

**Table S2** Summary of positions passing quality filters

<b>Filter</b>	<b>Passing Sites</b>	<b>Description</b>
Positions in Reference	2,460,580	unfiltered sequence
Regional Filtering	868,575	sites passing depth and MQ0 Ratio Filters
Site-Level Filtering	497,722	site-level filtering of heterozygote, MQ0 Ratio, and sites with missing genotypes
Second Depth	488,818	positions with extreme depth values
Indel Filter	484,924	+/- 5bp from indels