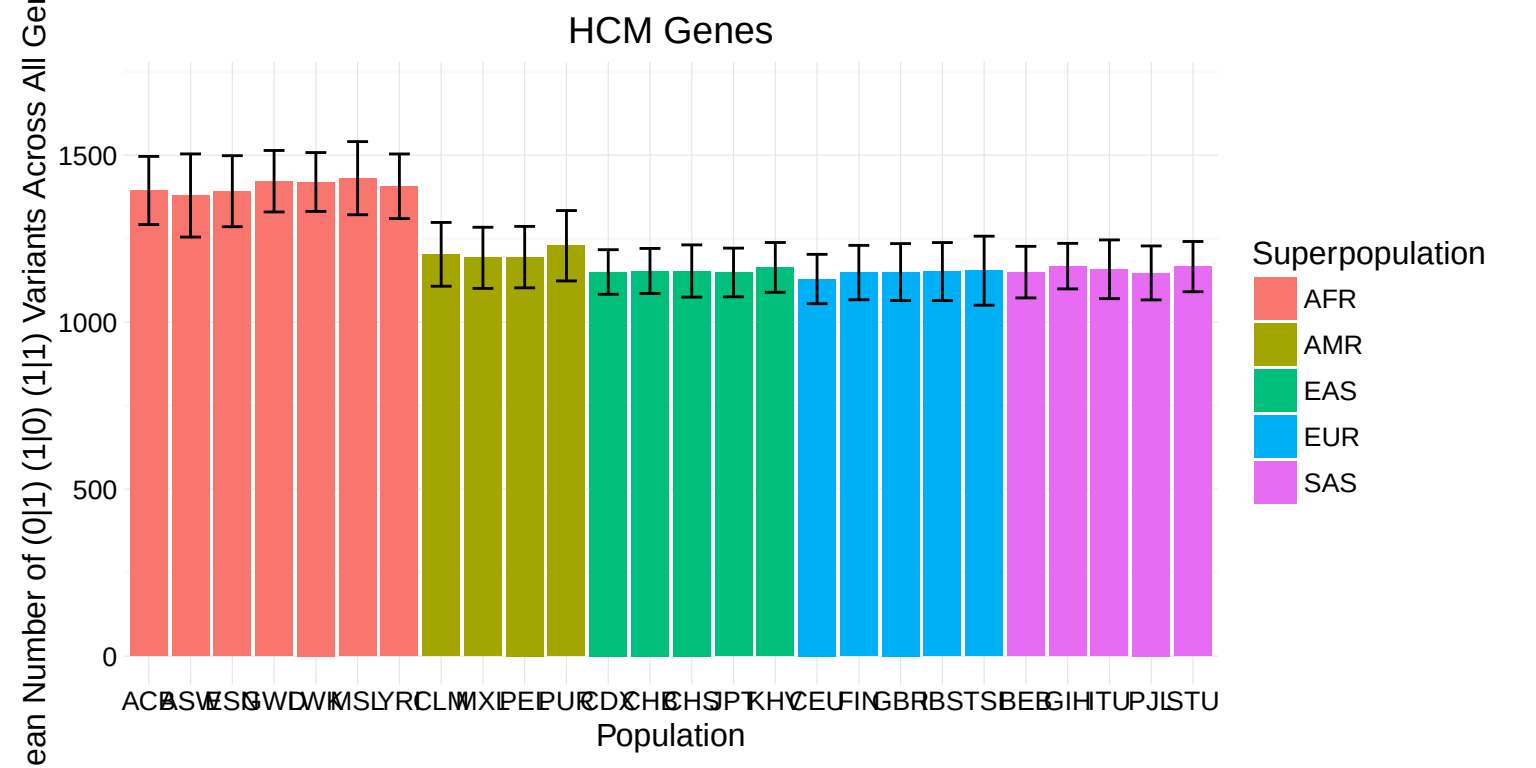
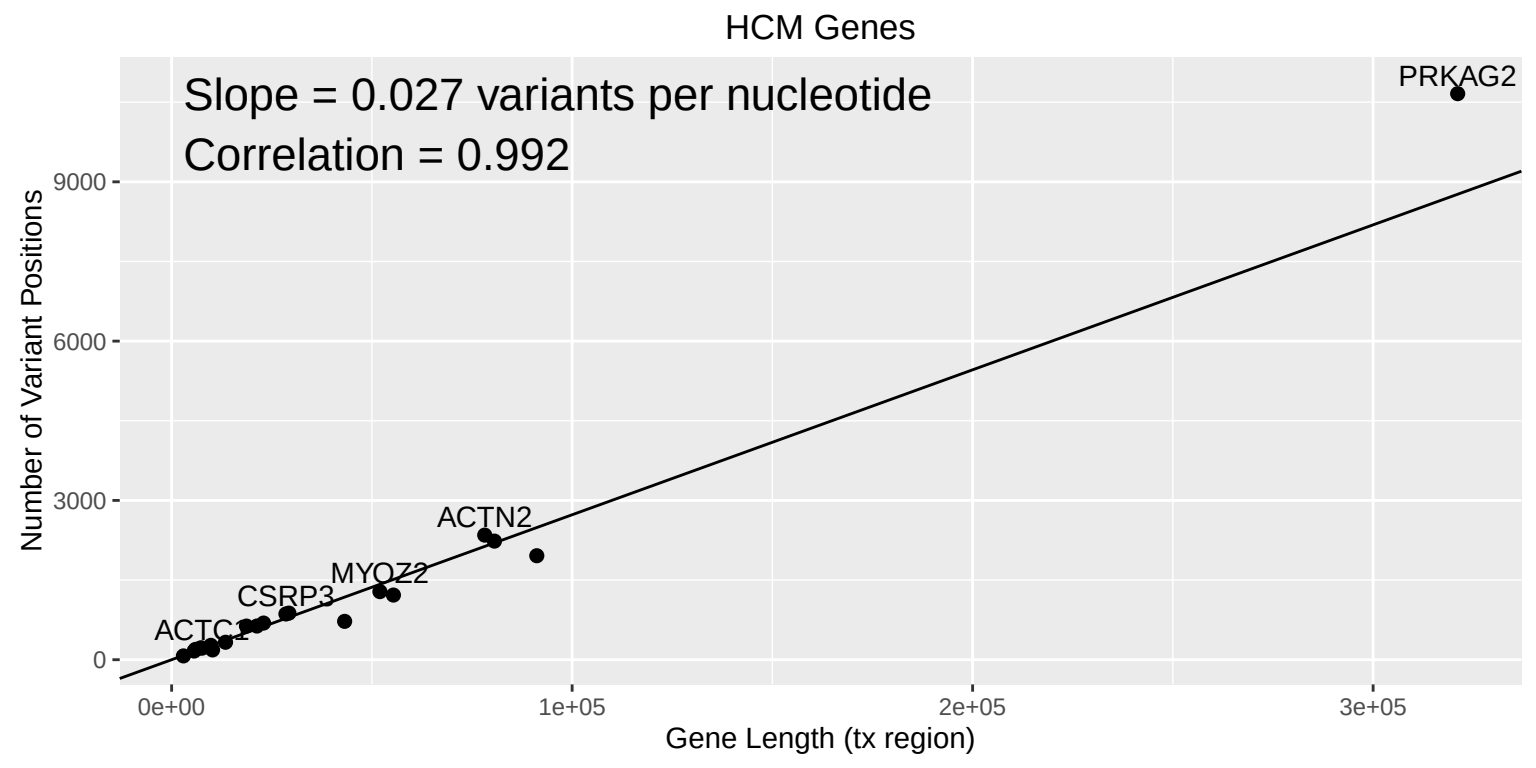

HCM Genes
Mean Number of (0|1) (1|0) (1|1) Variants Across All Genes
1500
1000
500
0
Population
Superpopulation
AFR
AMR
EAS
EUR
SAS


HCM Genes
Slope = 0.027 variants per nucleotide
Correlation = 0.992
PRKAG2
ACTN2
MYOZ2
CSRP3
ACTC1
Number of Variant Positions
9000
6000
3000
0
0e+00
1e+05
2e+05
3e+05
Gene Length (tx region)