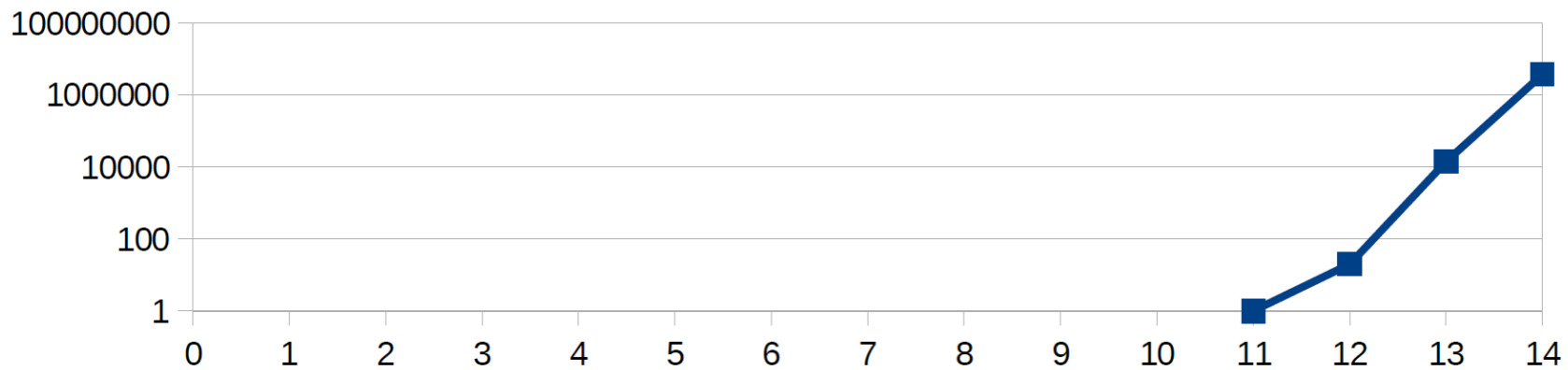
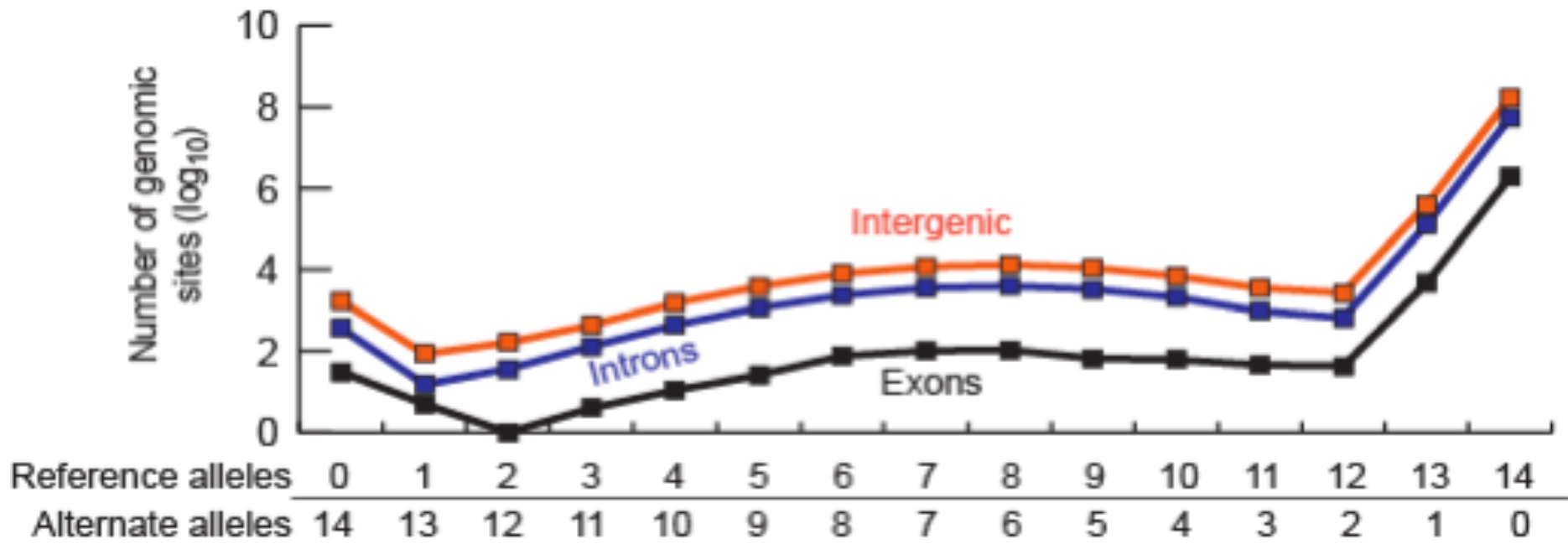


Heterozygosity



Notes: SW019 2x250 library against all Discover contigs > 50kb; ~50Mb
 -Q & -q > 30 in samtools
 Coverage = 14