

Table 1. Summary of the results of the analysis of the data collected from the 1000 Genomes Project									
Study	Population	Sample Size (n)	Genotyping Method	Genotyping Rate (%)	Genotyping Error Rate (%)	Genotyping Coverage (%)	Genotyping Accuracy (%)	Genotyping Precision (%)	Genotyping Recall (%)
1000 Genomes Project	Yoruba (YRI)	2,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CEU	2,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CHB	2,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CHS	2,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CHB+CHS	4,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
1000 Genomes Project	CHB+CHS	4,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CHB+CHS	4,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
	CHB+CHS	4,000	SNP arrays	99.9	0.1	99.9	99.9	99.9	99.9
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