling program completed over the remainder of Hines Hill, and E70/6136, are due shortly
- ❖ Further drilling is being planned for the quarter, with a full review of technical data to include upcoming results from the geochemical sampling program, and assaying of 1metre samples underway

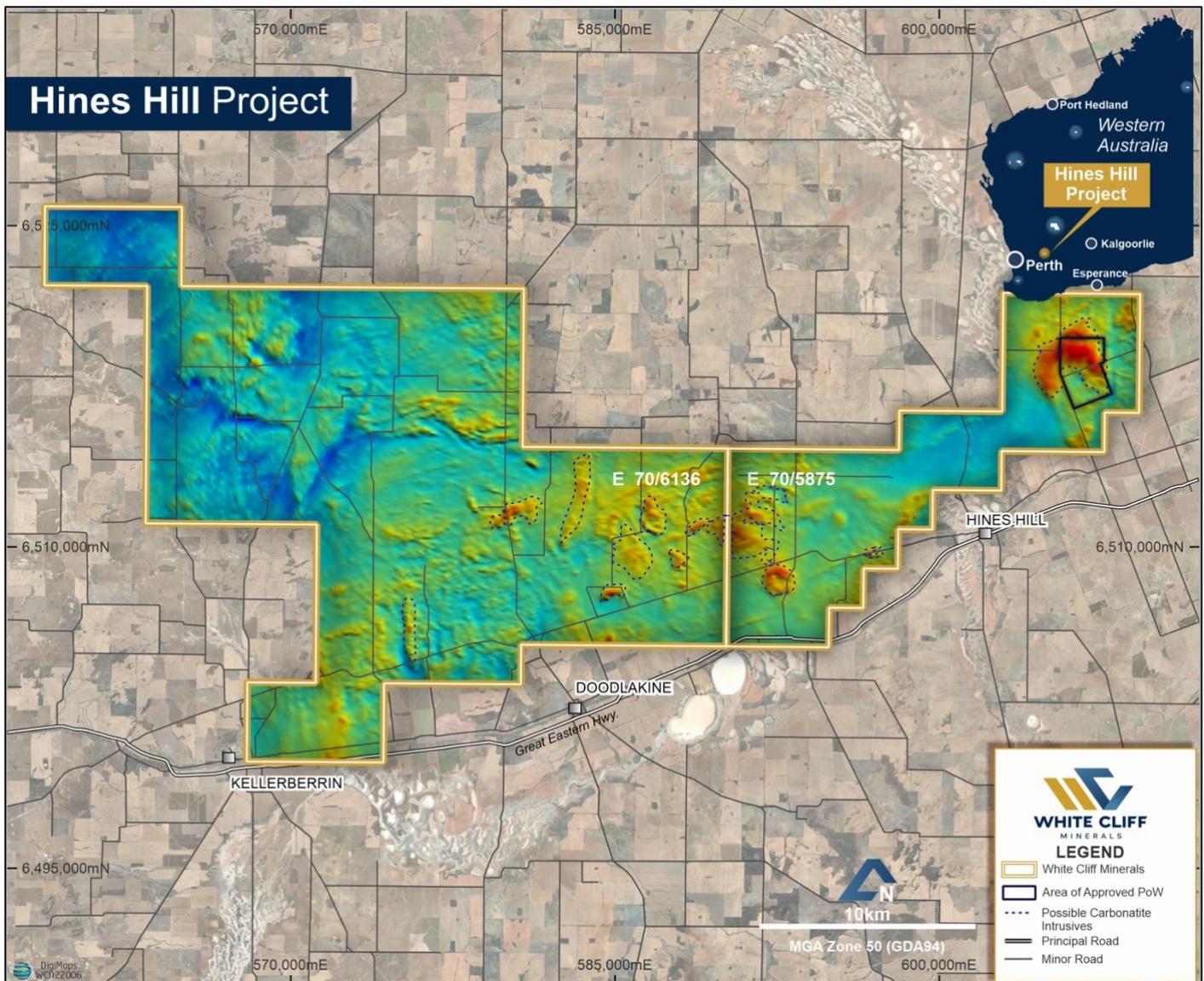
White Cliff Minerals Limited (**White Cliff** or the **Company**) is pleased to provide an update on the Company's Hines Hill REE project in the Wheatbelt region, WA (**Figure 1**), where maiden air core drilling has discovered mineralised REE clays from surface. The results are extremely positive (**Figure 2**) and further geochemical sampling over the remaining magnetic features has also been completed, to grow the project potential. The Company has also undertaken a maiden geochemical sampling program over E70/6136 an addition to the Hines Hill project area which is contiguous to the west and increases the project size to ~576km<sup>2</sup>.

Commenting on the progress, White Cliff Technical Director Ed Mead said:

"Our maiden drill program at Hines Hill has been extremely successful in demonstrating the potential of the project with the discovery of shallow mineralisation with intercepts of up to 1,602ppm TREO in clays from surface down through the oxidisation profile, and into the basement. It appears at this early stage that the granitoid basement is highly elevated in REE's and the REE Clays are the weathered granitoid. The Hines Hill project sits on salt affected broad acre farmland, which gives us the ability to work all year round; and with a land access agreement with the landowner, further encouraging results bodes well for progression of the project. We have also increased the scale of the project to ~576km<sup>2</sup> "

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**Figure 1:** Hines Hill REE Project location map in Western Australia, with the addition of tenement E70/6136(136 blocks) to original tenement E70/5875 (44 blocks increasing the project to ~576km<sup>2</sup> .

## Hines Hill - REE Project

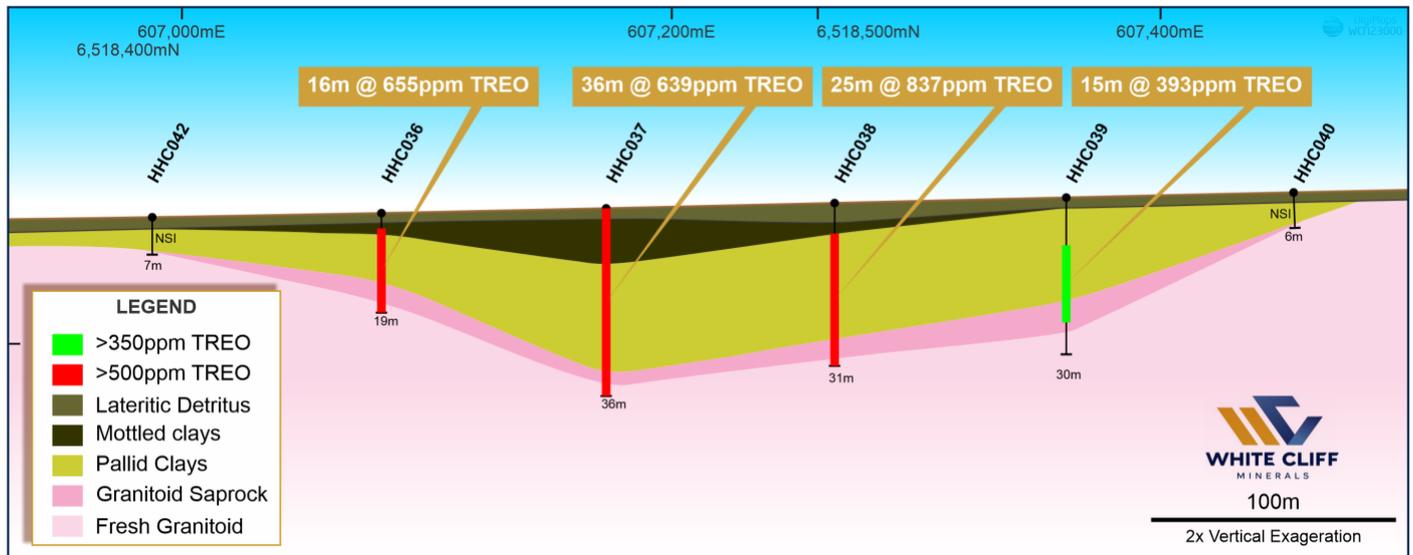
The Hines Hill REE project consists of two tenements (**Figure 1**), within the wheatbelt region, located about 200km east of Perth on the Great Eastern Highway. The tenement area of ~576Km<sup>2</sup> covers extensive broad acre grain growing properties.

Geochemical sampling by White Cliff has initially targeted two magnetic features tentatively interpreted to be carbonatite intrusives, although they may represent differential non-carbonatite intrusives (granitoids).

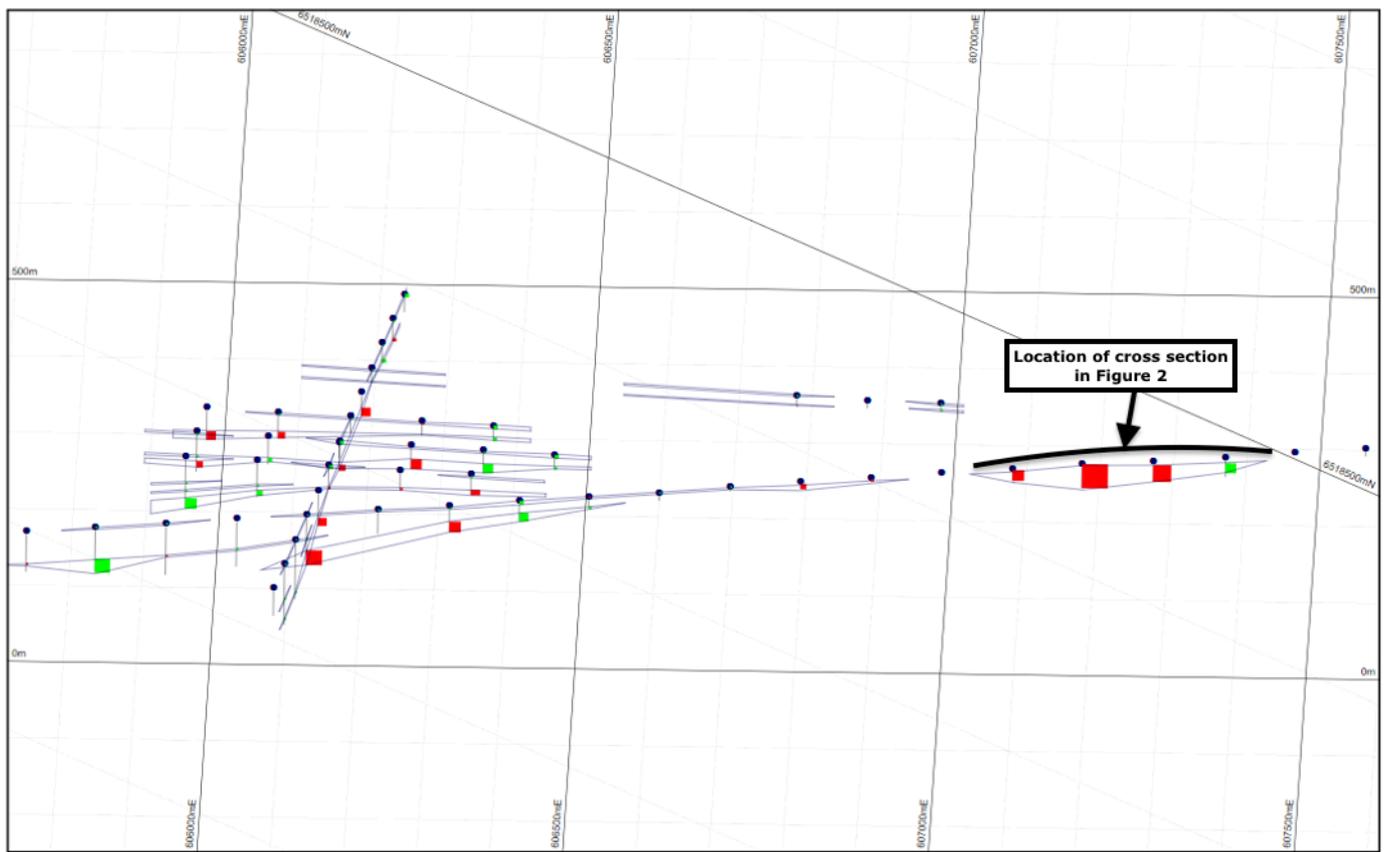
Drilling of the most north-eastern magnetic feature (**Figures 1 and 4**), where an approved POW allows drilling has intersected laterite, clays and basement geology (granitoid) with REE assays (**Figures 2 and 3**). The mineralised profile suggests a granitoid with high REE background is weathering and the resulting clays have equivalent or higher REE content due to reduction associated with weathering. Further drilling of the basement rocks will confirm the current interpretation of granitoid basement geology.

The maiden aircore drilling program totalled 49 drillholes for 1,861 metres (**Table 2**), ranging in depth from 8 metres to 92 metres. 716 (3 metre composite samples) were taken.

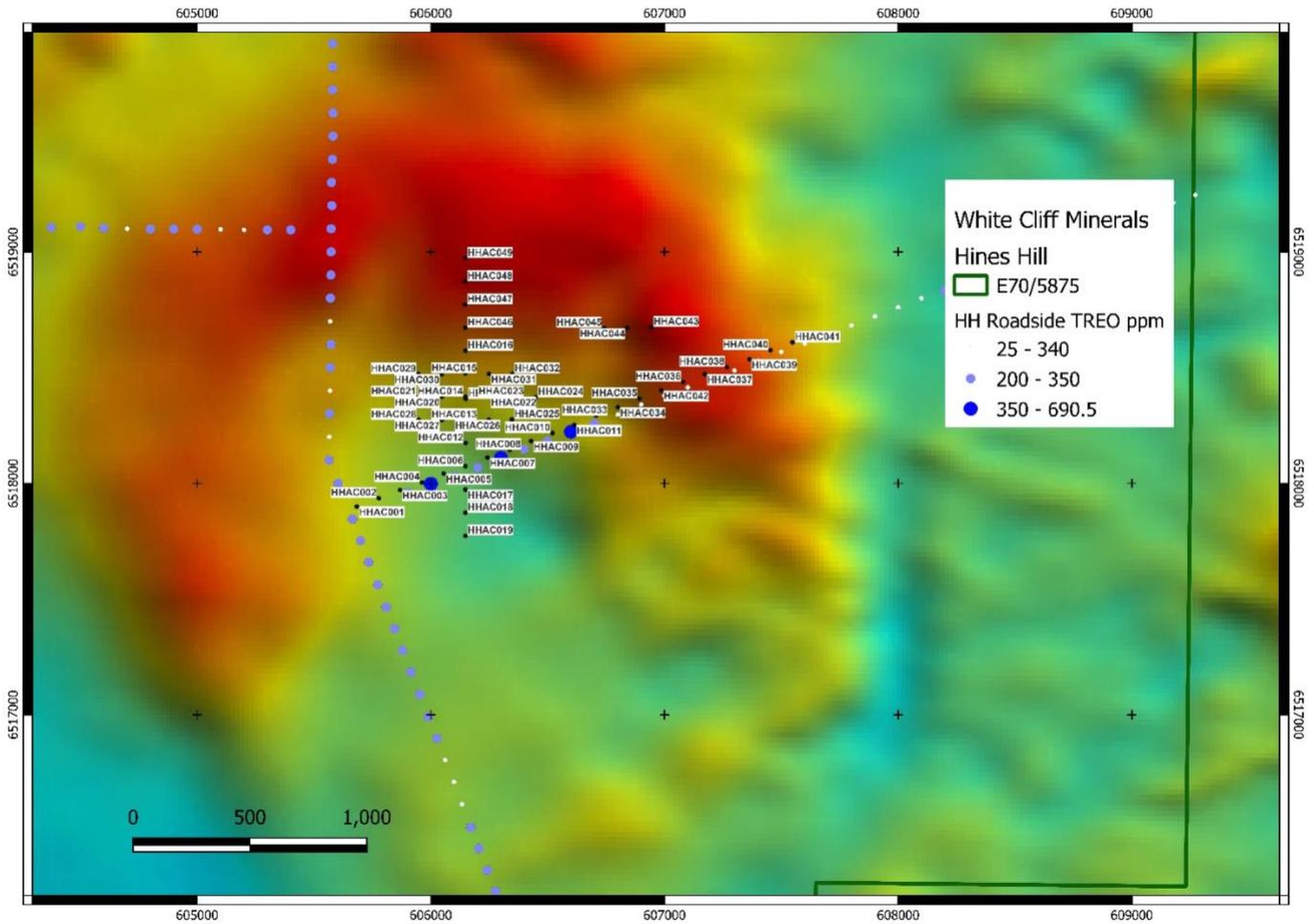
The results of the drilling (**Tables 1 and 3**) have returned positive intercepts for REE's with multiple zones of mineralisation intersected. The best and thickest zones of mineralisation appear to be associated with magnetic highs.



**Figure 2:** Hines Hill schematic drill section looking north. Refer Figure 3 for location of section.



**Figure 3:** Hines Hill drill hole locations and REE intercept schematic looking north. Red >500ppm TREO, Green < 500ppm TREO.



**Figure 4:** Hines Hill drill hole location plan from Aircore drill program.

Further roadside conventional geochemical sampling was completed on E70/5875, with 231 samples on a nominal 200m spacing.

ELA 70/6136 was applied for on 18 May 2022 and is contiguous to the west of Hines Hill. The application covers 136 blocks (~400km<sup>2</sup>) and is prospective for REE.

A first pass roadside conventional geochemical sampling program has been completed, with 813 samples on a nominal 200m spacing. These results are due shortly.

## Next Steps

The Company will review all technical data, that includes the geochemical samples which are due shortly, to determine the next suitable program which may include work to delineate a maiden resource on the magnetic feature area where drill holes HHC036 to 39 intersected wide zones of shallow REE mineralisation. These drill holes will have the 1 metre samples sent to ALS Laboratories for assay.

**Table 1: Significant drill intercepts from 3 metre composite samples**

Hole ID	From	To	Width	TREO ppm	MREO ppm	Intercept
HHC002	48	51	3	714	129	3m @ 714ppm TREO
HHC003	0	3	3	432	90	3m @ 432ppm TREO
HHC003	48	69	21	389	75	21m @ 389ppm TREO
HHC004	0	3	3	428	95	3m @ 428ppm TREO
HHC004	48	51	3	673	94	3m @ 673ppm TREO
HHC005	45	48	3	390	69	3m @ 390ppm TREO
HHC006	0	3	3	393	69	3m @ 393ppm TREO
HHC006	36	39	3	375	78	3m @ 375ppm TREO
HHC006	54	76	22	577	118	22m @ 577ppm TREO
HHC007	0	3	3	385	82	3m @ 385ppm TREO
HHC008	3	6	3	391	86	3m @ 391ppm TREO
HHC008	24	40	16	693	144	16m @ 693ppm TREO
HHC009	0	6	6	380	79	6m @ 380ppm TREO
HHC009	18	31	13	496	111	13m @ 496ppm TREO
HHC010	0	3	3	643	148	3m @ 643ppm TREO
HHC010	15	19	4	388	79	4m @ 388ppm TREO
HHC011	0	3	3	350	81	3m @ 350ppm TREO
HHC012	42	54	12	637	148	12m @ 637ppm TREO
HHC013	0	6	6	441	111	6m @ 441ppm TREO
HHC013	33	36	3	573	130	3m @ 573ppm TREO
HHC014	0	6	6	375	85	3m @ 375ppm TREO
HHC014A	33	42	9	503	107	9m @ 503ppm TREO
HHC015	3	6	3	357	82	3m @ 357ppm TREO
HHC016	24	37	13	534	120	13m @ 534ppm TREO
HHC017	0	3	3	480	104	3m @ 480ppm TREO
HHC017	78	81	3	490	98	3m @ 490ppm TREO
HHC018	0	3	3	409	96	3m @ 409ppm TREO
HHC018	51	54	3	486	89	3m @ 486ppm TREO
HHC018	81	84	3	386	77	3m @ 386ppm TREO
HHC020	33	39	6	481	108	6m @ 481ppm TREO
HHC021	3	6	3	408	98	3m @ 408ppm TREO
HHC021	36	39	3	412	93	3m @ 412ppm TREO
HHC021	45	54	9	655	127	9m @ 655ppm TREO
HHC022	3	6	3	349	85	3m @ 349ppm TREO
HHC022	21	36	15	518	110	15m @ 518ppm TREO
HHC023	3	6	3	427	107	3m @ 427ppm TREO
HHC023	21	35	14	492	99	14m @ 492ppm TREO
HHC024	0	6	6	350	79	6m @ 350ppm TREO
HHC024	18	22	4	459	98	4m @ 459ppm TREO
HHC025	3	6	3	473	103	3m @ 473ppm TREO
HHC025	24	37	13	587	131	13m @ 587ppm TREO
HHC026	27	31	4	600	125	4m @ 600ppm TREO
HHC027	36	39	3	372	93	3m @ 372ppm TREO
HHC027	45	53	8	472	100	8m @ 472ppm TREO
HHC028	39	42	3	436	90	3m @ 436ppm TREO
HHC028	51	54	3	472	96	3m @ 472ppm TREO
HHC028	63	79	16	441	100	16m @ 441ppm TREO
HHC029	36	49	13	604	111	13m @ 604ppm TREO
						Incl 4m @ 1182ppm TREO from 45m
HHC030	3	6	3	463	103	3m @ 463ppm TREO
HHC030	30	40	10	726	161	10m @ 726ppm TREO
						Incl 3m @ 1045ppm TREO from 30m
HHC031	3	6	3	621	183	3m @ 621ppm TREO
HHC031	21	22	1	471	117	1m @ 471ppm TREO
HHC032	0	6	6	395	89	6m @ 395ppm TREO
HHC032	18	23	5	398	81	5m @ 398ppm TREO
HHC033	0	3	3	478	98	3m @ 478ppm TREO

Hole ID	From	To	Width	TREO ppm	MREO ppm	Intercept
HHC034	3	11	8	642	149	8m @ 642ppm TREO
HHC035	3	7	4	855	205	4m @ 855ppm TREO
HHC036	3	19	16	655	143	16m @ 655ppm TREO
HHC037	0	36	36	639	131	36m @ 639ppm TREO
						Incl 3m @ 1126ppm TREO from 12m
HHC038	6	31	25	837	173	25m @ 837ppm TREO
						Incl 3m @ 1602ppm TREO from 6m
HHC039	9	24	15	393	75	15m @ 393ppm TREO
HHC043	0	3	3	364	72	3m @ 364ppm TREO
HHC043	9	12	3	390	81	3m @ 390ppm TREO
HHC045	0	3	3	400	82	3m @ 400ppm TREO
HHC045	15	17	2	612	126	2m @ 612ppm TREO
HHC046	3	6	3	395	88	3m @ 395ppm TREO
HHC046	21	23	2	478	118	2m @ 478ppm TREO
HHC047	3	6	3	442	109	3m @ 442ppm TREO
HHC047	24	30	6	393	78	6m @ 393ppm TREO
HHC048	3	6	3	418	105	3m @ 418ppm TREO
HHC048	30	35	5	701	142	5m @ 701ppm TREO
HHC049	0	6	6	357	79	6m @ 357ppm TREO

## UPCOMING NEWSFLOW

- January:** Hines Hill geochemical sampling results
- January:** High-resolution magnetics/radiometrics survey at Yinnetharra. Preliminary targets provided by SGC
- January:** Completion of AEM interpretation by SGC for Yinnetharra, Diemals, Hines Hill and North Kellerberrin, and Lake Tay
- January:** High-resolution magnetics/radiometrics survey at Yinnetharra, Final interpretation and targets provided by SGC
- January:** Diemals geochemical sampling results
- February:** Reconnaissance field trip at Lake Tay REE project
- February:** Planning for further drill program at Hines Hill

**ENDS**

## Further Information:

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*This announcement has been approved by the Board of White Cliff Minerals Limited.*

### **Competent Persons Statement**

The Information in this report that relates to exploration results, mineral resources or ore reserves is based on information compiled by Mr Allan Younger, who is a Member of the Australasian Institute of Mining and Metallurgy. Mr Younger is an employee of the company. Mr Younger has sufficient experience which is relevant to the style of mineralisation and type of deposits under consideration and to the activity that he is undertaking to qualify as a Competent Person as defined in the 2012 edition of the 'Australian Code for Reporting Exploration Results, Mineral Resources and Ore Reserves' (the JORC Code). Mr Younger consents to the inclusion of this information in the form and context in which it appears in this report.

### **Forward looking statements**

This announcement contains certain forward-looking statements and comments about future events, including the Company's expectations about the proposed transaction, the proposed tenements and the performance of its businesses. Forward looking statements can generally be identified by the use of forward-looking words such as 'expect', 'anticipate', 'likely', 'intend', 'should', 'could', 'may', 'predict', 'plan', 'propose', 'will', 'believe', 'forecast', 'estimate', 'target' and other similar expressions within the meaning of securities laws of applicable jurisdictions. Indications of, and guidance on, future earnings or financial position or performance are also forward-looking statements.

Forward looking statements involve inherent risks and uncertainties, both general and specific, and there is a risk that such predictions, forecasts, projections and other forward-looking statements will not be achieved. Forward looking statements are provided as a general guide only and should not be relied on as an indication or guarantee of future performance. Forward looking statements involve known and unknown risks, uncertainty and other factors which can cause the Company's actual results to differ materially from the plans, objectives, expectations, estimates and intentions expressed in such forward-looking statements and many of these factors are outside the control of the Company. As such, undue reliance should not be placed on any forward-looking statement. Past performance is not necessarily a guide to future performance and no representation or warranty is made by any person as to the likelihood of achievement or reasonableness of any forward-looking statements, forecast financial information or other forecast. Nothing contained in this announcement nor any information made available to you is, or shall be relied upon as, a promise, representation, warranty or guarantee as to the past, present or the future performance of the Company.

Except as required by law or the ASX Listing Rules, the Company assumes no obligation to provide any additional or updated information or to update any forward-looking statements, whether as a result of new information, future events or results, or otherwise.

**Table 2: Drilling Collars**

Hole ID	EAST	NORTH	RL	DEPTH M	Dip	Azimuth
HHAC001	605683	6517900	260	58	-90	0
HHAC002	605777	6517936	261	62	-90	0
HHAC003	605868	6517971	262	69	-90	0
HHAC004	605962	6518004	263	78	-90	0
HHAC005	606055	6518042	264	74	-90	0
HHAC006	606148	6518074	265	76	-90	0
HHAC007	606242	6518111	266	37	-90	0
HHAC008	606337	6518145	267	40	-90	0
HHAC009	606429	6518182	268	31	-90	0
HHAC010	606521	6518217	269	19	-90	0
HHAC011	606614	6518252	270	13	-90	0
HHAC012	606149	6518174	265	54	-90	0
HHAC013	606148	6518274	265	36	-90	0
HHAC014	606148	6518374	265	27	-90	0
HHAC014A	606149	6518365	265	42	-90	0
HHAC015	606148	6518474	265	26	-90	0
HHAC016	606148	6518573	265	37	-90	0
HHAC017	606148	6517973	265	90	-90	0
HHAC018	606148	6517873	265	92	-90	0
HHAC019	606148	6517773	265	42	-90	0
HHAC020	606048	6518374	265	43	-90	0
HHAC021	605948	6518374	265	61	-90	0
HHAC022	606248	6518374	265	36	-90	0
HHAC023	606349	6518374	265	35	-90	0
HHAC024	606449	6518372	265	22	-90	0
HHAC025	606347	6518276	265	32	-90	0
HHAC026	606248	6518274	265	31	-90	0
HHAC027	606048	6518274	265	55	-90	0
HHAC028	605948	6518274	265	79	-90	0
HHAC029	605947	6518472	265	49	-90	0
HHAC030	606047	6518472	265	40	-90	0
HHAC031	606248	6518473	265	22	-90	0
HHAC032	606348	6518473	271	23	-90	0
HHAC033	606707	6518293	272	8	-90	0
HHAC034	606800	6518329	273	11	-90	0
HHAC035	606893	6518366	274	7	-90	0
HHAC036	607080	6518438	275	19	-90	0
HHAC037	607172	6518472	276	36	-90	0
HHAC038	607267	6518502	277	31	-90	0
HHAC039	607363	6518536	278	30	-90	0
HHAC040	607454	6518575	279	6	-90	0
HHAC041	607548	6518609	280	13	-90	0
HHAC042	606986	6518402	266	7	-90	0
HHAC043	606943	6518675	266	12	-90	0
HHAC044	606841	6518671	266	13	-90	0
HHAC045	606742	6518671	266	17	-90	0
HHAC046	606147	6518672	266	23	-90	0
HHAC047	606146	6518772	266	30	-90	0
HHAC048	606146	6518872	266	35	-90	0
HHAC049	606147	6518972	266	28	-90	0

**Table 3: Assay Data**

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	PrO <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC001	0	3	74.07	35.54	2.33	1.49	0.85	3.25	0.53	0.18	24.96	7.88	4.58	0.47	0.13	1.31	15.49	173.05	35.64
HHC001	3	6	26.66	16.30	0.98	0.75	0.36	1.13	0.23	0.11	8.75	2.79	1.51	0.16	0.08	0.80	6.48	67.08	12.68
HHC001	6	9	23.83	15.83	0.77	0.56	0.25	0.76	0.16	0.09	6.65	2.21	1.19	0.13	0.08	0.67	4.95	58.15	9.76
HHC001	9	12	26.04	15.25	0.79	0.45	0.23	0.92	0.22	0.14	7.81	2.84	1.15	0.12	0.07	0.68	5.46	62.17	11.56
HHC001	12	15	19.90	12.43	0.63	0.49	0.17	0.68	0.16	0.10	5.72	2.11	1.02	0.13	0.11	0.55	4.83	49.04	8.59
HHC001	15	18	20.64	12.43	0.79	0.50	0.15	0.85	0.17	0.09	5.95	1.93	0.74	0.14	0.11	0.64	5.08	50.23	8.81
HHC001	18	21	21.62	14.07	0.86	0.48	0.14	0.78	0.16	0.08	6.30	2.02	1.01	0.12	0.02	0.60	5.33	53.60	9.29
HHC001	21	24	27.39	16.07	0.76	0.48	0.22	0.86	0.16	0.09	8.05	2.74	1.30	0.15	0.01	0.57	4.83	63.68	11.70
HHC001	24	27	29.36	16.77	0.65	0.54	0.19	0.80	0.17	0.10	8.16	2.96	1.03	0.15	0.01	0.66	5.33	66.89	11.93
HHC001	27	30	52.94	29.91	0.95	0.55	0.30	1.41	0.19	0.11	14.93	5.27	2.02	0.20	0.01	0.64	6.10	115.53	21.35
HHC001	30	33	23.83	15.25	0.61	0.47	0.25	0.86	0.18	0.08	7.46	2.33	1.00	0.11	0.01	0.55	4.06	57.06	10.51
HHC001	33	36	22.85	14.66	0.69	0.47	0.25	0.71	0.15	0.10	6.88	2.34	1.07	0.12	0.02	0.51	4.83	55.66	10.03
HHC001	36	39	18.67	11.26	0.55	0.41	0.19	0.66	0.11	0.08	5.37	1.91	0.96	0.12	0.02	0.52	3.94	44.77	7.94
HHC001	39	42	16.71	10.44	0.39	0.32	0.19	0.54	0.11	0.07	4.90	1.62	0.66	0.07	0.01	0.36	3.30	39.69	6.98
HHC001	42	45	31.45	17.24	0.70	0.46	0.20	0.93	0.16	0.10	9.21	2.84	1.70	0.15	0.06	0.61	4.57	70.39	12.91
HHC001	45	48	42.87	19.47	0.76	0.45	0.28	0.95	0.19	0.09	10.96	3.53	1.60	0.18	0.03	0.63	5.08	87.06	15.43
HHC001	48	51	88.81	44.33	1.40	0.83	0.53	2.11	0.30	0.15	21.93	7.50	3.33	0.26	0.07	0.73	8.13	180.41	31.09
HHC001	51	54	75.55	46.33	1.26	0.72	0.69	1.68	0.27	0.14	22.74	8.11	3.18	0.24	0.05	0.85	7.62	169.43	32.35
HHC001	54	58	148.64	75.29	2.27	1.26	1.01	3.61	0.45	0.17	39.77	13.47	5.39	0.48	0.13	1.33	14.22	307.49	56.00
HHC002	0	3	122.47	68.73	4.05	1.97	1.38	5.23	0.82	0.28	48.64	14.92	7.68	0.72	0.30	1.88	24.13	303.19	68.33
HHC002	3	6	36.11	23.10	1.06	0.81	0.42	1.36	0.29	0.13	11.66	3.66	1.88	0.18	0.10	0.75	7.87	89.38	16.56
HHC002	6	9	28.99	17.12	0.77	0.72	0.27	0.83	0.19	0.10	8.75	2.79	1.50	0.15	0.10	0.74	5.84	68.87	12.46
HHC002	9	12	20.64	12.55	0.67	0.56	0.22	0.65	0.17	0.09	6.18	1.99	0.89	0.13	0.08	0.55	4.70	50.06	8.97
HHC002	12	15	22.48	14.78	0.92	0.67	0.17	0.86	0.21	0.10	7.00	2.05	1.23	0.14	0.07	0.68	6.35	57.72	10.11
HHC002	15	18	20.39	11.61	0.69	0.47	0.19	0.67	0.17	0.08	5.83	1.79	0.88	0.11	0.11	0.56	5.21	48.75	8.41
HHC002	18	21	20.15	12.20	0.52	0.35	0.08	0.60	0.09	0.07	5.48	1.96	1.06	0.12	0.07	0.47	3.43	46.63	8.07
HHC002	21	24	21.62	12.43	0.67	0.40	0.27	0.68	0.14	0.09	6.18	2.14	1.08	0.11	0.01	0.60	4.19	50.60	9.09
HHC002	24	27	41.15	22.99	0.91	0.53	0.34	1.23	0.19	0.11	12.60	3.93	1.73	0.18	0.02	0.63	5.21	91.73	17.61
HHC002	27	30	48.40	28.85	1.07	0.62	0.28	1.24	0.22	0.14	14.00	4.89	2.32	0.21	0.03	0.79	7.11	110.16	20.17
HHC002	30	33	29.24	19.59	0.72	0.49	0.29	1.01	0.17	0.08	10.61	3.43	1.55	0.12	0.10	0.63	4.95	72.99	14.89
HHC002	33	36	20.39	12.43	0.56	0.38	0.15	0.71	0.14	0.08	6.30	2.11	1.07	0.09	0.11	0.43	3.68	48.65	9.07
HHC002	36	39	16.95	10.44	0.55	0.47	0.16	0.65	0.15	0.09	5.25	1.84	0.74	0.09	0.06	0.51	4.32	42.27	7.73
HHC002	39	42	19.90	10.44	0.64	0.53	0.16	0.63	0.17	0.13	6.42	1.97	0.89	0.11	0.08	0.74	4.83	47.63	9.13
HHC002	42	45	28.99	14.78	0.75	0.43	0.23	0.82	0.15	0.08	8.16	2.61	1.35	0.13	0.01	0.51	4.44	63.44	11.85
HHC002	45	48	83.04	38.47	1.15	0.71	0.53	1.73	0.23	0.13	20.30	6.37	3.04	0.25	0.06	0.74	7.75	164.47	28.06
HHC002	48	51	362.38	164.19	5.30	2.63	2.28	8.26	0.96	0.34	93.66	29.48	12.12	1.09	0.34	2.27	28.83	714.14	129.54
HHC002	51	54	157.24	73.42	2.34	1.03	1.16	3.48	0.33	0.17	45.84	14.26	6.67	0.44	0.08	1.00	11.94	319.38	62.87
HHC002	54	57	133.90	76.11	2.02	0.95	1.02	3.28	0.39	0.19	43.39	13.59	5.95	0.41	0.15	1.04	11.05	293.44	59.41
HHC002	57	60	119.03	61.69	2.09	1.14	0.94	3.00	0.39	0.22	35.69	11.18	4.71	0.42	0.13	1.10	11.68	253.41	49.38
HHC002	60	62	122.59	55.94	1.86	0.87	0.71	2.60	0.33	0.17	29.86	9.36	4.02	0.34	0.08	1.01	10.67	240.43	41.42
HHC003	0	3	200.23	83.74	5.68	2.84	1.76	6.86	1.04	0.36	63.80	19.15	10.29	1.09	0.37	2.44	32.38	432.02	89.73
HHC003	3	6	66.46	34.13	1.71	1.14	0.54	1.89	0.39	0.23	16.45	5.10	2.25	0.31	0.16	1.49	13.84	146.08	23.56
HHC003	6	9	28.62	17.01	0.86	0.70	0.35	1.03	0.22	0.14	8.40	2.63	1.16	0.15	0.03	0.74	6.73	68.76	12.05
HHC003	9	12	25.06	14.54	0.98	0.63	0.21	0.92	0.16	0.13	7.00	2.27	1.31	0.16	0.03	0.80	6.60	60.80	10.41
HHC003	12	15	21.13	12.90	0.80	0.58	0.23	0.84	0.21	0.11	6.18	1.99	0.97	0.15	0.10	0.67	6.48	53.36	9.13
HHC003	15	18	31.57	15.36	1.03	0.57	0.25	1.29	0.22	0.08	8.51	2.63	1.35	0.18	0.08	0.73	7.62	71.48	12.36
HHC003	18	21	21.50	14.07	0.65	0.35	0.15	0.80	0.11	0.10	6.53	2.27	0.81	0.14	0.05	0.44	4.95	52.94	9.60
HHC003	21	24	28.74	16.07	0.71	0.37	0.21	0.93	0.15	0.08	8.51	2.80	1.39	0.12	0.01	0.60	4.70	65.40	12.15
HHC003	24	27	26.04	14.89	0.65	0.53	0.24	0.80	0.18	0.10	7.46	2.33	1.41	0.13	0.03	0.59	4.83	60.23	10.58
HHC003	27	30	27.15	17.01	0.68	0.38	0.32	0.88	0.17	0.10	8.98	2.75	1.37	0.12	0.02	0.65	4.83	65.40	12.53
HHC003	30	33	21.99	12.78	0.56	0.40	0.25	0.68	0.15	0.09	6.53	1.95	1.04	0.12	0.03	0.63	4.19	51.40	9.16
HHC003	33	36	20.39	11.96	0.50	0.48	0.17	0.63	0.11	0.09	6.07	1.98	0.99	0.12	0.03	0.56	4.19	48.29	8.67
HHC003	36	39	34.15	22.28	0.87	0.53	0.29	1.24	0.19	0.14	12.48	3.95	1.75	0.16	0.11	0.56	5.59	84.30	17.47
HHC003	39	42	87.95	64.97	1.41	0.65	0.73	2.44	0.27	0.11	35.69	11.54	4.79	0.29	0.13	0.96	9.52	221.47	48.94
HHC003	42	45	84.51	58.29	1.42	0.69	0.60	2.31	0.25	0.11	30.79	9.61	4.04	0.32	0.08	0.79	8.38	202.18	42.14

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC003	45	48	119.52	75.18	1.86	0.74	0.90	3.07	0.33	0.15	40.82	12.93	5.65	0.39	0.09	0.95	10.29	272.86	56.00
HHC003	48	51	165.22	111.89	2.18	0.88	1.41	4.46	0.39	0.15	65.20	20.90	8.07	0.46	0.15	0.88	11.18	393.41	88.74
HHC003	51	54	227.87	127.25	2.16	0.93	1.70	4.58	0.37	0.14	74.77	24.65	9.95	0.54	0.17	1.04	11.94	488.03	102.11
HHC003	54	57	170.13	61.22	2.11	1.15	1.02	3.14	0.36	0.18	36.51	11.56	5.30	0.34	0.15	1.06	11.94	306.17	50.52
HHC003	57	60	318.16	150.70	4.35	1.92	2.11	7.28	0.71	0.24	90.98	28.39	12.00	0.84	0.25	1.58	22.86	642.37	124.56
HHC003	60	63	181.19	80.45	2.42	1.19	1.07	4.01	0.46	0.19	48.64	14.92	6.74	0.46	0.15	1.29	13.33	356.51	66.44
HHC003	63	66	86.23	53.01	1.35	0.80	0.52	1.94	0.27	0.14	23.79	8.16	2.64	0.26	0.16	0.87	8.00	188.15	33.56
HHC003	66	69	170.75	88.66	2.25	1.05	1.01	3.33	0.41	0.19	46.07	14.68	6.02	0.45	0.15	1.08	12.06	348.17	63.45
HHC004	0	3	186.72	82.10	6.63	3.24	1.99	7.75	1.20	0.45	67.18	20.06	10.61	1.14	0.55	3.40	35.30	428.33	95.02
HHC004	3	6	72.23	24.86	1.81	1.12	0.45	1.76	0.37	0.23	14.23	4.26	2.11	0.25	0.17	1.37	10.29	135.51	20.56
HHC004	6	9	26.78	15.83	0.76	0.51	0.21	0.91	0.15	0.10	7.93	2.53	1.28	0.14	0.08	0.65	5.21	63.06	11.36
HHC004	9	12	30.22	16.89	1.08	0.58	0.25	0.84	0.21	0.15	9.10	3.04	1.44	0.16	0.08	0.80	6.10	70.94	13.39
HHC004	12	15	19.29	10.79	0.63	0.45	0.13	0.62	0.13	0.11	5.60	1.64	0.94	0.11	0.03	0.71	4.44	45.61	7.98
HHC004	15	18	20.15	11.61	0.76	0.46	0.23	0.69	0.16	0.11	6.07	1.90	0.85	0.14	0.05	0.64	4.57	48.37	8.86
HHC004	18	21	20.39	11.38	0.63	0.41	0.20	0.73	0.14	0.10	6.18	1.99	0.67	0.14	0.05	0.54	4.32	47.86	8.95
HHC004	21	24	25.55	15.01	0.93	0.55	0.27	0.83	0.16	0.13	7.46	2.53	1.23	0.13	0.08	0.76	4.95	60.57	11.05
HHC004	24	27	26.04	15.13	0.81	0.57	0.21	0.74	0.15	0.07	7.35	2.42	0.87	0.13	0.07	0.60	4.06	59.22	10.71
HHC004	27	30	20.15	13.72	0.71	0.38	0.25	0.56	0.16	0.09	6.77	2.30	1.03	0.11	0.07	0.56	3.94	50.79	9.88
HHC004	30	33	33.54	22.40	0.75	0.45	0.30	1.12	0.15	0.08	11.78	3.73	1.25	0.12	0.07	0.57	4.32	80.61	16.38
HHC004	33	36	17.93	7.15	0.61	0.40	0.16	0.51	0.11	0.06	3.62	1.30	0.41	0.07	0.09	0.51	3.81	36.75	5.60
HHC004	36	39	122.72	107.19	2.17	0.87	1.44	3.99	0.61	0.13	58.90	18.85	7.33	0.45	0.11	1.08	11.43	337.26	80.37
HHC004	39	42	99.13	63.21	1.69	0.85	1.01	3.11	0.34	0.14	39.42	12.20	4.65	0.38	0.14	0.90	9.65	236.82	53.69
HHC004	42	45	54.17	30.49	1.01	0.47	0.56	1.52	0.19	0.09	19.13	6.22	2.61	0.18	0.10	0.67	5.21	122.62	26.54
HHC004	45	48	113.75	62.98	1.82	0.95	0.98	2.99	0.29	0.23	39.07	12.20	5.53	0.34	0.15	1.17	9.65	252.11	53.44
HHC004	48	51	421.34	129.59	2.26	1.17	1.34	4.37	0.38	0.15	68.47	22.77	7.48	0.51	0.18	1.02	12.32	673.35	94.01
HHC004	51	54	117.68	52.31	1.39	0.65	0.69	1.99	0.24	0.14	28.11	8.52	3.44	0.24	0.11	0.87	7.24	223.62	38.25
HHC004	54	57	125.30	62.51	1.47	0.71	0.73	2.16	0.24	0.13	32.31	10.80	3.56	0.32	0.09	0.88	7.87	249.07	44.90
HHC004	57	60	67.44	35.54	1.10	0.71	0.46	1.73	0.23	0.10	16.68	5.38	2.30	0.22	0.11	0.95	6.48	139.42	23.38
HHC004	60	63	43.85	25.57	0.90	0.54	0.36	1.04	0.21	0.10	11.20	3.82	1.53	0.14	0.10	0.67	5.46	95.48	16.05
HHC004	63	66	51.96	30.02	0.81	0.57	0.37	1.08	0.16	0.11	14.00	4.71	1.80	0.15	0.13	0.68	5.33	111.90	19.68
HHC004	66	69	183.03	81.86	1.40	0.69	0.96	2.19	0.26	0.10	41.06	13.35	4.37	0.33	0.09	0.72	7.87	338.29	56.14
HHC004	69	72	84.27	47.62	0.93	0.57	0.57	1.53	0.21	0.11	22.86	7.95	2.83	0.21	0.09	0.73	6.73	177.21	31.95
HHC004	72	75	83.65	48.44	1.25	0.62	0.51	1.65	0.18	0.09	24.14	7.83	3.17	0.22	0.09	0.76	6.60	179.21	33.45
HHC004	75	78	105.27	56.88	0.98	0.57	0.72	1.72	0.18	0.11	29.28	9.38	3.20	0.25	0.08	0.74	5.84	215.20	39.87
HHC005	0	3	130.82	69.66	4.14	2.52	1.54	5.79	0.86	0.35	53.07	15.40	8.34	0.76	0.31	2.29	29.97	325.83	73.38
HHC005	3	6	111.66	57.47	3.33	1.70	1.13	3.83	0.49	0.16	39.77	11.73	5.69	0.51	0.33	1.67	17.78	257.26	55.34
HHC005	6	9	26.16	16.30	1.01	0.64	0.37	0.98	0.17	0.07	8.86	2.69	1.04	0.11	0.16	0.68	5.46	64.72	12.67
HHC005	9	12	21.62	12.43	0.95	0.41	0.30	0.54	0.15	0.10	5.83	2.03	1.28	0.08	0.10	0.68	4.70	51.21	8.90
HHC005	12	15	20.88	12.43	1.08	0.65	0.23	0.74	0.16	0.14	6.42	1.97	0.95	0.15	0.13	0.76	5.59	52.28	9.62
HHC005	15	18	19.78	12.20	0.80	0.46	0.20	0.53	0.13	0.09	5.60	1.97	1.04	0.08	0.09	0.52	4.44	47.93	8.45
HHC005	18	21	27.88	15.83	0.79	0.46	0.23	0.80	0.13	0.10	8.16	2.43	1.22	0.08	0.07	0.58	4.44	63.21	11.47
HHC005	21	24	18.67	13.84	0.50	0.39	0.21	0.66	0.08	0.05	6.53	1.99	0.85	0.06	0.09	0.51	3.56	47.99	9.09
HHC005	24	27	34.64	23.57	0.63	0.49	0.25	0.85	0.11	0.11	9.91	3.42	1.08	0.08	0.09	0.38	4.19	79.83	14.05
HHC005	27	30	20.51	16.07	0.60	0.46	0.22	0.62	0.14	0.10	7.12	2.32	1.01	0.11	0.11	0.68	4.57	54.64	10.14
HHC005	30	33	7.74	5.39	0.34	0.32	0.15	0.32	0.08	0.05	2.45	0.79	0.67	0.04	0.05	0.38	3.05	21.81	3.61
HHC005	33	36	14.25	8.44	0.53	0.30	0.22	0.50	0.11	0.07	4.08	1.30	0.78	0.08	0.10	0.63	3.43	34.82	6.00
HHC005	36	39	75.06	28.85	1.14	0.55	0.44	1.34	0.16	0.15	15.40	5.10	2.28	0.16	0.15	0.81	6.22	137.80	21.80
HHC005	39	42	64.12	28.38	1.17	0.40	0.72	1.66	0.19	0.14	18.66	5.59	2.68	0.18	0.17	0.81	6.48	131.35	25.60
HHC005	42	45	66.95	29.09	1.30	0.75	0.51	1.49	0.19	0.14	17.50	5.20	2.52	0.27	0.14	0.74	6.98	133.75	24.26
HHC005	45	48	174.43	120.80	1.85	1.11	1.23	3.52	0.32	0.15	49.81	17.16	6.22	0.35	0.14	0.80	12.19	390.05	69.16
HHC005	48	51	105.89	65.44	1.35	1.03	0.83	2.56	0.26	0.22	31.73	10.12	4.30	0.26	0.21	0.90	9.40	234.50	43.46
HHC005	51	54	86.23	53.71	1.25	0.77	0.69	1.91	0.26	0.20	26.13	8.38	4.05	0.20	0.15	0.92	9.14	194.01	35.96
HHC005	54	57	46.68	27.91	0.83	0.53	0.38	1.23	0.17	0.13	13.06	4.72	1.99	0.12	0.14	0.68	6.10	104.67	18.73
HHC005	57	60	38.20	26.27	0.68	0.54	0.32	0.83	0.15	0.07	10.61	3.65	1.68	0.09	0.14	0.76	5.33	89.33	15.03
HHC005	60	63	51.59	28.97	0.70	0.71	0.34	1.21	0.16	0.14	14.23	4.55	1.87	0.14	0.17	0.65	5.59	111.01	19.63
HHC005	63	66	51.84	32.72	0.99	0.48	0.61	1.12	0.18	0.17	13.76	5.16	2.38	0.13	0.15	0.77	5.08	115.54	20.04
HHC005	66	69	94.59	70.49	1.14	0.59	0.98	1.90	0.18	0.13	34.76	11.22	4.79	0.22	0.18	0.71	6.35	228.23	47.34
HHC005	69	72	82.92	60.16	0.94	0.48	0.60	2.02	0.16	0.09	29.51	10.10	3.83	0.18	0.11	0.60	5.59	197.29	40.73
HHC005	72	74	120.14	66.62	1.37	0.50	0.96	2.04	0.18	0.08	34.53	11.92	4.75	0.20	0.21	0.72	5.59	249.80	48.02

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC006	0	3	162.76	86.67	5.05	2.53	1.53	6.30	0.90	0.40	61.82	19.09	10.10	0.87	0.42	2.40	32.26	393.11	86.83
HHC006	3	6	55.28	25.33	1.58	1.20	0.37	1.61	0.26	0.16	14.00	4.42	2.02	0.21	0.25	1.15	11.30	119.15	20.21
HHC006	6	9	32.55	21.34	1.21	0.74	0.36	0.98	0.22	0.10	10.85	3.47	1.61	0.20	0.14	0.71	7.87	82.35	15.72
HHC006	9	12	23.09	14.54	0.71	0.56	0.31	0.95	0.17	0.09	6.88	2.49	1.23	0.14	0.11	0.66	5.08	57.02	10.22
HHC006	12	15	15.97	9.97	0.63	0.46	0.24	0.53	0.13	0.13	5.02	1.62	1.00	0.11	0.11	0.47	3.56	39.93	7.37
HHC006	15	18	21.37	13.49	0.93	0.56	0.28	0.91	0.21	0.07	7.00	2.13	0.87	0.14	0.06	0.58	5.59	54.18	10.20
HHC006	18	21	21.87	13.02	0.70	0.39	0.25	0.76	0.17	0.08	7.00	2.30	0.92	0.13	0.06	0.69	4.19	52.52	10.12
HHC006	21	24	18.18	14.07	0.68	0.42	0.20	0.55	0.11	0.09	6.42	1.88	0.96	0.11	0.07	0.61	3.56	47.92	9.08
HHC006	24	27	19.78	13.25	0.69	0.63	0.22	0.88	0.19	0.10	6.18	2.20	1.10	0.12	0.06	0.54	5.46	51.39	9.19
HHC006	27	30	14.25	7.74	0.39	0.34	0.15	0.41	0.09	0.06	4.08	1.41	0.72	0.08	0.01	0.43	2.79	32.97	5.97
HHC006	30	33	26.04	10.56	0.52	0.39	0.30	0.70	0.13	0.07	6.88	2.02	1.06	0.09	0.01	0.50	3.94	53.19	9.51
HHC006	33	36	47.42	20.64	1.00	0.43	0.47	1.08	0.14	0.11	12.83	4.13	2.09	0.15	0.02	0.48	4.83	95.83	18.11
HHC006	36	39	150.48	117.87	2.48	0.97	1.24	4.17	0.46	0.14	55.17	19.63	6.74	0.46	0.17	1.28	13.97	375.22	77.74
HHC006	39	42	104.78	65.79	1.89	0.99	0.94	2.50	0.31	0.18	33.36	10.52	4.72	0.34	0.08	1.10	9.14	236.67	46.12
HHC006	42	45	97.53	51.02	1.48	0.77	0.76	1.99	0.26	0.15	26.71	8.61	3.40	0.26	0.05	0.91	6.60	200.51	37.06
HHC006	45	48	73.46	40.81	1.32	0.65	0.63	1.96	0.29	0.17	24.61	7.53	3.08	0.22	0.08	0.92	6.48	162.21	33.68
HHC006	48	51	34.40	21.58	0.96	0.48	0.43	1.20	0.23	0.14	11.90	3.43	1.35	0.19	0.14	0.87	5.59	82.86	16.48
HHC006	51	54	41.27	18.53	0.91	0.64	0.32	0.76	0.14	0.13	9.56	2.97	1.61	0.14	0.10	0.80	5.33	83.22	13.58
HHC006	54	57	187.95	86.67	1.86	0.93	1.16	3.93	0.39	0.16	49.46	16.01	6.31	0.41	0.16	0.97	10.92	367.27	67.73
HHC006	57	60	261.65	144.84	4.26	1.88	2.92	7.73	0.82	0.31	96.69	30.45	13.28	0.82	0.27	1.96	23.49	591.37	132.22
HHC006	60	63	287.45	161.26	3.94	1.97	2.43	7.24	0.79	0.31	96.46	29.84	13.10	0.82	0.29	1.74	25.78	633.41	131.06
HHC006	63	66	316.93	182.37	5.31	2.81	2.52	9.12	0.96	0.45	99.96	32.14	13.39	0.95	0.42	2.58	37.97	707.90	138.37
HHC006	66	69	287.45	168.30	5.62	3.28	2.34	9.08	1.15	0.41	100.43	30.21	13.97	1.11	0.40	3.21	40.89	667.84	137.36
HHC006	69	72	260.42	149.53	4.67	2.20	2.04	7.23	0.86	0.28	87.60	28.03	11.41	0.86	0.34	1.96	27.56	584.98	121.16
HHC006	72	76	230.94	131.94	3.44	1.76	2.01	5.88	0.64	0.17	77.22	22.96	10.56	0.67	0.25	1.30	18.79	508.54	104.29
HHC007	0	3	163.99	83.15	5.20	2.50	1.77	6.96	1.03	0.36	57.62	17.88	10.01	0.86	0.37	2.47	31.37	385.54	81.56
HHC007	3	6	152.94	55.00	3.81	1.88	1.03	4.39	0.68	0.23	37.32	11.41	6.25	0.64	0.30	1.62	21.84	299.32	53.18
HHC007	6	9	42.38	24.98	1.14	0.72	0.41	1.43	0.26	0.10	13.30	4.51	1.70	0.20	0.15	0.87	7.75	99.89	19.14
HHC007	9	12	20.88	12.90	0.81	0.47	0.21	0.62	0.14	0.07	6.65	1.95	1.07	0.14	0.13	0.54	4.95	51.52	9.55
HHC007	12	15	29.85	17.71	1.19	0.69	0.24	1.03	0.26	0.14	9.68	2.97	1.15	0.20	0.10	0.76	8.00	73.98	14.05
HHC007	15	18	16.95	11.02	0.61	0.55	0.17	0.62	0.16	0.10	5.95	1.84	1.15	0.13	0.15	0.58	4.44	44.43	8.52
HHC007	18	21	17.07	11.61	0.78	0.42	0.24	0.62	0.13	0.10	5.83	1.69	0.81	0.09	0.09	0.54	4.19	44.23	8.40
HHC007	21	24	19.65	14.19	0.73	0.42	0.23	0.74	0.15	0.09	6.42	2.02	0.86	0.12	0.11	0.68	5.46	51.88	9.29
HHC007	24	27	24.69	13.37	0.69	0.53	0.27	0.84	0.16	0.08	8.40	2.32	1.16	0.11	0.09	0.72	4.19	57.61	11.51
HHC007	27	30	37.34	14.89	0.71	0.50	0.28	0.96	0.11	0.11	9.68	3.09	1.28	0.13	0.08	0.61	4.44	74.23	13.62
HHC007	30	33	57.61	40.93	1.40	0.91	0.76	2.50	0.29	0.11	30.21	8.80	4.34	0.29	0.08	0.93	8.76	157.94	40.70
HHC007	33	37	114.73	81.51	2.44	1.17	1.24	4.00	0.38	0.18	51.32	15.28	7.06	0.35	0.21	1.24	12.45	293.56	69.40
HHC008	0	3	142.49	66.97	4.84	2.74	1.61	6.69	0.89	0.31	54.00	14.44	9.14	0.80	0.40	2.33	28.32	335.98	74.09
HHC008	3	6	167.68	80.57	5.67	2.89	1.68	7.07	1.03	0.33	62.40	17.22	11.65	1.04	0.42	2.51	28.95	391.11	86.32
HHC008	6	9	32.80	19.47	1.03	0.71	0.29	1.21	0.17	0.09	10.85	3.46	1.74	0.15	0.15	0.68	5.84	78.64	15.49
HHC008	9	12	25.43	15.36	0.81	0.57	0.39	0.89	0.17	0.09	7.81	2.43	1.70	0.12	0.18	0.61	5.59	62.17	11.18
HHC008	12	15	24.94	14.89	0.88	0.87	0.23	0.93	0.22	0.13	8.16	2.42	1.30	0.16	0.11	0.71	5.84	61.80	11.63
HHC008	15	18	18.18	11.26	0.80	0.73	0.23	0.62	0.18	0.14	5.83	1.88	0.92	0.14	0.21	0.65	5.71	47.49	8.66
HHC008	18	21	24.20	17.59	0.57	0.37	0.22	0.81	0.17	0.10	9.10	2.68	1.37	0.13	0.16	0.68	4.06	62.22	12.48
HHC008	21	24	12.28	7.74	0.54	0.41	0.20	0.36	0.14	0.08	4.55	1.23	0.75	0.08	0.05	0.44	3.43	32.28	6.40
HHC008	24	27	189.17	96.29	2.65	1.42	1.22	4.54	0.46	0.18	51.55	15.59	6.83	0.61	0.19	1.21	11.05	382.96	70.40
HHC008	27	30	395.54	224.00	6.07	2.56	2.98	10.22	0.97	0.30	128.30	37.45	16.29	1.18	0.38	2.20	27.05	855.50	173.01
HHC008	30	33	428.71	235.73	7.00	3.74	3.93	13.25	1.17	0.36	160.38	44.46	22.67	1.42	0.45	2.82	34.80	960.90	213.27
HHC008	33	36	323.07	190.58	5.10	2.61	2.59	9.58	0.93	0.26	109.64	33.59	13.63	1.12	0.40	1.82	28.19	723.10	149.44
HHC008	36	40	264.11	150.70	4.18	1.83	2.07	7.04	0.69	0.27	89.23	26.58	10.37	0.86	0.25	1.59	22.10	581.87	120.85
HHC009	0	3	182.42	92.89	4.82	2.58	1.75	6.54	0.88	0.39	64.62	18.55	10.39	0.82	0.43	2.08	27.43	416.59	88.81
HHC009	3	6	162.15	70.02	3.93	1.80	1.35	4.89	0.64	0.24	49.81	14.56	7.91	0.76	0.37	1.91	22.73	343.05	69.05
HHC009	6	9	33.04	19.94	1.02	0.65	0.37	0.95	0.21	0.11	11.20	3.25	1.91	0.16	0.17	0.77	6.73	80.49	15.63
HHC009	9	12	42.75	30.02	1.43	0.95	0.34	1.44	0.29	0.16	15.28	4.65	2.06	0.22	0.21	0.82	9.27	109.89	21.59
HHC009	12	15	36.11	22.17	1.45	0.78	0.25	1.33	0.25	0.17	10.50	3.54	1.92	0.20	0.13	0.66	8.00	87.46	15.68
HHC009	15	18	40.91	21.58	1.19	0.74	0.36	1.36	0.26	0.13	12.60	4.04	1.72	0.18	0.16	0.69	6.22	92.13	18.00
HHC009	18	21	289.90	187.65	4.87	2.36	2.91	9.08	0.86	0.26	125.97	39.15	17.10	1.04	0.29	1.71	22.10	705.23	171.02
HHC009	21	24	234.01	134.29	3.60	1.98	2.13	6.40	0.72	0.25	84.80	26.34	11.07	0.73	0.27	1.51	21.72	529.82	115.47
HHC009	24	27	214.36	121.38	3.11	1.32	1.82	5.49	0.50	0.15	77.68	21.93	9.61	0.62	0.23	0.99	17.27	476.46	103.34

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC009	27	31	144.95	79.05	2.98	1.88	0.89	3.90	0.56	0.48	48.64	15.46	6.84	0.49	0.31	2.54	20.06	329.04	67.58
HHC010	0	3	262.88	143.08	8.16	4.49	3.11	11.48	1.67	0.55	106.84	31.53	17.68	1.47	0.49	3.69	45.59	642.73	148.01
HHC010	3	6	48.40	27.33	1.66	0.90	0.46	1.49	0.34	0.13	14.11	4.92	2.13	0.20	0.17	0.81	9.78	112.83	20.89
HHC010	6	9	29.11	17.24	0.86	0.55	0.31	1.07	0.17	0.13	8.86	2.79	1.36	0.13	0.03	0.73	5.46	68.81	12.65
HHC010	9	12	32.18	19.23	1.00	0.77	0.14	1.23	0.19	0.14	9.45	3.12	1.75	0.16	0.07	0.66	6.98	77.08	13.73
HHC010	12	15	31.45	22.75	1.18	0.63	0.46	1.23	0.27	0.22	12.83	4.25	1.84	0.19	0.16	0.91	7.11	85.50	18.45
HHC010	15	19	173.20	105.90	2.71	1.26	1.44	4.67	0.46	0.17	57.04	18.55	6.48	0.59	0.11	1.01	13.97	387.56	78.88
HHC011	0	3	139.42	77.64	5.08	2.78	1.78	6.64	0.84	0.38	57.97	16.91	9.03	0.87	0.24	2.52	28.19	350.30	80.84
HHC011	3	6	41.15	26.04	1.31	0.81	0.39	1.57	0.25	0.19	12.83	4.25	2.06	0.22	0.14	0.95	9.27	101.44	18.62
HHC011	6	9	32.18	19.82	0.77	0.54	0.29	1.12	0.11	0.09	10.61	3.55	1.60	0.12	0.21	0.72	4.06	75.79	15.05
HHC011	9	13	26.29	17.01	1.14	0.59	0.22	0.82	0.17	0.15	8.05	2.54	1.18	0.16	0.16	0.76	6.10	65.33	11.89
HHC012	0	3	109.57	53.71	4.27	2.50	1.23	5.39	0.66	0.32	38.72	11.51	6.78	0.67	0.39	2.06	23.37	261.17	55.18
HHC012	3	6	133.28	67.55	4.57	2.04	1.44	5.30	0.86	0.26	46.07	13.95	8.26	0.72	0.30	2.06	25.52	312.18	65.31
HHC012	6	9	28.01	17.59	1.38	0.77	0.35	1.26	0.26	0.11	8.75	2.67	1.23	0.19	0.13	0.72	7.24	70.64	12.98
HHC012	9	12	24.08	15.25	1.07	0.86	0.29	1.12	0.18	0.14	7.23	2.48	1.30	0.11	0.15	0.80	6.60	61.64	10.88
HHC012	12	15	20.51	12.90	1.11	0.64	0.28	0.84	0.18	0.13	6.30	1.86	1.00	0.15	0.09	0.67	5.71	52.38	9.43
HHC012	15	18	18.30	11.49	0.81	0.43	0.15	0.67	0.17	0.10	5.25	1.58	0.80	0.13	0.02	0.50	5.08	45.50	7.78
HHC012	18	21	21.74	13.49	0.84	0.62	0.28	1.05	0.14	0.09	7.46	2.40	0.83	0.12	0.06	0.67	5.08	54.87	10.82
HHC012	21	24	20.27	15.36	0.78	0.57	0.23	0.70	0.17	0.11	6.65	2.26	1.10	0.12	0.02	0.68	4.57	53.61	9.81
HHC012	24	27	13.02	9.73	0.71	0.49	0.23	0.48	0.09	0.09	4.32	1.52	0.65	0.06	0.05	0.42	3.43	35.30	6.61
HHC012	27	30	21.62	8.33	0.57	0.41	0.21	0.66	0.10	0.09	6.18	1.70	0.92	0.09	0.01	0.46	3.17	44.53	8.55
HHC012	30	33	43.61	19.59	1.02	0.63	0.46	1.21	0.15	0.15	11.78	3.32	1.72	0.18	0.17	0.91	4.19	89.08	16.30
HHC012	33	36	41.03	20.99	1.06	0.74	0.44	1.29	0.22	0.10	11.90	3.87	2.23	0.16	0.11	0.82	5.46	90.42	16.98
HHC012	36	39	120.87	61.57	1.89	0.95	0.80	2.85	0.26	0.17	37.91	11.62	5.01	0.35	0.13	0.95	9.14	254.48	51.78
HHC012	39	42	105.03	72.95	1.88	0.85	1.01	2.97	0.36	0.17	42.92	13.41	5.76	0.38	0.15	1.04	10.41	259.28	58.59
HHC012	42	45	242.61	118.45	5.50	2.33	2.81	9.22	0.89	0.36	102.76	30.57	14.73	1.00	0.43	2.36	28.57	562.60	139.82
HHC012	45	48	196.54	91.48	5.31	2.22	2.62	8.39	0.78	0.35	90.28	25.37	14.26	0.96	0.34	2.13	25.02	466.06	121.93
HHC012	48	51	316.93	205.83	5.78	2.33	2.88	9.67	0.89	0.35	127.72	39.51	16.76	1.16	0.33	2.22	27.18	759.55	174.18
HHC012	51	54	339.04	212.86	4.65	2.14	2.70	8.09	0.72	0.33	111.27	37.82	14.67	0.95	0.35	1.95	21.33	758.88	154.69
HHC013	0	3	152.32	77.17	4.71	2.15	1.46	6.33	0.86	0.27	53.89	16.49	8.19	0.84	0.37	2.22	26.03	353.29	75.92
HHC013	3	6	160.31	149.53	6.84	3.54	2.69	10.12	1.07	0.38	105.09	33.71	15.94	1.26	0.41	2.92	34.92	528.72	146.90
HHC013	6	9	38.33	24.28	1.19	0.83	0.45	1.48	0.22	0.11	14.46	4.33	2.20	0.20	0.11	0.98	8.38	97.56	20.18
HHC013	9	12	27.02	16.54	1.25	0.89	0.30	0.82	0.24	0.11	8.75	2.49	1.07	0.18	0.14	0.74	7.37	67.90	12.66
HHC013	12	15	29.36	18.88	1.17	0.59	0.34	1.15	0.26	0.17	11.55	3.17	1.83	0.18	0.13	0.80	6.73	76.30	16.06
HHC013	15	18	19.04	11.61	0.80	0.61	0.10	0.75	0.21	0.14	5.83	1.75	1.07	0.12	0.09	0.83	4.95	47.90	8.50
HHC013	18	21	24.69	15.95	0.72	0.64	0.21	0.48	0.16	0.10	8.05	2.28	1.18	0.13	0.08	0.63	4.44	59.75	11.18
HHC013	21	24	23.34	18.30	0.65	0.47	0.29	0.80	0.17	0.16	8.40	2.50	1.28	0.14	0.10	0.83	4.70	62.12	11.69
HHC013	24	27	16.09	11.49	0.62	0.47	0.28	0.65	0.10	0.03	6.30	1.95	0.86	0.12	0.03	0.65	3.68	43.32	8.98
HHC013	27	30	32.06	21.58	1.04	0.54	0.46	0.90	0.18	0.09	13.30	3.64	2.05	0.12	0.08	0.58	4.57	81.20	18.10
HHC013	30	33	83.90	63.21	1.33	0.64	0.76	2.47	0.22	0.14	31.96	10.25	4.00	0.29	0.09	0.84	7.11	207.22	43.83
HHC013	33	36	228.48	166.54	4.07	2.46	2.04	6.57	0.77	0.27	97.28	27.43	11.65	0.84	0.29	1.86	22.48	573.01	129.61
HHC014	0	3	153.55	88.66	4.57	2.34	1.63	6.11	0.72	0.31	61.59	18.12	9.82	0.75	0.15	2.05	25.91	376.28	85.03
HHC014	3	6	149.86	87.14	5.49	2.79	1.47	6.50	0.82	0.31	60.07	17.64	8.78	0.85	0.21	2.17	29.08	373.18	84.04
HHC014	6	9	27.15	16.89	0.88	0.65	0.31	0.98	0.17	0.14	9.10	2.78	1.17	0.12	0.01	0.63	6.22	67.19	12.88
HHC014	9	12	23.34	15.01	0.86	0.66	0.23	0.74	0.16	0.10	7.93	2.05	1.21	0.12	0.01	0.59	6.10	59.11	10.96
HHC014	12	15	24.57	14.54	0.79	0.50	0.21	0.82	0.16	0.08	7.35	2.30	1.19	0.12	0.06	0.64	4.70	58.02	10.55
HHC014	15	18	21.99	12.31	0.57	0.45	0.10	0.82	0.13	0.10	5.95	1.97	0.94	0.12	0.01	0.52	4.44	50.42	8.61
HHC014	18	21	26.41	15.48	0.76	0.59	0.23	0.82	0.14	0.14	7.70	2.49	1.37	0.13	0.01	0.73	4.95	61.95	11.07
HHC014	21	24	27.02	19.00	0.69	0.49	0.22	0.81	0.14	0.10	8.75	2.89	1.47	0.12	0.01	0.44	4.57	66.72	12.44
HHC014	24	27	32.43	20.52	0.99	0.58	0.29	1.13	0.19	0.14	12.60	3.64	1.88	0.15	0.05	0.79	5.84	81.21	17.37
HHC014A	0	3	107.49	49.14	3.45	1.77	1.08	4.59	0.61	0.26	38.84	11.02	6.68	0.69	0.29	1.89	19.94	247.73	54.01
HHC014A	3	6	110.43	51.96	2.82	1.43	0.85	3.70	0.53	0.25	34.18	10.58	6.02	0.52	0.27	1.49	16.89	241.91	48.10
HHC014A	6	9	28.50	17.12	0.87	0.63	0.37	1.15	0.24	0.14	9.21	2.61	1.41	0.21	0.14	0.71	6.98	70.30	12.91
HHC014A	9	12	24.45	13.60	0.79	0.62	0.28	0.88	0.18	0.13	8.40	2.27	1.37	0.14	0.14	0.73	5.71	59.68	11.60
HHC014A	12	15	21.62	12.90	0.79	0.57	0.20	0.86	0.16	0.09	7.35	1.81	1.24	0.13	0.10	0.51	4.95	53.30	10.08
HHC014A	15	18	27.27	13.96	0.85	0.55	0.21	0.97	0.16	0.10	8.16	2.32	1.37	0.15	0.07	0.76	4.44	61.35	11.49
HHC014A	18	21	25.55	15.25	0.87	0.48	0.17	0.74	0.13	0.14	7.35	2.14	1.51	0.16	0.14	0.59	5.21	60.42	10.52
HHC014A	21	24	20.39	15.72	0.65	0.41	0.23	0.66	0.08	0.08	7.00	2.05	1.03	0.14	0.13	0.58	3.94	53.09	9.85
HHC014A	24	27	37.59	26.04	0.99	0.56	0.35	1.29	0.18	0.13	15.28	4.34	1.69	0.16	0.11	0.60	5.21	94.52	20.77

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC014A	27	30	138.81	71.54	2.36	1.23	1.10	4.00	0.36	0.13	45.26	13.11	6.60	0.44	0.19	1.23	12.32	298.67	61.16
HHC014A	30	33	25.18	13.84	0.50	0.51	0.16	0.62	0.11	0.15	7.58	2.45	1.00	0.07	0.10	0.66	5.08	58.03	10.61
HHC014A	33	36	227.25	115.76	3.65	1.84	1.84	6.36	0.58	0.25	75.58	21.63	10.58	0.71	0.23	1.51	19.56	487.33	101.56
HHC014A	36	39	240.77	120.21	4.14	1.90	2.11	6.70	0.65	0.19	81.53	23.80	11.05	0.76	0.33	1.54	21.33	517.02	110.24
HHC014A	39	42	235.85	112.47	3.97	1.75	1.92	6.96	0.66	0.22	80.25	23.50	11.24	0.84	0.30	1.58	22.60	504.11	108.55
HHC015	0	3	83.04	57.82	3.41	1.76	1.08	4.63	0.62	0.16	42.92	11.08	6.83	0.61	0.33	1.58	22.60	238.48	58.02
HHC015	3	6	140.65	84.68	4.97	2.42	1.67	6.30	0.90	0.35	60.19	15.95	8.88	0.84	0.38	2.00	27.18	357.36	81.94
HHC015	6	9	47.54	29.20	1.12	0.77	0.46	1.86	0.24	0.15	16.68	4.81	2.41	0.21	0.11	1.05	8.64	115.25	22.82
HHC015	9	12	18.30	11.26	0.54	0.37	0.25	0.56	0.11	0.09	6.77	1.68	0.79	0.09	0.05	0.56	4.19	45.61	9.08
HHC015	12	15	24.94	15.13	0.76	0.64	0.20	1.18	0.16	0.13	8.05	2.43	1.51	0.14	0.06	0.67	6.10	62.07	11.38
HHC015	15	18	28.38	20.29	0.91	0.67	0.17	1.06	0.23	0.08	10.73	2.73	1.16	0.16	0.03	0.68	6.48	73.77	14.53
HHC015	18	21	26.90	18.53	0.76	0.50	0.29	0.78	0.19	0.10	10.03	2.82	1.51	0.18	0.09	0.77	4.95	68.41	13.78
HHC015	21	24	29.24	21.11	0.87	0.40	0.32	0.90	0.14	0.13	10.26	3.11	1.41	0.16	0.21	0.55	5.08	73.89	14.41
HHC015	24	26	65.47	43.39	1.06	0.58	0.53	1.71	0.25	0.13	24.96	7.48	2.54	0.21	0.23	0.96	7.11	156.61	33.71
HHC016	0	3	80.71	44.92	3.08	1.58	0.80	3.41	0.47	0.23	33.13	8.82	5.09	0.48	0.23	1.40	17.27	201.60	45.50
HHC016	3	6	105.64	65.44	3.94	2.37	1.19	5.33	0.79	0.38	47.59	12.50	7.28	0.69	0.39	1.71	25.02	280.26	64.72
HHC016	6	9	31.69	20.05	1.17	0.90	0.30	1.45	0.25	0.14	10.61	3.06	1.52	0.16	0.14	0.98	7.62	80.05	15.01
HHC016	9	12	29.11	21.70	1.30	0.94	0.28	1.19	0.27	0.15	8.40	2.78	1.32	0.22	0.10	0.73	8.00	76.49	12.70
HHC016	12	15	20.15	12.43	0.99	0.55	0.19	0.75	0.16	0.11	6.30	2.02	0.82	0.12	0.07	0.64	5.33	50.62	9.42
HHC016	15	18	24.94	15.48	1.06	0.71	0.22	0.91	0.15	0.09	7.81	2.25	1.22	0.13	0.21	0.98	5.46	61.61	11.25
HHC016	18	21	71.62	44.92	1.57	0.98	0.64	2.42	0.36	0.19	26.13	8.08	3.54	0.36	0.13	1.21	11.81	173.95	36.15
HHC016	21	24	51.59	25.92	1.01	0.63	0.43	1.49	0.16	0.13	13.76	4.19	1.76	0.19	0.08	0.64	5.46	107.44	19.15
HHC016	24	27	192.24	131.94	3.55	1.57	1.67	5.64	0.63	0.26	79.90	23.86	10.77	0.78	0.27	1.94	19.94	474.95	108.08
HHC016	27	30	272.70	194.10	6.45	3.10	3.07	10.04	1.08	0.41	130.64	37.09	18.96	1.32	0.50	2.79	35.43	717.67	175.50
HHC016	30	33	243.84	146.60	4.01	2.20	2.06	6.39	0.70	0.28	86.31	26.22	11.04	0.74	0.32	1.92	22.35	554.97	117.28
HHC016	33	37	188.56	102.50	3.51	1.92	1.48	5.39	0.71	0.35	67.30	19.27	9.23	0.80	0.26	2.20	22.10	425.59	90.88
HHC017	0	3	213.74	99.34	6.28	3.38	1.89	8.45	1.21	0.44	75.82	21.32	11.09	1.09	0.43	2.41	33.27	480.17	104.51
HHC017	3	6	51.96	29.55	1.66	1.11	0.52	1.98	0.30	0.14	17.03	4.91	2.49	0.26	0.18	0.99	10.03	123.12	23.86
HHC017	6	9	28.62	17.12	1.18	0.86	0.31	1.13	0.23	0.14	9.45	2.71	1.84	0.22	0.18	0.82	8.25	73.07	13.56
HHC017	9	12	25.67	15.25	1.01	0.64	0.25	1.06	0.22	0.14	8.40	2.28	1.73	0.11	0.10	0.88	6.73	64.46	11.80
HHC017	12	15	17.32	10.20	0.72	0.48	0.17	0.67	0.16	0.09	4.90	1.61	0.92	0.14	0.08	0.55	4.32	42.33	7.37
HHC017	15	18	20.15	12.43	1.10	0.72	0.21	0.98	0.17	0.10	5.48	1.80	0.94	0.15	0.09	0.81	5.59	50.72	8.54
HHC017	18	21	22.73	16.77	1.21	0.74	0.21	0.91	0.23	0.08	7.81	2.07	1.04	0.14	0.15	0.71	6.86	61.65	11.23
HHC017	21	24	26.16	25.22	0.91	0.61	0.29	0.97	0.23	0.11	13.53	4.01	1.53	0.14	0.10	0.71	4.95	79.47	18.59
HHC017	24	27	24.45	17.71	0.52	0.34	0.27	0.78	0.11	0.08	10.38	2.60	1.12	0.13	0.08	0.42	3.17	62.17	13.62
HHC017	27	30	16.46	10.91	0.39	0.31	0.21	0.63	0.10	0.08	6.77	1.85	0.95	0.09	0.15	0.51	3.68	43.09	9.10
HHC017	30	33	58.72	32.13	1.09	0.50	0.54	1.72	0.15	0.08	18.66	5.69	2.60	0.20	0.05	0.60	5.33	128.07	25.64
HHC017	33	36	54.79	20.64	1.00	0.45	0.32	1.41	0.15	0.09	14.11	3.76	1.95	0.12	0.05	0.58	3.81	103.22	18.99
HHC017	36	39	60.44	26.04	1.09	0.61	0.36	1.29	0.21	0.11	17.96	5.01	2.27	0.22	0.09	0.69	5.84	122.24	24.29
HHC017	39	42	84.02	36.00	0.91	0.46	0.66	1.60	0.18	0.13	22.86	6.37	2.73	0.18	0.15	0.80	5.71	162.75	30.31
HHC017	42	45	88.57	53.71	1.47	0.82	0.86	2.17	0.26	0.11	27.53	8.77	3.84	0.28	0.09	0.74	7.87	197.10	38.05
HHC017	45	48	74.07	51.13	1.63	0.72	0.63	2.52	0.26	0.11	26.13	7.53	2.82	0.24	0.13	0.72	9.02	177.65	35.52
HHC017	48	51	97.29	48.44	1.27	0.88	0.65	1.74	0.22	0.11	26.24	8.07	3.40	0.22	0.18	0.59	7.24	196.55	35.81
HHC017	51	54	55.52	26.04	0.99	0.59	0.42	1.24	0.16	0.10	14.46	4.08	2.11	0.14	0.03	0.59	4.70	111.19	19.68
HHC017	54	57	42.01	20.41	1.16	0.77	0.30	1.09	0.19	0.16	11.90	3.41	1.69	0.16	0.07	0.91	5.21	89.44	16.63
HHC017	57	60	35.62	19.12	0.98	0.67	0.25	1.20	0.24	0.14	10.26	3.03	1.40	0.12	0.11	0.84	6.86	80.85	14.39
HHC017	60	63	46.31	24.04	1.23	0.67	0.41	1.21	0.19	0.14	12.48	4.01	1.90	0.16	0.17	0.75	6.10	99.78	17.88
HHC017	63	66	93.73	37.30	1.29	0.61	0.58	1.84	0.18	0.14	21.93	6.51	2.75	0.19	0.16	0.76	6.73	174.69	29.91
HHC017	66	69	108.96	47.15	1.78	0.88	0.73	2.46	0.27	0.17	25.78	7.87	3.97	0.33	0.15	1.33	8.13	209.94	35.75
HHC017	69	72	97.29	40.34	1.31	0.81	0.64	2.11	0.21	0.10	24.84	7.04	3.79	0.26	0.15	0.83	7.49	187.22	33.46
HHC017	72	75	122.59	54.18	1.64	0.86	0.80	2.88	0.32	0.20	34.29	10.51	4.01	0.34	0.13	0.82	8.25	241.84	46.79
HHC017	75	78	113.01	74.59	1.47	0.80	0.93	2.93	0.27	0.14	42.81	12.87	5.36	0.26	0.09	0.81	8.64	264.96	57.40
HHC017	78	81	235.85	124.90	2.81	1.20	1.56	4.54	0.39	0.16	73.25	21.87	9.14	0.46	0.22	1.31	12.57	490.23	98.39
HHC017	81	84	113.50	60.40	2.02	1.30	0.69	2.77	0.37	0.25	34.88	10.10	4.42	0.41	0.15	1.34	12.06	244.67	47.41
HHC017	84	87	83.04	47.62	2.28	1.32	0.66	2.73	0.42	0.20	25.31	7.76	4.24	0.34	0.17	1.31	13.08	190.49	35.69
HHC017	87	90	84.88	44.10	3.52	2.44	0.80	3.91	0.62	0.39	27.88	8.16	4.85	0.61	0.30	2.31	20.70	205.45	40.17
HHC018	0	3	164.61	87.96	5.45	3.09	1.85	7.48	1.04	0.43	69.63	19.81	10.58	1.01	0.56	2.43	32.76	408.70	95.91
HHC018	3	6	67.19	33.78	2.26	1.41	0.74	2.66	0.48	0.23	22.04	6.36	3.83	0.45	0.19	1.48	12.70	155.80	31.11
HHC018	6	9	24.20	16.77	1.03	0.65	0.32	1.01	0.19	0.11	8.16	2.32	1.26	0.18	0.08	0.73	6.73	63.77	11.69

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC018	9	12	18.30	10.91	0.70	0.48	0.24	0.59	0.07	0.07	5.25	1.74	0.90	0.09	0.01	0.56	3.94	43.85	7.78
HHC018	12	15	19.53	11.61	0.87	0.67	0.22	0.78	0.17	0.16	6.42	1.93	1.03	0.16	0.13	0.84	4.95	49.49	9.39
HHC018	15	18	15.48	9.15	0.65	0.54	0.20	0.55	0.09	0.07	4.90	1.43	0.85	0.06	0.05	0.59	4.57	39.17	7.04
HHC018	18	21	18.55	10.67	0.76	0.50	0.20	0.62	0.09	0.07	5.83	1.66	0.89	0.07	0.06	0.44	4.06	44.48	8.32
HHC018	21	24	25.18	16.07	0.94	0.62	0.24	0.84	0.14	0.10	7.58	2.25	1.15	0.11	0.09	0.71	4.83	60.84	10.88
HHC018	24	27	18.30	9.97	0.67	0.41	0.23	0.68	0.06	0.11	5.48	1.23	0.75	0.06	0.07	0.56	3.56	42.14	7.44
HHC018	27	30	9.09	6.22	0.39	0.23	0.13	0.29	0.06	0.05	3.62	1.01	0.48	0.05	0.02	0.40	2.54	24.56	5.07
HHC018	30	33	17.44	12.08	0.72	0.46	0.19	0.63	0.11	0.10	6.77	2.10	0.97	0.08	0.06	0.55	3.94	46.20	9.67
HHC018	33	36	21.13	10.67	0.59	0.34	0.27	0.61	0.07	0.07	7.46	2.10	1.01	0.07	0.06	0.44	3.30	48.19	10.22
HHC018	36	39	52.94	33.89	1.17	0.62	0.63	1.87	0.16	0.11	22.04	6.62	3.35	0.20	0.10	0.64	5.84	130.19	30.04
HHC018	39	42	66.58	55.47	1.79	0.74	0.95	2.95	0.24	0.14	45.84	12.50	5.96	0.35	0.15	0.90	8.38	202.95	60.49
HHC018	42	45	84.64	62.39	1.66	0.61	1.02	2.79	0.25	0.13	45.14	13.05	6.31	0.31	0.14	0.96	8.13	227.51	60.16
HHC018	45	48	59.45	39.64	1.00	0.63	0.54	1.54	0.14	0.13	21.81	6.42	2.96	0.20	0.17	0.66	5.46	140.75	29.43
HHC018	48	51	127.75	66.26	1.50	0.79	1.03	2.89	0.25	0.14	40.12	11.76	5.66	0.36	0.15	0.89	8.00	267.56	53.75
HHC018	51	54	249.37	117.87	2.44	1.22	1.53	4.25	0.38	0.15	66.48	19.69	9.38	0.46	0.18	0.93	11.43	485.77	89.08
HHC018	54	57	44.47	23.34	0.67	0.39	0.31	0.92	0.11	0.11	14.35	4.26	2.08	0.13	0.05	0.57	3.17	94.93	19.41
HHC018	57	60	138.20	65.32	1.65	0.88	0.80	2.70	0.32	0.11	37.32	11.36	5.09	0.26	0.21	0.95	8.38	273.55	50.59
HHC018	60	63	77.51	42.69	1.04	0.74	0.49	1.48	0.21	0.13	21.81	6.51	2.49	0.18	0.11	0.85	6.60	162.85	29.54
HHC018	63	66	35.01	22.28	0.91	0.58	0.38	0.89	0.15	0.11	11.43	3.20	1.73	0.12	0.10	0.65	5.33	82.88	15.66
HHC018	66	69	35.87	21.23	0.95	0.53	0.32	0.98	0.16	0.10	11.31	3.42	1.73	0.13	0.11	0.60	5.71	83.16	15.82
HHC018	69	72	43.98	25.68	0.96	0.57	0.36	1.12	0.14	0.08	14.35	4.19	1.91	0.15	0.14	0.57	5.21	99.41	19.66
HHC018	72	75	93.36	58.29	1.40	0.59	0.52	1.96	0.16	0.11	28.11	8.78	3.66	0.20	0.18	0.61	7.11	205.06	38.49
HHC018	75	78	96.80	62.04	1.40	0.63	0.78	2.05	0.21	0.08	31.73	9.99	3.85	0.21	0.11	0.79	6.60	217.26	43.33
HHC018	78	81	135.12	81.51	1.80	0.78	0.89	2.82	0.25	0.14	41.64	12.57	5.50	0.34	0.14	0.74	8.51	292.75	56.35
HHC018	81	84	175.66	107.55	2.07	1.10	1.15	3.63	0.34	0.14	56.92	17.22	8.05	0.46	0.16	0.83	11.18	386.44	76.66
HHC018	84	87	93.48	58.17	1.50	0.66	0.67	2.31	0.22	0.14	31.84	9.91	5.07	0.22	0.14	0.80	7.49	212.62	43.48
HHC018	87	90	43.24	23.69	1.18	0.54	0.38	1.50	0.18	0.08	15.40	4.60	2.06	0.19	0.14	0.58	6.35	100.11	21.37
HHC018	90	92	47.78	29.55	0.98	0.59	0.36	1.43	0.18	0.10	16.68	4.64	2.28	0.16	0.10	0.75	5.46	111.07	22.46
HHC019	0	3	103.80	43.51	2.63	1.66	0.90	3.16	0.46	0.25	30.91	9.17	5.52	0.47	0.25	1.70	15.87	220.26	43.18
HHC019	3	6	35.87	20.17	1.47	0.93	0.36	1.34	0.22	0.17	11.78	3.14	1.83	0.21	0.24	0.98	7.75	86.45	16.60
HHC019	6	9	31.08	19.47	0.98	0.77	0.30	1.00	0.19	0.11	10.61	2.95	1.58	0.16	0.19	0.72	6.86	76.97	14.70
HHC019	9	12	28.13	18.06	0.91	0.50	0.34	0.71	0.16	0.08	8.98	2.49	1.25	0.13	0.17	0.65	5.08	67.64	12.51
HHC019	12	15	21.37	12.90	0.85	0.48	0.25	0.66	0.13	0.09	6.77	1.98	0.88	0.12	0.14	0.63	3.68	50.92	9.71
HHC019	15	18	12.90	7.62	0.40	0.27	0.08	0.52	0.11	0.05	3.97	1.22	0.60	0.08	0.09	0.39	2.92	31.23	5.67
HHC019	18	21	17.20	10.67	0.61	0.34	0.20	0.58	0.14	0.06	4.32	1.61	0.96	0.08	0.10	0.60	4.19	41.65	6.61
HHC019	21	24	22.23	14.19	0.60	0.38	0.23	0.63	0.10	0.08	6.42	2.19	0.88	0.09	0.15	0.52	3.43	52.13	9.29
HHC019	24	27	23.71	15.36	0.57	0.50	0.16	0.74	0.13	0.11	8.40	2.54	1.12	0.12	0.15	0.44	4.70	58.76	11.63
HHC019	27	30	18.30	11.14	0.41	0.35	0.23	0.56	0.08	0.09	5.95	1.82	0.75	0.09	0.15	0.54	2.67	43.15	8.28
HHC019	30	33	17.44	12.08	0.44	0.40	0.20	0.62	0.13	0.09	6.07	1.72	0.86	0.11	0.16	0.54	3.68	44.52	8.32
HHC019	33	36	9.34	6.57	0.37	0.16	0.15	0.28	0.07	0.07	3.50	1.05	0.46	0.08	0.10	0.43	2.79	25.42	5.00
HHC019	36	39	25.43	13.84	0.70	0.51	0.21	0.99	0.11	0.09	7.81	2.84	1.81	0.08	0.07	0.52	3.81	58.83	11.44
HHC019	39	42	41.27	28.85	1.09	0.61	0.53	1.76	0.19	0.08	19.01	5.71	2.84	0.26	0.14	0.73	5.71	108.80	26.08
HHC020	0	3	111.54	58.52	3.64	1.74	1.30	4.58	0.70	0.24	42.11	11.97	6.34	0.65	0.33	1.65	22.48	267.78	58.37
HHC020	3	6	106.38	69.78	3.62	1.53	1.12	4.58	0.62	0.19	46.31	13.47	7.29	0.59	0.21	1.48	17.14	274.31	63.98
HHC020	6	9	26.16	16.54	1.16	0.74	0.31	0.97	0.19	0.13	8.16	2.61	1.16	0.15	0.10	0.77	6.73	65.90	12.09
HHC020	9	12	19.65	12.67	0.87	0.62	0.31	0.85	0.13	0.10	6.42	1.84	1.03	0.14	0.06	0.58	5.59	50.85	9.27
HHC020	12	15	18.79	12.43	0.77	0.49	0.20	0.67	0.15	0.06	6.30	1.80	0.65	0.12	0.07	0.50	4.70	47.69	8.99
HHC020	15	18	26.90	15.72	0.67	0.38	0.16	0.85	0.16	0.07	8.86	2.49	1.12	0.08	0.09	0.50	4.19	62.25	12.10
HHC020	18	21	27.39	17.12	0.72	0.64	0.21	0.82	0.17	0.09	8.05	2.51	1.18	0.13	0.05	0.65	5.21	64.94	11.41
HHC020	21	24	35.13	24.16	0.79	0.50	0.25	1.18	0.18	0.10	10.61	3.49	1.30	0.13	0.07	0.72	4.83	83.45	15.03
HHC020	24	27	16.95	13.25	0.56	0.31	0.22	0.62	0.11	0.10	6.07	1.93	1.18	0.07	0.11	0.46	3.05	45.00	8.63
HHC020	27	30	68.42	61.57	1.78	0.86	0.93	2.70	0.38	0.19	33.83	9.69	4.43	0.33	0.21	1.14	12.57	199.02	45.62
HHC020	30	33	80.83	48.91	1.40	0.71	0.81	2.33	0.27	0.10	29.04	8.63	3.68	0.31	0.14	0.93	7.24	185.32	39.38
HHC020	33	36	214.97	160.09	4.69	2.13	2.32	7.92	0.77	0.28	98.56	29.84	13.57	0.86	0.33	1.87	24.13	562.32	133.96
HHC020	36	39	180.57	106.84	2.75	1.15	1.35	4.39	0.41	0.11	59.49	18.61	7.93	0.54	0.22	0.98	14.35	399.71	81.39
HHC020	39	43	143.72	85.85	2.44	1.29	1.20	3.82	0.46	0.16	46.07	14.74	6.09	0.44	0.24	0.87	13.84	321.23	63.69
HHC021	0	3	97.29	52.42	3.50	1.97	1.23	4.60	0.65	0.25	38.96	10.73	5.74	0.65	0.30	1.53	19.56	239.36	53.83
HHC021	3	6	156.01	102.15	5.07	2.42	1.78	6.86	1.00	0.35	70.80	20.84	10.41	0.93	0.26	1.73	27.43	408.05	97.64
HHC021	6	9	25.80	16.42	1.19	0.64	0.29	0.97	0.29	0.13	8.16	2.46	1.51	0.18	0.17	0.81	7.87	66.89	12.00

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC021	9	12	19.78	12.78	1.02	0.56	0.28	0.88	0.14	0.07	5.48	1.85	0.96	0.13	0.08	0.76	6.22	50.99	8.48
HHC021	12	15	20.76	12.78	1.11	0.58	0.17	0.78	0.17	0.08	6.53	2.07	0.66	0.14	0.13	0.52	5.21	51.70	9.85
HHC021	15	18	23.22	17.01	0.65	0.56	0.19	0.78	0.15	0.07	7.12	2.14	1.44	0.14	0.09	0.56	5.46	59.57	10.05
HHC021	18	21	20.39	12.43	0.61	0.51	0.23	0.71	0.14	0.08	5.72	1.97	1.08	0.09	0.08	0.40	3.94	48.38	8.39
HHC021	21	24	27.76	18.06	0.81	0.49	0.29	0.82	0.18	0.08	8.98	3.08	1.38	0.13	0.09	0.55	5.21	67.92	13.01
HHC021	24	27	67.32	41.40	1.12	0.55	0.46	1.84	0.25	0.14	22.16	6.91	2.99	0.20	0.11	0.81	8.00	154.27	30.40
HHC021	27	30	50.73	45.04	1.02	0.53	0.44	1.66	0.19	0.14	19.13	6.19	2.18	0.22	0.07	0.77	6.35	134.66	26.56
HHC021	30	33	17.69	11.49	0.54	0.29	0.16	0.48	0.10	0.09	6.18	1.72	0.95	0.06	0.06	0.34	3.43	43.58	8.50
HHC021	33	36	41.77	20.29	1.01	0.61	0.28	1.09	0.18	0.13	11.08	3.43	1.65	0.19	0.14	0.55	5.97	88.35	15.71
HHC021	36	39	162.15	112.35	3.53	1.93	1.68	5.42	0.64	0.27	69.40	19.75	9.04	0.65	0.31	1.80	22.73	411.67	93.34
HHC021	39	42	NS																
HHC021	42	45	99.62	64.39	1.76	1.18	0.95	3.22	0.33	0.19	42.11	12.93	5.10	0.35	0.13	1.48	11.18	244.91	57.14
HHC021	45	48	195.93	110.48	1.87	0.89	1.17	3.18	0.32	0.11	60.30	19.09	6.97	0.35	0.15	0.79	8.76	410.37	81.62
HHC021	48	51	355.01	192.93	1.95	0.93	1.49	3.71	0.30	0.10	89.11	31.78	9.43	0.41	0.18	0.72	8.51	696.55	123.25
HHC021	51	54	438.54	208.76	2.65	0.86	2.51	5.49	0.36	0.10	128.89	44.10	15.13	0.61	0.18	0.84	8.76	857.78	176.25
HHC021	54	57	75.79	35.30	1.39	1.18	0.59	1.89	0.29	0.28	24.38	7.52	3.72	0.27	0.18	1.21	11.05	165.04	33.55
HHC021	57	60	80.09	42.92	1.46	0.71	0.49	2.09	0.33	0.20	26.36	8.28	3.62	0.28	0.15	0.97	10.54	178.49	36.38
HHC022	0	3	120.75	60.05	3.70	2.15	1.27	4.60	0.69	0.26	41.64	12.44	6.81	0.69	0.31	1.84	22.99	280.19	58.47
HHC022	3	6	126.53	81.86	5.21	2.66	1.84	7.11	0.95	0.34	61.59	16.85	9.06	0.85	0.45	2.39	31.24	348.93	84.50
HHC022	6	9	38.82	22.75	1.31	0.88	0.35	1.39	0.25	0.09	11.66	3.54	1.77	0.19	0.09	0.81	7.87	91.78	16.70
HHC022	9	12	26.66	16.18	1.01	0.82	0.35	1.05	0.22	0.15	8.63	2.40	1.12	0.12	0.10	0.72	7.37	66.90	12.16
HHC022	12	15	24.94	15.13	0.99	0.61	0.25	0.88	0.15	0.14	7.58	2.51	1.03	0.15	0.16	0.75	6.10	61.36	11.23
HHC022	15	18	22.73	12.67	0.94	0.70	0.16	0.71	0.18	0.14	7.00	2.13	1.29	0.14	0.10	0.79	5.71	55.38	10.21
HHC022	18	21	26.41	17.59	0.81	0.62	0.29	0.77	0.15	0.07	8.86	2.54	1.19	0.12	0.14	0.71	4.70	64.97	12.33
HHC022	21	24	155.39	121.97	2.64	1.26	1.44	4.81	0.47	0.13	67.88	19.03	8.59	0.58	0.21	1.00	14.73	400.12	90.13
HHC022	24	27	238.92	136.63	4.05	1.98	2.13	7.30	0.76	0.30	83.40	24.53	11.89	0.79	0.33	1.98	25.65	540.63	112.76
HHC022	27	30	249.37	141.32	4.05	1.90	2.20	6.39	0.64	0.26	80.48	24.41	10.01	0.76	0.22	1.75	21.33	545.09	109.70
HHC022	30	33	244.45	131.35	3.96	1.49	2.67	7.84	0.70	0.23	89.93	25.37	12.23	0.89	0.29	1.48	21.59	544.47	120.16
HHC022	33	36	254.28	135.46	3.88	1.94	2.22	7.10	0.71	0.19	89.81	25.37	12.00	0.80	0.24	1.46	22.10	557.57	119.86
HHC023	0	3	117.56	59.46	4.24	2.20	1.26	4.83	0.85	0.24	46.31	12.99	7.60	0.71	0.37	1.89	24.64	285.11	64.23
HHC023	3	6	146.79	105.67	5.98	3.34	2.05	8.32	1.05	0.31	78.97	20.72	12.87	1.04	0.45	2.36	36.70	426.61	106.70
HHC023	6	9	41.64	19.59	1.31	0.69	0.44	1.24	0.24	0.13	10.73	3.37	1.58	0.18	0.15	0.98	7.24	89.49	15.59
HHC023	9	12	23.59	13.84	1.01	0.73	0.30	1.03	0.22	0.11	7.12	2.17	1.22	0.16	0.17	0.67	6.10	58.44	10.46
HHC023	12	15	25.31	15.25	1.17	0.69	0.29	1.12	0.23	0.10	7.93	2.45	1.08	0.16	0.14	0.68	6.48	63.07	11.72
HHC023	15	18	20.51	12.43	0.72	0.49	0.17	0.59	0.17	0.10	5.60	1.88	0.77	0.13	0.11	0.65	4.83	49.16	8.34
HHC023	18	21	34.52	26.15	0.72	0.53	0.45	1.03	0.13	0.08	13.76	3.89	1.76	0.16	0.21	0.60	5.33	89.33	18.54
HHC023	21	24	205.14	120.80	3.99	1.98	1.69	6.35	0.71	0.28	69.87	21.08	10.39	0.80	0.35	2.07	23.75	469.26	95.74
HHC023	24	27	200.84	114.23	3.12	1.59	1.56	5.69	0.53	0.19	68.12	19.75	8.49	0.68	0.31	1.64	20.45	447.20	91.68
HHC023	27	30	246.91	138.98	3.81	1.89	1.76	6.55	0.62	0.19	80.02	24.77	11.83	0.69	0.34	1.41	21.21	540.97	109.29
HHC023	30	33	230.33	130.77	3.35	1.40	1.91	5.69	0.62	0.19	76.05	22.77	10.01	0.67	0.26	1.33	19.30	504.65	102.85
HHC023	33	35	230.94	132.53	3.23	1.59	1.69	6.07	0.57	0.24	70.45	21.14	8.60	0.67	0.35	1.32	18.29	497.69	95.49
HHC024	0	3	152.94	76.11	5.14	2.81	1.57	6.89	0.89	0.28	56.69	16.13	9.23	0.95	0.46	2.40	27.81	360.32	78.91
HHC024	3	6	126.53	79.16	5.05	2.66	1.55	6.64	0.96	0.31	57.15	16.73	9.58	0.86	0.47	2.14	28.83	338.62	79.80
HHC024	6	9	28.13	17.94	1.11	0.82	0.28	1.03	0.24	0.10	9.21	2.61	1.48	0.14	0.19	0.90	6.73	70.93	13.08
HHC024	9	12	54.30	31.31	1.21	0.80	0.34	1.58	0.27	0.14	17.38	5.32	2.26	0.16	0.18	0.87	7.87	123.98	24.07
HHC024	12	15	25.92	15.72	0.95	0.69	0.23	0.75	0.17	0.09	8.40	2.73	1.39	0.16	0.17	0.69	6.48	64.54	12.25
HHC024	15	18	89.55	64.50	1.56	0.99	0.65	2.16	0.25	0.16	29.74	9.29	3.55	0.26	0.21	1.08	9.02	212.97	40.85
HHC024	18	22	207.60	114.70	3.47	1.53	1.68	5.54	0.57	0.17	72.78	21.02	10.17	0.60	0.27	1.49	17.65	459.26	97.87
HHC025	0	3	138.20	68.26	4.28	2.28	1.38	6.03	0.84	0.28	51.79	15.22	8.79	0.86	0.32	2.05	25.02	325.58	72.15
HHC025	3	6	213.13	92.18	6.32	3.20	2.28	8.28	1.15	0.39	74.30	21.45	12.93	1.18	0.43	2.43	33.14	472.78	103.25
HHC025	6	9	49.26	20.05	1.42	0.80	0.39	1.28	0.30	0.14	11.55	3.52	1.67	0.24	0.14	0.97	8.51	100.23	16.72
HHC025	9	12	29.24	16.77	1.04	0.83	0.30	1.05	0.26	0.15	8.98	2.82	1.51	0.16	0.13	0.97	8.13	72.34	13.01
HHC025	12	15	22.48	12.67	0.90	0.63	0.23	0.81	0.18	0.14	6.53	2.08	0.97	0.14	0.10	0.81	5.84	54.51	9.65
HHC025	15	18	20.76	11.61	0.61	0.39	0.14	0.68	0.13	0.14	4.90	1.80	0.92	0.09	0.08	0.65	3.68	46.57	7.40
HHC025	18	21	18.55	12.90	0.57	0.38	0.22	0.73	0.11	0.10	5.95	1.93	0.77	0.09	0.08	0.63	3.56	46.57	8.55
HHC025	21	24	37.71	30.02	0.83	0.45	0.52	1.09	0.15	0.11	15.05	4.92	1.95	0.16	0.07	0.59	4.19	97.81	20.95
HHC025	24	27	380.80	215.80	5.99	2.65	3.49	11.39	1.08	0.40	158.63	46.88	22.44	1.32	0.40	2.48	27.68	881.42	212.82
HHC025	27	30	272.70	158.91	4.64	2.13	2.29	7.72	0.76	0.31	96.69	29.48	13.51	0.89	0.38	1.95	25.78	618.14	131.71
HHC025	30	33	128.37	72.48	1.63	0.95	0.90	3.42	0.34	0.09	41.64	12.69	5.51	0.33	0.15	0.95	10.41	279.86	56.29

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC025	33	37	264.11	140.15	3.91	1.86	2.26	7.01	0.68	0.22	92.03	26.82	11.32	0.76	0.32	1.55	21.59	574.58	123.53
HHC026	0	3	150.48	72.24	4.77	2.46	1.45	6.06	0.79	0.27	53.19	15.34	8.00	0.78	0.40	2.28	25.14	343.66	74.08
HHC026	3	6	124.68	73.30	4.75	2.47	1.51	5.97	0.86	0.26	53.89	14.44	8.64	0.80	0.37	2.12	29.97	324.02	73.88
HHC026	6	9	31.82	18.53	1.19	0.99	0.42	1.29	0.23	0.15	9.21	3.19	1.65	0.21	0.15	0.95	7.87	77.85	13.81
HHC026	9	12	24.81	15.25	0.77	0.78	0.28	0.86	0.18	0.11	7.00	2.42	1.30	0.13	0.14	0.68	5.84	60.55	10.31
HHC026	12	15	24.32	13.84	0.99	0.69	0.22	0.90	0.22	0.10	8.28	2.39	1.23	0.13	0.22	0.65	5.59	59.76	11.79
HHC026	15	18	56.63	31.20	1.92	1.12	0.59	2.62	0.33	0.19	20.06	6.23	2.76	0.33	0.32	1.13	11.68	137.11	28.54
HHC026	18	21	22.73	19.70	1.18	0.67	0.16	1.15	0.17	0.14	6.88	2.34	0.77	0.19	0.19	0.89	7.62	64.79	10.60
HHC026	21	24	14.50	9.97	0.76	0.55	0.23	0.65	0.14	0.11	4.55	1.57	0.82	0.08	0.11	0.72	5.33	40.09	6.96
HHC026	24	27	39.55	29.20	0.92	0.51	0.46	1.13	0.17	0.16	15.16	4.31	2.24	0.15	0.11	0.76	4.95	99.81	20.55
HHC026	27	31	267.79	150.12	4.73	2.53	2.48	7.56	0.77	0.30	92.26	27.18	14.90	0.85	0.35	2.16	26.54	600.52	125.02
HHC027	0	3	123.45	61.92	4.32	2.02	1.39	5.07	0.79	0.27	46.07	13.35	8.15	0.71	0.27	2.07	25.52	295.39	64.44
HHC027	3	6	119.03	48.20	3.27	1.49	0.90	3.41	0.56	0.23	33.01	9.54	6.19	0.45	0.32	1.57	17.40	245.58	46.27
HHC027	6	9	31.69	18.53	1.31	0.79	0.34	1.16	0.22	0.15	10.26	2.92	1.47	0.18	0.16	0.97	7.49	77.64	14.67
HHC027	9	12	26.66	16.30	1.17	0.85	0.36	1.09	0.24	0.14	8.51	2.67	1.51	0.15	0.13	1.07	7.11	67.96	12.51
HHC027	12	15	23.09	12.90	0.73	0.61	0.20	0.82	0.15	0.09	6.88	2.14	1.18	0.14	0.11	0.75	4.95	54.75	9.90
HHC027	15	18	19.16	11.49	0.64	0.56	0.20	0.46	0.14	0.13	5.95	1.93	1.04	0.07	0.11	0.46	4.19	46.54	8.60
HHC027	18	21	17.93	11.02	0.63	0.50	0.15	0.61	0.15	0.07	5.83	1.73	0.93	0.11	0.15	0.54	3.81	44.16	8.30
HHC027	21	24	32.06	25.92	0.77	0.50	0.42	0.81	0.15	0.09	14.00	3.65	1.77	0.09	0.10	0.60	4.83	85.76	18.51
HHC027	24	27	18.43	14.66	0.60	0.37	0.22	0.67	0.11	0.07	6.30	1.90	1.29	0.11	0.11	0.35	3.94	49.11	8.90
HHC027	27	30	13.76	10.91	0.59	0.49	0.15	0.62	0.15	0.06	4.20	1.26	0.93	0.12	0.08	0.41	3.94	37.65	6.16
HHC027	30	33	27.52	17.71	0.70	0.56	0.29	0.96	0.13	0.11	8.98	2.78	1.23	0.13	0.11	0.57	4.57	66.35	12.59
HHC027	33	36	75.67	32.60	1.37	0.64	0.43	1.61	0.19	0.10	19.36	5.88	3.25	0.19	0.13	0.69	6.22	148.34	26.80
HHC027	36	39	140.04	105.55	2.69	1.52	1.57	4.78	0.49	0.17	68.93	20.78	9.46	0.53	0.18	1.46	14.10	372.26	92.93
HHC027	39	42	68.54	48.32	1.43	0.81	0.97	2.47	0.23	0.16	35.69	9.61	4.65	0.27	0.14	1.01	7.37	181.67	47.00
HHC027	42	45	43.85	28.26	1.15	0.73	0.50	1.81	0.18	0.13	18.66	5.35	3.14	0.19	0.18	0.68	7.62	112.44	25.35
HHC027	45	48	228.48	137.22	5.04	2.29	2.13	7.71	0.80	0.30	89.11	25.98	13.51	0.89	0.39	2.39	28.57	544.81	121.02
HHC027	48	51	169.52	96.17	3.45	1.58	1.44	4.97	0.53	0.23	58.55	17.10	9.18	0.60	0.30	1.57	21.97	387.15	79.70
HHC027	51	53	222.95	124.32	3.66	1.91	1.81	5.94	0.62	0.19	73.25	22.59	9.71	0.59	0.31	1.51	19.81	489.17	100.09
HHC028	0	3	117.68	64.86	4.29	2.25	1.20	5.49	0.71	0.18	47.47	13.23	8.48	0.71	0.31	1.82	22.86	291.54	65.70
HHC028	3	6	85.74	34.95	1.97	1.07	0.63	2.49	0.32	0.19	20.18	6.31	3.85	0.35	0.26	1.14	11.43	170.89	28.81
HHC028	6	9	28.62	17.36	0.99	0.66	0.31	1.08	0.24	0.10	9.80	2.86	1.41	0.13	0.14	0.99	7.37	72.07	13.78
HHC028	9	12	24.08	14.89	0.93	0.71	0.20	0.86	0.15	0.09	7.70	2.15	1.16	0.14	0.13	0.84	6.60	60.63	10.92
HHC028	12	15	28.13	16.42	0.88	0.62	0.20	0.84	0.16	0.10	8.40	2.68	1.08	0.12	0.17	0.75	5.97	66.52	12.08
HHC028	15	18	29.48	16.42	0.77	0.49	0.30	0.84	0.15	0.09	8.40	2.62	1.58	0.13	0.06	0.58	5.08	66.99	11.92
HHC028	18	21	28.38	15.48	0.77	0.47	0.22	0.88	0.15	0.11	9.33	2.72	1.55	0.14	0.16	0.72	5.08	66.15	12.96
HHC028	21	24	32.31	20.05	0.95	0.61	0.29	1.05	0.14	0.11	10.26	3.19	1.55	0.13	0.10	0.64	5.33	76.72	14.54
HHC028	24	27	35.87	22.64	1.14	0.75	0.36	1.29	0.24	0.15	11.90	3.73	2.02	0.15	0.14	0.85	6.73	87.96	16.92
HHC028	27	30	43.73	34.60	0.86	0.55	0.35	1.35	0.15	0.11	16.80	5.04	2.52	0.19	0.07	0.72	5.97	112.99	22.88
HHC028	30	33	35.87	22.52	0.87	0.35	0.19	1.14	0.15	0.14	10.38	3.26	1.60	0.12	0.14	0.50	4.57	81.80	14.63
HHC028	33	36	29.60	17.59	0.65	0.46	0.28	0.77	0.14	0.08	9.56	2.95	1.32	0.07	0.08	0.59	3.68	67.84	13.24
HHC028	36	39	87.22	36.47	1.14	0.71	0.60	1.96	0.19	0.13	25.08	7.16	3.97	0.24	0.11	0.73	7.37	173.07	33.61
HHC028	39	42	205.14	108.95	2.58	1.28	1.47	4.41	0.48	0.14	66.95	20.30	9.16	0.49	0.16	1.13	13.71	436.37	90.33
HHC028	42	45	79.85	46.56	1.73	0.89	0.69	2.28	0.27	0.11	30.09	8.55	4.16	0.28	0.14	0.96	9.65	186.23	40.66
HHC028	45	48	116.82	57.12	2.57	1.43	0.97	3.20	0.40	0.18	37.56	10.72	5.45	0.39	0.18	1.28	12.19	250.46	51.23
HHC028	48	51	114.98	64.74	2.54	1.19	1.37	4.35	0.48	0.24	53.19	14.86	7.86	0.46	0.29	1.31	13.21	281.05	71.04
HHC028	51	54	234.62	106.14	3.06	1.52	1.71	5.24	0.52	0.20	71.27	20.66	9.94	0.60	0.25	1.53	15.11	472.38	95.59
HHC028	54	57	114.73	55.36	1.71	0.96	0.78	2.56	0.27	0.11	37.56	10.62	5.68	0.26	0.15	0.92	8.25	239.93	50.15
HHC028	57	60	73.34	39.76	1.26	0.74	0.44	1.54	0.26	0.09	24.84	7.48	3.24	0.15	0.21	0.71	6.10	160.16	33.74
HHC028	60	63	144.95	81.04	1.65	0.88	0.94	2.67	0.24	0.14	46.19	14.80	6.76	0.28	0.21	0.85	6.98	308.59	62.92
HHC028	63	66	325.53	167.12	4.05	1.56	2.76	7.40	0.56	0.18	127.72	36.61	17.28	0.76	0.26	1.50	17.40	710.69	169.15
HHC028	66	69	198.39	91.24	3.37	1.75	2.11	5.97	0.56	0.20	82.46	23.50	12.00	0.71	0.27	1.57	17.02	441.13	110.04
HHC028	69	72	34.15	17.94	0.72	0.69	0.20	1.09	0.15	0.18	12.48	3.64	1.92	0.14	0.15	0.77	7.24	81.47	16.98
HHC028	72	75	200.84	104.50	3.27	1.64	1.51	5.24	0.47	0.17	68.47	19.87	10.08	0.55	0.25	1.13	17.27	435.26	92.17
HHC028	75	79	236.47	124.90	3.72	1.74	1.64	5.99	0.61	0.20	79.55	23.20	11.39	0.59	0.24	1.66	19.81	511.71	107.05
HHC029	0	3	116.08	55.83	3.73	2.34	1.23	4.86	0.66	0.20	44.44	12.38	7.73	0.62	0.31	1.65	19.68	271.77	61.18
HHC029	3	6	130.21	71.19	4.03	2.31	1.35	5.08	0.71	0.27	51.67	14.56	7.94	0.72	0.31	1.80	22.48	314.63	70.98
HHC029	6	9	26.78	15.60	1.04	0.45	0.27	0.98	0.17	0.11	8.40	2.55	1.12	0.13	0.11	0.64	5.84	64.19	12.12
HHC029	9	12	21.87	12.67	0.76	0.45	0.21	0.66	0.15	0.08	6.77	1.91	1.44	0.09	0.15	0.63	5.08	52.89	9.53

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC029	12	15	22.85	11.85	0.91	0.62	0.20	0.85	0.17	0.08	7.23	2.03	1.28	0.09	0.11	0.65	5.71	54.63	10.26
HHC029	15	18	34.64	17.47	0.91	0.58	0.24	0.90	0.16	0.06	10.15	3.08	1.79	0.13	0.08	0.58	4.95	75.72	14.26
HHC029	18	21	22.36	12.78	0.63	0.33	0.15	0.51	0.13	0.01	6.42	2.07	0.99	0.08	0.11	0.30	4.06	50.92	9.19
HHC029	21	24	38.69	22.87	1.01	0.59	0.20	1.18	0.15	0.10	12.36	3.60	1.75	0.13	0.11	0.49	5.33	88.57	17.10
HHC029	24	27	53.07	34.83	1.19	0.89	0.34	1.90	0.22	0.08	19.83	6.05	2.75	0.24	0.17	0.82	8.00	130.38	27.31
HHC029	27	30	43.36	29.20	1.04	0.62	0.24	1.27	0.16	0.09	16.10	4.76	2.27	0.15	0.13	0.68	5.59	105.67	22.05
HHC029	30	33	97.66	40.81	1.43	0.98	0.53	2.25	0.25	0.07	24.38	7.09	3.47	0.22	0.11	0.92	7.37	187.55	33.13
HHC029	33	36	99.13	30.61	1.11	0.43	0.32	1.19	0.21	0.06	16.33	4.68	2.42	0.12	0.15	0.58	6.48	163.82	22.24
HHC029	36	39	255.51	116.22	3.41	1.94	1.41	4.38	0.63	0.28	71.62	21.99	9.50	0.64	0.23	1.87	19.94	509.56	97.65
HHC029	39	42	122.23	68.02	5.22	3.88	0.81	4.33	1.11	0.73	35.69	11.72	5.50	0.66	0.62	4.60	37.34	302.45	53.29
HHC029	42	45	80.95	56.88	4.06	3.01	0.61	3.24	0.82	0.45	26.36	8.75	4.19	0.59	0.43	3.04	33.27	226.66	39.76
HHC029	45	49	595.77	314.31	4.58	1.93	2.73	7.56	0.66	0.20	158.05	54.97	18.44	0.81	0.27	1.71	20.70	1182.71	218.41
HHC030	0	3	141.88	72.01	4.11	2.08	1.52	4.91	0.60	0.25	48.64	14.86	7.34	0.64	0.18	1.94	22.60	323.55	68.24
HHC030	3	6	191.02	113.88	5.03	2.63	1.98	7.13	0.81	0.30	73.48	23.50	10.47	0.87	0.30	2.07	29.72	463.18	102.88
HHC030	6	9	62.16	35.65	1.88	1.09	0.37	1.88	0.30	0.17	19.13	6.81	3.04	0.26	0.15	0.92	11.43	145.24	28.08
HHC030	9	12	37.83	22.28	1.00	0.63	0.31	1.34	0.22	0.11	12.13	3.83	1.73	0.19	0.14	0.82	7.75	90.31	17.15
HHC030	12	15	28.13	15.72	1.26	0.69	0.28	0.89	0.19	0.13	9.33	2.74	1.31	0.20	0.13	0.81	6.98	68.78	13.54
HHC030	15	18	26.04	14.07	0.75	0.61	0.17	0.84	0.16	0.08	7.00	2.39	0.86	0.11	0.08	0.58	5.59	59.33	10.24
HHC030	18	21	33.41	20.05	0.77	0.59	0.31	0.92	0.17	0.13	9.33	3.13	1.45	0.15	0.01	0.69	4.95	76.08	13.38
HHC030	21	24	24.94	16.54	0.62	0.53	0.16	0.68	0.11	0.13	8.75	2.72	1.35	0.07	0.06	0.51	4.06	61.22	12.16
HHC030	24	27	32.43	27.44	0.94	0.51	0.29	1.23	0.16	0.09	11.78	4.12	1.62	0.16	0.10	0.58	5.97	87.44	17.01
HHC030	27	30	33.66	19.47	0.87	0.96	0.27	0.99	0.16	0.13	10.26	3.07	1.92	0.18	0.13	0.89	5.71	78.67	14.38
HHC030	30	33	151.71	105.79	2.33	1.09	1.17	2.99	0.34	0.17	44.67	15.53	5.37	0.39	0.14	1.23	10.92	343.82	62.92
HHC030	33	36	418.88	294.37	7.59	3.33	3.72	13.20	1.19	0.33	182.54	53.64	22.55	1.41	0.37	2.68	39.87	1045.67	245.18
HHC030	36	40	334.12	202.31	5.46	2.76	2.89	9.52	0.97	0.25	125.97	39.15	16.70	1.01	0.33	1.86	29.84	773.15	171.59
HHC031	0	3	118.91	61.81	3.73	2.31	1.25	4.56	0.71	0.26	41.99	12.50	7.06	0.64	0.33	2.00	20.95	279.02	58.86
HHC031	3	6	140.65	211.69	8.92	3.61	3.20	13.66	1.37	0.34	132.97	39.51	21.45	1.65	0.43	2.84	39.11	621.40	183.04
HHC031	6	9	35.75	21.46	1.42	0.77	0.29	1.27	0.23	0.14	10.38	3.25	1.46	0.20	0.16	0.88	8.13	85.78	15.25
HHC031	9	12	33.04	22.40	1.12	0.74	0.29	1.16	0.26	0.10	11.78	3.46	1.47	0.16	0.05	0.76	7.75	84.56	16.53
HHC031	12	15	28.38	16.18	0.94	0.74	0.28	1.24	0.21	0.15	8.51	2.68	1.77	0.19	0.16	0.93	6.60	68.98	12.33
HHC031	15	18	26.90	13.72	0.80	0.59	0.23	0.84	0.16	0.06	6.88	2.59	1.18	0.14	0.03	0.61	4.83	59.58	10.41
HHC031	18	21	148.02	80.92	1.79	1.02	0.85	3.09	0.30	0.15	38.49	11.95	5.73	0.39	0.18	0.85	10.54	304.27	52.62
HHC031	21	22	168.29	146.01	3.11	1.50	1.77	5.33	0.52	0.17	87.71	25.73	12.06	0.66	0.22	1.10	17.27	471.45	117.22
HHC032	0	3	144.34	78.46	4.79	2.46	1.48	6.34	0.99	0.30	57.97	17.22	9.60	0.85	0.32	2.25	29.46	356.82	80.82
HHC032	3	6	182.42	99.10	5.23	2.74	1.85	7.05	0.86	0.34	69.52	20.78	9.86	0.93	0.31	2.10	29.21	432.30	96.46
HHC032	6	9	34.40	19.94	1.02	0.65	0.31	1.08	0.27	0.10	10.61	3.43	1.86	0.19	0.08	0.84	6.48	81.27	15.26
HHC032	9	12	26.90	16.54	0.91	0.66	0.39	1.07	0.17	0.13	8.05	2.71	1.17	0.18	0.11	0.74	6.10	65.82	11.84
HHC032	12	15	23.22	14.43	1.06	0.80	0.30	0.89	0.19	0.14	6.88	2.11	1.23	0.13	0.17	0.85	6.73	59.13	10.18
HHC032	15	18	25.55	16.42	0.87	0.62	0.28	0.85	0.18	0.10	6.88	2.37	1.10	0.11	0.06	0.59	5.21	61.19	10.23
HHC032	18	21	135.74	89.37	1.79	1.02	1.23	3.50	0.31	0.14	49.81	16.55	6.95	0.51	0.08	0.68	10.16	317.82	68.65
HHC032	21	23	238.92	137.80	3.31	1.62	1.81	5.94	0.55	0.23	71.62	24.41	10.33	0.68	0.22	1.37	19.05	517.85	100.01
HHC033	0	3	215.58	105.08	5.26	2.98	1.99	7.37	0.95	0.38	70.10	21.93	11.34	1.05	0.38	2.29	31.87	478.55	98.33
HHC033	3	6	118.91	40.34	3.32	1.86	0.96	3.79	0.63	0.30	27.53	8.61	4.72	0.55	0.32	1.91	18.16	231.92	40.01
HHC033	6	8	69.28	42.10	1.65	0.90	0.59	2.34	0.30	0.15	24.14	7.73	3.54	0.34	0.16	1.01	9.27	163.52	33.87
HHC034	0	3	114.00	46.68	3.93	2.66	0.98	4.45	0.65	0.36	33.36	9.63	6.20	0.67	0.40	2.32	22.73	249.03	47.58
HHC034	3	6	156.62	131.94	3.26	1.42	1.75	5.13	0.54	0.22	80.71	26.34	11.27	0.64	0.24	1.18	17.27	438.53	110.95
HHC034	6	9	353.78	249.81	7.20	3.56	3.54	12.56	1.26	0.38	152.80	47.36	21.74	1.41	0.39	2.97	38.99	897.74	208.77
HHC034	9	11	253.05	144.25	3.87	1.64	1.83	6.83	0.64	0.22	85.96	27.43	12.47	0.87	0.22	1.48	21.33	562.09	118.13
HHC035	0	3	117.93	58.05	4.06	2.14	1.33	4.96	0.69	0.32	37.44	11.07	5.81	0.66	0.37	1.88	23.11	269.81	53.23
HHC035	3	7	334.12	227.52	7.39	3.70	3.75	12.33	1.42	0.39	150.47	46.15	22.55	1.45	0.49	2.93	40.76	855.44	205.46
HHC036	0	3	105.03	56.65	2.20	1.49	1.17	3.37	0.45	0.22	35.34	9.97	5.08	0.44	0.19	1.13	11.43	234.14	47.95
HHC036	3	6	176.89	105.90	2.81	0.95	1.55	5.58	0.42	0.16	67.53	19.87	9.01	0.60	0.16	0.91	10.16	402.52	90.82
HHC036	6	9	389.40	214.04	6.82	2.70	3.80	12.33	1.07	0.32	141.13	43.74	19.89	1.42	0.31	2.36	31.62	870.94	193.11
HHC036	9	12	340.27	177.68	6.35	2.60	2.93	10.75	1.02	0.36	124.80	37.70	18.03	1.24	0.27	2.41	33.65	760.06	170.08
HHC036	12	15	265.33	141.32	8.18	4.54	2.83	11.70	1.44	0.48	100.43	29.60	15.60	1.43	0.53	3.75	51.94	639.09	139.65
HHC036	15	19	273.93	154.22	4.51	2.66	2.35	8.05	1.01	0.28	90.98	29.60	12.70	0.95	0.19	1.90	33.02	616.36	126.04
HHC037	0	3	284.99	111.65	3.16	1.68	1.31	4.52	0.52	0.18	54.59	17.88	7.87	0.60	0.19	1.36	14.86	505.35	76.22
HHC037	3	6	280.08	151.88	2.97	1.43	1.46	5.04	0.48	0.16	72.43	23.20	9.02	0.56	0.23	1.00	15.62	565.56	99.17
HHC037	6	9	113.14	68.96	1.57	0.78	0.63	2.37	0.27	0.11	28.58	9.03	3.66	0.31	0.11	0.99	9.40	239.91	39.48

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC037	9	12	232.78	116.81	2.40	1.46	1.47	4.71	0.45	0.23	69.87	21.57	9.83	0.52	0.25	1.42	13.21	476.98	94.35
<b>HHC037</b>	<b>12</b>	<b>15</b>	<b>524.53</b>	<b>262.71</b>	<b>6.86</b>	<b>2.90</b>	<b>4.49</b>	<b>14.18</b>	<b>1.03</b>	<b>0.36</b>	<b>193.04</b>	<b>53.64</b>	<b>27.25</b>	<b>1.43</b>	<b>0.43</b>	<b>2.64</b>	<b>31.11</b>	<b>1126.62</b>	<b>254.98</b>
HHC037	15	18	350.09	172.99	5.15	2.22	3.05	10.26	0.78	0.32	124.22	35.28	17.86	1.20	0.40	2.11	25.27	751.19	165.85
HHC037	18	21	266.56	132.53	6.90	3.13	3.06	12.39	1.21	0.36	99.49	26.82	16.99	1.53	0.45	2.78	31.87	606.08	134.74
HHC037	21	24	310.79	154.81	6.21	2.96	2.43	9.87	1.02	0.40	99.84	29.96	14.73	1.24	0.45	3.04	33.27	671.01	137.25
HHC037	24	27	310.79	151.88	5.52	3.19	2.80	9.69	1.08	0.43	108.24	31.41	15.31	1.04	0.47	2.98	38.10	682.92	146.21
HHC037	27	30	312.01	157.16	4.74	2.38	2.52	7.98	0.81	0.32	101.94	30.08	13.80	0.88	0.34	2.14	28.32	665.43	137.65
HHC037	30	33	357.46	186.48	5.68	2.74	2.92	9.92	0.88	0.36	120.14	34.80	16.35	1.12	0.41	2.30	29.33	770.90	161.73
HHC037	33	36	280.08	142.50	4.42	2.18	2.35	7.75	0.73	0.31	93.78	25.98	13.68	0.94	0.34	1.99	25.27	602.29	125.11
HHC038	0	3	64.49	33.78	2.08	1.14	0.63	2.48	0.30	0.16	22.39	6.26	3.61	0.38	0.22	1.07	11.94	150.91	31.11
HHC038	3	6	61.17	37.85	1.42	0.79	0.42	1.86	0.27	0.14	18.43	5.94	2.49	0.28	0.16	0.73	8.00	139.76	26.08
<b>HHC038</b>	<b>6</b>	<b>9</b>	<b>733.35</b>	<b>443.32</b>	<b>8.94</b>	<b>3.80</b>	<b>5.34</b>	<b>17.87</b>	<b>1.41</b>	<b>0.34</b>	<b>241.44</b>	<b>73.70</b>	<b>31.43</b>	<b>1.95</b>	<b>0.46</b>	<b>2.71</b>	<b>35.81</b>	<b>1601.86</b>	<b>326.04</b>
HHC038	9	12	459.42	248.63	6.42	2.30	3.00	10.67	1.01	0.26	139.97	43.86	19.66	1.18	0.26	1.59	24.89	963.11	191.42
HHC038	12	15	359.92	198.20	5.93	2.70	3.22	10.74	1.04	0.26	132.39	38.54	19.54	1.16	0.27	2.16	29.59	805.68	178.03
HHC038	15	18	325.53	184.13	5.15	2.57	2.54	9.12	0.89	0.26	102.06	31.90	13.97	1.05	0.33	1.94	26.79	708.23	140.16
HHC038	18	21	309.56	165.36	4.64	2.06	2.17	7.64	0.78	0.17	93.43	28.63	12.00	0.86	0.37	1.55	25.02	654.23	127.56
HHC038	21	24	357.46	186.48	6.00	2.95	2.79	10.00	1.02	0.33	124.80	36.85	17.28	1.22	0.41	2.55	34.54	784.70	168.88
HHC038	24	27	296.04	154.81	6.66	3.50	2.42	9.44	1.20	0.52	105.09	30.93	15.13	1.15	0.48	3.48	45.84	676.71	143.83
HHC038	27	31	264.11	140.74	4.61	2.37	2.35	8.05	0.84	0.31	91.80	26.70	13.92	0.92	0.27	1.89	27.68	586.54	124.03
HHC039	0	3	61.05	22.64	1.56	0.90	0.49	1.87	0.32	0.19	15.63	4.46	2.63	0.26	0.17	0.96	10.67	123.79	21.91
HHC039	3	6	36.61	21.58	0.64	0.38	0.25	0.89	0.10	0.05	10.26	3.36	1.12	0.12	0.02	0.41	3.56	79.35	14.38
HHC039	6	9	58.35	39.76	1.06	0.50	0.43	1.28	0.17	0.06	16.56	5.61	2.15	0.19	0.01	0.64	4.83	131.58	23.41
HHC039	9	12	170.75	129.59	2.23	0.85	0.78	2.95	0.31	0.13	45.14	17.10	4.88	0.35	0.16	0.92	8.38	384.51	64.82
HHC039	12	15	107.85	67.55	1.49	0.55	0.43	1.43	0.21	0.10	25.78	9.39	2.95	0.21	0.09	0.69	6.48	225.20	36.87
HHC039	15	18	205.76	120.80	2.18	0.85	0.82	3.38	0.32	0.15	58.67	20.66	6.15	0.40	0.11	0.95	10.16	431.34	81.91
HHC039	18	21	224.18	122.56	2.26	1.17	1.49	5.05	0.40	0.20	75.35	24.04	9.33	0.49	0.16	0.88	12.70	480.27	102.15
HHC039	21	24	211.28	108.84	2.94	1.37	1.48	4.51	0.52	0.17	64.04	22.35	8.19	0.62	0.19	1.31	15.87	443.68	89.95
HHC039	24	27	122.23	62.51	2.73	1.40	1.11	3.38	0.41	0.15	38.49	12.32	5.25	0.45	0.22	1.39	15.11	267.14	53.99
HHC039	27	30	129.60	67.20	2.55	1.19	1.07	3.05	0.40	0.24	41.29	13.23	5.68	0.49	0.17	1.24	13.46	280.86	57.56
HHC040	3	6	87.34	46.44	2.32	1.18	0.96	2.97	0.46	0.16	28.69	8.69	4.48	0.46	0.17	1.21	14.73	200.25	40.16
HHC040	0	3	99.62	33.19	1.78	1.29	0.50	2.43	0.39	0.22	20.76	6.68	3.49	0.29	0.24	1.65	12.70	185.24	29.52
HHC041	0	3	59.45	32.13	1.40	1.12	0.66	1.83	0.34	0.15	18.08	5.96	2.79	0.26	0.14	1.10	10.92	136.35	25.69
HHC041	3	6	101.59	54.65	2.09	1.33	1.23	2.64	0.37	0.23	31.84	9.92	5.10	0.36	0.18	1.15	12.70	225.38	44.22
HHC041	6	9	100.73	53.95	2.17	1.28	0.94	2.67	0.31	0.18	30.56	10.40	4.93	0.32	0.26	1.18	12.45	222.33	43.45
HHC041	9	13	127.14	66.50	2.41	1.38	1.04	3.50	0.46	0.11	37.91	12.38	5.84	0.47	0.22	1.39	15.75	276.51	53.17
HHC042	0	3	96.68	47.50	2.52	1.46	0.97	3.39	0.48	0.23	29.63	9.23	4.84	0.51	0.23	1.50	16.25	215.42	41.89
HHC042	3	7	89.67	55.83	1.69	0.75	0.67	2.12	0.30	0.16	28.23	8.78	3.43	0.27	0.13	0.98	11.18	204.18	38.97
HHC043	0	3	163.38	75.41	4.94	2.78	1.59	6.01	0.84	0.45	50.74	15.22	8.72	0.74	0.42	2.73	29.97	363.93	71.64
HHC043	3	6	35.75	21.11	1.35	0.89	0.27	1.24	0.26	0.14	11.31	3.40	1.75	0.16	0.08	0.72	8.00	86.44	16.23
HHC043	6	9	33.04	19.12	1.02	0.66	0.22	1.05	0.18	0.13	10.03	3.07	1.44	0.16	0.13	0.81	5.59	76.65	14.29
HHC043	9	12	178.12	94.41	3.10	1.72	1.52	5.15	0.48	0.20	58.20	18.67	8.21	0.60	0.18	1.49	17.65	389.70	80.57
HHC044	0	3	121.98	59.58	3.27	1.61	1.30	4.39	0.57	0.19	40.24	12.69	6.01	0.59	0.30	1.87	20.19	274.77	56.79
HHC044	3	6	40.29	21.81	1.30	0.73	0.36	1.41	0.26	0.15	10.96	3.70	1.69	0.24	0.21	0.95	7.24	91.29	16.19
HHC044	6	9	22.97	14.07	1.00	0.66	0.19	0.84	0.24	0.14	7.70	2.37	1.30	0.19	0.24	0.83	7.24	59.97	11.25
HHC044	9	13	102.08	62.86	2.03	1.21	0.88	3.47	0.38	0.16	38.26	11.71	5.93	0.39	0.21	0.95	10.03	240.53	52.38
HHC045	0	3	184.87	80.22	4.94	2.57	1.69	6.95	0.85	0.33	57.50	18.24	10.42	0.94	0.39	2.03	28.45	400.39	81.62
HHC045	3	6	33.41	18.76	1.19	0.79	0.42	1.08	0.24	0.14	10.73	3.55	1.72	0.15	0.13	0.81	6.98	80.11	15.63
HHC045	6	9	26.90	15.83	1.06	0.70	0.43	1.04	0.16	0.09	7.93	2.54	1.84	0.13	0.11	0.68	5.46	64.91	11.65
HHC045	9	12	38.57	20.17	1.35	1.19	0.30	1.34	0.32	0.16	8.98	3.32	1.67	0.25	0.21	1.37	10.03	89.23	13.91
HHC045	12	15	93.48	47.97	1.71	0.77	0.56	2.52	0.31	0.15	25.78	8.66	4.05	0.28	0.18	0.85	8.00	195.27	36.43
HHC045	15	17	287.45	150.70	3.88	1.92	1.88	7.09	0.71	0.15	91.80	29.48	12.18	0.81	0.32	1.70	21.84	611.89	125.97
HHC046	0	3	105.15	51.49	3.95	2.13	1.02	4.84	0.79	0.22	38.49	11.96	6.80	0.67	0.32	1.97	23.87	253.66	55.07
HHC046	3	6	167.06	90.31	5.36	2.50	1.59	6.78	0.99	0.26	62.52	19.33	8.96	0.92	0.35	1.99	25.65	394.57	88.13
HHC046	6	9	37.83	19.23	1.11	0.61	0.34	1.29	0.22	0.11	9.21	3.65	1.41	0.16	0.09	0.87	6.60	82.75	14.14
HHC046	9	12	21.50	13.60	0.83	0.69	0.28	0.98	0.24	0.11	6.88	2.13	1.18	0.15	0.13	0.80	6.98	56.48	9.99
HHC046	12	15	21.62	13.02	0.85	0.61	0.22	0.89	0.14	0.06	6.53	1.90	0.72	0.07	0.07	0.59	5.21	52.48	9.35
HHC046	15	18	26.78	14.19	0.86	0.51	0.21	0.88	0.15	0.06	7.35	2.25	1.35	0.11	0.08	0.51	4.83	60.10	10.56
HHC046	18	21	31.57	17.94	1.08	0.67	0.27	0.98	0.19	0.13	9.21	2.69	1.28	0.12	0.14	0.77	6.73	73.78	13.11
HHC046	21	23	162.15	153.05	3.81	1.77	1.71	5.87	0.61	0.24	87.25	26.22	9.96	0.72	0.18	1.24	22.86	477.63	117.99

Hole ID	From	To	CeO <sub>2</sub>	La <sub>2</sub> O <sub>3</sub>	Dy <sub>2</sub> O <sub>3</sub>	Er <sub>2</sub> O <sub>3</sub>	Eu <sub>2</sub> O <sub>3</sub>	Gd <sub>2</sub> O <sub>3</sub>	Ho <sub>2</sub> O <sub>3</sub>	Lu <sub>2</sub> O <sub>3</sub>	Nd <sub>2</sub> O <sub>3</sub>	Pr <sub>6</sub> O <sub>11</sub>	Sm <sub>2</sub> O <sub>3</sub>	Tb <sub>4</sub> O <sub>7</sub>	Tm <sub>2</sub> O <sub>3</sub>	Yb <sub>2</sub> O <sub>3</sub>	Y <sub>2</sub> O <sub>3</sub>	TREO	MREO
HHC047	0	3	122.10	79.98	4.95	2.41	1.64	6.11	0.82	0.28	56.45	16.43	9.23	0.85	0.37	2.20	29.46	333.30	78.68
HHC047	3	6	152.94	118.45	5.98	2.97	1.91	8.59	0.99	0.25	79.20	22.71	12.12	1.08	0.35	2.04	32.89	442.47	108.97
HHC047	6	9	37.10	20.64	1.09	0.72	0.27	1.38	0.22	0.15	10.26	3.25	1.35	0.20	0.07	0.66	6.48	83.83	14.80
HHC047	9	12	24.69	15.36	0.72	0.59	0.22	0.80	0.15	0.11	7.81	2.38	0.95	0.15	0.03	0.76	5.46	60.21	11.07
HHC047	12	15	27.15	16.65	1.10	0.71	0.13	1.00	0.15	0.06	8.63	2.60	1.60	0.11	0.07	0.61	5.59	66.15	12.44
HHC047	15	18	27.52	16.07	0.91	0.56	0.15	0.77	0.16	0.05	7.70	2.46	1.11	0.12	0.21	0.58	5.46	63.82	11.19
HHC047	18	21	24.08	15.01	0.69	0.50	0.35	0.82	0.16	0.08	7.81	2.21	1.37	0.14	0.09	0.54	4.70	58.55	10.86
HHC047	21	24	42.01	29.91	1.17	0.67	0.44	1.43	0.22	0.16	13.53	4.65	2.15	0.22	0.13	0.95	7.24	104.87	19.58
HHC047	24	27	90.16	63.21	1.96	0.99	0.69	2.86	0.37	0.18	33.71	10.54	4.67	0.32	0.18	0.96	10.79	221.61	46.52
HHC047	27	30	179.96	98.98	3.05	1.56	1.45	5.14	0.54	0.14	57.39	17.22	8.93	0.66	0.17	1.32	16.76	393.26	78.32
HHC048	0	3	119.65	67.79	4.66	2.04	1.49	5.80	0.77	0.27	47.82	13.23	8.30	0.69	0.29	1.89	24.64	299.32	66.41
HHC048	3	6	125.91	119.04	6.20	3.00	1.97	8.58	1.00	0.27	75.58	22.47	12.23	1.06	0.33	2.47	38.35	418.46	105.31
HHC048	6	9	29.48	18.06	1.21	0.69	0.36	1.27	0.26	0.10	9.56	2.97	1.35	0.16	0.03	0.80	7.11	73.42	13.91
HHC048	9	12	28.01	16.89	1.27	0.74	0.27	0.98	0.21	0.13	8.28	2.63	1.02	0.15	0.11	0.84	7.24	68.77	12.34
HHC048	12	15	28.13	15.01	0.94	0.53	0.25	1.03	0.19	0.10	7.70	2.30	1.02	0.14	0.08	0.51	6.35	64.28	11.08
HHC048	15	18	31.94	17.83	0.80	0.42	0.19	0.81	0.15	0.09	9.21	2.86	1.35	0.11	0.07	0.50	4.57	70.89	12.99
HHC048	18	21	31.82	19.12	1.08	0.46	0.17	1.06	0.17	0.07	9.80	2.97	1.30	0.15	0.05	0.50	4.44	73.16	14.00
HHC048	21	24	67.68	41.75	1.24	0.58	0.39	2.21	0.23	0.13	21.70	7.33	2.75	0.22	0.06	0.77	7.11	154.16	30.49
HHC048	24	27	60.31	41.52	1.15	0.66	0.43	1.96	0.22	0.10	20.30	5.90	2.50	0.20	0.02	0.66	8.13	144.06	27.54
HHC048	27	30	59.21	34.48	1.22	0.63	0.44	1.82	0.21	0.10	17.03	5.47	2.53	0.25	0.10	0.65	6.22	130.36	23.97
HHC048	30	33	330.44	131.94	5.78	3.13	2.25	7.78	1.05	0.44	89.81	24.89	12.12	0.89	0.49	3.09	36.70	650.81	121.38
HHC048	33	35	339.04	192.34	6.17	2.54	2.83	10.83	1.03	0.33	128.30	37.09	18.15	1.19	0.40	2.62	34.41	777.28	172.76
HHC049	0	3	163.38	84.09	4.88	2.30	1.62	6.49	0.87	0.36	61.24	16.67	9.18	0.80	0.32	2.14	29.72	384.06	83.59
HHC049	3	6	128.98	76.23	4.35	2.29	1.48	6.51	0.88	0.39	53.77	14.80	8.42	0.73	0.34	2.14	28.95	330.27	73.65
HHC049	6	9	57.12	31.43	1.61	0.89	0.53	1.84	0.23	0.14	17.96	4.78	2.73	0.24	0.06	1.02	9.27	129.85	24.59
HHC049	9	12	24.81	14.66	0.67	0.40	0.20	0.80	0.11	0.06	7.12	2.31	0.83	0.07	0.03	0.48	4.19	56.73	10.16
HHC049	12	15	26.29	16.54	0.78	0.57	0.22	0.67	0.17	0.13	8.40	2.40	1.16	0.14	0.06	0.57	4.83	62.92	11.72
HHC049	15	18	85.13	45.74	1.18	0.77	0.24	1.68	0.22	0.08	25.19	7.55	2.77	0.19	0.15	0.61	5.33	176.84	34.12
HHC049	18	21	33.29	20.52	0.91	0.51	0.17	1.24	0.17	0.07	10.38	2.91	1.39	0.16	0.10	0.63	6.10	78.57	14.36
HHC049	21	24	51.10	35.18	1.27	0.75	0.38	1.75	0.26	0.14	17.96	5.53	3.08	0.26	0.13	0.84	7.37	126.02	25.03
HHC049	24	28	40.29	36.36	1.01	0.74	0.50	1.68	0.15	0.07	18.90	5.78	2.03	0.16	0.15	0.40	6.98	115.20	25.85

## APPENDIX 1.

The following Tables are provided to ensure compliance with the JORC Code (2012 Edition) requirements for the reporting of Exploration Results at Hines Hill

### Section 1: Sampling Techniques and Data

(Criteria in this section applies to all succeeding sections)

Criteria	JORC Code explanation	Commentary
Sampling techniques	<i>Nature and quality of sampling (eg cut channels, random chips, or specific specialised industry standard measurement tools appropriate to the minerals under investigation, such as down hole gamma sondes, or handheld XRF instruments, etc). These examples should not be taken as limiting the broad meaning of sampling.</i>	Every metre drilled was sampled at the drill rig using a rig mounted static cone splitter to collect 2 – 3kg sub samples.  3m composites were collected using the pipe/spear method of sampling the coarse reject sample collected in standard green bags, which remain at the drill site.
	<i>Include reference to measures taken to ensure sample representivity and the appropriate calibration of any measurement tools or systems used.</i>	Standard reference material, sample duplicates were automatically collected at 25m sample intervals from the cone splitter  Where a duplicate, produced from the cone splitter, wasn't sampled due to it being in a non-mineralised zone, a 4m composite field duplicate was obtained using the pipe/spear method from the sample reject bag. This method maintained a ~25m duplicate and standard insertion rate throughout the entire program.
	<i>Aspects of the determination of mineralisation that are Material to the Public Report. In cases where 'industry standard' work has been done this would be relatively simple (eg 'reverse circulation drilling was used to obtain 1 m samples from which 3 kg was pulverised to produce a 30 g charge for fire assay'). In other cases more explanation may be required, such as where there is coarse gold that has inherent sampling problems. Unusual commodities or mineralisation types (eg submarine nodules) may warrant disclosure of detailed information.</i>	1m samples were taken for each metre drilled and stored. 3m composite samples were taken for each drill hole and sent to the laboratory for crushing, splitting and analysis.
Drilling techniques	<i>Drill type (e.g. core, reverse circulation, open-hole hammer, rotary air blast, auger, Bangka, sonic etc) and details (e.g. core diameter, triple or standard tube, depth of diamond tails, face-sampling bit or other type, whether core is orientated and if so, by what method, etc).</i>	The drilling was undertaken by Strike Drilling with their X 350 Air Core tracked rig with 3.5inch RC/AC capability. Clay profile was drilled with blade to refusal and hammer drilled for basement sample
Drill sample recovery	<i>Method of recording and assessing core and chip sample recoveries and results assessed.</i>	Drill recovery was routinely recorded via estimation of the comparative percentage of the volume of the sample bag by the company geologist. The sample recovery was deemed adequate for representative assays.
	<i>Measures taken to maximise sample recovery and ensure representative nature of the samples.</i>	A qualitative estimate of sample weight was undertaken to ensure consistency of sample size and to monitor sample recoveries at the time of drilling.
	<i>Whether a relationship exists between sample recovery and grade and whether sample bias may have occurred due to preferential loss/gain of fine/coarse material.</i>	Drill sample recovery and quality is considered to be adequate for the drilling technique employed.
Criteria	JORC Code explanation	Commentary

<b>Logging</b>	<i>Whether core and chip samples have been geologically and geotechnically logged to a level of detail to support appropriate Mineral Resource estimation, mining studies and metallurgical studies.</i>	All holes have been geologically logged for lithology, mineralisation and weathering. A brief description of each drilling sample was recorded and a permanent record has been collected and stored in chip trays for reference.
	<i>The total length and percentage of the relevant intersections logged.</i>	All intersections logged 100% as all lengths are relevant at the current stage of exploration.
<b>Sub-sampling techniques and sample preparation</b>	<i>If core, whether cut or sawn and whether quarter, half or all core taken.</i>	A sub sample from the RC drill rig of approximately 2-4kg was taken from the sample splitter off the cyclone.
	<i>If non-core, whether riffled, tube sampled, rotary split, etc and whether sampled wet or dry.</i>	
	<i>For all sample types, the nature, quality and appropriateness of the sample preparation technique.</i>	
	<i>Quality control procedures adopted for all sub-sampling stages to maximise representivity of samples.</i>	No sub-sampling has been undertaken.
	<i>Measures taken to ensure that the sampling is representative of the in-situ material collected, including for instance results for field duplicate/second-half sampling.</i>	No sub-sampling has been undertaken.
	<i>Whether sample sizes are appropriate to the grain size of the material being sampled.</i>	The sample size of 2-4 kilograms is appropriate and representative of the grain size and mineralisation style of the deposit.
<b>Quality of assay data and laboratory tests</b>	<i>The nature, quality and appropriateness of the assaying and laboratory procedures used and whether the technique is considered partial or total. For geophysical tools, spectrometers, handheld XRF instruments, etc, the parameters used in determining the analysis including instrument make and model, reading times, calibrations factors applied and their derivation, etc.</i>	Samples were submitted to ALS Laboratories for analysis.  Elements were analysed using MS81L-REE™ : Ba, Ce, Cr, Cs, Dy, Er, Eu, Ga, Gd, Hf, Ho, La, Lu, Nb, Nd, Pr, Rb, Sc, Sm, Sn, Sr, Ta, Tb, Te, Th, Ti, Tm, U, V, W, Y, Yb, Zr.
	<i>Nature of quality control procedures adopted (eg standards, blanks, duplicates, external laboratory checks) and whether acceptable levels of accuracy (ie lack of bias) and precision have been established.</i>	CRM & field duplicated samples were inserted every 25 samples for QA/QC control.
<b>Verification of sampling and assaying</b>	<i>The verification of significant intersections by either independent or alternative company personnel.</i>	Significant intercepts are reviewed by 2 or more company geologists.
	<i>The use of twinned holes.</i>	No twinned drill holes.
	<i>Documentation of primary data, data entry procedures, data verification, data storage (physical and electronic) protocols.</i>	All field data were collected manually and transferred to spreadsheets. Sample location coordinates were determined and recorded using a handheld GPS.
	<i>Discuss any adjustment to assay data.</i>	The REE assay data were converted from reported elemental assays to the equivalent oxide compound as applicable to rare earth oxides. The oxides were calculated from the element according to the following factors: <ul style="list-style-type: none"> <li>• CeO<sub>2</sub> 1.1526</li> <li>• La<sub>2</sub>O<sub>3</sub> 1.1728</li> <li>• Nd<sub>2</sub>O<sub>3</sub> 1.1664</li> <li>• Pr<sub>6</sub>O<sub>11</sub> 1.2082</li> </ul> <ul style="list-style-type: none"> <li>• Ho<sub>2</sub>O<sub>3</sub> 1.1455</li> <li>• Lu<sub>2</sub>O<sub>3</sub> 1.1371</li> <li>• Sm<sub>2</sub>O<sub>3</sub> 1.1596</li> <li>• Tb<sub>2</sub>O<sub>3</sub> 1.1762</li> <li>• Tm<sub>2</sub>O<sub>3</sub> 1.1421</li> <li>• Y<sub>2</sub>O<sub>3</sub> 1.2699</li> <li>• Yb<sub>2</sub>O<sub>3</sub> 1.1387</li> <li>• Dy<sub>2</sub>O<sub>3</sub> 1.1477</li> <li>• Er<sub>2</sub>O<sub>3</sub> 1.1435</li> <li>• Eu<sub>2</sub>O<sub>3</sub> 1.1579</li> <li>• Gd<sub>2</sub>O<sub>3</sub> 1.1526</li> </ul>
<b>Criteria</b>	<b>JORC Code explanation</b>	<b>Commentary</b>
<b>Location of data points</b>	<i>Accuracy and quality of surveys used to locate drill holes (collar and down-hole surveys), trenches, mine workings and other locations used in Mineral Resource estimation.</i>	All locations determined by handheld GPS using GDA94 datum in UTM Zone 50.

	<i>Specification of the grid system used.</i>	
	<i>Quality and adequacy of topographic control.</i>	
<b>Data spacing and distribution</b>	<i>Data spacing for reporting of Exploration Results.</i>	Sample spacing was 100m.
	<i>Whether the data spacing and distribution is sufficient to establish the degree of geological and grade continuity appropriate for the Mineral Resource and Ore Reserve estimation procedure(s) and classifications applied.</i>	Maiden first pass drilling is not designed for an MRE and is too coarse. The drill spacing is intended to identify REE mineralisation, and will have reduced spacing in future programs.
	<i>Whether sample compositing has been applied.</i>	3m sample compositing has been applied.
<b>Orientation of data in relation to geological structure</b>	<i>Whether the orientation of sampling achieves unbiased sampling of possible structures and the extent to which this is known, considering the deposit type.</i>	Drill holes are vertical and suitable for clay profile drilling.
	<i>If the relationship between the drilling orientation and the orientation of key mineralised structures is considered to have introduced a sampling bias, this should be assessed and reported if material.</i>	No bias is seen in the orientation of drilling
<b>Sample security</b>	<i>The measures taken to ensure sample security.</i>	All samples were placed in plastic or calico bags, taken to Perth and delivered to ALS laboratory by White Cliff staff.
<b>Audits or reviews</b>	<i>The results of any audits or reviews of sampling techniques and data.</i>	Data is validated upon up-loading into the master database. Any validation issues identified are investigated prior to reporting of results.

## Section 2: Reporting of Exploration Results

(Criteria listed in the preceding section also apply to this section)

Criteria	JORC Code explanation	Commentary
<b>Mineral tenement and land tenure status</b>	<i>Type, reference name/number, location and ownership including agreements or material issues with third parties such as joint ventures, partnerships, overriding royalties, native title interests, historical sites, wilderness or national park and environmental settings.</i>	The Hines Hill exploration license E70/5875 is held 100% by Magnet Resource Company Pty Ltd, a 100% subsidiary of White Cliff Minerals Limited. The tenement was granted on 21/10/21, has annual expenditure of \$44,000 and the tenement is in good standing.
	<i>The security of the tenure held at the time of reporting along with any known impediments to obtaining a licence to operate in the area.</i>	A land access agreement has been signed with the landowners.
<b>Exploration done by other parties</b>	<i>Acknowledgment and appraisal of exploration by other parties.</i>	No previous exploration.
<b>Geology</b>	<i>Deposit type, geological setting and style of mineralisation.</i>	Potential Carbonatite within Archean terrane.
<b>Drill hole information</b>	<i>A summary of all information material to the understanding of the exploration results including a tabulation of the following information for all Material drill holes:</i>	A summary of all exploration drilling information and sampling is contained in tabulated data within this announcement.

Criteria	JORC Code explanation	Commentary
	<p>easting and northing of the drill hole collar elevation or RL (Reduced Level – elevation above sea level in metres) of the drill hole collar</p> <p>dip and azimuth of the hole</p> <p>down hole length and interception depth</p> <p>hole length.</p> <p>If the exclusion of this information is justified on the basis that the information is not Material and this exclusion does not detract from the understanding of the report, the Competent Person should clearly explain why this is the case.</p>	
<b>Data aggregation methods</b>	<p>In reporting Exploration Results, weighting averaging techniques, maximum and/or minimum grade truncations (eg. cutting of high grades) and cut-off grades are usually Material and should be stated.</p>	<p>Intersections have been calculated generally using a 350ppm cut off and internal waste of up to 2m thickness with total intercepts greater than 350ppm.</p> <p>No upper cut off has been applied to intersections or samples.</p>
	<p>Where aggregate intercepts incorporate short lengths of high-grade results and longer lengths of low grade results, the procedure used for such aggregation should be stated and some typical examples of such aggregations should be shown in detail.</p>	<p>Only relevant elements (REE) are reported here. However, the samples underwent multi element assay as industry standard.</p>
	<p>The assumptions used for any reporting of metal equivalent values should be clearly stated.</p>	<p>No metal equivalent values are being used.</p>
<b>Relationship between mineralisation widths and intercept lengths</b>	<p>These relationships are particularly important in the reporting of Exploration Results. If the geometry of the mineralisation with respect to the drill hole angle is known, its nature should be reported. If it is not known and only the down hole lengths are reported, there should be a clear statement to this effect (eg 'down hole length, true width not known').</p>	<p>Drill holes have been drilled vertical, which is generally perpendicular to the horizontal orebody within the clay profile.</p>
<b>Diagrams</b>	<p>Appropriate maps and sections (with scales) and tabulations of intercepts should be included for any significant discovery being reported. These should include, but not be limited to a plan view of drill hole collar locations and appropriate sectional views.</p>	<p>Location maps of projects within the release with relevant exploration information contained.</p>
<b>Balanced reporting</b>	<p>Where comprehensive reporting of all Exploration Results is not practicable, representative reporting of both low and high grades and/or widths should be practiced to avoid misleading reporting of Exploration Results.</p>	<p>The reporting of exploration results is considered balanced by the competent person.</p>
<b>Other substantive exploration data</b>	<p>Other exploration data, if meaningful and material, should be reported including (but not limited to): geological observations; geophysical survey results; geochemical survey results; bulk samples – size and method of treatment; metallurgical test results; bulk density, groundwater, geotechnical and rock characteristics; potential deleterious or contaminating substances.</p>	<p>No other exploration to report.</p>
<b>Further work</b>	<p>The nature and scale of planned further work (eg. tests for lateral extensions or depth extensions or large-scale step-out drilling).</p> <p>Diagrams clearly highlighting the areas of possible extensions, including the main geological interpretations and future drilling areas, provided this information is not commercially sensitive.</p>	<p>Full technical review which includes upcoming geochemical sampling results, assay of 1m samples within the ore zone and then further drilling.</p>