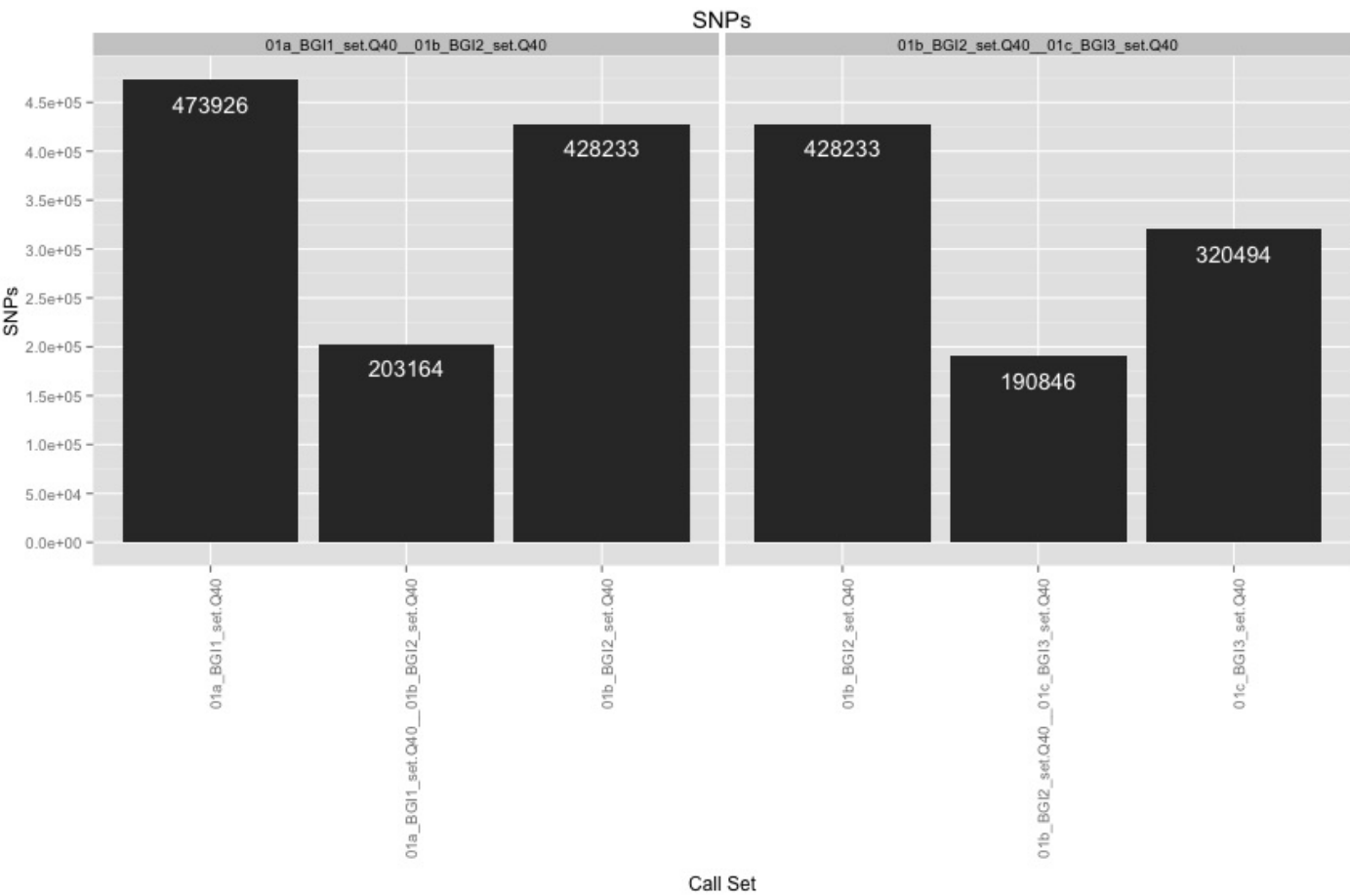


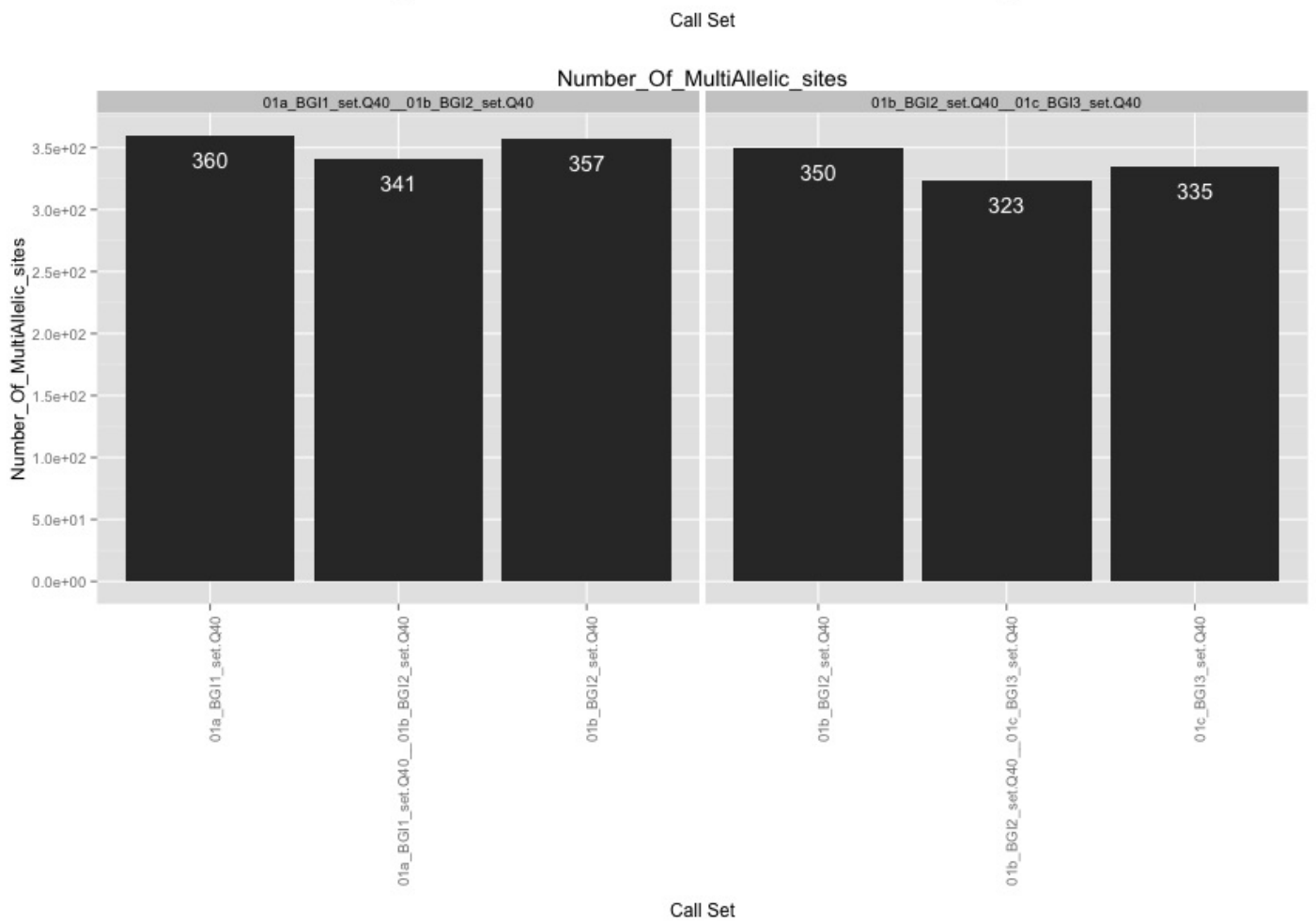
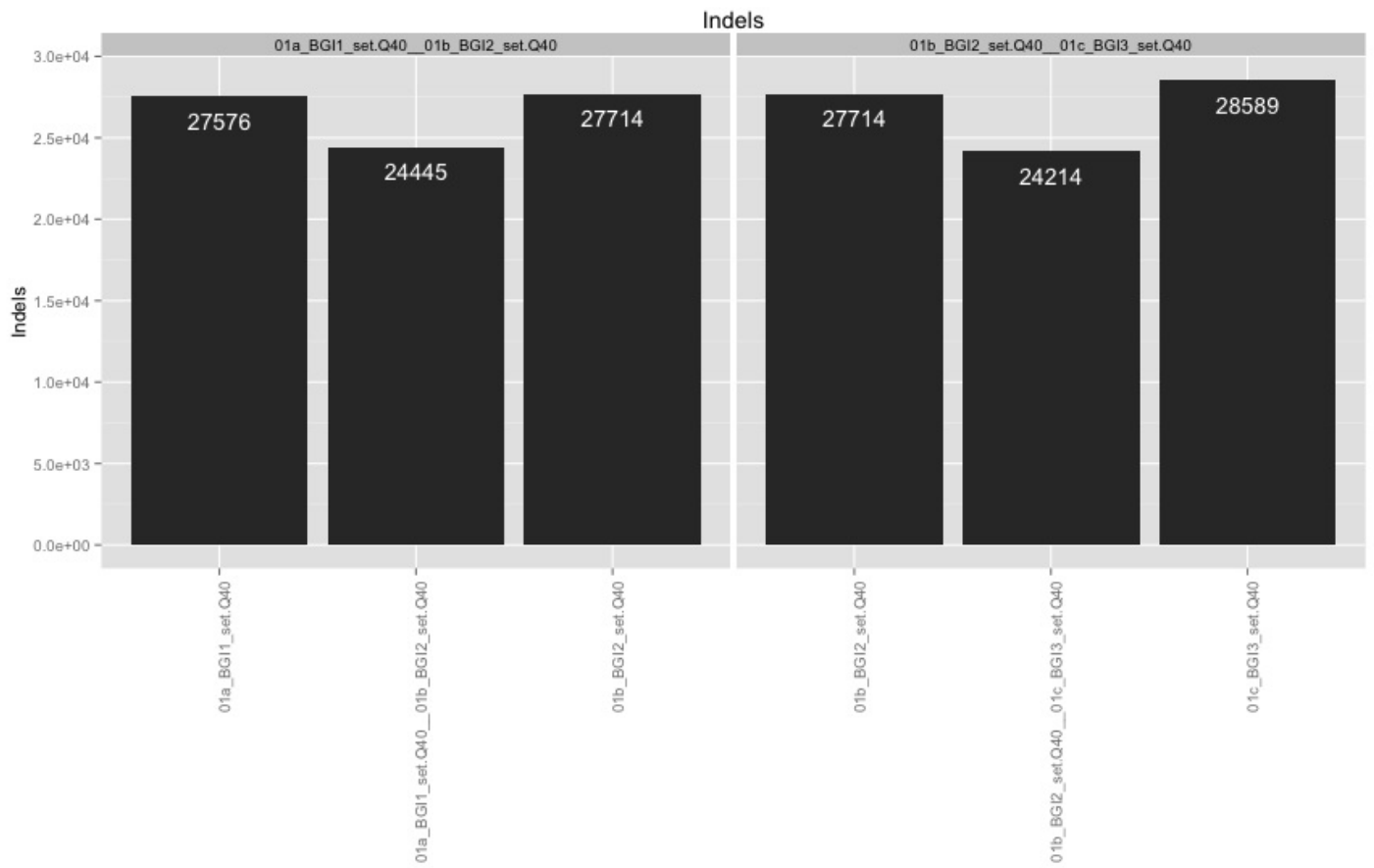
# VCF Compare Script

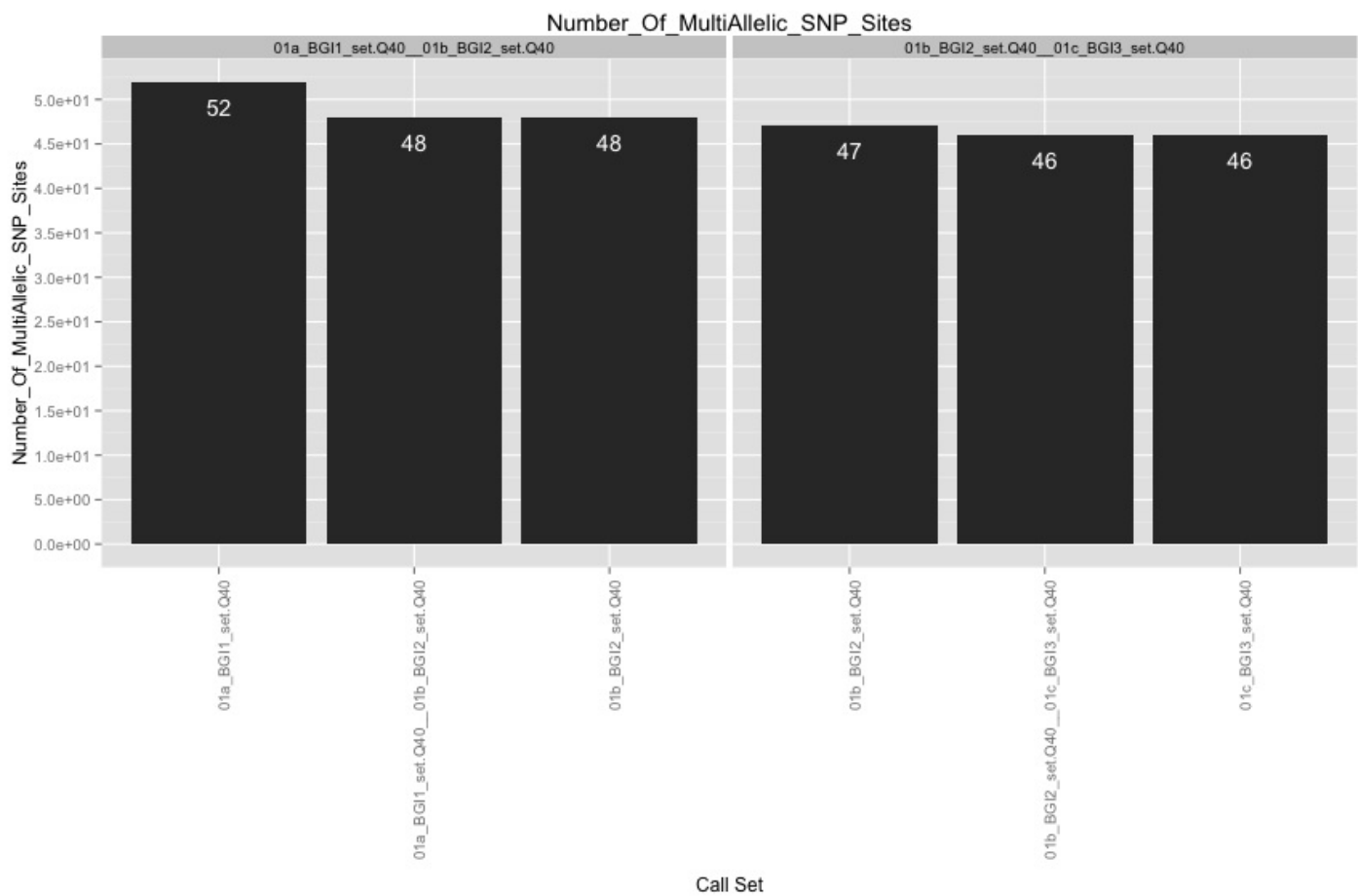
## Files

[1] "01aBGI1set.txt.Q40.vcf.gz" "01bBGI2set.txt.Q40.vcf.gz" [3] "01cBGI3set.txt.Q40.vcf.gz"

## Individual VCF Results





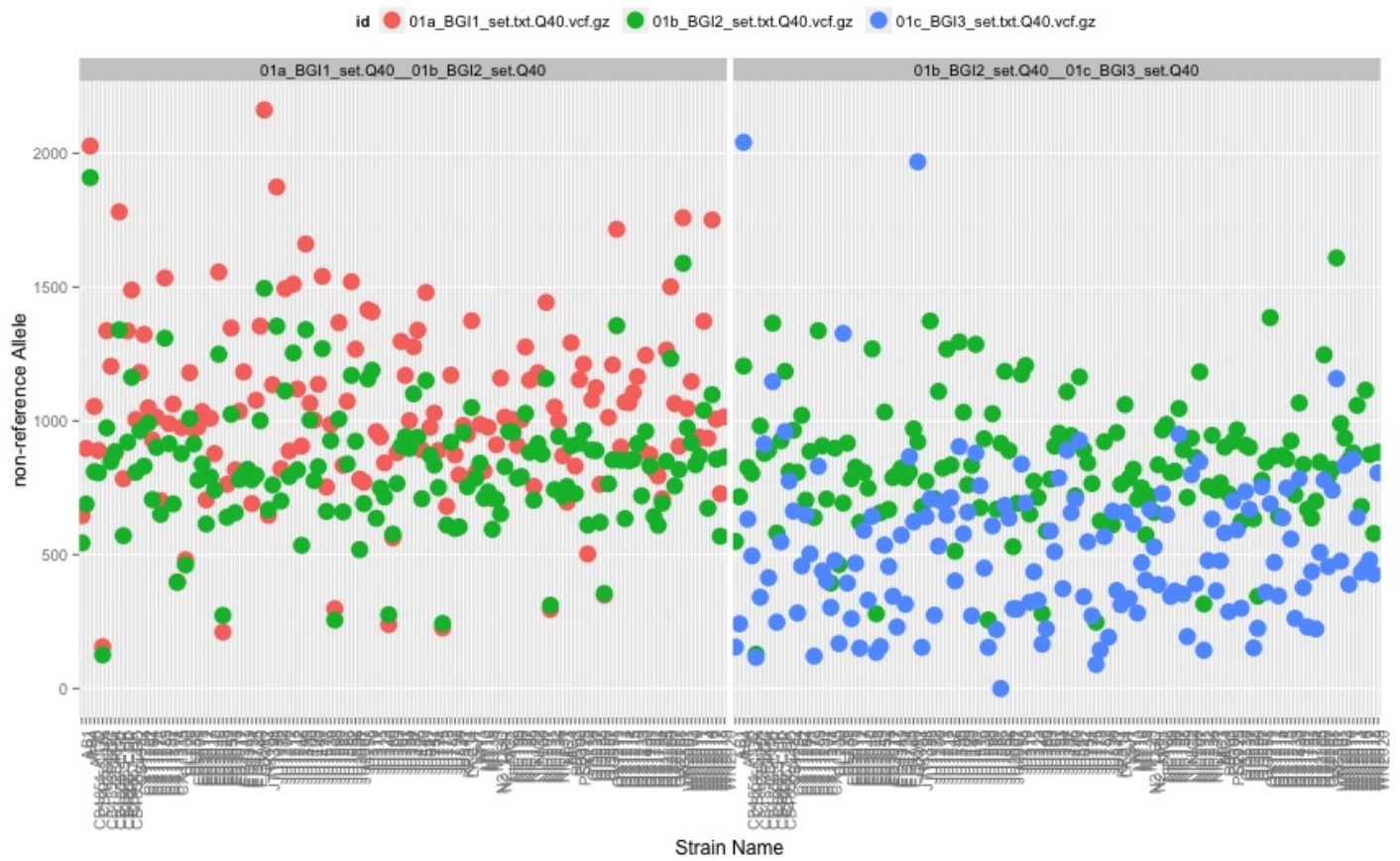


## Number of SNPs/Strain



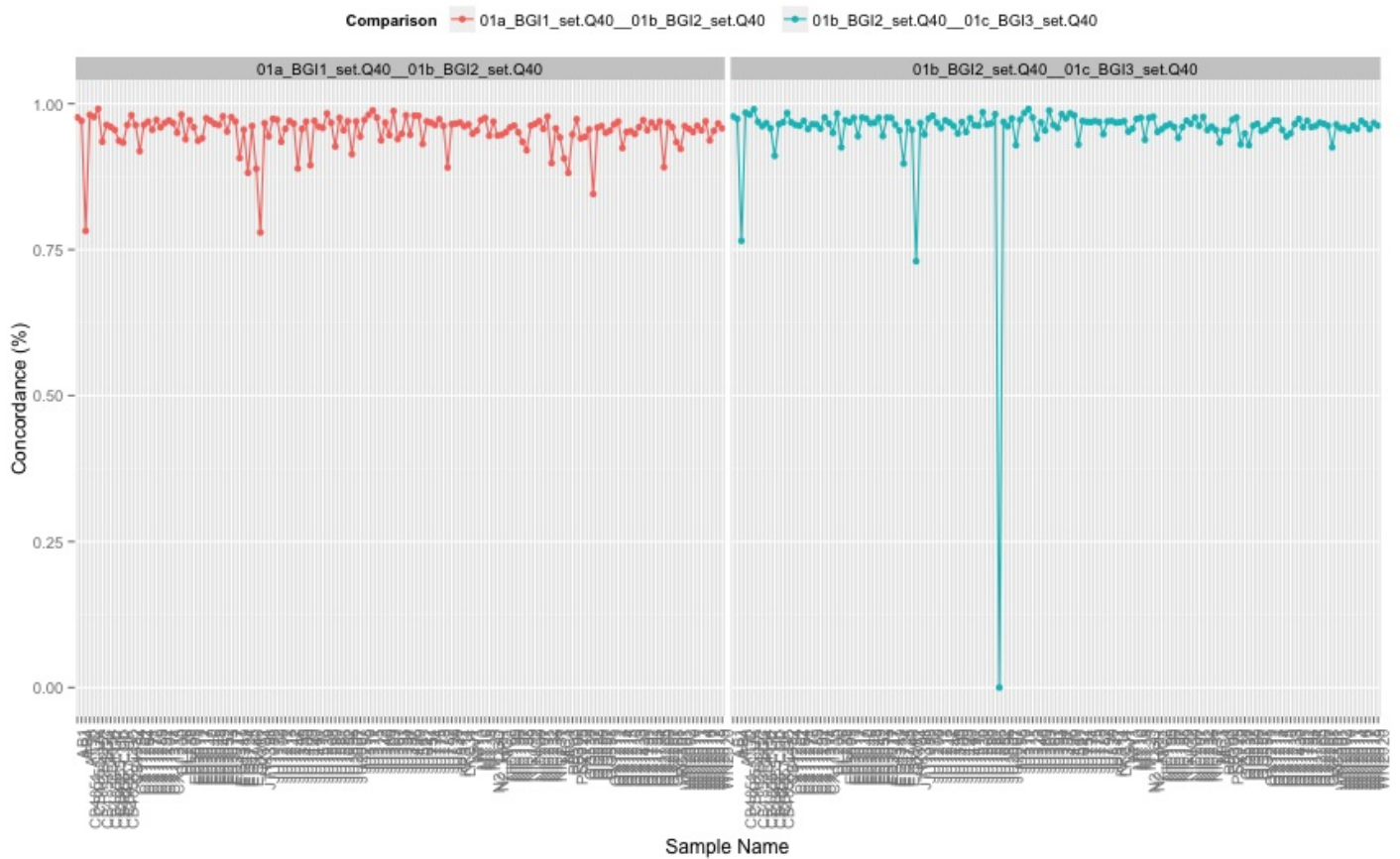
## Singletons

## Singletons



## Ind. Sample Concordance

### Concordance by Strain



## Pairwise Concordance

[[1]]



