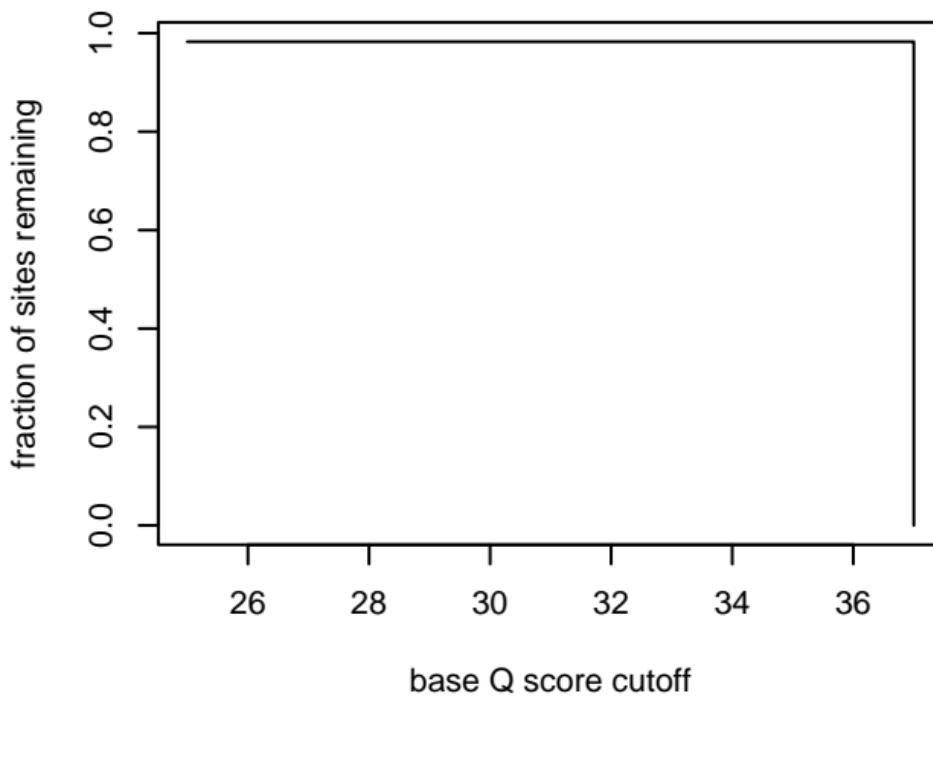
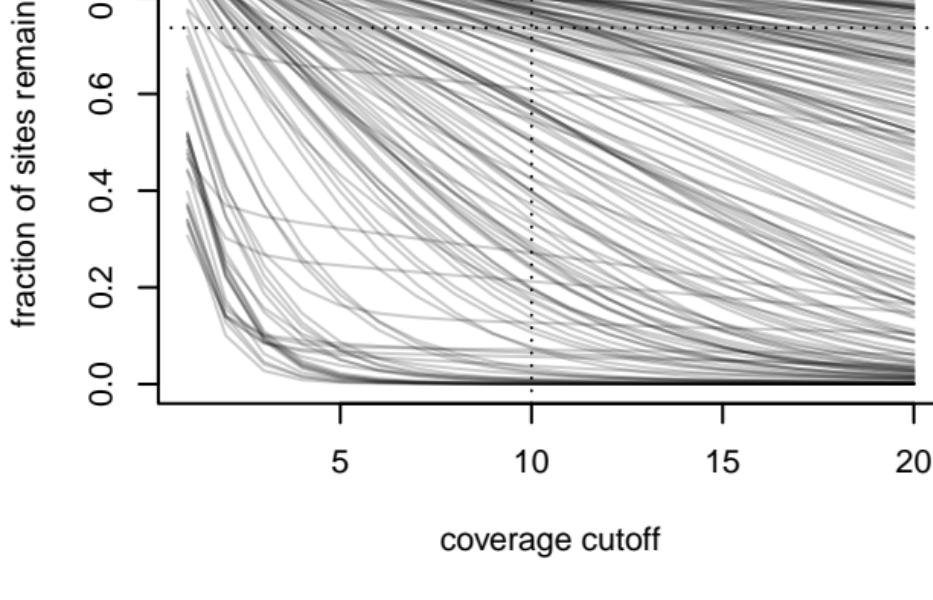


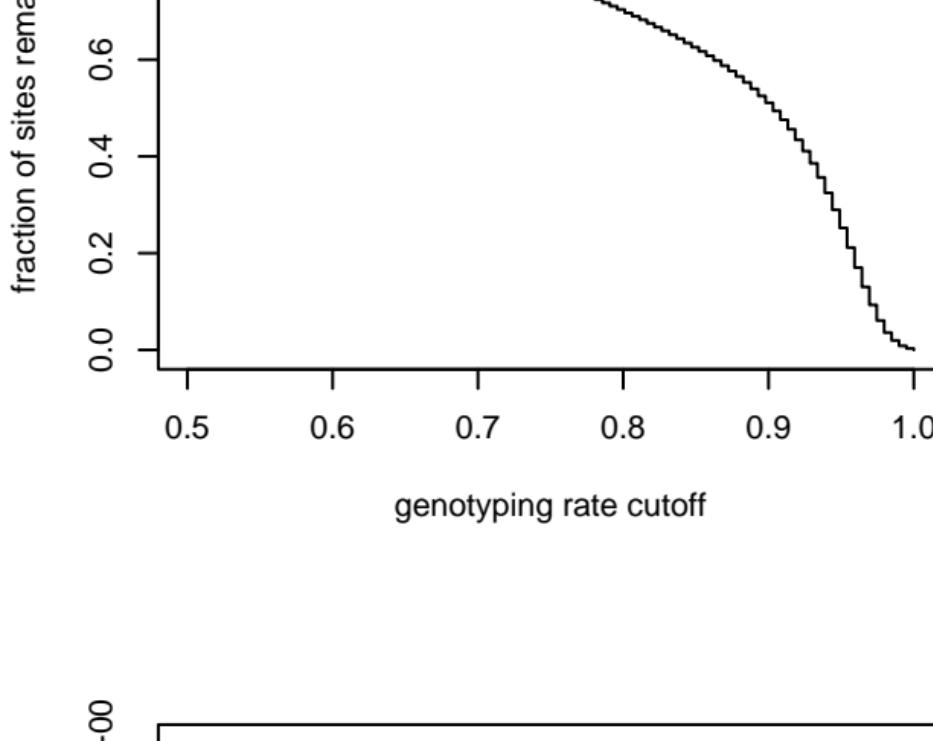
1781183 sites



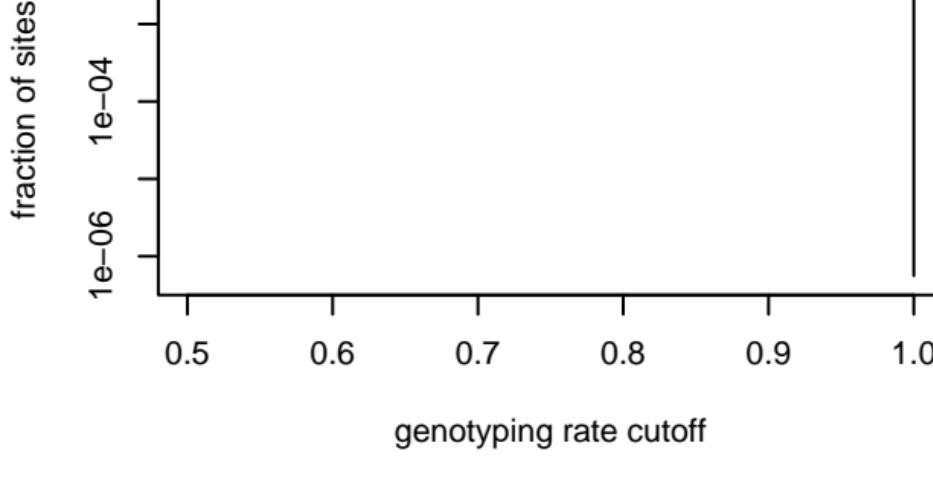
base Q score cutoff



coverage cutoff



genotyping rate cutoff



genotyping rate cutoff