

Table S2 Proportion of each genotype in the PAG2014 dataset (Sanger sequencing) as called by the 1000 Genomes. The diagonal shows the proportion of correctly called genotypes. ALT = alternative allele; REF = reference allele.

		1000 Genomes		
		ALT/ALT	ALT/REF	REF/REF
PAG2014	ALT/ALT	0.699	0.227	0.074
	ALT/REF	0.029	0.681	0.290
	REF/REF	0.001	0.069	0.930