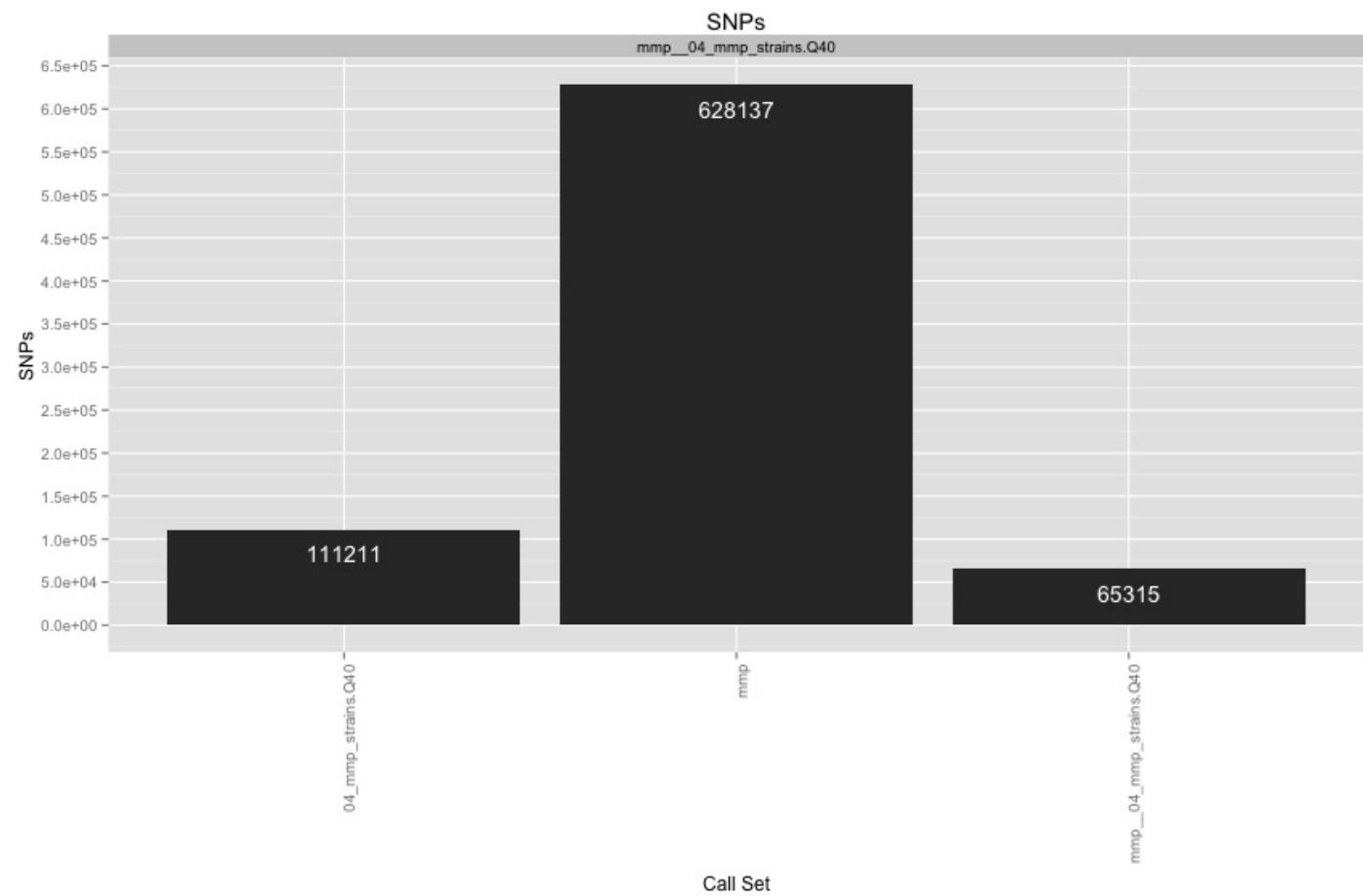


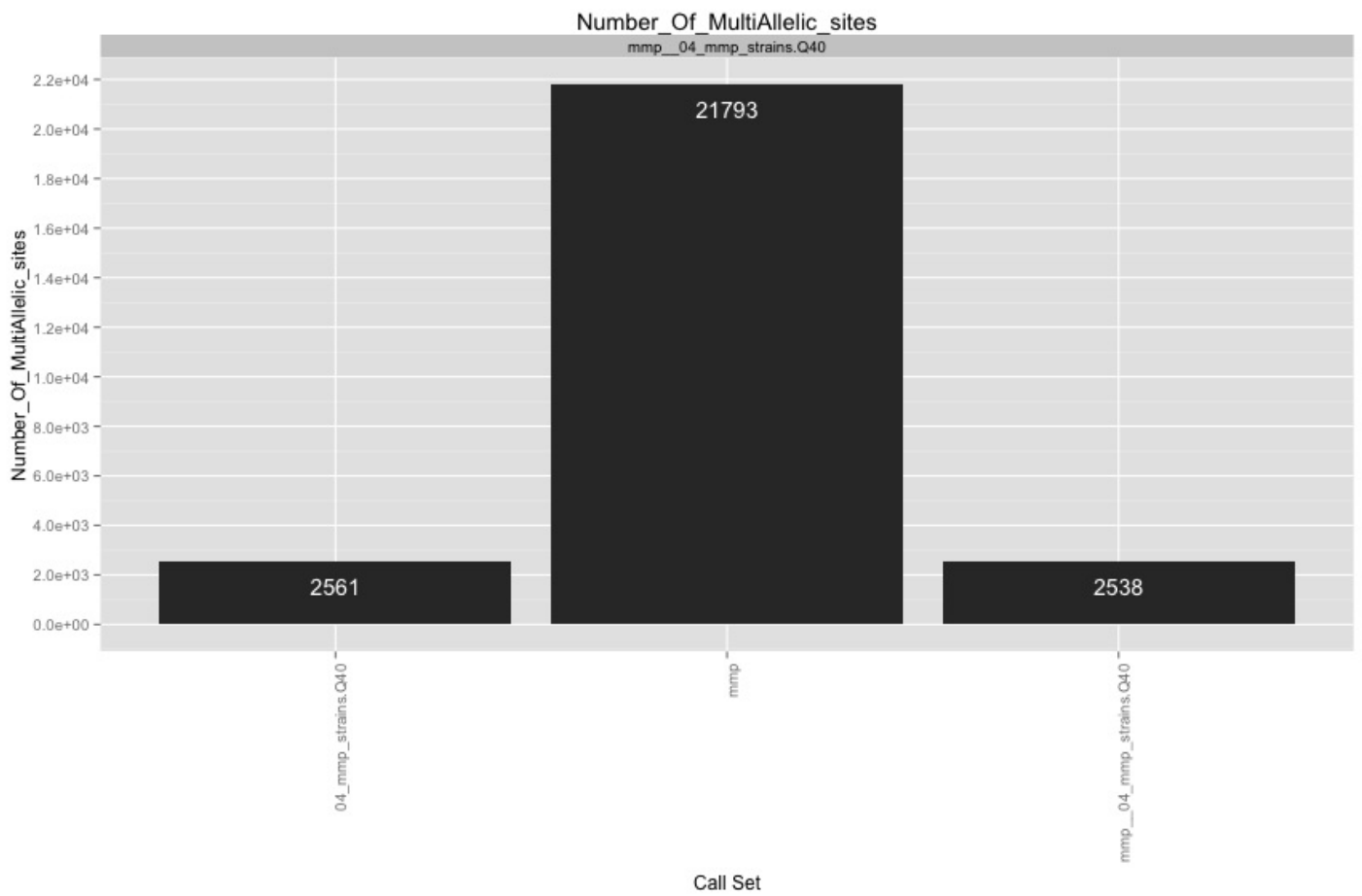
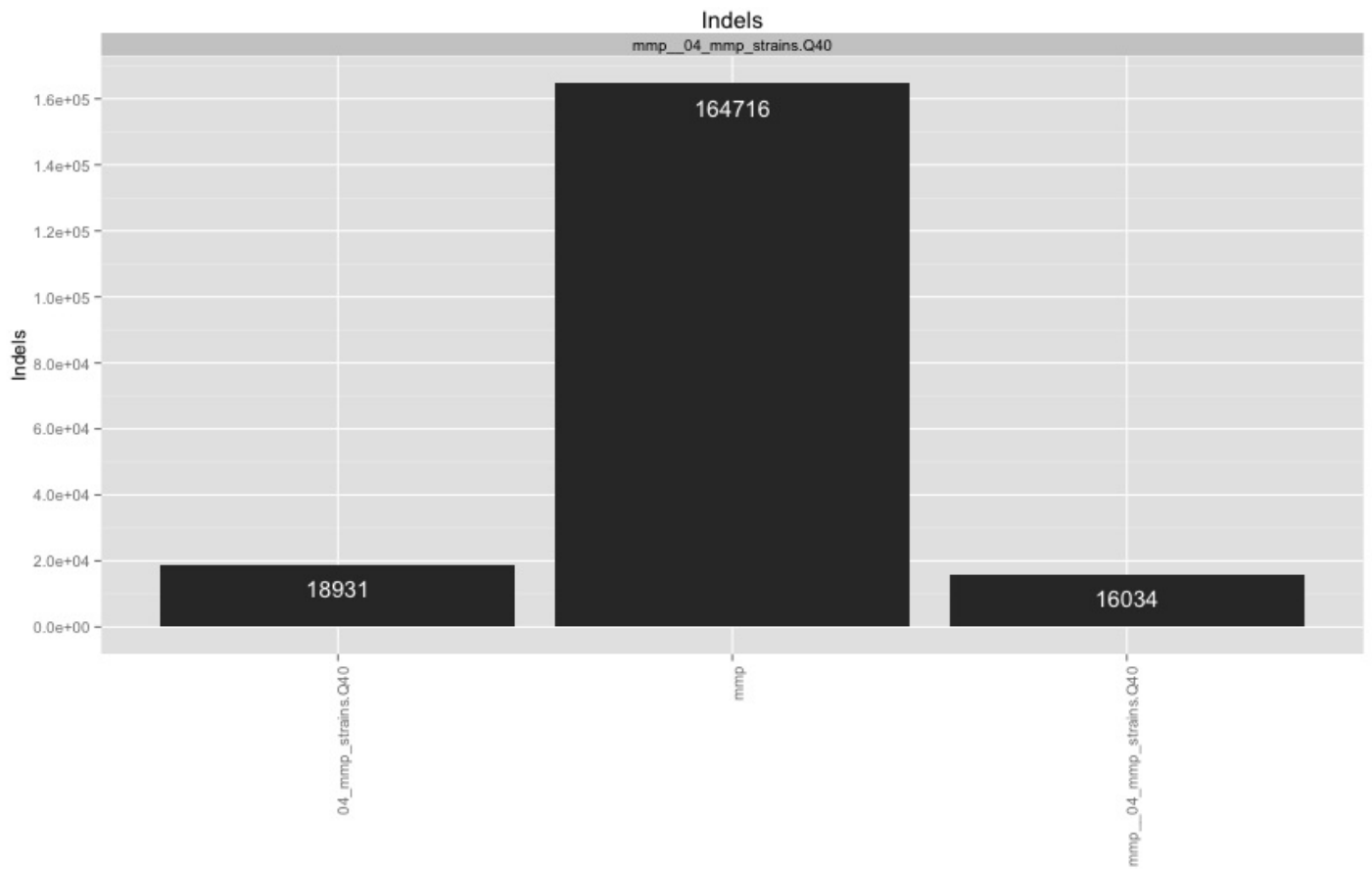
VCF Compare Script

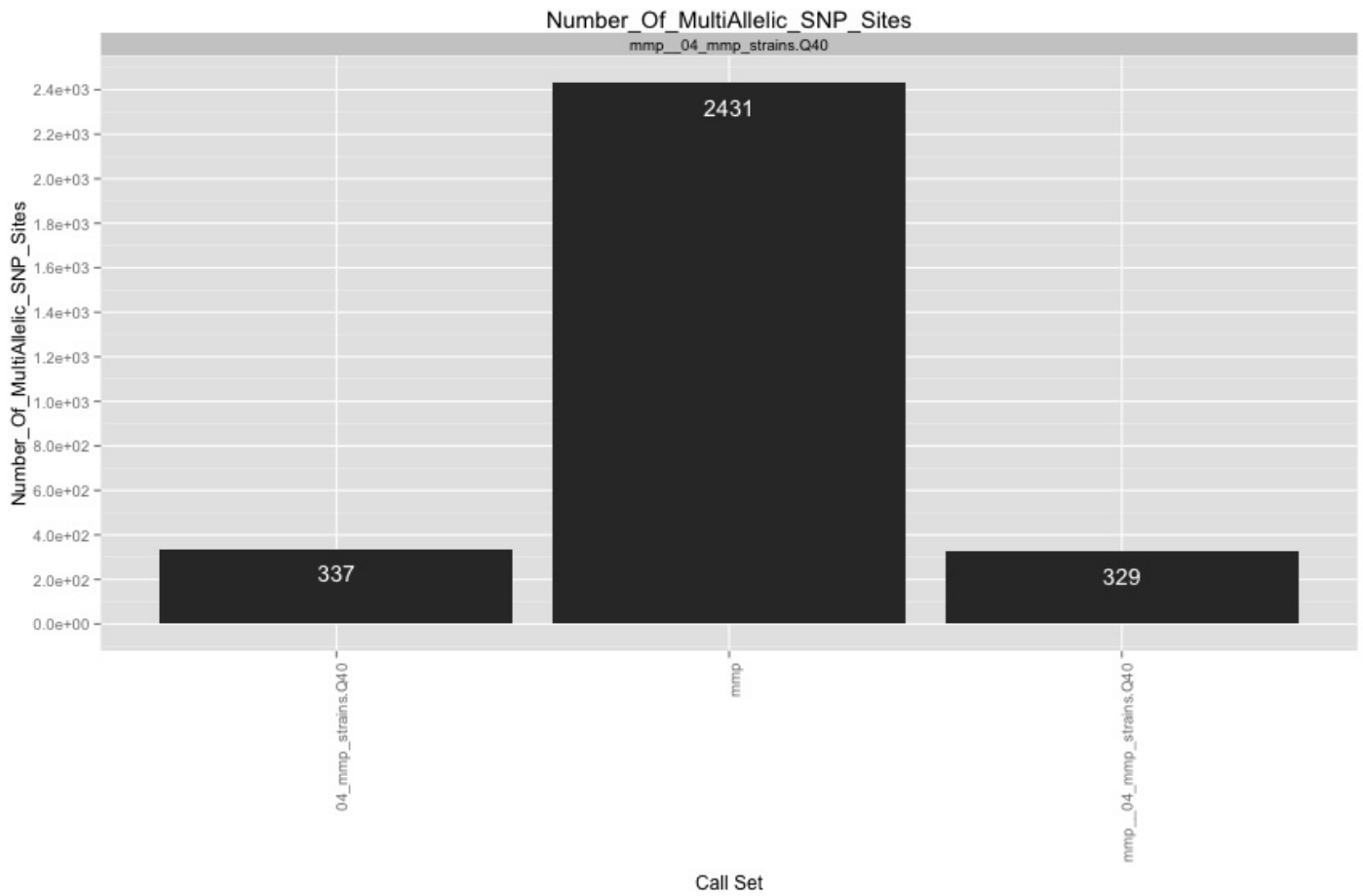
Files

[1] "mmp.vcf.gz" "04mmpstrains.txt.Q40.vcf.gz"

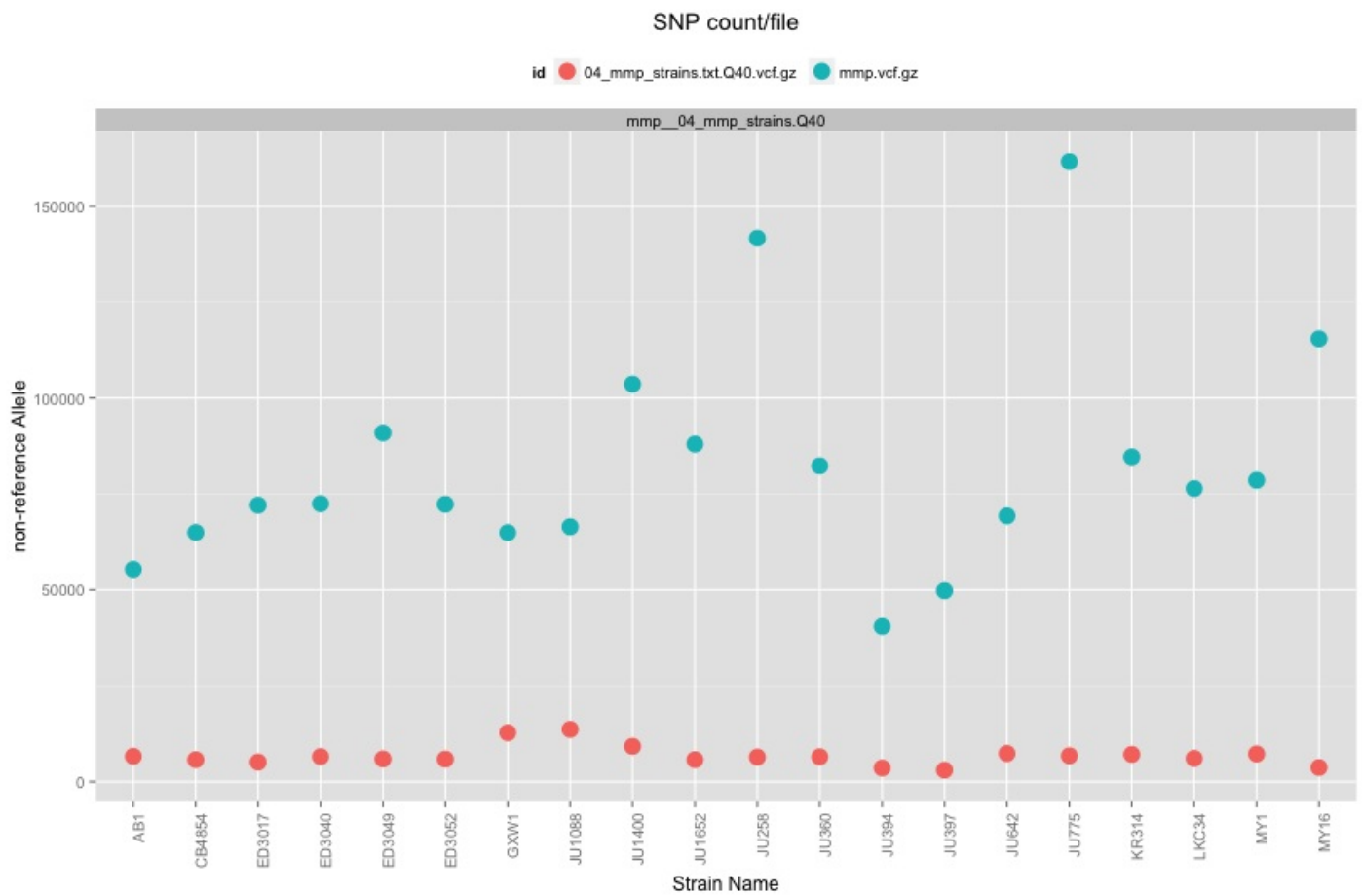
Individual VCF Results



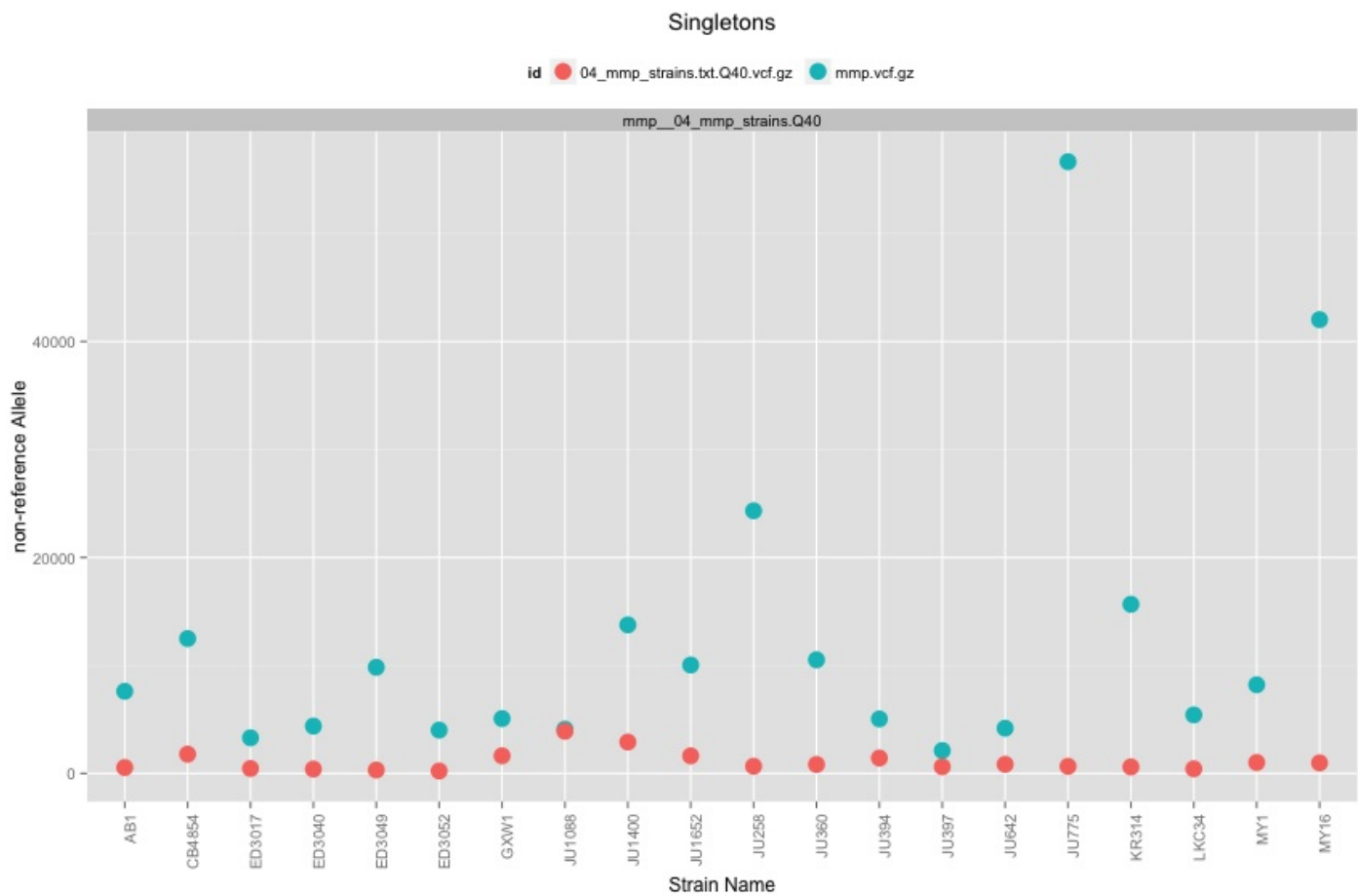




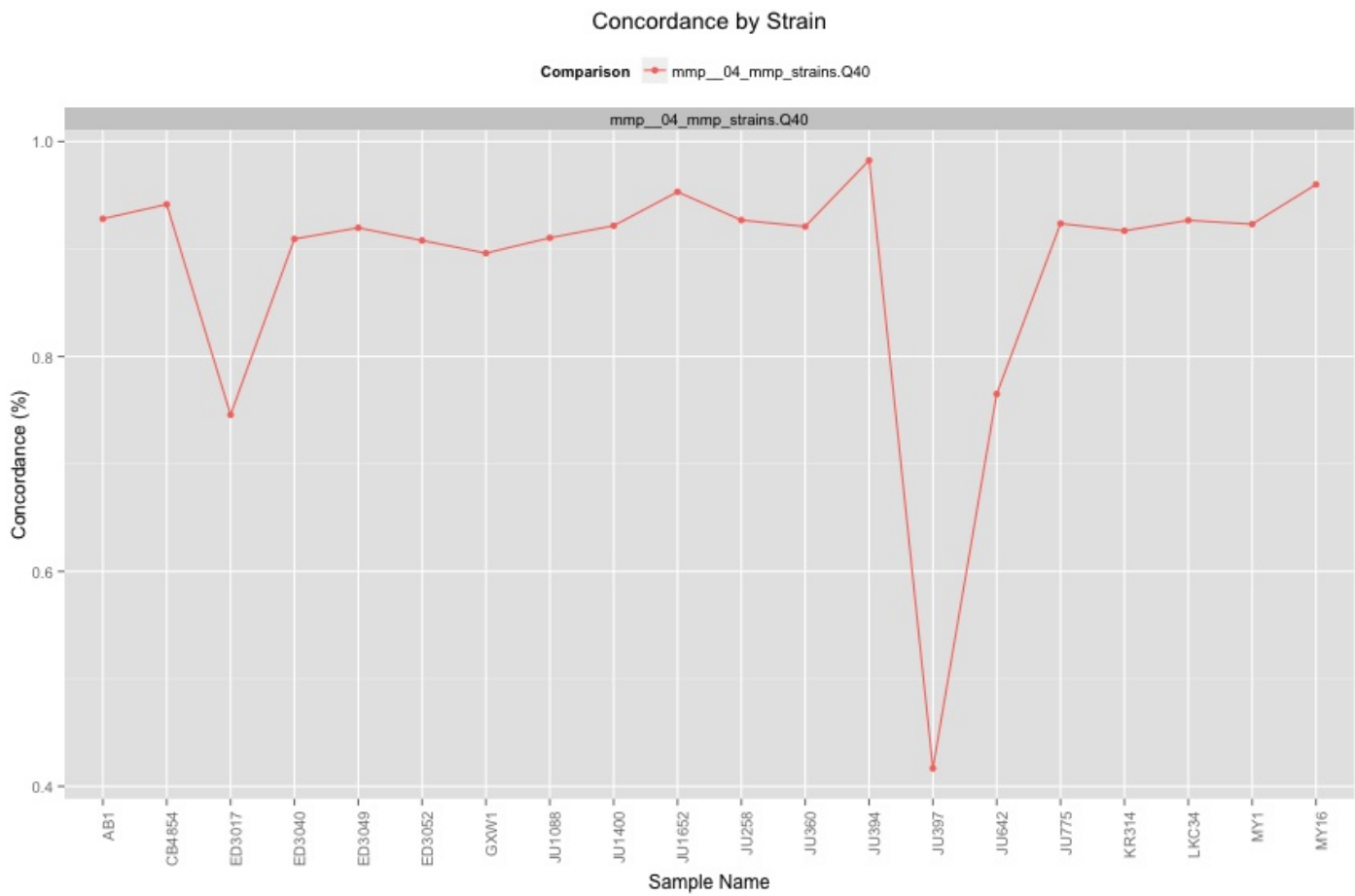
Number of SNPs/Strain



Singletons



Ind. Sample Concordance



Pairwise Concordance

[[1]]

