

Appendix 3. U–Pb age. U–Pb 年代.
Table A1. Sample KRH. 試料KRH.

Labels	$^{206}\text{Pb}_c^{(1)}$ (%)	U (ppm)	Th (ppm)	Th/U	$^{238}\text{U}/^{206}\text{Pb}^*^{(1)}$	$^{207}\text{Pb}^*/^{206}\text{Pb}^*^{(1)}$	$^{238}\text{U}/^{206}\text{Pb}^* \text{ age}^{(1)}$ (Ma)	$^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age}^{(1)}$ (Ma)	Disc ⁽³⁾ (%)
KRH_001	0.46	461	259	0.58	101.58 ± 1.80	0.0456 ± 0.0053	63.1 ± 1.1	63.1 ± 1.1	
KRH_002	1.26	242	147	0.62	107.69 ± 2.43	0.0383 ± 0.0070	59.6 ± 1.3	59.6 ± 1.3	
KRH_003	0.00	232	100	0.44	25.94 ± 0.39	0.0520 ± 0.0019	243.9 ± 3.6	243.9 ± 3.6	
KRH_004	0.46	251	184	0.75	97.80 ± 2.34	0.0537 ± 0.0071	65.6 ± 1.6	65.6 ± 1.6	
KRH_005	0.00	334	321	0.99	99.57 ± 1.65	0.0524 ± 0.0030	64.4 ± 1.1	64.4 ± 1.1	
KRH_006	2.02	217	88	0.42	103.11 ± 2.61	0.0342 ± 0.0071	62.2 ± 1.6	62.2 ± 1.6	
KRH_007	0.43	257	194	0.77	96.51 ± 2.06	0.0454 ± 0.0068	66.5 ± 1.4	66.5 ± 1.4	
KRH_008	0.31	189	115	0.62	68.07 ± 1.45	0.0521 ± 0.0063	94.0 ± 2.0	94.0 ± 2.0	
KRH_009	1.43	280	218	0.80	106.48 ± 2.89	0.0516 ± 0.0081	60.3 ± 1.6	60.3 ± 1.6	
KRH_010	0.34	305	220	0.74	103.56 ± 2.10	0.0523 ± 0.0069	61.9 ± 1.3	61.9 ± 1.3	
KRH_011	0.89	221	158	0.73	104.49 ± 2.50	0.0343 ± 0.0077	61.4 ± 1.5	61.4 ± 1.5	
KRH_012	0.00	251	131	0.53	100.41 ± 2.09	0.0497 ± 0.0042	63.9 ± 1.3	63.9 ± 1.3	
KRH_013	0.08	339	221	0.67	100.94 ± 1.99	0.0441 ± 0.0059	63.6 ± 1.2	63.6 ± 1.2	
KRH_014	0.17	373	159	0.44	104.65 ± 1.87	0.0462 ± 0.0047	61.3 ± 1.1	61.3 ± 1.1	
KRH_015	0.00	234	177	0.78	106.09 ± 2.20	0.0535 ± 0.0041	60.5 ± 1.2	60.5 ± 1.2	
KRH_016	0.08	197	147	0.77	93.91 ± 2.48	0.0473 ± 0.0090	68.3 ± 1.8	68.3 ± 1.8	
KRH_017	0.96	249	200	0.82	100.49 ± 2.46	0.0452 ± 0.0086	63.8 ± 1.6	63.8 ± 1.6	
KRH_018	0.00	244	189	0.79	98.05 ± 2.03	0.0521 ± 0.0039	65.4 ± 1.3	65.4 ± 1.3	
KRH_019	0.71	153	69	0.46	96.23 ± 3.03	0.0503 ± 0.0091	66.6 ± 2.1	66.6 ± 2.1	
KRH_020	1.09	203	190	0.96	97.89 ± 2.81	0.0400 ± 0.0112	65.5 ± 1.9	65.5 ± 1.9	
KRH_021	0.00	245	234	0.98	100.31 ± 2.22	0.0502 ± 0.0038	63.9 ± 1.4	63.9 ± 1.4	
KRH_022	0.00	309	212	0.71	99.33 ± 1.91	0.0442 ± 0.0029	64.6 ± 1.2	64.6 ± 1.2	
KRH_023	0.67	254	142	0.58	99.95 ± 2.27	0.0432 ± 0.0076	64.2 ± 1.5	64.2 ± 1.5	
KRH_024	0.00	283	285	1.03	96.94 ± 2.07	0.0498 ± 0.0037	66.2 ± 1.4	66.2 ± 1.4	
KRH_025	0.00	270	208	0.79	98.59 ± 2.17	0.0448 ± 0.0030	65.1 ± 1.4	65.1 ± 1.4	
KRH_026	0.64	355	276	0.80	102.97 ± 2.08	0.0424 ± 0.0064	62.3 ± 1.3	62.3 ± 1.3	
KRH_027	0.72	251	161	0.66	94.73 ± 2.33	0.0449 ± 0.0071	67.7 ± 1.7	67.7 ± 1.7	
KRH_028	1.11	125	76	0.63	99.19 ± 2.94	0.0457 ± 0.0101	64.7 ± 1.9	64.7 ± 1.9	
KRH_029	0.48	227	177	0.80	96.57 ± 2.18	0.0439 ± 0.0074	66.4 ± 1.5	66.4 ± 1.5	
KRH_030	0.77	306	221	0.74	101.74 ± 2.24	0.0363 ± 0.0067	63.1 ± 1.4	63.1 ± 1.4	
KRH_031	1.06	306	270	0.91	105.72 ± 2.54	0.0348 ± 0.0077	60.7 ± 1.5	60.7 ± 1.5	
KRH_032	0.55	373	205	0.56	102.80 ± 2.13	0.0446 ± 0.0058	62.4 ± 1.3	62.4 ± 1.3	
KRH_033	0.02	241	180	0.77	98.20 ± 2.14	0.0493 ± 0.0077	65.3 ± 1.4	65.3 ± 1.4	
KRH_034	0.00	245	169	0.71	102.72 ± 2.49	0.0434 ± 0.0034	62.5 ± 1.5	62.5 ± 1.5	
KRH_035	1.23	255	184	0.74	104.30 ± 2.71	0.0383 ± 0.0059	61.5 ± 1.6	61.5 ± 1.6	
KRH_036	0.00	272	210	0.79	99.56 ± 1.97	0.0465 ± 0.0066	64.4 ± 1.3	64.4 ± 1.3	
KRH_037	0.48	284	222	0.80	98.63 ± 2.49	0.0459 ± 0.0080	65.0 ± 1.6	65.0 ± 1.6	
KRH_038	0.33	343	202	0.60	101.61 ± 2.20	0.0394 ± 0.0059	63.1 ± 1.4	63.1 ± 1.4	
KRH_039	0.10	421	248	0.60	99.33 ± 1.89	0.0501 ± 0.0056	64.6 ± 1.2	64.6 ± 1.2	
KRH_040	0.63	277	180	0.66	91.22 ± 1.84	0.0396 ± 0.0059	70.3 ± 1.4	70.3 ± 1.4	
KRH_041	0.45	309	232	0.77	102.58 ± 2.31	0.0484 ± 0.0072	62.5 ± 1.4	62.5 ± 1.4	
KRH_042	1.40	226	150	0.68	99.91 ± 2.65	0.0394 ± 0.0088	64.2 ± 1.7	64.2 ± 1.7	
KRH_043	4.62	169	126	0.76	102.71 ± 2.92	0.0306 ± 0.0122	62.5 ± 1.8	62.5 ± 1.8	
KRH_044	0.23	345	197	0.59	101.24 ± 2.31	0.0499 ± 0.0061	63.4 ± 1.4	63.4 ± 1.4	
KRH_045	1.15	244	162	0.68	97.58 ± 2.51	0.0485 ± 0.0080	65.7 ± 1.7	65.7 ± 1.7	
KRH_046	0.88	316	178	0.58	99.37 ± 2.18	0.0424 ± 0.0058	64.5 ± 1.4	64.5 ± 1.4	
KRH_047	0.18	210	120	0.59	97.17 ± 2.61	0.0472 ± 0.0087	66.0 ± 1.8	66.0 ± 1.8	
KRH_048	0.00	192	132	0.71	96.38 ± 2.16	0.0502 ± 0.0040	66.5 ± 1.5	66.5 ± 1.5	
KRH_049	0.63	277	186	0.69	100.63 ± 2.32	0.0431 ± 0.0074	63.7 ± 1.5	63.7 ± 1.5	
KRH_050	1.25	171	115	0.69	96.64 ± 2.76	0.0374 ± 0.0088	66.4 ± 1.9	66.4 ± 1.9	
KRH_051	0.00	185	121	0.67	93.88 ± 2.44	0.0426 ± 0.0038	68.3 ± 1.8	68.3 ± 1.8	
KRH_052	0.00	409	272	0.68	98.54 ± 2.09	0.0465 ± 0.0026	65.1 ± 1.4	65.1 ± 1.4	
KRH_053	0.00	221	138	0.64	81.64 ± 1.76	0.0481 ± 0.0038	78.5 ± 1.7	78.5 ± 1.7	
KRH_054	0.89	193	142	0.76	100.42 ± 3.00	0.0462 ± 0.0095	63.9 ± 1.9	63.9 ± 1.9	
KRH_055	0.00	253	162	0.66	100.16 ± 2.38	0.0483 ± 0.0038	64.0 ± 1.5	64.0 ± 1.5	
KRH_056#	0.00	191	127	0.68	97.50 ± 2.23	0.0570 ± 0.0047	65.8 ± 1.5	65.8 ± 1.5	
KRH_057	0.66	238	200	0.86	98.94 ± 2.44	0.0539 ± 0.0100	64.8 ± 1.6	64.8 ± 1.6	
KRH_058	0.05	180	128	0.73	101.17 ± 2.37	0.0513 ± 0.0076	63.4 ± 1.5	63.4 ± 1.5	
KRH_059	0.00	312	168	0.55	95.74 ± 1.90	0.0453 ± 0.0030	67.0 ± 1.3	67.0 ± 1.3	
KRH_060	0.47	267	235	0.90	95.76 ± 2.35	0.0475 ± 0.0083	67.0 ± 1.6	67.0 ± 1.6	
KRH_061	0.00	253	146	0.59	97.67 ± 2.34	0.0442 ± 0.0037	65.7 ± 1.6	65.7 ± 1.6	
KRH_062	0.26	231	151	0.67	96.16 ± 2.53	0.0448 ± 0.0074	66.7 ± 1.7	66.7 ± 1.7	
KRH_063	0.00	305	290	0.98	96.34 ± 2.07	0.0448 ± 0.0032	66.6 ± 1.4	66.6 ± 1.4	
KRH_064	0.06	251	171	0.70	96.67 ± 2.50	0.0519 ± 0.0066	66.3 ± 1.7	66.3 ± 1.7	
KRH_065	0.00	256	133	0.53	97.98 ± 2.14	0.0547 ± 0.0039	65.5 ± 1.4	65.5 ± 1.4	
KRH_066	2.30	255	159	0.64	100.41 ± 2.25	0.0516 ± 0.0081	63.9 ± 1.4	63.9 ± 1.4	
KRH_067	1.09	170	117	0.71	98.56 ± 2.99	0.0420 ± 0.0109	65.1 ± 2.0	65.1 ± 2.0	
KRH_068	0.00	243	184	0.78	100.45 ± 2.64	0.0469 ± 0.0044	63.9 ± 1.7	63.9 ± 1.7	
KRH_069	0.00	279	172	0.63	98.58 ± 2.04	0.0458 ± 0.0037	65.1 ± 1.3	65.1 ± 1.3	
KRH_070	0.00	388	211	0.56	89.99 ± 1.75	0.0477 ± 0.0031	71.2 ± 1.4	71.2 ± 1.4	
KRH_071	0.00	334	193	0.59	100.17 ± 2.31	0.0481 ± 0.0033	64.0 ± 1.5	64.0 ± 1.5	
KRH_072	0.00	257	185	0.74	98.98 ± 2.15	0.0547 ± 0.0041	64.8 ± 1.4	64.8 ± 1.4	

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb–Pb and U–Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.
Table A2. Sample MJK. 試料MJK.

Labels	²⁰⁶ Pb _c ⁽¹⁾	U	Th	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾	Disc ⁽²⁾
	(%)	(ppm)	(ppm)				(Ma)	(Ma)	(%)
MJK_001	2.57	112	76	0.69	91.14 ± 3.40	0.0502 ± 0.0154	70.3 ± 2.6		
MJK_002	0.00	266	101	0.39	25.25 ± 0.40	0.0497 ± 0.0025	250.4 ± 3.9		
MJK_003	0.00	159	197	1.27	29.43 ± 0.62	0.0472 ± 0.0037	215.4 ± 4.5		
MJK_004	0.00	245	131	0.55	64.44 ± 1.70	0.0559 ± 0.0042	99.3 ± 2.6		
MJK_005	0.00	80	60	0.78	89.75 ± 4.06	0.0559 ± 0.0117	71.4 ± 3.2		
MJK_006	0.15	447	334	0.77	64.34 ± 1.41	0.0533 ± 0.0064	99.4 ± 2.2		
MJK_007	2.15	317	177	0.57	89.99 ± 2.62	0.0322 ± 0.0081	71.2 ± 2.1		
MJK_008	0.25	355	197	0.57	77.18 ± 1.86	0.0510 ± 0.0069	83.0 ± 2.0		
MJK_009	1.06	1021	1004	1.01	85.69 ± 1.66	0.0429 ± 0.0066	74.8 ± 1.4		
MJK_010	0.00	279	192	0.71	66.38 ± 1.82	0.0416 ± 0.0040	96.4 ± 2.6		
MJK_011	0.13	1071	279	0.27	78.17 ± 1.25	0.0462 ± 0.0030	81.9 ± 1.3		
MJK_012	0.00	122	83	0.70	94.25 ± 4.09	0.0389 ± 0.0068	68.0 ± 2.9		
MJK_013	0.00	338	130	0.39	67.21 ± 1.53	0.0425 ± 0.0035	95.2 ± 2.2		
MJK_014	0.00	190	131	0.71	88.83 ± 3.15	0.0514 ± 0.0065	72.2 ± 2.5		
MJK_015	0.00	299	310	1.06	12.55 ± 0.21	0.0544 ± 0.0016	494.1 ± 7.9		
MJK_016	0.88	446	482	1.11	94.47 ± 2.71	0.0375 ± 0.0092	67.9 ± 1.9		
MJK_017	0.00	465	218	0.48	3.26 ± 0.04	0.1118 ± 0.0015	1724.7 ± 20.2	1830 ± 24	5.75
MJK_018	0.00	378	348	0.94	39.14 ± 0.78	0.0446 ± 0.0027	162.6 ± 3.2		
MJK_019	0.00	428	221	0.53	97.22 ± 2.46	0.0540 ± 0.0043	66.0 ± 1.7		
MJK_020	0.51	716	295	0.42	77.13 ± 1.60	0.0388 ± 0.0045	83.0 ± 1.7		
MJK_021#	0.00	1261	479	0.39	68.80 ± 1.14	0.0549 ± 0.0022	93.0 ± 1.5		
MJK_022	1.40	124	77	0.64	99.27 ± 4.07	0.0252 ± 0.0158	64.6 ± 2.6		
MJK_023	0.00	141	130	0.94	86.12 ± 3.88	0.0362 ± 0.0062	74.4 ± 3.3		
MJK_024#	0.00	222	169	0.78	87.16 ± 2.80	0.0638 ± 0.0070	73.5 ± 2.3		
MJK_025	0.97	191	123	0.66	99.72 ± 4.18	0.0331 ± 0.0113	64.3 ± 2.7		
MJK_026	1.09	399	209	0.54	87.13 ± 2.65	0.0400 ± 0.0079	73.6 ± 2.2		
MJK_027	0.00	66	56	0.87	78.81 ± 5.06	0.0403 ± 0.0088	81.3 ± 5.2		
MJK_028	0.00	160	81	0.52	70.60 ± 2.38	0.0415 ± 0.0061	90.7 ± 3.0		
MJK_029	0.98	512	350	0.70	99.77 ± 2.90	0.0369 ± 0.0092	64.3 ± 1.9		
MJK_030	0.00	777	272	0.36	63.40 ± 1.15	0.0465 ± 0.0025	100.9 ± 1.8		
MJK_031	0.68	454	463	1.05	97.97 ± 2.63	0.0480 ± 0.0100	65.5 ± 1.8		
MJK_032	2.11	235	150	0.65	73.16 ± 2.15	0.0294 ± 0.0101	87.5 ± 2.6		
MJK_033	0.10	168	86	0.52	80.36 ± 3.28	0.0419 ± 0.0105	79.7 ± 3.2		
MJK_034	0.00	366	183	0.51	82.94 ± 2.45	0.0527 ± 0.0044	77.3 ± 2.3		
MJK_035	0.00	2144	1711	0.82	2.61 ± 0.03	0.1357 ± 0.0012	2091.8 ± 22.2	2175 ± 15	3.83
MJK_036	0.96	139	75	0.55	72.38 ± 3.11	0.0292 ± 0.0113	88.5 ± 3.8		
MJK_037	0.27	316	116	0.38	55.53 ± 1.59	0.0442 ± 0.0057	115.0 ± 3.3		
MJK_038	0.00	270	127	0.48	61.42 ± 1.63	0.0468 ± 0.0039	104.1 ± 2.7		
MJK_039	0.00	176	80	0.47	72.62 ± 2.27	0.0455 ± 0.0051	88.2 ± 2.7		
MJK_040	0.46	461	131	0.29	64.07 ± 1.52	0.0506 ± 0.0050	99.8 ± 2.3		
MJK_041	0.46	295	291	1.01	95.16 ± 3.01	0.0339 ± 0.0128	67.4 ± 2.1		
MJK_042	0.00	508	310	0.63	83.79 ± 1.88	0.0419 ± 0.0030	76.5 ± 1.7		
MJK_043	0.00	448	184	0.42	85.06 ± 1.92	0.0477 ± 0.0037	75.3 ± 1.7		
MJK_044	0.00	235	155	0.68	25.94 ± 0.50	0.0539 ± 0.0027	243.9 ± 4.7		
MJK_045	0.15	1132	608	0.55	86.33 ± 1.48	0.0478 ± 0.0040	74.2 ± 1.3		
MJK_046	0.00	161	56	0.35	71.04 ± 2.05	0.0548 ± 0.0065	90.1 ± 2.6		
MJK_047	0.00	370	440	1.22	83.95 ± 2.00	0.0529 ± 0.0044	76.3 ± 1.8		
MJK_048	0.25	615	232	0.39	64.65 ± 1.13	0.0453 ± 0.0041	98.9 ± 1.7		
MJK_049	0.00	365	181	0.51	70.93 ± 1.43	0.0556 ± 0.0042	90.2 ± 1.8		
MJK_050	0.00	389	295	0.78	93.34 ± 2.11	0.0493 ± 0.0039	68.7 ± 1.5		
MJK_051	0.56	55	48	0.89	12.51 ± 0.31	0.0608 ± 0.0092	495.8 ± 11.8		
MJK_052	0.24	215	126	0.60	74.48 ± 2.17	0.0458 ± 0.0083	86.0 ± 2.5		
MJK_053	0.00	133	31	0.24	27.10 ± 0.65	0.0512 ± 0.0045	233.6 ± 5.5		
MJK_054	0.00	138	102	0.76	32.06 ± 0.95	0.0495 ± 0.0044	198.0 ± 5.8		
MJK_055	0.00	703	268	0.39	60.00 ± 1.07	0.0489 ± 0.0024	106.5 ± 1.9		
MJK_056#	0.00	387	178	0.47	84.85 ± 2.42	0.0635 ± 0.0054	75.5 ± 2.1		
MJK_057	1.16	386	290	0.77	99.54 ± 2.45	0.0410 ± 0.0078	64.4 ± 1.6		
MJK_058	0.14	118	58	0.50	24.13 ± 0.64	0.0536 ± 0.0065	261.8 ± 6.8		
MJK_059	0.08	462	328	0.73	61.04 ± 1.42	0.0508 ± 0.0066	104.8 ± 2.4		
MJK_060	0.00	510	199	0.40	84.62 ± 1.87	0.0505 ± 0.0037	75.7 ± 1.7		
MJK_061	0.47	365	270	0.76	34.88 ± 0.62	0.0484 ± 0.0059	182.2 ± 3.2		
MJK_062	0.00	1358	574	0.43	68.94 ± 1.07	0.0468 ± 0.0018	92.8 ± 1.4		
MJK_063	0.00	280	90	0.33	2.95 ± 0.04	0.1151 ± 0.0018	1880.9 ± 23.4	1883 ± 28	0.11
MJK_064	1.07	88	64	0.74	94.88 ± 5.27	0.0514 ± 0.0211	67.6 ± 3.7		
MJK_065	1.83	99	45	0.47	92.76 ± 4.92	0.0539 ± 0.0178	69.1 ± 3.6		
MJK_066	1.30	817	768	0.96	89.89 ± 1.97	0.0407 ± 0.0072	71.3 ± 1.6		
MJK_067	1.06	234	127	0.56	75.03 ± 2.13	0.0447 ± 0.0084	85.3 ± 2.4		
MJK_068	0.00	87	74	0.88	83.36 ± 3.61	0.0572 ± 0.0129	76.9 ± 3.3		
MJK_069	0.00	231	116	0.52	84.44 ± 2.38	0.0492 ± 0.0052	75.9 ± 2.1		
MJK_070	0.36	160	68	0.43	37.04 ± 0.93	0.0437 ± 0.0067	171.7 ± 4.3		
MJK_071	0.00	182	134	0.75	77.05 ± 2.87	0.0565 ± 0.0059	83.1 ± 3.1		
MJK_072	2.07	223	170	0.78	85.80 ± 2.94	0.0252 ± 0.0115	74.7 ± 2.5		
MJK_073	0.69	233	108	0.48	69.74 ± 2.04	0.0368 ± 0.0069	91.8 ± 2.7		
MJK_074	0.00	207	174	0.86	85.08 ± 2.71	0.0460 ± 0.0064	75.3 ± 2.4		
MJK_075	1.34	201	94	0.48	64.02 ± 2.05	0.0312 ± 0.0084	99.9 ± 3.2		

MJK_076	0.05	296	73	0.25	3.12 ± 0.04	0.1189 ±	0.0020	1791.9 ± 20.6			
MJK_077	0.00	292	124	0.43	82.43 ± 2.43	0.0456 ±	0.0044	77.7 ± 2.3			
MJK_078#	0.18	525	132	0.26	3.08 ± 0.04	0.1429 ±	0.0020	1813.9 ± 22.5			
MJK_079	0.01	282	156	0.57	87.53 ± 2.60	0.0422 ±	0.0091	73.2 ± 2.2			
MJK_080	0.00	296	234	0.81	35.31 ± 0.76	0.0528 ±	0.0036	180.0 ± 3.8			
MJK_081	0.00	112	80	0.74	12.38 ± 0.24	0.0634 ±	0.0039	500.7 ± 9.2			
MJK_082	0.01	253	221	0.90	3.05 ± 0.04	0.1151 ±	0.0029	1829.9 ± 21.3	1883 ± 44		2.82
MJK_083	0.00	188	185	1.01	84.54 ± 3.02	0.0510 ±	0.0069	75.8 ± 2.7			
MJK_084	0.00	475	233	0.50	60.58 ± 1.23	0.0435 ±	0.0031	105.5 ± 2.1			
MJK_085	0.00	913	521	0.59	60.55 ± 0.99	0.0446 ±	0.0018	105.6 ± 1.7			
MJK_086	0.83	463	580	1.28	81.50 ± 1.97	0.0433 ±	0.0096	78.6 ± 1.9			
MJK_087#	0.00	434	163	0.39	73.23 ± 1.96	0.0602 ±	0.0051	87.4 ± 2.3			
MJK_088	0.32	360	140	0.40	23.73 ± 0.46	0.0492 ±	0.0036	266.0 ± 5.0			
MJK_089	0.54	602	207	0.35	80.19 ± 1.77	0.0441 ±	0.0050	79.9 ± 1.8			
MJK_090	0.12	598	543	0.93	29.31 ± 0.50	0.0478 ±	0.0042	216.3 ± 3.7			
MJK_091	0.07	217	42	0.20	3.00 ± 0.05	0.1171 ±	0.0022	1857.1 ± 24.7	1913 ± 33		2.92
MJK_092	0.45	53	41	0.80	2.20 ± 0.05	0.1488 ±	0.0050	2412.9 ± 42.6	2333 ± 57		-3.42
MJK_093	0.00	187	121	0.66	62.05 ± 2.18	0.0497 ±	0.0049	103.1 ± 3.6			
MJK_094	0.38	669	231	0.35	66.21 ± 1.19	0.0437 ±	0.0042	96.6 ± 1.7			
MJK_095	0.00	218	132	0.62	77.30 ± 2.62	0.0561 ±	0.0061	82.9 ± 2.8			
MJK_096	0.00	497	188	0.39	62.28 ± 1.26	0.0499 ±	0.0028	102.7 ± 2.1			
MJK_097	0.00	334	118	0.36	36.89 ± 0.77	0.0528 ±	0.0034	172.4 ± 3.5			
MJK_098	0.00	440	25	0.06	2.04 ± 0.03	0.1931 ±	0.0022	2570.0 ± 28.9	2770 ± 19		7.22
MJK_099	0.00	67	40	0.61	24.49 ± 0.78	0.0453 ±	0.0048	258.0 ± 8.0			
MJK_100	2.38	730	385	0.54	79.33 ± 1.70	0.0429 ±	0.0066	80.8 ± 1.7			
MJK_101	0.16	502	300	0.61	84.96 ± 1.88	0.0517 ±	0.0069	75.4 ± 1.7			
MJK_102	0.11	412	343	0.85	79.49 ± 1.90	0.0439 ±	0.0078	80.6 ± 1.9			
MJK_103	1.24	110	76	0.71	94.08 ± 4.35	0.0329 ±	0.0150	68.2 ± 3.1			
MJK_104	0.00	169	104	0.63	64.59 ± 2.00	0.0396 ±	0.0049	99.0 ± 3.0			
MJK_105	0.31	612	454	0.76	32.92 ± 0.65	0.0467 ±	0.0042	192.9 ± 3.7			
MJK_106	0.58	463	242	0.54	91.13 ± 2.24	0.0406 ±	0.0071	70.4 ± 1.7			
MJK_107	0.00	394	145	0.38	62.72 ± 1.87	0.0420 ±	0.0040	102.0 ± 3.0			
MJK_108	0.00	359	151	0.43	63.10 ± 1.70	0.0408 ±	0.0037	101.4 ± 2.7			
MJK_109	0.00	465	283	0.63	81.29 ± 2.27	0.0471 ±	0.0035	78.8 ± 2.2			
MJK_110	0.00	402	190	0.49	2.82 ± 0.06	0.1168 ±	0.0018	1957.8 ± 34.3	1909 ± 27		-2.56
MJK_111	0.22	734	396	0.55	75.16 ± 1.75	0.0482 ±	0.0059	85.2 ± 2.0			
MJK_112	0.00	74	55	0.76	97.65 ± 5.25	0.0334 ±	0.0105	65.7 ± 3.5			
MJK_113	0.00	52	53	1.05	12.48 ± 0.37	0.0534 ±	0.0039	497.0 ± 14.3			
MJK_114	0.00	462	141	0.31	85.50 ± 2.11	0.0446 ±	0.0038	75.0 ± 1.8			
MJK_115	0.14	86	20	0.24	12.31 ± 0.25	0.0566 ±	0.0044	503.6 ± 9.7			
MJK_116	0.00	209	187	0.92	82.99 ± 2.49	0.0420 ±	0.0045	77.2 ± 2.3			
MJK_117	0.00	228	217	0.98	26.40 ± 0.61	0.0458 ±	0.0031	239.7 ± 5.5			
MJK_118	0.00	399	199	0.51	25.44 ± 0.49	0.0495 ±	0.0023	248.6 ± 4.7			
MJK_119	0.00	170	66	0.40	72.55 ± 2.41	0.0451 ±	0.0072	88.3 ± 2.9			
MJK_120	0.55	565	285	0.52	68.03 ± 1.35	0.0452 ±	0.0055	94.1 ± 1.9			
MJK_121	0.00	1745	910	0.53	69.64 ± 1.26	0.0497 ±	0.0017	91.9 ± 1.7			
MJK_122	0.00	608	205	0.35	12.54 ± 0.18	0.0567 ±	0.0015	494.8 ± 6.8			
MJK_123	0.00	619	192	0.32	2.96 ± 0.04	0.1139 ±	0.0013	1877.5 ± 24.6	1863 ± 21		-0.78
MJK_124	0.00	486	193	0.41	64.97 ± 1.41	0.0522 ±	0.0040	98.5 ± 2.1			
MJK_125	0.00	167	84	0.52	85.61 ± 3.42	0.0409 ±	0.0058	74.9 ± 3.0			
MJK_126	0.19	586	119	0.21	3.03 ± 0.04	0.1154 ±	0.0019	1840.7 ± 22.4	1887 ± 30		2.45
MJK_127	0.00	537	232	0.44	65.65 ± 1.45	0.0464 ±	0.0029	97.5 ± 2.1			
MJK_128	1.00	168	131	0.80	95.45 ± 3.33	0.0501 ±	0.0131	67.2 ± 2.3			
MJK_129	0.00	166	115	0.71	25.37 ± 0.56	0.0497 ±	0.0035	249.2 ± 5.4			
MJK_130	0.00	201	77	0.39	2.75 ± 0.04	0.1314 ±	0.0020	2000.3 ± 25.4	2118 ± 27		5.56
MJK_131	0.33	1615	740	0.47	38.54 ± 0.56	0.0502 ±	0.0023	165.1 ± 2.4			
MJK_132	0.00	287	51	0.18	2.96 ± 0.04	0.1145 ±	0.0018	1877.9 ± 22.9	1872 ± 28		-0.31
MJK_133	0.00	325	176	0.56	25.40 ± 0.47	0.0472 ±	0.0021	248.9 ± 4.5			
MJK_134	0.00	208	78	0.39	84.10 ± 2.50	0.0454 ±	0.0053	76.2 ± 2.2			
MJK_135	0.70	191	215	1.15	30.59 ± 0.65	0.0459 ±	0.0077	207.4 ± 4.3			
MJK_136	0.00	237	136	0.59	89.16 ± 2.74	0.0463 ±	0.0052	71.9 ± 2.2			
MJK_137	0.00	290	257	0.91	28.93 ± 0.51	0.0460 ±	0.0026	219.1 ± 3.8			
MJK_138	0.00	369	145	0.40	78.15 ± 1.75	0.0437 ±	0.0029	82.0 ± 1.8			
MJK_139	0.11	583	291	0.51	78.09 ± 1.67	0.0459 ±	0.0049	82.0 ± 1.7			
MJK_140	1.41	295	121	0.42	84.58 ± 2.28	0.0429 ±	0.0075	75.8 ± 2.0			
MJK_141	0.00	212	58	0.28	3.01 ± 0.04	0.1125 ±	0.0017	1850.9 ± 21.3	1842 ± 27		-0.48
MJK_142	0.00	402	128	0.33	24.93 ± 0.42	0.0539 ±	0.0021	253.5 ± 4.1			
MJK_143	0.00	243	58	0.25	3.01 ± 0.04	0.1147 ±	0.0017	1851.6 ± 19.7	1876 ± 27		1.30

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.
Table A3. Sample SGD. 試料SGD.

Labels	²⁰⁶ Pb _c ⁽¹⁾	U	Th	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾	Disc ⁽²⁾
	(%)	(ppm)	(ppm)				(Ma)	(Ma)	(%)
SGD_001	0.00	272	353	1.33	91.01 ± 2.46	0.0443 ± 0.0044	70.4 ± 1.9		
SGD_002#	0.36	466	263	0.58	62.62 ± 1.04	0.0650 ± 0.0056	102.1 ± 1.7		
SGD_003	1.52	248	301	1.24	98.80 ± 3.04	0.0397 ± 0.0130	64.9 ± 2.0		
SGD_004	0.78	438	186	0.44	81.85 ± 1.71	0.0452 ± 0.0057	78.3 ± 1.6		
SGD_005	0.00	400	159	0.41	77.57 ± 1.70	0.0541 ± 0.0035	82.6 ± 1.8		
SGD_006	0.95	141	68	0.49	87.10 ± 3.14	0.0289 ± 0.0105	73.6 ± 2.6		
SGD_007	3.75	225	171	0.78	75.99 ± 2.08	0.0505 ± 0.0113	84.3 ± 2.3		
SGD_008	1.81	294	570	1.99	108.33 ± 3.57	0.0349 ± 0.0208	59.2 ± 1.9		
SGD_009	0.00	391	447	1.17	81.22 ± 1.61	0.0444 ± 0.0035	78.9 ± 1.6		
SGD_010	0.00	130	173	1.36	101.09 ± 4.18	0.0604 ± 0.0094	63.5 ± 2.6		
SGD_011#	0.00	547	296	0.55	74.80 ± 1.53	0.0553 ± 0.0029	85.6 ± 1.7		
SGD_012	0.00	402	449	1.15	79.98 ± 1.52	0.0540 ± 0.0037	80.1 ± 1.5		
SGD_013	1.05	464	466	1.03	103.84 ± 2.47	0.0350 ± 0.0092	61.8 ± 1.5		
SGD_014	1.15	439	211	0.49	88.22 ± 1.74	0.0422 ± 0.0063	72.7 ± 1.4		
SGD_015#	0.55	517	135	0.27	4.15 ± 0.05	0.1078 ± 0.0015	1392.7 ± 15.6	1764 ± 24	21.05
SGD_016	0.00	1320	721	0.56	86.19 ± 1.30	0.0487 ± 0.0020	74.4 ± 1.1		
SGD_017	0.00	85	82	0.99	83.72 ± 4.22	0.0280 ± 0.0100	76.5 ± 3.8		
SGD_018	0.15	570	315	0.57	84.66 ± 1.71	0.0472 ± 0.0049	75.7 ± 1.5		
SGD_019	0.36	329	140	0.44	82.59 ± 2.04	0.0472 ± 0.0061	77.6 ± 1.9		
SGD_020	1.42	68	71	1.08	98.40 ± 6.60	0.0269 ± 0.0261	65.2 ± 4.4		
SGD_021	0.00	97	96	1.02	110.38 ± 5.68	0.0422 ± 0.0108	58.1 ± 3.0		
SGD_022	0.00	93	130	1.44	112.28 ± 5.88	0.0326 ± 0.0091	57.2 ± 3.0		
SGD_023	0.00	80	55	0.71	92.25 ± 4.68	0.0459 ± 0.0099	69.5 ± 3.5		
SGD_024	0.53	611	255	0.43	88.52 ± 1.86	0.0446 ± 0.0051	72.4 ± 1.5		
SGD_025	0.42	443	209	0.48	82.65 ± 1.87	0.0431 ± 0.0053	77.5 ± 1.7		
SGD_026	1.12	304	319	1.08	88.38 ± 2.48	0.0325 ± 0.0108	72.5 ± 2.0		
SGD_027	0.00	119	127	1.09	102.46 ± 4.30	0.0369 ± 0.0078	62.6 ± 2.6		
SGD_028	0.00	389	154	0.41	2.91 ± 0.04	0.1259 ± 0.0015	1903.2 ± 21.5	2042 ± 21	6.79
SGD_029	0.59	115	115	1.03	107.34 ± 5.68	0.0395 ± 0.0211	59.8 ± 3.2		
SGD_030	0.00	210	127	0.62	3.13 ± 0.04	0.1129 ± 0.0018	1789.3 ± 18.2	1848 ± 29	3.17
SGD_031	0.12	738	422	0.59	85.50 ± 1.66	0.0431 ± 0.0050	75.0 ± 1.4		
SGD_032	0.00	486	401	0.85	83.66 ± 1.72	0.0509 ± 0.0037	76.6 ± 1.6		
SGD_033	0.00	146	190	1.33	97.48 ± 3.94	0.0577 ± 0.0093	65.8 ± 2.6		
SGD_034	1.64	215	136	0.65	84.68 ± 2.69	0.0319 ± 0.0116	75.7 ± 2.4		
SGD_035	0.00	840	412	0.50	82.17 ± 1.47	0.0436 ± 0.0023	78.0 ± 1.4		
SGD_036	2.64	158	227	1.47	106.95 ± 4.61	0.0342 ± 0.0221	60.0 ± 2.6		
SGD_037#	2.41	256	82	0.33	80.97 ± 2.97	0.0676 ± 0.0099	79.1 ± 2.9		
SGD_038	0.00	220	182	0.85	81.40 ± 2.48	0.0550 ± 0.0056	78.7 ± 2.4		
SGD_039	0.00	218	125	0.59	98.95 ± 2.58	0.0414 ± 0.0047	64.8 ± 1.7		
SGD_040	0.00	143	147	1.05	99.45 ± 4.07	0.0474 ± 0.0083	64.5 ± 2.6		
SGD_041	1.61	209	328	1.61	97.65 ± 4.04	0.0359 ± 0.0222	65.7 ± 2.7		
SGD_042	0.17	94	99	1.08	97.25 ± 4.90	0.0399 ± 0.0212	65.9 ± 3.3		
SGD_043	0.00	391	247	0.65	100.82 ± 2.50	0.0511 ± 0.0042	63.6 ± 1.6		
SGD_044	0.00	231	125	0.55	2.98 ± 0.04	0.1166 ± 0.0020	1864.6 ± 20.8	1906 ± 31	2.17
SGD_045	0.00	310	284	0.94	73.07 ± 1.92	0.0479 ± 0.0046	87.6 ± 2.3		
SGD_046	1.94	188	222	1.21	103.61 ± 4.36	0.0365 ± 0.0175	61.9 ± 2.6		
SGD_047	0.00	105	121	1.19	101.42 ± 4.57	0.0503 ± 0.0095	63.3 ± 2.8		
SGD_048	0.61	374	263	0.72	85.28 ± 2.34	0.0437 ± 0.0082	75.2 ± 2.1		
SGD_049	2.31	123	104	0.86	98.30 ± 3.92	0.0318 ± 0.0171	65.2 ± 2.6		
SGD_050	1.40	486	477	1.01	89.51 ± 2.27	0.0394 ± 0.0093	71.6 ± 1.8		
SGD_051	2.39	137	124	0.93	90.29 ± 3.71	0.0333 ± 0.0185	71.0 ± 2.9		
SGD_052	0.00	218	117	0.55	72.28 ± 2.17	0.0425 ± 0.0048	88.6 ± 2.6		
SGD_053	0.64	295	435	1.51	110.63 ± 3.83	0.0488 ± 0.0170	58.0 ± 2.0		
SGD_054	0.00	474	434	0.94	83.03 ± 1.57	0.0441 ± 0.0032	77.2 ± 1.4		
SGD_055	0.32	160	126	0.81	75.77 ± 2.71	0.0433 ± 0.0123	84.5 ± 3.0		
SGD_056	0.27	197	296	1.54	101.00 ± 3.91	0.0576 ± 0.0198	63.5 ± 2.4		
SGD_057	0.02	534	81	0.16	3.06 ± 0.03	0.1157 ± 0.0014	1824.0 ± 18.1	1892 ± 23	3.59
SGD_058	0.00	89	63	0.72	23.57 ± 0.64	0.0594 ± 0.0048	267.9 ± 7.2		
SGD_059	0.70	447	295	0.68	75.11 ± 1.71	0.0429 ± 0.0069	85.3 ± 1.9		
SGD_060	0.00	278	348	1.28	105.22 ± 3.29	0.0402 ± 0.0044	61.0 ± 1.9		
SGD_061	2.47	236	102	0.44	105.55 ± 3.38	0.0344 ± 0.0108	60.8 ± 1.9		
SGD_062	0.00	289	59	0.21	3.07 ± 0.04	0.1157 ± 0.0017	1816.8 ± 18.6	1892 ± 26	3.98
SGD_063	1.75	256	225	0.90	89.04 ± 2.84	0.0339 ± 0.0125	72.0 ± 2.3		
SGD_064	0.00	113	111	1.01	89.13 ± 4.11	0.0521 ± 0.0084	71.9 ± 3.3		
SGD_065	2.78	340	140	0.42	87.49 ± 2.30	0.0462 ± 0.0087	73.3 ± 1.9		
SGD_066	0.00	962	961	1.02	81.41 ± 1.24	0.0497 ± 0.0024	78.7 ± 1.2		
SGD_067	1.18	384	236	0.63	84.77 ± 2.22	0.0417 ± 0.0085	75.6 ± 2.0		
SGD_068#	0.00	495	531	1.10	82.12 ± 1.77	0.0555 ± 0.0034	78.0 ± 1.7		
SGD_069	1.30	104	105	1.04	102.10 ± 4.86	0.0445 ± 0.0215	62.8 ± 3.0		
SGD_070	1.80	205	267	1.34	100.96 ± 4.03	0.0234 ± 0.0196	63.5 ± 2.5		
SGD_071	0.38	80	82	1.05	93.05 ± 5.01	0.0383 ± 0.0244	68.9 ± 3.7		
SGD_072	0.00	581	373	0.66	66.88 ± 1.38	0.0478 ± 0.0031	95.7 ± 2.0		
SGD_073	0.00	173	202	1.20	98.43 ± 3.09	0.0446 ± 0.0063	65.2 ± 2.0		
SGD_074	0.00	955	499	0.54	82.52 ± 1.34	0.0499 ± 0.0024	77.7 ± 1.2		
SGD_075	0.00	860	346	0.41	61.27 ± 0.94	0.0483 ± 0.0019	104.4 ± 1.6		

SGD_076	1.59	289	204	0.72	101.80 ± 3.21	0.0407 ± 0.0116	63.0 ± 2.0				
SGD_077	0.00	360	341	0.97	97.08 ± 2.34	0.0539 ± 0.0047	66.1 ± 1.6				
SGD_078	0.00	321	246	0.79	77.76 ± 2.14	0.0438 ± 0.0037	82.4 ± 2.3				
SGD_079	0.39	450	344	0.78	77.77 ± 1.90	0.0443 ± 0.0085	82.4 ± 2.0				
SGD_080	0.35	358	284	0.81	102.88 ± 2.83	0.0430 ± 0.0101	62.4 ± 1.7				
SGD_081	0.00	217	180	0.85	76.89 ± 2.04	0.0478 ± 0.0049	83.3 ± 2.2				
SGD_082	0.05	160	93	0.59	102.11 ± 3.68	0.0631 ± 0.0129	62.8 ± 2.3				
SGD_083	1.81	184	201	1.12	77.94 ± 2.91	0.0693 ± 0.0170	82.2 ± 3.0				
SGD_084	0.00	207	288	1.43	92.87 ± 3.08	0.0439 ± 0.0063	69.0 ± 2.3				
SGD_085	0.10	106	60	0.58	2.47 ± 0.04	0.1347 ± 0.0032	2189.7 ± 30.6	2161 ± 42			-1.33
SGD_086	1.04	100	57	0.58	88.61 ± 4.27	0.0450 ± 0.0154	72.3 ± 3.5				
SGD_087	0.00	251	130	0.53	85.57 ± 2.48	0.0399 ± 0.0041	74.9 ± 2.2				
SGD_088	0.97	265	188	0.73	83.52 ± 2.48	0.0393 ± 0.0097	76.7 ± 2.3				
SGD_089	0.00	551	366	0.68	3.16 ± 0.04	0.1165 ± 0.0014	1773.1 ± 18.6	1904 ± 21			6.88
SGD_090	1.23	129	97	0.77	76.01 ± 3.44	0.0453 ± 0.0172	84.3 ± 3.8				
SGD_091	0.00	120	61	0.53	3.23 ± 0.05	0.1183 ± 0.0025	1738.5 ± 24.1	1932 ± 37			10.01
SGD_092	0.00	87	113	1.33	85.24 ± 3.80	0.0470 ± 0.0108	75.2 ± 3.3				
SGD_093	0.32	95	94	1.01	106.55 ± 6.38	0.0366 ± 0.0217	60.2 ± 3.6				
SGD_094	2.71	138	86	0.64	89.79 ± 3.78	0.0296 ± 0.0149	71.4 ± 3.0				
SGD_095	0.42	346	124	0.37	79.97 ± 1.95	0.0489 ± 0.0059	80.1 ± 1.9				
SGD_096	0.33	218	256	1.20	107.31 ± 3.70	0.0632 ± 0.0144	59.8 ± 2.1				
SGD_097	0.57	56	56	1.02	2.29 ± 0.05	0.1568 ± 0.0066	2338.3 ± 42.0	2423 ± 70			3.50
SGD_098	0.00	203	158	0.80	77.31 ± 1.97	0.0419 ± 0.0046	82.9 ± 2.1				
SGD_099	0.54	117	120	1.05	108.49 ± 5.02	0.0502 ± 0.0190	59.1 ± 2.7				
SGD_100	0.00	146	149	1.05	93.32 ± 3.31	0.0413 ± 0.0061	68.7 ± 2.4				
SGD_101	0.00	321	256	0.82	87.37 ± 2.06	0.0545 ± 0.0049	73.4 ± 1.7				
SGD_102	3.90	171	112	0.67	86.92 ± 2.64	0.0473 ± 0.0161	73.7 ± 2.2				
SGD_103	0.00	930	457	0.50	73.66 ± 1.35	0.0484 ± 0.0020	86.9 ± 1.6				
SGD_104	0.00	346	403	1.19	83.15 ± 1.98	0.0416 ± 0.0036	77.1 ± 1.8				
SGD_105	0.03	1226	447	0.37	77.53 ± 1.24	0.0521 ± 0.0030	82.6 ± 1.3				
SGD_106	0.00	326	178	0.56	102.35 ± 2.36	0.0450 ± 0.0042	62.7 ± 1.4				
SGD_107	0.22	286	145	0.52	86.58 ± 2.32	0.0528 ± 0.0080	74.0 ± 2.0				
SGD_108	0.00	168	132	0.81	80.23 ± 2.61	0.0459 ± 0.0062	79.9 ± 2.6				
SGD_109	2.45	183	173	0.97	87.57 ± 3.54	0.0472 ± 0.0156	73.2 ± 2.9				
SGD_110	1.84	508	335	0.68	76.60 ± 1.56	0.0347 ± 0.0066	83.6 ± 1.7				
SGD_111	0.00	361	163	0.46	75.20 ± 1.60	0.0426 ± 0.0032	85.2 ± 1.8				
SGD_112	0.00	292	79	0.28	80.19 ± 2.23	0.0497 ± 0.0036	79.9 ± 2.2				
SGD_113	1.51	262	256	1.00	109.95 ± 3.78	0.0312 ± 0.0119	58.4 ± 2.0				
SGD_114	0.73	712	66	0.09	3.68 ± 0.07	0.1081 ± 0.0023	1548.4 ± 25.6	1768 ± 39			12.42
SGD_115	0.35	46	30	0.67	12.42 ± 0.36	0.0512 ± 0.0083	499.3 ± 13.8				
SGD_116	0.00	161	102	0.65	88.93 ± 2.89	0.0615 ± 0.0073	72.1 ± 2.3				
SGD_117#	0.00	282	663	2.41	96.45 ± 2.87	0.0631 ± 0.0055	66.5 ± 2.0				
SGD_118	0.00	856	449	0.54	79.84 ± 1.36	0.0462 ± 0.0021	80.2 ± 1.4				
SGD_119	0.00	323	240	0.76	104.87 ± 2.55	0.0432 ± 0.0043	61.2 ± 1.5				
SGD_120	0.00	263	154	0.60	94.93 ± 2.58	0.0430 ± 0.0044	67.6 ± 1.8				
SGD_121	0.23	1050	395	0.39	88.40 ± 1.42	0.0484 ± 0.0037	72.5 ± 1.2				
SGD_122	0.00	95	84	0.90	93.69 ± 5.07	0.0523 ± 0.0135	68.4 ± 3.7				
SGD_123	1.83	110	104	0.96	101.31 ± 5.01	0.0468 ± 0.0227	63.3 ± 3.1				
SGD_124	0.00	397	216	0.56	78.86 ± 1.66	0.0471 ± 0.0041	81.2 ± 1.7				
SGD_125	1.97	279	349	1.28	77.80 ± 2.59	0.0462 ± 0.0131	82.3 ± 2.7				
SGD_126	0.00	947	929	1.01	75.62 ± 1.35	0.0442 ± 0.0021	84.7 ± 1.5				
SGD_127	0.00	193	195	1.03	96.81 ± 2.98	0.0448 ± 0.0072	66.2 ± 2.0				
SGD_128	1.94	349	348	1.02	95.91 ± 3.05	0.0242 ± 0.0126	66.9 ± 2.1				
SGD_129	0.00	364	147	0.42	77.68 ± 1.78	0.0476 ± 0.0038	82.5 ± 1.9				
SGD_130	1.79	506	317	0.64	103.86 ± 2.63	0.0378 ± 0.0080	61.8 ± 1.6				
SGD_131	0.49	312	251	0.83	75.76 ± 1.96	0.0396 ± 0.0096	84.5 ± 2.2				
SGD_132#	0.00	338	415	1.26	95.97 ± 2.64	0.0586 ± 0.0052	66.8 ± 1.8				
SGD_133	0.00	85	32	0.38	82.27 ± 3.74	0.0357 ± 0.0091	77.9 ± 3.5				
SGD_134	0.00	650	507	0.80	64.71 ± 1.39	0.0434 ± 0.0025	98.9 ± 2.1				
SGD_135	0.00	201	311	1.59	97.84 ± 3.28	0.0576 ± 0.0070	65.6 ± 2.2				
SGD_136	2.87	296	241	0.83	92.94 ± 2.77	0.0393 ± 0.0121	69.0 ± 2.0				
SGD_137	1.13	153	117	0.78	68.62 ± 2.42	0.0338 ± 0.0119	93.3 ± 3.3				
SGD_138#	0.00	759	506	0.68	87.03 ± 1.59	0.0526 ± 0.0025	73.6 ± 1.3				
SGD_139	0.39	520	237	0.47	83.04 ± 1.59	0.0496 ± 0.0057	77.2 ± 1.5				
SGD_140	0.00	620	28	0.05	3.11 ± 0.04	0.1177 ± 0.0014	1798.4 ± 19.5	1923 ± 21			6.48
SGD_141#	0.00	614	534	0.89	96.35 ± 2.02	0.0557 ± 0.0038	66.6 ± 1.4				
SGD_142	0.86	530	218	0.42	78.21 ± 1.63	0.0534 ± 0.0057	81.9 ± 1.7				
SGD_143#	0.00	610	229	0.38	64.16 ± 1.14	0.0559 ± 0.0030	99.7 ± 1.8				
SGD_144	0.00	244	284	1.20	97.84 ± 3.40	0.0460 ± 0.0051	65.6 ± 2.3				

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U} - ^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1 - (238\text{U}/206\text{Pb}^* \text{ age}) / (207\text{Pb}^*/206\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.

Table A4. Sample NTK. 試料NTK.

Labels	²⁰⁶ Pb _c ⁽¹⁾	U	Th	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾	Disc ⁽³⁾
	(%)	(ppm)	(ppm)				(Ma)	(Ma)	(%)
NTK_001	2.45	403	209	0.53	60.99 ± 1.22	0.0450 ± 0.0068	104.8 ± 2.1		
NTK_002	0.00	663	777	1.20	76.59 ± 1.60	0.0476 ± 0.0020	83.6 ± 1.7		
NTK_003	0.31	529	308	0.60	76.23 ± 1.78	0.0467 ± 0.0057	84.0 ± 2.0		
NTK_004	0.00	167	98	0.60	100.15 ± 3.47	0.0541 ± 0.0058	64.0 ± 2.2		
NTK_005	1.57	164	98	0.61	97.49 ± 3.18	0.0512 ± 0.0135	65.8 ± 2.1		
NTK_006	1.13	384	232	0.62	92.21 ± 2.09	0.0426 ± 0.0065	69.5 ± 1.6		
NTK_007	0.58	160	59	0.38	97.37 ± 3.22	0.0444 ± 0.0088	65.9 ± 2.2		
NTK_008	1.23	178	111	0.64	77.96 ± 2.16	0.0371 ± 0.0094	82.2 ± 2.3		
NTK_009	0.99	221	103	0.48	99.19 ± 2.60	0.0325 ± 0.0080	64.7 ± 1.7		
NTK_010	0.59	203	116	0.58	98.72 ± 2.94	0.0422 ± 0.0091	65.0 ± 1.9		
NTK_011	0.00	957	868	0.93	66.56 ± 0.90	0.0481 ± 0.0018	96.1 ± 1.3		
NTK_012	1.09	144	54	0.38	97.59 ± 3.62	0.0369 ± 0.0100	65.7 ± 2.4		
NTK_013	0.82	1074	587	0.56	65.89 ± 1.22	0.0448 ± 0.0038	97.1 ± 1.8		
NTK_014	0.20	1053	700	0.68	91.17 ± 1.45	0.0424 ± 0.0036	70.3 ± 1.1		
NTK_015	0.73	414	242	0.60	71.93 ± 1.61	0.0449 ± 0.0063	89.0 ± 2.0		
NTK_016	0.11	242	150	0.64	74.59 ± 2.10	0.0440 ± 0.0081	85.8 ± 2.4		
NTK_017	0.00	246	134	0.56	97.82 ± 2.60	0.0531 ± 0.0047	65.6 ± 1.7		
NTK_018	1.01	181	153	0.87	95.54 ± 2.89	0.0342 ± 0.0112	67.1 ± 2.0		
NTK_019	0.74	226	129	0.59	90.39 ± 2.51	0.0456 ± 0.0084	70.9 ± 2.0		
NTK_020	0.00	259	152	0.60	93.62 ± 2.76	0.0547 ± 0.0049	68.5 ± 2.0		
NTK_021	0.45	290	166	0.59	99.58 ± 2.73	0.0383 ± 0.0083	64.4 ± 1.8		
NTK_022	0.39	105	32	0.31	101.40 ± 4.35	0.0451 ± 0.0129	63.3 ± 2.7		
NTK_023	1.17	192	81	0.43	94.63 ± 2.95	0.0374 ± 0.0095	67.8 ± 2.1		
NTK_024	0.00	767	292	0.39	66.03 ± 1.06	0.0465 ± 0.0021	96.9 ± 1.5		
NTK_025	0.00	227	117	0.53	86.13 ± 2.24	0.0500 ± 0.0052	74.4 ± 1.9		
NTK_026	0.00	69	41	0.61	90.09 ± 4.07	0.0568 ± 0.0100	71.2 ± 3.2		
NTK_027	0.81	245	101	0.42	98.69 ± 3.00	0.0428 ± 0.0077	65.0 ± 2.0		
NTK_028	0.00	143	36	0.26	96.37 ± 3.22	0.0371 ± 0.0062	66.5 ± 2.2		
NTK_029	0.19	390	198	0.52	73.72 ± 1.55	0.0479 ± 0.0056	86.9 ± 1.8		
NTK_030	0.16	728	377	0.53	63.72 ± 1.19	0.0479 ± 0.0037	100.4 ± 1.9		
NTK_031	1.59	279	177	0.65	97.69 ± 2.92	0.0375 ± 0.0094	65.6 ± 1.9		
NTK_032	1.06	306	158	0.53	82.92 ± 1.90	0.0457 ± 0.0074	77.3 ± 1.8		
NTK_033	4.58	277	138	0.51	88.61 ± 2.51	0.0421 ± 0.0128	72.3 ± 2.0		
NTK_034	0.00	153	46	0.31	99.39 ± 3.33	0.0514 ± 0.0063	64.5 ± 2.2		
NTK_035	0.00	860	680	0.81	65.76 ± 1.05	0.0491 ± 0.0018	97.3 ± 1.5		
NTK_036	0.56	648	263	0.42	79.93 ± 1.42	0.0413 ± 0.0039	80.2 ± 1.4		
NTK_037	0.00	297	166	0.57	68.09 ± 1.29	0.0491 ± 0.0031	94.0 ± 1.8		
NTK_038	0.00	42	23	0.56	96.88 ± 5.12	0.0434 ± 0.0168	66.2 ± 3.5		
NTK_039	0.88	189	85	0.46	99.22 ± 2.94	0.0571 ± 0.0099	64.6 ± 1.9		
NTK_040	0.00	152	38	0.26	91.50 ± 3.20	0.0430 ± 0.0051	70.1 ± 2.4		
NTK_041	1.43	256	100	0.40	69.12 ± 1.72	0.0381 ± 0.0060	92.6 ± 2.3		
NTK_042	0.00	157	50	0.33	96.53 ± 3.09	0.0368 ± 0.0058	66.4 ± 2.1		
NTK_043	0.00	124	58	0.48	93.99 ± 3.55	0.0356 ± 0.0060	68.2 ± 2.6		
NTK_044	0.00	131	90	0.71	97.80 ± 4.41	0.0404 ± 0.0095	65.6 ± 2.9		
NTK_045	0.39	518	209	0.41	76.00 ± 1.38	0.0516 ± 0.0046	84.3 ± 1.5		
NTK_046	0.00	87	21	0.25	93.48 ± 4.44	0.0391 ± 0.0079	68.6 ± 3.2		
NTK_047	0.00	670	433	0.66	82.13 ± 1.57	0.0472 ± 0.0025	78.0 ± 1.5		
NTK_048	1.05	128	132	1.06	35.64 ± 0.86	0.0350 ± 0.0091	178.4 ± 4.3		
NTK_049	0.19	174	95	0.56	90.91 ± 2.92	0.0508 ± 0.0104	70.5 ± 2.3		
NTK_050	0.00	88	39	0.46	97.93 ± 4.05	0.0372 ± 0.0089	65.5 ± 2.7		
NTK_051	0.71	356	382	1.10	69.33 ± 1.56	0.0487 ± 0.0081	92.3 ± 2.1		
NTK_052	0.00	372	183	0.50	76.58 ± 1.78	0.0458 ± 0.0031	83.6 ± 1.9		
NTK_053	1.16	208	144	0.71	93.02 ± 2.57	0.0358 ± 0.0093	68.9 ± 1.9		
NTK_054	0.00	110	48	0.44	93.69 ± 3.23	0.0355 ± 0.0076	68.4 ± 2.3		
NTK_055	0.00	60	26	0.44	89.76 ± 4.17	0.0437 ± 0.0082	71.4 ± 3.3		
NTK_056	1.39	308	263	0.88	92.88 ± 1.97	0.0343 ± 0.0085	69.0 ± 1.5		
NTK_057	0.00	455	250	0.57	66.50 ± 1.27	0.0468 ± 0.0026	96.2 ± 1.8		
NTK_058	0.00	225	110	0.50	82.38 ± 1.82	0.0509 ± 0.0037	77.8 ± 1.7		
NTK_059	0.00	187	89	0.49	99.38 ± 2.70	0.0555 ± 0.0048	64.5 ± 1.7		
NTK_060	0.00	179	60	0.35	90.03 ± 2.28	0.0483 ± 0.0051	71.2 ± 1.8		
NTK_061	1.12	305	202	0.68	82.16 ± 1.97	0.0386 ± 0.0070	78.0 ± 1.9		
NTK_062	0.00	73	46	0.64	81.89 ± 2.95	0.0342 ± 0.0073	78.2 ± 2.8		
NTK_063	0.45	332	189	0.58	83.10 ± 2.07	0.0478 ± 0.0075	77.1 ± 1.9		
NTK_064	0.16	120	46	0.39	93.46 ± 3.21	0.0410 ± 0.0098	68.6 ± 2.3		
NTK_065	2.49	147	100	0.70	93.68 ± 2.99	0.0555 ± 0.0135	68.5 ± 2.2		
NTK_066	0.13	401	244	0.62	79.86 ± 1.56	0.0516 ± 0.0058	80.2 ± 1.6		
NTK_067	0.00	112	41	0.37	93.59 ± 3.38	0.0441 ± 0.0058	68.5 ± 2.5		
NTK_068	1.02	268	172	0.66	78.93 ± 1.92	0.0453 ± 0.0076	81.2 ± 2.0		
NTK_069	1.27	151	71	0.48	96.49 ± 3.21	0.0457 ± 0.0094	66.5 ± 2.2		
NTK_070	0.00	424	361	0.87	95.71 ± 1.86	0.0493 ± 0.0031	67.0 ± 1.3		
NTK_071	0.00	144	43	0.30	95.11 ± 2.78	0.0477 ± 0.0050	67.4 ± 2.0		
NTK_072	0.00	827	433	0.54	89.76 ± 1.33	0.0496 ± 0.0021	71.4 ± 1.1		
NTK_073	0.11	416	431	1.06	89.60 ± 1.90	0.0493 ± 0.0077	71.5 ± 1.5		

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}$ - $^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.
Table A5. Sample TKO2. 試料TKO2.

Labels	²⁰⁶ Pb _c ⁽¹⁾	U	Th	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾		²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾	Disc ⁽²⁾
	(%)	(ppm)	(ppm)					(Ma)	(Ma)	(%)
TAK2_001	0.00	1132	741	0.67	86.31 ± 1.18	0.0507 ±	0.0019	74.3 ± 1.0		
TAK2_002	0.28	574	286	0.51	92.20 ± 1.39	0.0447 ±	0.0043	69.5 ± 1.0		
TAK2_003	0.00	382	298	0.80	83.45 ± 1.53	0.0507 ±	0.0029	76.8 ± 1.4		
TAK2_004	0.00	242	147	0.62	67.92 ± 1.51	0.0501 ±	0.0038	94.2 ± 2.1		
TAK2_005	0.00	340	134	0.40	84.65 ± 1.54	0.0525 ±	0.0027	75.7 ± 1.4		
TAK2_006	0.14	204	108	0.55	84.84 ± 2.27	0.0469 ±	0.0076	75.5 ± 2.0		
TAK2_007	0.00	767	336	0.45	71.71 ± 0.93	0.0485 ±	0.0019	89.3 ± 1.1		
TAK2_008	0.88	213	37	0.18	3.09 ± 0.03	0.1174 ±	0.0017	1807.8 ± 16.2	1919 ± 26	5.79
TAK2_009	0.14	302	171	0.58	63.82 ± 1.31	0.0458 ±	0.0050	100.2 ± 2.0		
TAK2_010	0.00	272	226	0.85	78.31 ± 1.79	0.0499 ±	0.0033	81.8 ± 1.9		
TAK2_011	0.00	272	193	0.73	93.26 ± 2.04	0.0458 ±	0.0029	68.8 ± 1.5		
TAK2_012	0.00	132	78	0.61	91.41 ± 2.83	0.0524 ±	0.0051	70.1 ± 2.2		
TAK2_013#	0.00	287	245	0.88	89.58 ± 1.90	0.0543 ±	0.0033	71.6 ± 1.5		
TAK2_014	0.07	196	90	0.47	96.86 ± 2.40	0.0452 ±	0.0077	66.2 ± 1.6		
TAK2_015	0.03	288	180	0.64	89.19 ± 1.80	0.0485 ±	0.0070	71.9 ± 1.4		
TAK2_016	0.00	298	173	0.60	90.99 ± 1.90	0.0471 ±	0.0035	70.5 ± 1.5		
TAK2_017#	0.00	298	110	0.38	63.05 ± 1.22	0.0537 ±	0.0027	101.4 ± 1.9		
TAK2_018	0.00	48	23	0.50	85.29 ± 3.96	0.0463 ±	0.0117	75.1 ± 3.5		
TAK2_019	0.00	337	168	0.51	90.61 ± 1.76	0.0485 ±	0.0034	70.8 ± 1.4		
TAK2_020	0.31	307	130	0.43	61.78 ± 1.08	0.0487 ±	0.0049	103.5 ± 1.8		
TAK2_021	4.28	293	263	0.92	62.29 ± 1.52	0.0498 ±	0.0141	102.7 ± 2.5		
TAK2_022	0.00	361	251	0.72	99.58 ± 2.05	0.0445 ±	0.0029	64.4 ± 1.3		
TAK2_023	0.00	730	170	0.24	68.24 ± 0.95	0.0480 ±	0.0021	93.8 ± 1.3		
TAK2_024	1.11	210	128	0.63	72.34 ± 1.68	0.0465 ±	0.0079	88.5 ± 2.0		
TAK2_025	0.00	416	204	0.50	66.44 ± 1.05	0.0439 ±	0.0024	96.3 ± 1.5		
TAK2_026	0.00	67	39	0.61	91.13 ± 3.88	0.0584 ±	0.0099	70.4 ± 3.0		
TAK2_027	0.00	365	170	0.48	74.17 ± 1.16	0.0521 ±	0.0032	86.3 ± 1.3		
TAK2_028	0.38	287	151	0.54	66.77 ± 1.24	0.0420 ±	0.0057	95.8 ± 1.8		
TAK2_029	0.58	63	32	0.52	96.59 ± 3.99	0.0447 ±	0.0149	66.4 ± 2.7		
TAK2_030	1.15	282	182	0.66	91.55 ± 2.07	0.0458 ±	0.0074	70.0 ± 1.6		
TAK2_031	0.00	243	217	0.92	85.27 ± 1.83	0.0464 ±	0.0039	75.2 ± 1.6		
TAK2_032	0.00	561	870	1.59	73.75 ± 1.15	0.0491 ±	0.0025	86.8 ± 1.3		
TAK2_033	0.00	34	17	0.52	90.97 ± 4.91	0.0464 ±	0.0126	70.5 ± 3.8		
TAK2_034	0.17	876	374	0.44	25.01 ± 0.31	0.0488 ±	0.0018	252.7 ± 3.1		
TAK2_035	0.00	712	598	0.86	89.28 ± 1.43	0.0504 ±	0.0025	71.8 ± 1.1		
TAK2_036	2.34	160	135	0.87	25.14 ± 0.47	0.0472 ±	0.0069	251.4 ± 4.6		
TAK2_037	0.00	436	473	1.11	94.80 ± 1.77	0.0445 ±	0.0025	67.6 ± 1.3		
TAK2_038	0.56	208	133	0.65	92.52 ± 2.25	0.0491 ±	0.0090	69.3 ± 1.7		
TAK2_039	0.23	753	229	0.31	60.97 ± 0.81	0.0433 ±	0.0027	104.9 ± 1.4		
TAK2_040	0.00	59	34	0.59	89.54 ± 4.07	0.0554 ±	0.0092	71.6 ± 3.2		
TAK2_041	0.00	579	323	0.57	92.14 ± 1.47	0.0500 ±	0.0027	69.6 ± 1.1		
TAK2_042	0.00	954	705	0.76	91.44 ± 1.26	0.0488 ±	0.0018	70.1 ± 1.0		
TAK2_043	0.00	319	269	0.87	68.01 ± 1.11	0.0483 ±	0.0032	94.1 ± 1.5		
TAK2_044	0.00	205	121	0.61	94.03 ± 2.29	0.0483 ±	0.0041	68.2 ± 1.7		
TAK2_045	0.00	275	218	0.81	81.20 ± 1.76	0.0469 ±	0.0038	78.9 ± 1.7		
TAK2_046	0.52	546	353	0.66	75.71 ± 1.26	0.0435 ±	0.0055	84.6 ± 1.4		
TAK2_047	0.00	722	326	0.46	67.30 ± 0.77	0.0488 ±	0.0020	95.1 ± 1.1		
TAK2_048	0.26	239	89	0.38	63.66 ± 1.25	0.0471 ±	0.0055	100.5 ± 2.0		
TAK2_049	3.66	129	71	0.56	94.84 ± 3.12	0.0456 ±	0.0146	67.6 ± 2.2		
TAK2_050	0.46	142	82	0.59	79.45 ± 2.65	0.0492 ±	0.0103	80.6 ± 2.7		
TAK2_051	0.45	167	106	0.65	88.48 ± 2.59	0.0507 ±	0.0093	72.5 ± 2.1		
TAK2_052	0.00	172	68	0.40	64.85 ± 1.63	0.0448 ±	0.0039	98.6 ± 2.5		
TAK2_053	0.00	378	223	0.60	94.02 ± 1.88	0.0541 ±	0.0036	68.2 ± 1.4		
TAK2_054	0.00	117	59	0.52	75.12 ± 2.03	0.0391 ±	0.0045	85.3 ± 2.3		
TAK2_055	0.00	426	217	0.52	94.33 ± 1.66	0.0427 ±	0.0032	68.0 ± 1.2		
TAK2_056	0.00	569	293	0.53	63.97 ± 0.98	0.0464 ±	0.0020	100.0 ± 1.5		
TAK2_057	0.00	411	135	0.34	61.24 ± 0.97	0.0496 ±	0.0023	104.4 ± 1.6		
TAK2_058	0.59	194	77	0.41	70.59 ± 1.63	0.0371 ±	0.0065	90.7 ± 2.1		
TAK2_059#	0.00	322	165	0.52	87.39 ± 1.85	0.0580 ±	0.0036	73.3 ± 1.5		
TAK2_060	1.45	201	53	0.27	98.98 ± 2.57	0.0442 ±	0.0077	64.8 ± 1.7		
TAK2_061	0.73	152	70	0.47	93.25 ± 2.88	0.0358 ±	0.0091	68.8 ± 2.1		
TAK2_062	0.00	355	216	0.62	84.32 ± 1.45	0.0497 ±	0.0028	76.0 ± 1.3		
TAK2_063	0.00	289	234	0.83	94.53 ± 1.90	0.0449 ±	0.0039	67.8 ± 1.4		
TAK2_064	0.17	317	169	0.55	95.20 ± 2.40	0.0379 ±	0.0062	67.4 ± 1.7		
TAK2_065	0.48	235	137	0.60	95.94 ± 2.30	0.0430 ±	0.0074	66.8 ± 1.6		
TAK2_066#	0.00	169	136	0.82	95.68 ± 2.74	0.0282 ±	0.0042	67.0 ± 1.9		
TAK2_067	0.00	444	244	0.56	74.11 ± 1.35	0.0438 ±	0.0028	86.4 ± 1.6		
TAK2_068	0.00	456	179	0.40	62.66 ± 1.00	0.0474 ±	0.0028	102.1 ± 1.6		
TAK2_069	0.00	166	91	0.56	97.95 ± 3.08	0.0550 ±	0.0056	65.5 ± 2.0		
TAK2_070	0.00	240	90	0.39	28.60 ± 0.45	0.0485 ±	0.0022	221.5 ± 3.5		
TAK2_071	0.00	307	131	0.44	82.29 ± 1.71	0.0492 ±	0.0038	77.9 ± 1.6		
TAK2_072	0.00	313	136	0.45	61.33 ± 1.03	0.0492 ±	0.0032	104.3 ± 1.7		
TAK2_073	0.00	263	78	0.30	60.22 ± 1.21	0.0510 ±	0.0036	106.2 ± 2.1		
TAK2_074	0.00	177	118	0.68	86.01 ± 2.45	0.0515 ±	0.0051	74.5 ± 2.1		
TAK2_075	0.46	254	224	0.90	88.32 ± 2.30	0.0491 ±	0.0088	72.6 ± 1.9		

TAK2_076	0.07	220	156	0.73	86.89 ± 2.37	0.0460 ±	0.0082	73.8 ± 2.0
TAK2_077#	0.00	245	221	0.93	80.60 ± 2.18	0.0581 ±	0.0048	79.5 ± 2.1
TAK2_078	0.03	222	156	0.72	85.95 ± 2.21	0.0425 ±	0.0096	74.6 ± 1.9
TAK2_079	0.00	471	327	0.71	90.37 ± 1.66	0.0463 ±	0.0035	70.9 ± 1.3
TAK2_080	0.00	676	490	0.74	91.66 ± 1.47	0.0439 ±	0.0023	69.9 ± 1.1
TAK2_081	0.00	159	87	0.56	89.15 ± 2.89	0.0432 ±	0.0043	71.9 ± 2.3
TAK2_082	0.00	291	137	0.48	86.63 ± 2.07	0.0406 ±	0.0045	74.0 ± 1.8
TAK2_083	0.00	535	244	0.47	90.01 ± 1.56	0.0454 ±	0.0027	71.2 ± 1.2
TAK2_084	0.00	1103	352	0.33	63.16 ± 0.72	0.0483 ±	0.0016	101.3 ± 1.1
TAK2_085	0.00	65	37	0.59	94.31 ± 4.19	0.0390 ±	0.0084	68.0 ± 3.0
TAK2_086	0.00	150	47	0.32	73.26 ± 1.94	0.0521 ±	0.0059	87.4 ± 2.3
TAK2_087#	0.00	612	301	0.50	76.24 ± 1.03	0.0527 ±	0.0022	84.0 ± 1.1
TAK2_088#	0.00	80	68	0.87	90.11 ± 3.69	0.0659 ±	0.0088	71.1 ± 2.9
TAK2_089	0.00	369	155	0.43	86.68 ± 1.35	0.0485 ±	0.0028	73.9 ± 1.1
TAK2_090	0.00	518	405	0.80	83.71 ± 1.48	0.0486 ±	0.0029	76.5 ± 1.3
TAK2_091	0.07	553	266	0.49	94.69 ± 1.65	0.0390 ±	0.0045	67.7 ± 1.2
TAK2_092	0.37	688	391	0.58	92.88 ± 1.61	0.0451 ±	0.0043	69.0 ± 1.2
TAK2_093	0.00	70	40	0.59	93.96 ± 3.79	0.0458 ±	0.0086	68.2 ± 2.7
TAK2_094	0.49	355	232	0.67	76.91 ± 1.65	0.0400 ±	0.0065	83.3 ± 1.8
TAK2_095	0.00	446	289	0.66	94.13 ± 1.69	0.0479 ±	0.0035	68.1 ± 1.2

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(238\text{U}/206\text{Pb}^* \text{ age})/(207\text{Pb}^*/206\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.
Table A6. Sample SB1. 試料SB1.

Labels	²⁰⁶ Pb _c ⁽¹⁾ (%)	U (ppm)	Th (ppm)	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾ (Ma)	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾ Disc ⁽²⁾ (Ma)	(%)
SHB1_001	0.00	760	372	0.50	82.87 ± 1.38	0.0441 ± 0.0025	77.3 ± 1.3		
SHB1_002	0.13	440	255	0.59	80.14 ± 1.74	0.0527 ± 0.0073	79.9 ± 1.7		
SHB1_003	0.87	479	281	0.60	74.81 ± 1.50	0.0581 ± 0.0076	85.6 ± 1.7		
SHB1_004	2.92	387	118	0.31	63.37 ± 1.19	0.0520 ± 0.0066	100.9 ± 1.9		
SHB1_005	0.00	393	292	0.76	2.47 ± 0.03	0.1433 ± 0.0016	2192.6 ± 20.7	2269 ± 20	3.37
SHB1_006	0.00	305	158	0.53	60.52 ± 1.36	0.0569 ± 0.0046	105.6 ± 2.3		
SHB1_007	1.36	147	112	0.78	89.19 ± 3.56	0.0239 ± 0.0162	71.9 ± 2.9		
SHB1_008	0.70	540	308	0.58	90.67 ± 2.15	0.0448 ± 0.0069	70.7 ± 1.7		
SHB1_009	1.67	222	154	0.71	97.29 ± 3.27	0.0439 ± 0.0136	65.9 ± 2.2		
SHB1_010	0.04	347	157	0.47	85.62 ± 1.85	0.0400 ± 0.0061	74.9 ± 1.6		
SHB1_011	0.00	380	289	0.78	92.94 ± 2.15	0.0419 ± 0.0041	69.0 ± 1.6		
SHB1_012	0.00	371	302	0.83	90.26 ± 2.38	0.0429 ± 0.0039	71.0 ± 1.9		
SHB1_013	0.00	184	102	0.57	64.62 ± 1.86	0.0569 ± 0.0060	99.0 ± 2.8		
SHB1_014	1.75	456	521	1.17	90.54 ± 2.16	0.0366 ± 0.0096	70.8 ± 1.7		
SHB1_015	2.05	344	161	0.48	57.38 ± 1.18	0.0367 ± 0.0077	111.4 ± 2.3		
SHB1_016	0.14	836	311	0.38	64.70 ± 1.07	0.0415 ± 0.0037	98.9 ± 1.6		
SHB1_017	0.00	179	101	0.58	90.64 ± 3.50	0.0472 ± 0.0067	70.7 ± 2.7		
SHB1_018	0.00	272	256	0.96	89.17 ± 2.52	0.0473 ± 0.0055	71.9 ± 2.0		
SHB1_019	0.00	455	323	0.73	66.51 ± 1.50	0.0428 ± 0.0034	96.2 ± 2.1		
SHB1_020	0.59	380	220	0.59	90.74 ± 2.51	0.0405 ± 0.0077	70.7 ± 1.9		
SHB1_021	0.98	445	210	0.48	86.43 ± 2.23	0.0391 ± 0.0073	74.2 ± 1.9		
SHB1_022	1.86	756	324	0.44	80.05 ± 1.49	0.0382 ± 0.0051	80.0 ± 1.5		
SHB1_023	0.00	292	113	0.40	72.92 ± 2.11	0.0497 ± 0.0052	87.8 ± 2.5		
SHB1_024	0.00	94	29	0.32	99.62 ± 5.14	0.0476 ± 0.0144	64.4 ± 3.3		
SHB1_025#	4.42	251	310	1.26	2.92 ± 0.05	0.0822 ± 0.0107	1899.1 ± 26.6	1251 ± 236	-51.80
SHB1_026	1.17	345	192	0.57	81.44 ± 2.09	0.0429 ± 0.0082	78.7 ± 2.0		
SHB1_027	0.21	288	247	0.88	88.09 ± 2.67	0.0463 ± 0.0116	72.8 ± 2.2		
SHB1_028	0.00	75	28	0.38	104.42 ± 6.71	0.0486 ± 0.0172	61.4 ± 3.9		
SHB1_029	0.00	227	128	0.58	90.42 ± 2.81	0.0487 ± 0.0063	70.9 ± 2.2		
SHB1_030	0.00	560	311	0.57	25.05 ± 0.31	0.0515 ± 0.0018	252.4 ± 3.1		
SHB1_031	0.00	120	37	0.32	88.42 ± 4.33	0.0486 ± 0.0087	72.5 ± 3.5		
SHB1_032	0.00	255	190	0.76	89.98 ± 2.72	0.0482 ± 0.0061	71.2 ± 2.1		
SHB1_033	0.03	344	218	0.65	26.06 ± 0.50	0.0525 ± 0.0055	242.7 ± 4.6		
SHB1_034	0.00	234	147	0.64	70.70 ± 2.05	0.0566 ± 0.0065	90.5 ± 2.6		
SHB1_035	1.44	265	130	0.50	70.38 ± 2.02	0.0539 ± 0.0089	90.9 ± 2.6		
SHB1_036	0.00	68	36	0.54	91.62 ± 6.00	0.0269 ± 0.0104	70.0 ± 4.6		
SHB1_037	0.00	149	102	0.70	81.46 ± 3.12	0.0512 ± 0.0074	78.7 ± 3.0		
SHB1_038	0.46	872	270	0.32	36.88 ± 0.52	0.0452 ± 0.0025	172.5 ± 2.4		
SHB1_039	1.64	319	292	0.94	87.54 ± 2.57	0.0516 ± 0.0124	73.2 ± 2.1		
SHB1_040	0.61	318	258	0.83	90.27 ± 2.78	0.0427 ± 0.0112	71.0 ± 2.2		
SHB1_041	2.35	205	143	0.72	81.10 ± 2.97	0.0620 ± 0.0125	79.0 ± 2.9		
SHB1_042	0.00	250	116	0.47	82.47 ± 2.33	0.0384 ± 0.0051	77.7 ± 2.2		
SHB1_043	0.39	466	150	0.33	62.89 ± 1.19	0.0513 ± 0.0055	101.7 ± 1.9		
SHB1_044#	0.00	265	132	0.51	76.09 ± 1.85	0.0585 ± 0.0053	84.2 ± 2.0		
SHB1_045	0.00	273	180	0.68	35.17 ± 0.62	0.0439 ± 0.0034	180.7 ± 3.1		
SHB1_046	1.18	551	463	0.86	87.38 ± 2.18	0.0372 ± 0.0074	73.4 ± 1.8		
SHB1_047	0.51	939	493	0.54	81.39 ± 1.30	0.0438 ± 0.0048	78.7 ± 1.2		
SHB1_048	0.00	240	183	0.78	86.17 ± 2.43	0.0491 ± 0.0062	74.4 ± 2.1		
SHB1_049	0.35	215	67	0.32	59.94 ± 1.73	0.0441 ± 0.0086	106.7 ± 3.1		
SHB1_050	0.26	200	78	0.40	95.60 ± 3.54	0.0382 ± 0.0107	67.1 ± 2.5		
SHB1_051	0.39	165	142	0.89	68.01 ± 2.39	0.0507 ± 0.0136	94.1 ± 3.3		
SHB1_052	0.48	655	592	0.93	95.68 ± 2.35	0.0450 ± 0.0090	67.0 ± 1.6		
SHB1_053	0.00	237	207	0.90	72.71 ± 2.05	0.0476 ± 0.0050	88.1 ± 2.5		
SHB1_054	0.00	174	116	0.68	90.01 ± 3.24	0.0439 ± 0.0079	71.2 ± 2.5		
SHB1_055	2.51	221	170	0.79	74.02 ± 2.63	0.0208 ± 0.0139	86.5 ± 3.0		
SHB1_056	0.00	194	146	0.77	68.50 ± 2.34	0.0476 ± 0.0072	93.4 ± 3.2		
SHB1_057	0.29	586	293	0.51	91.35 ± 2.24	0.0467 ± 0.0057	70.2 ± 1.7		
SHB1_058	0.00	61	40	0.67	33.22 ± 1.45	0.0387 ± 0.0067	191.2 ± 8.2		
SHB1_059	0.00	331	284	0.88	27.53 ± 0.56	0.0478 ± 0.0034	230.0 ± 4.6		
SHB1_060	0.53	223	105	0.48	79.92 ± 2.48	0.0396 ± 0.0102	80.2 ± 2.5		
SHB1_061	0.00	307	215	0.72	94.00 ± 2.91	0.0518 ± 0.0063	68.2 ± 2.1		
SHB1_062	0.00	566	266	0.48	79.08 ± 1.61	0.0458 ± 0.0060	81.0 ± 1.6		
SHB1_063	0.00	814	299	0.38	66.98 ± 1.38	0.0428 ± 0.0029	95.5 ± 2.0		
SHB1_064	0.00	152	99	0.67	2.06 ± 0.03	0.1861 ± 0.0029	2548.5 ± 27.2	2710 ± 25	5.96
SHB1_065	1.02	211	138	0.67	85.09 ± 2.74	0.0474 ± 0.0137	75.3 ± 2.4		
SHB1_066	0.88	228	169	0.76	93.68 ± 3.42	0.0301 ± 0.0135	68.5 ± 2.5		
SHB1_067#	0.00	2093	1001	0.49	91.18 ± 1.12	0.0523 ± 0.0021	70.3 ± 0.9		
SHB1_068	0.00	571	340	0.61	80.82 ± 1.92	0.0505 ± 0.0036	79.3 ± 1.9		
SHB1_069	0.18	276	174	0.65	63.67 ± 1.67	0.0511 ± 0.0088	100.5 ± 2.6		
SHB1_070	1.65	229	152	0.68	96.48 ± 3.51	0.0309 ± 0.0144	66.5 ± 2.4		
SHB1_071	1.42	527	245	0.48	101.17 ± 3.07	0.0414 ± 0.0100	63.4 ± 1.9		
SHB1_072	0.00	111	50	0.46	87.78 ± 3.49	0.0656 ± 0.0117	73.0 ± 2.9		
SHB1_073	0.00	56	29	0.54	98.55 ± 6.93	0.0610 ± 0.0266	65.1 ± 4.6		
SHB1_074	0.00	166	87	0.54	93.64 ± 3.67	0.0363 ± 0.0068	68.5 ± 2.7		
SHB1_075	0.00	259	257	1.02	95.91 ± 2.84	0.0559 ± 0.0072	66.9 ± 2.0		

SHB1_076	0.00	72	44	0.64	100.86 ± 5.99	0.0164 ±	0.0248	63.6 ± 3.8				
SHB1_077	0.00	204	155	0.78	89.79 ± 3.48	0.0568 ±	0.0075	71.4 ± 2.7				
SHB1_078	0.00	119	82	0.71	91.50 ± 4.06	0.0433 ±	0.0099	70.1 ± 3.1				
SHB1_079	0.00	353	205	0.60	88.66 ± 2.22	0.0484 ±	0.0049	72.3 ± 1.8				
SHB1_080	0.00	1390	1018	0.75	2.96 ± 0.02	0.1290 ±	0.0011	1877.9 ± 13.1	2085 ± 15		9.93	
SHB1_081	0.00	551	371	0.69	70.89 ± 1.39	0.0494 ±	0.0035	90.3 ± 1.8				
SHB1_082	0.42	823	456	0.57	66.07 ± 1.02	0.0468 ±	0.0049	96.8 ± 1.5				
SHB1_083	0.00	335	453	1.39	28.60 ± 0.52	0.0515 ±	0.0029	221.5 ± 4.0				
SHB1_084	0.81	584	406	0.71	98.02 ± 2.39	0.0418 ±	0.0084	65.4 ± 1.6				
SHB1_085	0.00	586	740	1.30	30.50 ± 0.45	0.0492 ±	0.0022	208.0 ± 3.0				
SHB1_086	0.55	69	34	0.50	92.47 ± 5.75	0.0320 ±	0.0286	69.3 ± 4.3				
SHB1_087	0.75	534	291	0.56	73.83 ± 1.60	0.0551 ±	0.0070	86.7 ± 1.9				
SHB1_088	0.00	978	306	0.32	68.11 ± 1.10	0.0509 ±	0.0027	94.0 ± 1.5				

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}$ - $^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.

Table A7. Sample SB2. 試料SB2.

Labels	²⁰⁶ Pb _c ⁽¹⁾ (%)	U (ppm)	Th (ppm)	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾ (Ma)	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾ (Ma)	Disc ⁽²⁾ (%)
SHB2_001	0.00	390	236	0.62	89.20 ± 2.07	0.0500 ± 0.0044	71.9 ± 1.7		
SHB2_002	1.11	426	244	0.59	90.24 ± 2.37	0.0593 ± 0.0085	71.0 ± 1.9		
SHB2_003	0.72	301	228	0.78	93.66 ± 2.90	0.0453 ± 0.0113	68.5 ± 2.1		
SHB2_004	0.00	158	78	0.51	76.48 ± 2.76	0.0400 ± 0.0067	83.7 ± 3.0		
SHB2_005	2.37	286	144	0.52	93.62 ± 2.65	0.0369 ± 0.0092	68.5 ± 1.9		
SHB2_006	0.53	576	403	0.72	90.45 ± 2.19	0.0466 ± 0.0069	70.9 ± 1.7		
SHB2_007	0.03	357	172	0.49	91.61 ± 2.28	0.0590 ± 0.0079	70.0 ± 1.7		
SHB2_008#	0.00	302	142	0.48	87.18 ± 2.03	0.0605 ± 0.0058	73.5 ± 1.7		
SHB2_009	0.00	90	62	0.71	85.55 ± 3.80	0.0713 ± 0.0123	74.9 ± 3.3		
SHB2_010	0.31	63	56	0.90	2.82 ± 0.06	0.1082 ± 0.0044	1954.0 ± 34.1	1771 ± 72	-10.33
SHB2_011	0.35	763	329	0.44	95.26 ± 1.83	0.0487 ± 0.0053	67.3 ± 1.3		
SHB2_012#	0.00	336	160	0.49	60.84 ± 1.32	0.0565 ± 0.0037	105.1 ± 2.3		
SHB2_013	0.00	439	311	0.73	91.00 ± 2.04	0.0430 ± 0.0044	70.5 ± 1.6		
SHB2_014	0.34	519	443	0.88	90.93 ± 2.09	0.0404 ± 0.0085	70.5 ± 1.6		
SHB2_015	0.00	417	228	0.56	67.95 ± 1.42	0.0481 ± 0.0038	94.2 ± 2.0		
SHB2_016	0.91	540	406	0.77	91.34 ± 2.17	0.0483 ± 0.0081	70.2 ± 1.7		
SHB2_017	2.28	492	264	0.55	95.07 ± 2.25	0.0469 ± 0.0088	67.5 ± 1.6		
SHB2_018	2.42	915	446	0.50	81.78 ± 1.48	0.0562 ± 0.0060	78.3 ± 1.4		
SHB2_019	1.96	405	248	0.63	73.48 ± 1.64	0.0594 ± 0.0097	87.1 ± 1.9		
SHB2_020	3.18	474	248	0.54	84.29 ± 2.07	0.0483 ± 0.0081	76.0 ± 1.9		
SHB2_021	0.00	437	262	0.62	87.62 ± 2.17	0.0543 ± 0.0045	73.2 ± 1.8		
SHB2_022	0.49	554	307	0.57	89.93 ± 1.96	0.0495 ± 0.0069	71.3 ± 1.5		
SHB2_023	1.43	366	116	0.33	72.73 ± 1.63	0.0597 ± 0.0070	88.0 ± 2.0		
SHB2_024	0.00	479	297	0.64	94.31 ± 1.84	0.0529 ± 0.0038	68.0 ± 1.3		
SHB2_025	0.00	710	349	0.50	71.80 ± 1.31	0.0498 ± 0.0025	89.2 ± 1.6		
SHB2_026	0.03	285	161	0.58	13.25 ± 0.18	0.0596 ± 0.0035	468.9 ± 6.2		
SHB2_027	0.00	76	47	0.63	89.61 ± 4.77	0.0637 ± 0.0142	71.5 ± 3.8		
SHB2_028	0.62	520	206	0.41	64.46 ± 1.21	0.0523 ± 0.0049	99.2 ± 1.8		
SHB2_029	0.00	736	788	1.10	87.71 ± 1.71	0.0472 ± 0.0031	73.1 ± 1.4		
SHB2_030	2.89	603	529	0.90	104.73 ± 2.91	0.0465 ± 0.0115	61.3 ± 1.7		
SHB2_031	0.00	205	190	0.95	84.70 ± 2.76	0.0543 ± 0.0059	75.7 ± 2.5		
SHB2_032	0.19	259	144	0.57	94.48 ± 3.28	0.0615 ± 0.0103	67.9 ± 2.3		
SHB2_033	0.00	163	92	0.58	62.38 ± 2.06	0.0440 ± 0.0059	102.5 ± 3.4		
SHB2_034	0.00	357	135	0.39	61.23 ± 1.33	0.0425 ± 0.0033	104.4 ± 2.3		
SHB2_035	0.38	166	73	0.45	64.33 ± 2.38	0.0509 ± 0.0106	99.4 ± 3.6		
SHB2_036	0.00	797	511	0.66	70.13 ± 1.10	0.0477 ± 0.0027	91.3 ± 1.4		
SHB2_037	0.00	116	75	0.66	86.70 ± 3.64	0.0408 ± 0.0091	73.9 ± 3.1		
SHB2_038	0.00	322	290	0.92	85.98 ± 2.40	0.0465 ± 0.0044	74.5 ± 2.1		
SHB2_039	1.90	772	435	0.58	24.78 ± 0.30	0.0515 ± 0.0032	255.0 ± 3.1		
SHB2_040	0.00	341	255	0.77	24.16 ± 0.39	0.0533 ± 0.0025	261.4 ± 4.1		
SHB2_041	0.38	199	152	0.78	82.92 ± 2.90	0.0523 ± 0.0112	77.3 ± 2.7		
SHB2_042	0.87	356	202	0.58	89.06 ± 2.16	0.0435 ± 0.0080	72.0 ± 1.7		
SHB2_043	0.00	372	256	0.70	89.77 ± 2.27	0.0457 ± 0.0040	71.4 ± 1.8		
SHB2_044	0.00	156	128	0.84	78.81 ± 2.66	0.0539 ± 0.0075	81.3 ± 2.7		
SHB2_045#	0.62	409	75	0.19	4.29 ± 0.06	0.1245 ± 0.0020	1350.8 ± 16.4	2023 ± 27	33.23
SHB2_046	1.96	314	276	0.90	89.12 ± 2.76	0.0456 ± 0.0120	71.9 ± 2.2		
SHB2_047	0.00	315	214	0.70	81.96 ± 2.00	0.0540 ± 0.0049	78.2 ± 1.9		
SHB2_048	0.00	394	312	0.81	89.15 ± 2.25	0.0470 ± 0.0039	71.9 ± 1.8		
SHB2_049	0.14	489	413	0.87	88.21 ± 2.02	0.0446 ± 0.0080	72.7 ± 1.7		
SHB2_050	0.00	432	391	0.93	26.46 ± 0.34	0.0513 ± 0.0030	239.1 ± 3.0		
SHB2_051	0.00	208	116	0.57	86.77 ± 2.43	0.0374 ± 0.0055	73.9 ± 2.1		
SHB2_052#	0.28	152	72	0.48	2.69 ± 0.04	0.1658 ± 0.0032	2034.8 ± 22.7	2517 ± 32	19.16
SHB2_053	0.00	1033	674	0.67	67.85 ± 1.24	0.0518 ± 0.0021	94.3 ± 1.7		
SHB2_054	2.48	383	276	0.74	95.70 ± 2.82	0.0442 ± 0.0114	67.0 ± 2.0		
SHB2_055	1.04	521	355	0.70	90.91 ± 2.08	0.0446 ± 0.0082	70.5 ± 1.6		
SHB2_056	0.00	782	902	1.18	94.80 ± 1.38	0.0480 ± 0.0034	67.6 ± 1.0		
SHB2_057	0.00	374	205	0.56	90.97 ± 2.25	0.0439 ± 0.0049	70.5 ± 1.7		
SHB2_058	0.64	244	153	0.64	85.60 ± 2.57	0.0328 ± 0.0104	74.9 ± 2.2		
SHB2_059	1.67	287	179	0.64	88.47 ± 2.30	0.0306 ± 0.0088	72.5 ± 1.9		
SHB2_060	1.22	463	276	0.61	88.60 ± 2.46	0.0465 ± 0.0098	72.4 ± 2.0		
SHB2_061	0.62	478	280	0.60	76.98 ± 1.62	0.0422 ± 0.0069	83.2 ± 1.7		
SHB2_062	0.00	320	171	0.55	90.24 ± 2.12	0.0473 ± 0.0045	71.0 ± 1.7		
SHB2_063	0.18	470	234	0.51	67.50 ± 1.37	0.0482 ± 0.0057	94.8 ± 1.9		
SHB2_064	0.01	235	150	0.65	87.94 ± 2.77	0.0487 ± 0.0110	72.9 ± 2.3		
SHB2_065	2.53	452	260	0.59	96.61 ± 2.53	0.0401 ± 0.0092	66.4 ± 1.7		
SHB2_066	0.00	558	397	0.73	91.51 ± 1.75	0.0443 ± 0.0040	70.1 ± 1.3		
SHB2_067	2.04	579	608	1.08	60.11 ± 1.37	0.0360 ± 0.0089	106.4 ± 2.4		
SHB2_068	0.21	508	233	0.47	2.75 ± 0.03	0.1398 ± 0.0019	2002.1 ± 18.0	2226 ± 23	10.06
SHB2_069	0.00	301	212	0.72	91.01 ± 2.27	0.0474 ± 0.0050	70.4 ± 1.8		
SHB2_070	0.73	284	129	0.47	92.24 ± 2.97	0.0398 ± 0.0094	69.5 ± 2.2		
SHB2_071	0.43	148	73	0.51	93.83 ± 4.25	0.0347 ± 0.0139	68.3 ± 3.1		
SHB2_072	0.76	297	159	0.55	90.87 ± 2.51	0.0337 ± 0.0086	70.6 ± 1.9		
SHB2_073#	0.00	451	482	1.10	90.09 ± 1.97	0.0626 ± 0.0055	71.2 ± 1.5		
SHB2_074	2.53	139	82	0.61	91.87 ± 4.12	0.0266 ± 0.0177	69.8 ± 3.1		
SHB2_075	0.84	155	93	0.62	89.00 ± 3.71	0.0459 ± 0.0158	72.0 ± 3.0		

SHB2_076	0.00	203	122	0.62	92.83 ± 2.58	0.0536 ±	0.0068	69.1 ± 1.9
SHB2_077	0.43	476	229	0.49	89.77 ± 2.02	0.0512 ±	0.0064	71.4 ± 1.6
SHB2_078	1.30	348	319	0.94	91.73 ± 2.35	0.0359 ±	0.0113	69.9 ± 1.8
SHB2_079	2.23	249	140	0.57	89.76 ± 2.79	0.0377 ±	0.0106	71.4 ± 2.2
SHB2_080	1.33	906	1166	1.32	92.58 ± 1.80	0.0567 ±	0.0097	69.3 ± 1.3
SHB2_081	0.00	404	210	0.53	76.46 ± 1.58	0.0406 ±	0.0036	83.8 ± 1.7
SHB2_082	0.00	476	356	0.77	75.36 ± 1.46	0.0461 ±	0.0030	85.0 ± 1.6
SHB2_083	0.24	237	189	0.82	88.72 ± 2.73	0.0513 ±	0.0114	72.3 ± 2.2
SHB2_084	1.17	245	166	0.70	86.37 ± 2.75	0.0296 ±	0.0119	74.2 ± 2.4
SHB2_085	2.34	595	518	0.89	86.44 ± 2.00	0.0339 ±	0.0095	74.2 ± 1.7
SHB2_086	0.00	361	349	0.99	86.97 ± 2.00	0.0449 ±	0.0051	73.7 ± 1.7
SHB2_087	0.25	464	263	0.58	75.04 ± 1.51	0.0393 ±	0.0061	85.3 ± 1.7
SHB2_088	0.30	512	306	0.61	90.01 ± 2.20	0.0450 ±	0.0071	71.2 ± 1.7
SHB2_089	0.00	447	266	0.61	88.80 ± 1.88	0.0451 ±	0.0047	72.2 ± 1.5
SHB2_090	0.00	360	281	0.80	89.37 ± 2.26	0.0497 ±	0.0046	71.7 ± 1.8
SHB2_091	0.33	254	201	0.81	86.39 ± 2.44	0.0479 ±	0.0109	74.2 ± 2.1
SHB2_092	0.95	293	211	0.74	84.79 ± 2.35	0.0432 ±	0.0102	75.6 ± 2.1
SHB2_093	0.00	301	123	0.42	2.40 ± 0.03	0.1637 ±	0.0019	2243.4 ± 23.3 2496 ± 19 10.12

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.
 "#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(238\text{U}/206\text{Pb}^* \text{ age})/(207\text{Pb}^*/206\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.
Table A8. Sample CCO. 試料CCO.

Labels	²⁰⁶ Pb _c ⁽¹⁾ (%)	U (ppm)	Th (ppm)	Th/U	²³⁸ U/ ²⁰⁶ Pb* ⁽¹⁾	²⁰⁷ Pb*/ ²⁰⁶ Pb* ⁽¹⁾	²³⁸ U/ ²⁰⁶ Pb* age ⁽¹⁾ (Ma)	²⁰⁷ Pb*/ ²⁰⁶ Pb* age ⁽¹⁾ (Ma)	Disc ⁽²⁾ (%)
CCO_001	0.00	79	59	0.77	30.62 ± 0.95	0.0468 ± 0.0044	207.1 ± 6.3		
CCO_002	0.00	166	127	0.78	26.28 ± 0.54	0.0502 ± 0.0031	240.7 ± 4.9		
CCO_003	1.26	335	315	0.96	85.15 ± 2.32	0.0429 ± 0.0095	75.3 ± 2.0		
CCO_004	0.00	230	116	0.52	2.98 ± 0.04	0.1135 ± 0.0016	1864.5 ± 22.5	1857 ± 26	-0.41
CCO_005	0.35	465	161	0.36	31.28 ± 0.58	0.0469 ± 0.0034	202.8 ± 3.7		
CCO_006	0.51	253	123	0.50	85.29 ± 2.74	0.0556 ± 0.0091	75.1 ± 2.4		
CCO_007	0.14	430	91	0.22	3.00 ± 0.05	0.1211 ± 0.0017	1852.6 ± 25.5	1973 ± 24	6.10
CCO_008	0.00	233	169	0.74	65.33 ± 1.62	0.0423 ± 0.0039	97.9 ± 2.4		
CCO_009	0.13	334	273	0.84	28.83 ± 0.56	0.0490 ± 0.0048	219.8 ± 4.2		
CCO_010	0.00	275	197	0.74	89.15 ± 2.43	0.0435 ± 0.0041	71.9 ± 2.0		
CCO_011	0.00	138	119	0.89	32.27 ± 0.95	0.0443 ± 0.0039	196.7 ± 5.7		
CCO_012	0.71	494	295	0.61	79.42 ± 1.93	0.0578 ± 0.0077	80.7 ± 2.0		
CCO_013	0.00	151	142	0.96	2.69 ± 0.04	0.1377 ± 0.0024	2036.8 ± 24.3	2199 ± 30	7.38
CCO_014	0.00	181	95	0.54	87.53 ± 2.75	0.0467 ± 0.0055	73.2 ± 2.3		
CCO_015	0.93	302	199	0.68	88.71 ± 2.58	0.0416 ± 0.0089	72.3 ± 2.1		
CCO_016	0.00	243	94	0.39	3.13 ± 0.05	0.1155 ± 0.0018	1786.7 ± 22.5	1889 ± 28	5.42
CCO_017	1.16	798	449	0.58	88.83 ± 1.96	0.0427 ± 0.0056	72.2 ± 1.6		
CCO_018	0.04	202	79	0.40	2.83 ± 0.05	0.1362 ± 0.0026	1947.7 ± 29.0	2181 ± 33	10.70
CCO_019	0.00	717	501	0.72	28.31 ± 0.51	0.0500 ± 0.0019	223.8 ± 4.0		
CCO_020	1.17	96	66	0.70	86.56 ± 5.09	0.0456 ± 0.0188	74.1 ± 4.3		
CCO_021	0.00	55	41	0.76	26.08 ± 0.87	0.0533 ± 0.0061	242.6 ± 8.0		
CCO_022	0.00	335	239	0.73	84.08 ± 2.51	0.0465 ± 0.0046	76.2 ± 2.3		
CCO_023	0.00	209	120	0.59	89.32 ± 2.80	0.0382 ± 0.0048	71.8 ± 2.2		
CCO_024	0.95	363	230	0.65	88.75 ± 2.55	0.0305 ± 0.0087	72.2 ± 2.1		
CCO_025	0.07	228	43	0.20	3.12 ± 0.05	0.1103 ± 0.0020	1793.1 ± 23.1	1806 ± 32	0.71
CCO_026	0.31	225	189	0.86	65.21 ± 2.14	0.0345 ± 0.0116	98.1 ± 3.2		
CCO_027	0.00	1181	754	0.65	89.15 ± 1.65	0.0497 ± 0.0024	71.9 ± 1.3		
CCO_028	0.83	68	54	0.82	26.03 ± 1.07	0.0335 ± 0.0135	243.0 ± 9.8		
CCO_029	0.00	161	48	0.31	2.46 ± 0.04	0.1480 ± 0.0024	2202.4 ± 29.5	2325 ± 27	5.27
CCO_030	0.00	147	79	0.55	84.98 ± 3.15	0.0390 ± 0.0072	75.4 ± 2.8		
CCO_031	0.00	1158	540	0.48	85.84 ± 1.48	0.0514 ± 0.0024	74.7 ± 1.3		
CCO_032#	0.00	458	131	0.29	3.64 ± 0.05	0.1128 ± 0.0014	1566.1 ± 17.3	1847 ± 22	15.21
CCO_033	0.75	204	301	1.51	2.85 ± 0.04	0.1176 ± 0.0046	1936.4 ± 23.5	1921 ± 69	-0.80
CCO_034	0.00	285	32	0.11	3.10 ± 0.04	0.1138 ± 0.0016	1801.0 ± 20.7	1863 ± 25	3.33
CCO_035	0.00	913	528	0.59	3.03 ± 0.03	0.1130 ± 0.0012	1839.4 ± 18.5	1850 ± 18	0.57
CCO_036	0.00	302	171	0.58	84.03 ± 2.14	0.0505 ± 0.0042	76.3 ± 1.9		
CCO_037	0.00	167	123	0.75	87.54 ± 2.89	0.0507 ± 0.0066	73.2 ± 2.4		
CCO_038	0.88	628	359	0.59	91.36 ± 1.87	0.0373 ± 0.0060	70.2 ± 1.4		
CCO_039	0.00	168	138	0.84	86.16 ± 2.91	0.0381 ± 0.0050	74.4 ± 2.5		
CCO_040	0.00	231	122	0.54	83.07 ± 2.15	0.0396 ± 0.0043	77.1 ± 2.0		
CCO_041	0.00	131	79	0.62	87.01 ± 3.28	0.0474 ± 0.0067	73.7 ± 2.8		
CCO_042#	0.00	209	117	0.58	3.17 ± 0.05	0.1137 ± 0.0018	1767.3 ± 22.4	1861 ± 28	5.03
CCO_043	2.41	290	140	0.50	3.03 ± 0.05	0.1457 ± 0.0026	1840.2 ± 24.7	2297 ± 31	19.89
CCO_044	0.00	479	401	0.86	40.05 ± 0.69	0.0457 ± 0.0024	159.0 ± 2.7		
CCO_045	0.00	352	150	0.44	84.77 ± 1.86	0.0411 ± 0.0038	75.6 ± 1.7		
CCO_046	1.96	168	137	0.84	89.52 ± 2.96	0.0388 ± 0.0140	71.6 ± 2.4		
CCO_047	0.37	183	14	0.08	28.93 ± 0.73	0.0472 ± 0.0041	219.0 ± 5.4		
CCO_048	0.00	98	59	0.62	35.33 ± 0.99	0.0437 ± 0.0042	179.9 ± 5.0		
CCO_049	0.00	60	54	0.91	23.92 ± 0.67	0.0541 ± 0.0057	264.0 ± 7.2		
CCO_050	0.00	209	57	0.28	2.99 ± 0.04	0.1190 ± 0.0017	1859.7 ± 24.1	1942 ± 26	4.24
CCO_051#	0.00	177	130	0.75	101.47 ± 3.28	0.0616 ± 0.0071	63.2 ± 2.0		
CCO_052	0.15	323	142	0.45	84.69 ± 2.34	0.0391 ± 0.0069	75.7 ± 2.1		
CCO_053	0.00	197	198	1.03	87.90 ± 2.78	0.0413 ± 0.0047	72.9 ± 2.3		
CCO_054	0.00	392	217	0.57	83.36 ± 1.75	0.0444 ± 0.0035	76.9 ± 1.6		
CCO_055	0.98	395	200	0.52	83.89 ± 2.16	0.0375 ± 0.0069	76.4 ± 2.0		
CCO_056	0.00	470	326	0.71	2.19 ± 0.03	0.1737 ± 0.0019	2427.4 ± 23.8	2595 ± 18	6.46
CCO_057	0.00	823	267	0.33	85.09 ± 1.40	0.0478 ± 0.0026	75.3 ± 1.2		
CCO_058	0.00	665	788	1.22	31.59 ± 0.50	0.0523 ± 0.0018	200.9 ± 3.1		
CCO_059	0.00	201	92	0.47	2.86 ± 0.04	0.1225 ± 0.0017	1934.7 ± 24.7	1994 ± 24	2.97
CCO_060	0.19	238	72	0.31	2.96 ± 0.04	0.1136 ± 0.0022	1878.8 ± 23.4	1858 ± 35	-1.12
CCO_061	0.00	795	416	0.54	83.51 ± 1.75	0.0490 ± 0.0026	76.7 ± 1.6		
CCO_062	0.97	70	95	1.40	34.43 ± 1.33	0.0500 ± 0.0177	184.6 ± 7.0		
CCO_063	0.00	181	207	1.17	3.11 ± 0.05	0.1121 ± 0.0022	1795.8 ± 23.0	1836 ± 34	2.19
CCO_064	0.00	432	188	0.45	82.59 ± 2.03	0.0462 ± 0.0031	77.6 ± 1.9		
CCO_065	0.71	105	48	0.47	25.34 ± 0.74	0.0431 ± 0.0055	249.5 ± 7.1		
CCO_066	0.12	147	86	0.60	87.96 ± 3.08	0.0674 ± 0.0118	72.9 ± 2.5		
CCO_067	0.15	871	297	0.35	24.80 ± 0.40	0.0534 ± 0.0021	254.8 ± 4.0		
CCO_068	0.00	533	40	0.08	2.85 ± 0.04	0.1143 ± 0.0014	1941.7 ± 21.8	1870 ± 22	-3.83
CCO_069	0.00	273	82	0.31	62.55 ± 1.55	0.0467 ± 0.0036	102.2 ± 2.5		
CCO_070	0.65	206	32	0.16	2.97 ± 0.05	0.1113 ± 0.0020	1868.5 ± 26.7	1821 ± 33	-2.61
CCO_071	0.66	626	540	0.89	36.15 ± 0.63	0.0445 ± 0.0047	175.9 ± 3.0		
CCO_072	0.75	150	120	0.82	82.34 ± 2.89	0.0402 ± 0.0130	77.8 ± 2.7		
CCO_073	0.00	1298	187	0.15	3.10 ± 0.05	0.1236 ± 0.0012	1801.6 ± 23.0	2009 ± 18	10.32
CCO_074	0.00	261	154	0.61	82.07 ± 2.35	0.0564 ± 0.0049	78.1 ± 2.2		
CCO_075#	0.00	235	192	0.84	83.15 ± 2.28	0.0624 ± 0.0051	77.1 ± 2.1		

CCO_076	0.00	327	49	0.16	3.07 ± 0.04	0.1145 ±	0.0017	1818.4 ± 19.9	1874 ± 26	2.97
CCO_077	0.00	453	176	0.40	3.04 ± 0.04	0.1157 ±	0.0014	1835.8 ± 20.9	1891 ± 22	2.92
CCO_078	0.00	232	97	0.43	88.21 ± 2.15	0.0499 ±	0.0045	72.7 ± 1.8		
CCO_079	0.00	520	49	0.10	3.31 ± 0.04	0.1174 ±	0.0014	1703.7 ± 20.0	1918 ± 20	11.17
CCO_080	0.00	259	116	0.46	3.12 ± 0.04	0.1177 ±	0.0015	1791.9 ± 19.3	1923 ± 23	6.82
CCO_081	0.00	704	450	0.66	84.89 ± 1.59	0.0465 ±	0.0025	75.5 ± 1.4		
CCO_082	2.41	140	97	0.71	90.61 ± 3.57	0.0289 ±	0.0129	70.8 ± 2.8		
CCO_083	2.65	208	215	1.06	89.65 ± 2.55	0.0241 ±	0.0123	71.5 ± 2.0		
CCO_084	0.00	916	220	0.25	22.99 ± 0.32	0.0512 ±	0.0015	274.5 ± 3.8		
CCO_085	0.00	339	246	0.75	84.87 ± 2.13	0.0417 ±	0.0034	75.5 ± 1.9		
CCO_086	0.00	352	240	0.70	85.47 ± 2.15	0.0447 ±	0.0064	75.0 ± 1.9		
CCO_087	0.01	480	151	0.32	3.08 ± 0.04	0.1123 ±	0.0015	1814.9 ± 22.7	1838 ± 23	1.26
CCO_088	0.00	231	125	0.55	3.21 ± 0.05	0.1141 ±	0.0017	1750.5 ± 23.2	1866 ± 27	6.19
CCO_089	1.31	207	146	0.72	92.15 ± 2.96	0.0512 ±	0.0128	69.6 ± 2.2		
CCO_090	0.66	451	146	0.33	20.19 ± 0.35	0.0495 ±	0.0029	311.6 ± 5.2		
CCO_091	0.00	319	224	0.72	88.06 ± 2.17	0.0411 ±	0.0034	72.8 ± 1.8		
CCO_092	0.00	186	120	0.66	1.62 ± 0.02	0.2941 ±	0.0032	3095.9 ± 34.0	3441 ± 17	10.03
CCO_093	2.40	143	65	0.47	84.01 ± 3.03	0.0552 ±	0.0141	76.3 ± 2.7		
CCO_094	0.00	91	87	0.98	15.37 ± 0.39	0.0595 ±	0.0082	406.2 ± 10.0		
CCO_095	0.65	108	41	0.39	23.76 ± 0.62	0.0420 ±	0.0061	265.8 ± 6.8		
CCO_096	0.00	323	284	0.90	2.94 ± 0.04	0.1147 ±	0.0016	1887.8 ± 20.1	1876 ± 25	-0.63
CCO_097	0.00	103	94	0.94	2.19 ± 0.03	0.1614 ±	0.0029	2424.9 ± 29.7	2472 ± 29	1.91
CCO_098	0.00	387	238	0.63	84.92 ± 1.73	0.0465 ±	0.0030	75.5 ± 1.5		
CCO_099	1.26	403	441	1.12	87.38 ± 2.22	0.0348 ±	0.0091	73.4 ± 1.9		
CCO_100	0.00	348	45	0.13	3.45 ± 0.04	0.1137 ±	0.0014	1640.8 ± 18.0	1861 ± 22	11.83
CCO_101	0.00	107	153	1.47	2.15 ± 0.04	0.1618 ±	0.0026	2457.7 ± 37.2	2476 ± 27	0.74
CCO_102	0.32	364	291	0.82	24.21 ± 0.49	0.0479 ±	0.0050	260.9 ± 5.2		
CCO_103	1.57	411	360	0.90	81.84 ± 1.99	0.0313 ±	0.0082	78.3 ± 1.9		
CCO_104	1.09	276	148	0.55	84.99 ± 2.57	0.0458 ±	0.0089	75.4 ± 2.3		
CCO_105	1.32	73	93	1.31	4.20 ± 0.08	0.0978 ±	0.0072	1375.5 ± 24.1	1584 ± 132	13.16
CCO_106	0.47	116	90	0.79	84.76 ± 3.49	0.0614 ±	0.0160	75.6 ± 3.1		
CCO_107	0.00	349	323	0.95	86.58 ± 2.13	0.0517 ±	0.0048	74.0 ± 1.8		
CCO_108	0.63	442	273	0.64	85.05 ± 1.98	0.0462 ±	0.0072	75.4 ± 1.7		
CCO_109	0.46	178	79	0.45	3.06 ± 0.05	0.1126 ±	0.0026	1824.2 ± 26.0	1843 ± 40	1.02
CCO_110	0.05	159	22	0.15	3.05 ± 0.05	0.1147 ±	0.0020	1826.4 ± 24.1	1876 ± 32	2.64
CCO_111	2.64	73	62	0.87	23.38 ± 0.78	0.0338 ±	0.0118	270.0 ± 8.9		
CCO_112	0.49	125	86	0.71	33.78 ± 0.92	0.0425 ±	0.0087	188.1 ± 5.1		
CCO_113	1.09	306	222	0.74	82.48 ± 2.50	0.0400 ±	0.0092	77.7 ± 2.3		
CCO_114	0.00	87	80	0.94	87.32 ± 3.94	0.0506 ±	0.0099	73.4 ± 3.3		
CCO_115	0.26	161	85	0.54	82.12 ± 2.50	0.0494 ±	0.0099	78.0 ± 2.4		
CCO_116	0.61	744	287	0.40	37.95 ± 0.68	0.0441 ±	0.0028	167.7 ± 3.0		
CCO_117	0.00	1035	487	0.48	91.01 ± 1.67	0.0500 ±	0.0024	70.4 ± 1.3		
CCO_118	3.89	250	181	0.74	87.42 ± 2.80	0.0346 ±	0.0118	73.3 ± 2.3		
CCO_119	0.17	200	69	0.35	23.62 ± 0.41	0.0462 ±	0.0039	267.3 ± 4.5		
CCO_120	0.00	1096	594	0.56	82.92 ± 1.46	0.0473 ±	0.0023	77.3 ± 1.4		
CCO_121	0.13	132	43	0.34	2.17 ± 0.03	0.1648 ±	0.0030	2447.2 ± 28.5	2507 ± 30	2.39
CCO_122	0.00	732	366	0.51	25.53 ± 0.38	0.0499 ±	0.0016	247.7 ± 3.6		
CCO_123	0.00	937	700	0.77	83.23 ± 1.46	0.0447 ±	0.0021	77.0 ± 1.3		
CCO_124	0.31	214	125	0.60	2.28 ± 0.04	0.1422 ±	0.0026	2342.0 ± 34.1	2255 ± 32	-3.86
CCO_125#	0.09	693	318	0.47	3.76 ± 0.06	0.1232 ±	0.0021	1520.2 ± 20.0	2004 ± 30	24.14
CCO_126	1.06	206	245	1.22	82.59 ± 2.74	0.0490 ±	0.0137	77.6 ± 2.6		
CCO_127	0.75	723	327	0.46	85.59 ± 1.89	0.0413 ±	0.0050	74.9 ± 1.6		
CCO_128	0.08	297	44	0.15	3.02 ± 0.04	0.1117 ±	0.0018	1841.6 ± 22.1	1829 ± 29	-0.69
CCO_129	0.00	895	466	0.53	77.91 ± 1.50	0.0436 ±	0.0025	82.2 ± 1.6		
CCO_130	0.11	250	120	0.49	3.15 ± 0.05	0.1070 ±	0.0018	1777.4 ± 23.0	1749 ± 32	-1.62
CCO_131	0.00	444	120	0.28	30.04 ± 0.50	0.0487 ±	0.0020	211.1 ± 3.5		
CCO_132	0.00	455	209	0.47	85.69 ± 2.01	0.0518 ±	0.0032	74.8 ± 1.7		
CCO_133	0.00	204	63	0.32	2.97 ± 0.04	0.1141 ±	0.0016	1873.0 ± 24.4	1866 ± 26	-0.38
CCO_134	0.00	127	123	0.99	36.12 ± 0.90	0.0516 ±	0.0045	176.0 ± 4.3		
CCO_135	0.00	1211	308	0.26	62.47 ± 0.85	0.0490 ±	0.0017	102.4 ± 1.4		
CCO_136	0.00	256	133	0.53	84.41 ± 2.29	0.0443 ±	0.0047	75.9 ± 2.0		
CCO_137	0.00	205	163	0.82	26.58 ± 0.50	0.0532 ±	0.0031	238.1 ± 4.4		
CCO_138	0.00	901	400	0.46	80.94 ± 1.29	0.0485 ±	0.0022	79.2 ± 1.3		
CCO_139	0.00	375	81	0.22	29.18 ± 0.45	0.0470 ±	0.0020	217.2 ± 3.3		
CCO_140	0.06	209	95	0.47	2.77 ± 0.04	0.1351 ±	0.0024	1989.8 ± 23.7	2166 ± 30	8.13
CCO_141	0.00	240	160	0.68	25.27 ± 0.48	0.0490 ±	0.0024	250.2 ± 4.7		
CCO_142	0.00	290	182	0.64	34.99 ± 0.69	0.0502 ±	0.0027	181.6 ± 3.5		
CCO_143	0.21	344	109	0.32	2.86 ± 0.04	0.1224 ±	0.0017	1934.9 ± 23.2	1992 ± 25	2.87
CCO_144#	0.80	289	108	0.38	3.61 ± 0.06	0.1247 ±	0.0020	1576.5 ± 22.3	2026 ± 28	22.19

Errors are 1-sigma; Pb_c and Pb* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. U-Pb age. U-Pb 年代.

Table A9. Sample SBR. 試料SBR.

Labels	$^{206}\text{Pb}_c^{(1)}$	U	Th	Th/U	$^{238}\text{U}/^{206}\text{Pb}^*^{(1)}$	$^{207}\text{Pb}^*/^{206}\text{Pb}^*^{(1)}$	$^{238}\text{U}/^{206}\text{Pb}^* \text{ age}^{(1)}$	$^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age}^{(1)}$	Disc ⁽³⁾
	(%)	(ppm)	(ppm)				(Ma)	(Ma)	(%)
SBR_001	0.00	700	354	0.52	64.21 ± 0.83	0.0503 ± 0.0016	99.6 ± 1.3		
SBR_002	5.20	446	342	0.79	81.13 ± 1.26	0.0556 ± 0.0089	79.0 ± 1.2		
SBR_003	0.56	846	462	0.56	80.73 ± 1.10	0.0421 ± 0.0030	79.4 ± 1.1		
SBR_004	0.00	578	335	0.59	81.06 ± 1.22	0.0456 ± 0.0019	79.0 ± 1.2		
SBR_005	0.16	407	267	0.67	78.39 ± 1.20	0.0467 ± 0.0042	81.7 ± 1.2		
SBR_006	0.00	434	260	0.62	73.47 ± 1.14	0.0476 ± 0.0022	87.1 ± 1.3		
SBR_007	0.18	211	162	0.79	83.14 ± 1.80	0.0406 ± 0.0066	77.1 ± 1.7		
SBR_008	0.00	232	145	0.64	77.52 ± 1.59	0.0460 ± 0.0031	82.6 ± 1.7		
SBR_009#	0.00	169	85	0.51	81.63 ± 1.85	0.0572 ± 0.0039	78.5 ± 1.8		
SBR_010	0.37	380	350	0.94	85.21 ± 1.58	0.0458 ± 0.0075	75.2 ± 1.4		
SBR_011	0.29	194	110	0.58	84.93 ± 2.02	0.0488 ± 0.0078	75.5 ± 1.8		
SBR_012	15.39	323	105	0.33	59.54 ± 1.00	0.0657 ± 0.0102	107.4 ± 1.8		
SBR_013	0.00	296	134	0.46	77.31 ± 1.57	0.0493 ± 0.0028	82.9 ± 1.7		
SBR_014	0.00	421	271	0.66	81.60 ± 1.50	0.0455 ± 0.0024	78.5 ± 1.4		
SBR_015	0.40	451	231	0.53	83.22 ± 1.58	0.0496 ± 0.0046	77.0 ± 1.5		
SBR_016	0.00	394	246	0.64	79.95 ± 1.26	0.0439 ± 0.0022	80.1 ± 1.3		
SBR_017	0.00	593	210	0.36	83.85 ± 1.31	0.0499 ± 0.0022	76.4 ± 1.2		
SBR_018	0.00	80	83	1.06	85.40 ± 2.63	0.0461 ± 0.0056	75.0 ± 2.3		
SBR_019	0.00	339	326	0.98	81.93 ± 1.46	0.0483 ± 0.0030	78.2 ± 1.4		
SBR_020	1.00	159	123	0.79	84.94 ± 2.36	0.0429 ± 0.0094	75.5 ± 2.1		
SBR_021	0.48	176	133	0.77	79.23 ± 2.19	0.0486 ± 0.0089	80.9 ± 2.2		
SBR_022	0.47	631	425	0.69	79.80 ± 2.14	0.0464 ± 0.0044	80.3 ± 2.1		
SBR_023	7.09	155	84	0.56	84.81 ± 3.01	0.0419 ± 0.0118	75.6 ± 2.7		
SBR_024	1.07	334	430	1.32	84.42 ± 2.39	0.0388 ± 0.0078	75.9 ± 2.1		
SBR_025	3.00	287	263	0.94	84.84 ± 2.70	0.0536 ± 0.0091	75.5 ± 2.4		
SBR_026	0.00	364	111	0.31	63.07 ± 1.77	0.0503 ± 0.0026	101.4 ± 2.8		
SBR_027	0.00	137	66	0.49	79.73 ± 2.95	0.0527 ± 0.0059	80.4 ± 3.0		
SBR_028	0.41	476	289	0.62	80.65 ± 2.28	0.0485 ± 0.0048	79.4 ± 2.2		
SBR_029	0.00	463	209	0.46	82.65 ± 2.22	0.0440 ± 0.0019	77.5 ± 2.1		
SBR_030	1.38	282	198	0.72	69.93 ± 2.53	0.0472 ± 0.0086	91.5 ± 3.3		
SBR_031	0.00	205	164	0.82	79.60 ± 2.94	0.0480 ± 0.0032	80.5 ± 3.0		
SBR_032	0.21	336	235	0.72	63.73 ± 2.24	0.0485 ± 0.0051	100.4 ± 3.5		
SBR_033	0.96	333	320	0.99	83.94 ± 2.88	0.0403 ± 0.0070	76.3 ± 2.6		
SBR_034	0.00	514	187	0.37	63.54 ± 2.10	0.0511 ± 0.0023	100.7 ± 3.3		
SBR_035	3.26	257	162	0.65	83.74 ± 2.99	0.0526 ± 0.0077	76.5 ± 2.7		
SBR_036	4.47	86	54	0.65	76.70 ± 2.93	0.0613 ± 0.0173	83.5 ± 3.2		
SBR_037	0.00	624	251	0.41	60.90 ± 0.92	0.0475 ± 0.0017	105.0 ± 1.6		
SBR_038	0.00	596	529	0.91	78.97 ± 1.28	0.0474 ± 0.0020	81.1 ± 1.3		
SBR_039	0.00	830	527	0.65	84.18 ± 1.30	0.0490 ± 0.0021	76.1 ± 1.2		
SBR_040	0.00	656	362	0.57	62.03 ± 0.91	0.0511 ± 0.0019	103.1 ± 1.5		
SBR_041	0.00	150	84	0.57	79.29 ± 1.80	0.0439 ± 0.0045	80.8 ± 1.8		
SBR_042	0.37	268	144	0.55	82.35 ± 1.79	0.0486 ± 0.0057	77.8 ± 1.7		
SBR_043	0.00	257	137	0.55	83.90 ± 1.93	0.0467 ± 0.0032	76.4 ± 1.7		
SBR_044	0.00	583	298	0.52	61.46 ± 0.84	0.0465 ± 0.0019	104.0 ± 1.4		
SBR_045	1.48	179	80	0.46	81.30 ± 1.99	0.0351 ± 0.0071	78.8 ± 1.9		
SBR_046	0.00	285	232	0.84	74.07 ± 1.53	0.0532 ± 0.0037	86.4 ± 1.8		
SBR_047	0.30	549	261	0.49	79.15 ± 1.24	0.0459 ± 0.0036	80.9 ± 1.3		
SBR_048	0.00	349	189	0.56	79.37 ± 1.56	0.0525 ± 0.0032	80.7 ± 1.6		
SBR_049	0.00	259	287	1.14	75.11 ± 1.48	0.0454 ± 0.0031	85.3 ± 1.7		
SBR_050	0.00	458	475	1.06	83.16 ± 1.33	0.0494 ± 0.0027	77.1 ± 1.2		
SBR_051	0.00	308	286	0.95	80.15 ± 1.51	0.0506 ± 0.0032	79.9 ± 1.5		
SBR_052	0.25	187	173	0.95	83.67 ± 2.22	0.0534 ± 0.0095	76.6 ± 2.0		
SBR_053	0.04	298	419	1.44	84.26 ± 2.01	0.0502 ± 0.0100	76.1 ± 1.8		
SBR_054	0.53	155	106	0.70	82.86 ± 2.00	0.0524 ± 0.0094	77.3 ± 1.9		
SBR_055	0.00	153	142	0.95	81.04 ± 2.03	0.0418 ± 0.0041	79.1 ± 2.0		
SBR_056	0.83	398	262	0.68	81.21 ± 1.48	0.0446 ± 0.0059	78.9 ± 1.4		
SBR_057	0.35	186	178	0.98	77.38 ± 2.00	0.0402 ± 0.0079	82.8 ± 2.1		
SBR_058	1.40	295	144	0.50	85.49 ± 1.54	0.0407 ± 0.0054	75.0 ± 1.3		
SBR_059	0.07	1014	662	0.67	82.18 ± 1.02	0.0511 ± 0.0035	78.0 ± 1.0		
SBR_060	0.24	302	293	0.99	84.32 ± 1.71	0.0442 ± 0.0065	76.0 ± 1.5		
SBR_061	0.00	323	299	0.95	82.02 ± 1.49	0.0486 ± 0.0030	78.1 ± 1.4		
SBR_062	0.00	383	308	0.83	81.70 ± 1.38	0.0512 ± 0.0027	78.4 ± 1.3		
SBR_063	0.15	324	105	0.33	82.14 ± 1.54	0.0449 ± 0.0044	78.0 ± 1.5		
SBR_064	0.00	419	535	1.31	82.97 ± 1.36	0.0501 ± 0.0024	77.2 ± 1.3		
SBR_065	0.00	483	251	0.53	81.86 ± 1.36	0.0447 ± 0.0022	78.3 ± 1.3		
SBR_066	0.41	210	129	0.63	84.40 ± 2.14	0.0416 ± 0.0078	75.9 ± 1.9		
SBR_067	0.00	379	188	0.51	79.61 ± 1.48	0.0450 ± 0.0027	80.5 ± 1.5		
SBR_068	0.00	397	289	0.75	80.48 ± 1.56	0.0465 ± 0.0025	79.6 ± 1.5		
SBR_069	0.72	265	321	1.24	80.75 ± 1.84	0.0470 ± 0.0091	79.3 ± 1.8		
SBR_070	0.00	296	178	0.62	82.81 ± 1.59	0.0514 ± 0.0031	77.4 ± 1.5		
SBR_071	0.00	305	234	0.79	88.09 ± 1.92	0.0454 ± 0.0025	72.8 ± 1.6		

Errors are 1-sigma; Pb_c and Pb^* indicate the common and radiogenic portions, respectively.

"#" with labels mean the data are discordant.

(1) Common Pb corrected by assuming $^{206}\text{Pb}/^{238}\text{U}-^{208}\text{Pb}/^{232}\text{Th}$ age-concordance

(2) The degree of discordance for an analyzed spot indicates the chronological difference between the two ages determined by Pb-Pb and U-Pb methods, and is defined as $\{1-(^{238}\text{U}/^{206}\text{Pb}^* \text{ age})/(^{207}\text{Pb}^*/^{206}\text{Pb}^* \text{ age})\} \times 100$ (%) (e.g., Song et al., 1996).

Appendix 3. O–kb age. O–kb 年代.

Table A10. Summary of the youngest age clusters for dated zircons in the nine sandstone samples (n = number of grains).

砂岩9試料中のジルコンが持つ最若年代集団のまとめ (n は粒子数).

sample name	number of data		YSG	YC1 σ		YC2 σ	
	All	Conc.	(Ma)	(Ma)	n	(Ma)	n
KRH (Kiriata)	72	71	59.6 \pm 1.3	61.1 \pm 0.5	10	63.2 \pm 0.2	42
MJK (Mijikano)	143	138	64.3 \pm 1.9	66.0 \pm 0.6	14	67.8 \pm 0.6	22
SGD (Sagadani)	144	134	57.2 \pm 3.0	60.1 \pm 0.6	16	62.9 \pm 0.4	45
NTK (Natekami)	73	73	63.3 \pm 2.7	65.8 \pm 0.4	26	67.6 \pm 0.4	43
TKO2 (Takeo)	95	88	64.4 \pm 1.3	66.1 \pm 0.6	10	67.8 \pm 0.3	26
SHB1 (Shoubu1)	88	85	61.4 \pm 3.9	65.5 \pm 0.8	11	69.3 \pm 0.5	36
SHB2 (Shoubu2)	93	88	61.3 \pm 1.7	68.0 \pm 0.5	14	66.7 \pm 1.0	9
CCO (Chichioni)	144	138	69.6 \pm 2.2	72.0 \pm 0.4	23	74.3 \pm 0.3	55
SBR (Sobura)	71	70	72.8 \pm 1.6	75.2 \pm 0.5	13	77.0 \pm 0.3	38

Age errors are 1 σ

tone samples (n = number of grains).