S2 Table. Cell lines in pooled set and sequencing quality

Normal HapMap cell line	Total reads	Uniquely aligned reads to hg19	Fraction of uniquely alinged reads (%)	On target bases	Mean target coverage	FractionTarget Covered (> ×100) (%)
NA19059	7,747,706	6,844,528	88	299,216,207	868	99
NA12767	9,685,732	6,413,560	66	276,595,345	803	99
NA19235	7,789,868	6,928,292	89	277,114,644	804	99
NA18595	6,975,830	6,224,850	89	271,406,072	788	99
NA19108	9,491,778	8,209,666	86	347,944,445	1010	99
NA19190	7,912,268	6,939,115	88	300,840,256	873	99
NA18618	5,961,450	5,328,755	89	233,253,852	677	99
NA19214	9,611,314	8,301,406	86	361,595,594	1049	99
NA18511	10,059,030	8,684,561	86	374,578,154	1087	99
XX	9,103,128	5,729,837	63	233,249,469	677	99