Table S1. The genotype concordance of 24 CHB samples derived from our sequencing results compared to ~80x WGS (Lan et al. 2017). The average genotype concordance for SNPs and Indels is 99.54% and 96.03%, respectively.

Samples	SNPs	Indels
HG00421	99.60%	96.28%
HG00428	99.65%	96.09%
HG00436	99.11%	96.11%
HG00442	99.22%	96.07%
HG00443	99.89%	96.46%
HG00472	99.75%	96.81%
HG00473	99.53%	96.38%
HG00478	99.71%	96.05%
HG00500	99.86%	96.07%
HG00619	99.54%	96.01%
HG00653	99.35%	95.54%
HG00683	99.64%	96.12%
HG00699	99.80%	96.07%
NA18532	99.76%	96.09%
NA18542	99.46%	95.96%
NA18547	99.75%	96.16%
NA18570	99.71%	96.33%
NA18572	99.74%	96.61%
NA18573	98.71%	94.25%
NA18577	99.55%	95.58%
NA18605	99.54%	95.79%
NA18611	99.21%	95.83%
NA18620	99.43%	96.04%
NA18624	99.43%	96.11%