

Supplementary Table S1.

Sample ID	Total number of sequenced reads	Total number of uniquely mapped non-duplicate reads	Total number of covered bases	Median coverage (and range) per base	Percentage of targeted bases with coverage >200
Osteoblastic					
OS					
1	78162786	57374475	2688927	1583(0-9562)	97.68
2	42343816	30624504	2688606	842(0-9226)	91.06
3	54553830	41293994	2688469	1033(0-10208)	98.23
4	60115584	46167862	2689512	1305(0-6212)	97.37
5	66068070	50590563	2688710	1281(0-17122)	96.67
6	52646996	38906717	2688555	1179(0-6407)	96.8
7	77888456	56638929	2688759	1472(0-18804)	97.52
8	53000830	42208802	2688479	1480(0-14042)	95.74
9	60926136	48145550	2689341	1424(0-6960)	97.25
10	52529878	38915874	2688603	1101(0-5596)	95.97
11	72629840	52018116	2689132	1514(0-9046)	98.06
12	76676114	54269584	2688839	1427(0-4985)	97.52
13	54839798	43117797	2688498	1288(0-5913)	97.34
14	76610172	56103756	2688743	1639(0-8551)	98.51
15	62740242	43083827	2689006	1171(0-4820)	96.98
16	80357252	59019469	2689259	1613(0-9111)	98.18
17	72385074	52810447	2689211	1362(0-9245)	97.27
18	67257524	44159544	2689484	1210(0-26469)	97.6
19	84875116	60547033	2687275	1879(0-11530)	98.35
20	75407976	55463257	2689405	1626(0-16995)	98.03
21	76521488	53764057	2688531	1474(0-7533)	97.3
Chondroblastic					
OS					
22	76236968	48305066	2688998	1627(0-8505)	97.38
23	76372648	54909260	2688785	1534(0-13621)	97.63
24	62857048	46781681	2689438	1222(0-30499)	97.63