RPL genes (n=94,678 loci; 47 heterozygous + 47 homozygous SNPs) 2x RPL +10% (n=189,356 loci; 94 heterozygous + 94 homozygous SNPs) 2x RPL +2% (n=189,356 loci; 94 heterozygous + 94 homozygous SNPs)

Accuracy = -10 log(1 - correct/total SNP calls); An accuracy of 30 means 99.9% of loci are called correctly