





Bat.117.1	-3.4E-5	101	0.1253	0.5	0.242	0.7	0.936	2.6	--	213	139	60.0	14.7	0.67	0.49	1827 ±96	1792 ±108	1786 ±106	2040 ±11	2267 ±131	+12	3.1	6.0	0.1258	0.6	5.68	6.1	0.328	6.0	0.99
Bat.23.1	-6.4E-5	163	0.1089	1.0	0.399	0.9	0.882	1.8	--	44	47	12.6	5.1	1.09	0.62	1836 ±97	1843 ±112	1759 ±115	1795 ±29	2320 ±136	-3	3.0	6.1	0.1097	1.6	4.99	6.3	0.330	6.1	0.97
Bat.49.1	-7.2E-5	246	0.1074	1.3	0.240	1.5	0.843	2.1	--	24	15	6.9	1.7	0.64	0.78	1838 ±98	1849 ±113	1789 ±107	1772 ±47	2398 ±156	-4	3.0	6.1	0.1084	2.6	4.93	6.6	0.330	6.1	0.92
Bat.10.1	-1.3E-4	57	0.1099	0.9	0.182	1.2	0.832	3.2	--	53	24	15.0	2.8	0.48	0.60	1851 ±97	1855 ±113	1811 ±104	1826 ±23	2468 ±148	-2	3.0	6.0	0.1116	1.3	5.12	6.2	0.333	6.0	0.98
Bat.105.1	-3.1E-5	208	0.1298	0.9	0.146	3.4	0.994	3.0	--	54	22	16.3	2.4	0.42	0.61	1938 ±101	1908 ±117	1916 ±108	2101 ±19	2331 ±158	+9	2.9	6.0	0.1302	1.1	6.30	6.1	0.351	6.0	0.98
Bat.71.1	-5.6E-5	57	0.1287	1.3	0.102	1.1	1.036	2.9	--	103	29	31.2	3.3	0.29	0.51	1947 ±103	1920 ±120	1929 ±108	2090 ±24	2428 ±146	+8	2.8	6.2	0.1294	1.4	6.29	6.3	0.353	6.2	0.98
Bat.108.1	-1.6E-4	42	0.1294	0.8	0.146	1.3	0.991	1.9	--	60	23	18.1	2.8	0.40	0.62	1952 ±102	1920 ±118	1920 ±108	2119 ±19	2538 ±154	+9	2.8	6.1	0.1315	1.1	6.41	6.1	0.354	6.1	0.99
Bat.61.1	-3.3E-5	131	0.1265	0.7	0.103	1.2	1.015	3.0	--	77	21	23.6	2.5	0.29	0.54	1960 ±102	1942 ±119	1941 ±106	2056 ±14	2473 ±148	+5	2.8	6.0	0.1269	0.8	6.22	6.1	0.355	6.0	0.99
Bat.77.1	-7.6E-5	80	0.1274	0.8	0.404	5.3	0.982	4.3	--	69	82	21.0	8.6	1.23	0.57	1962 ±102	1941 ±119	1911 ±125	2077 ±17	2244 ±171	+6	2.8	6.0	0.1284	1.0	6.30	6.1	0.356	6.0	0.99
Bat.67.1	-3.0E-4	47	0.1245	1.0	0.281	1.2	0.951	3.7	--	47	37	14.6	4.3	0.80	0.71	1974 ±104	1953 ±122	1916 ±117	2078 ±31	2487 ±153	+6	2.8	6.1	0.1286	1.8	6.35	6.4	0.358	6.1	0.96
Bat.13.1	-1.0E-5	190	0.1269	0.7	0.121	0.7	0.918	4.5	--	269	84	83.6	10.3	0.32	0.41	1989 ±108	1976 ±127	1964 ±113	2057 ±14	2582 ±154	+4	2.8	6.3	0.1270	0.8	6.33	6.3	0.361	6.3	0.99
Bat.80.1	-2.8E-4	32	0.1264	1.0	0.158	3.7	0.968	3.1	--	45	19	14.1	2.4	0.44	0.71	2014 ±105	1997 ±124	1976 ±111	2098 ±23	2670 ±186	+5	2.7	6.1	0.1300	1.3	6.57	6.2	0.367	6.1	0.98
Bat.58.1	-2.4E-5	45	0.1268	0.3	0.099	0.6	0.996	2.9	--	401	105	130.0	13.1	0.27	0.67	2066 ±108	2067 ±131	2046 ±112	2059 ±6	2635 ±153	-0	2.6	6.1	0.1272	0.3	6.62	6.1	0.378	6.1	1.00
Bat.41.1	9.2E-6	189	0.1286	1.1	0.133	0.6	0.959	2.8	0.02	246	87	80.0	10.7	0.36	0.39	2074 ±106	2073 ±129	2048 ±112	2078 ±20	2610 ±148	+0	2.6	6.0	0.1285	1.1	6.72	6.1	0.379	6.0	0.98
Bat.39.1	-9.3E-5	73	0.1242	4.3	0.168	2.5	1.034	1.4	--	88	39	28.8	5.0	0.45	1.96	2082 ±114	2093 ±141	2044 ±122	2035 ±76	2727 ±187	-3	2.6	6.4	0.1254	4.3	6.60	7.7	0.381	6.4	0.83
Bat.31.1	-1.9E-5	153	0.1277	1.1	0.106	0.9	0.874	4.9	--	180	50	59.2	6.4	0.29	0.45	2088 ±107	2092 ±131	2066 ±112	2070 ±20	2680 ±155	-1	2.6	6.0	0.1280	1.2	6.75	6.1	0.383	6.0	0.98
Bat.60.1	-9.3E-5	42	0.1262	0.6	0.147	0.9	0.986	3.5	--	85	34	28.4	4.3	0.41	0.49	2115 ±109	2127 ±135	2085 ±115	2063 ±13	2683 ±156	-3	2.6	6.0	0.1275	0.7	6.83	6.1	0.388	6.0	0.99
Bat.52.1	-3.6E-5	98	0.1279	0.6	0.128	1.0	0.979	3.0	--	85	30	28.5	3.7	0.36	0.49	2122 ±109	2132 ±135	2097 ±114	2076 ±12	2646 ±154	-3	2.6	6.0	0.1284	0.7	6.90	6.1	0.390	6.0	0.99
Bat.40.1	3.2E-5	102	0.1272	1.9	0.202	0.6	1.029	3.2	0.06	188	100	63.0	12.8	0.55	0.45	2126 ±109	2143 ±136	2086 ±118	2053 ±33	2684 ±153	-4	2.6	6.0	0.1267	1.9	6.83	6.3	0.391	6.0	0.95
Bat.47.1	2.6E-5	84	0.1919	0.7	0.107	1.1	0.952	4.9	0.05	240	53	81.0	8.7	0.23	1.93	2136 ±113	2099 ±117	2099 ±117	2756 ±12	3409 ±209	+26	2.5	6.2	0.1916	0.7	10.38	6.3	0.393	6.2	0.99
Bat.45.1	-4.0E-5	68	0.1295	0.5	0.106	0.9	0.997	1.9	--	126	35	42.5	4.6	0.28	0.44	2139 ±109	2148 ±136	2114 ±114	2099 ±10	2814 ±162	-2	2.5	6.0	0.1301	0.6	7.06	6.0	0.393	6.0	1.00
Bat.17.1	-6.5E-5	25	0.1276	0.8	0.105	0.7	0.971	2.4	--	228	64	77.4	8.4	0.29	0.41	2147 ±110	2163 ±137	2124 ±114	2077 ±15	2766 ±157	-4	2.5	6.0	0.1285	0.9	7.00	6.1	0.395	6.0	0.99
Bat.1.1	-1.7E-4	55	0.1256	1.0	0.206	1.2	0.982	4.0	--	38	20	12.7	2.7	0.56	0.66	2147 ±111	2166 ±139	2096 ±120	2069 ±24	2842 ±171	-4	2.5	6.1	0.1279	1.4	6.97	6.2	0.395	6.1	0.98
Bat.26.1	-1.9E-5	173	0.1258	1.5	0.172	2.0	1.010	2.2	--	147	68	50.1	8.8	0.48	1.30	2148 ±110	2172 ±139	2113 ±118	2044 ±27	2697 ±165	-6	2.5	6.0	0.1260	1.5	6.87	6.2	0.395	6.0	0.97
Bat.98.1	-6.8E-5	46	0.1426	0.8	0.129	1.4	1.068	4.2	--	210	75	72.0	9.6	0.37	0.46	2165 ±110	2138 ±135	2140 ±116	2270 ±15	2692 ±157	+5	2.5	6.0	0.1435	0.9	7.89	6.1	0.399	6.0	0.99
Bat.46.1	1.6E-5	111	0.1281	0.4	0.126	0.7	0.991	3.2	0.03	214	72	74.1	9.4	0.35	0.78	2179 ±115	2206 ±147	2154 ±121	2069 ±8	2735 ±162	-6	2.5	6.2	0.1279	0.4	7.09	6.3	0.402	6.2	1.00
Bat.22.1	4.0E-6	1022	0.1342	1.0	0.129	0.9	1.035	2.1	0.01	115	38	40.4	5.2	0.34	0.47	2205 ±112	2219 ±143	2175 ±118	2153 ±18	2889 ±167	-3	2.5	6.0	0.1341	1.0	7.54	6.1	0.408	6.0	0.99

Errors are 1-sigma; Pb<sub>c</sub> and Pb<sub>r</sub> indicate the common and radiogenic portions, respectively.

Error in Standard calibration was 0.35% (not included in above errors but required when comparing data from different mounts).

(1) Common Pb corrected using measured <sup>204</sup>Pb.

(2) Common Pb corrected by assuming <sup>206</sup>Pb/<sup>238</sup>U-<sup>207</sup>Pb/<sup>235</sup>U age-concordance

(3) Common Pb corrected by assuming <sup>206</sup>Pb/<sup>238</sup>U-<sup>207</sup>Pb/<sup>232</sup>Th age-concordance





