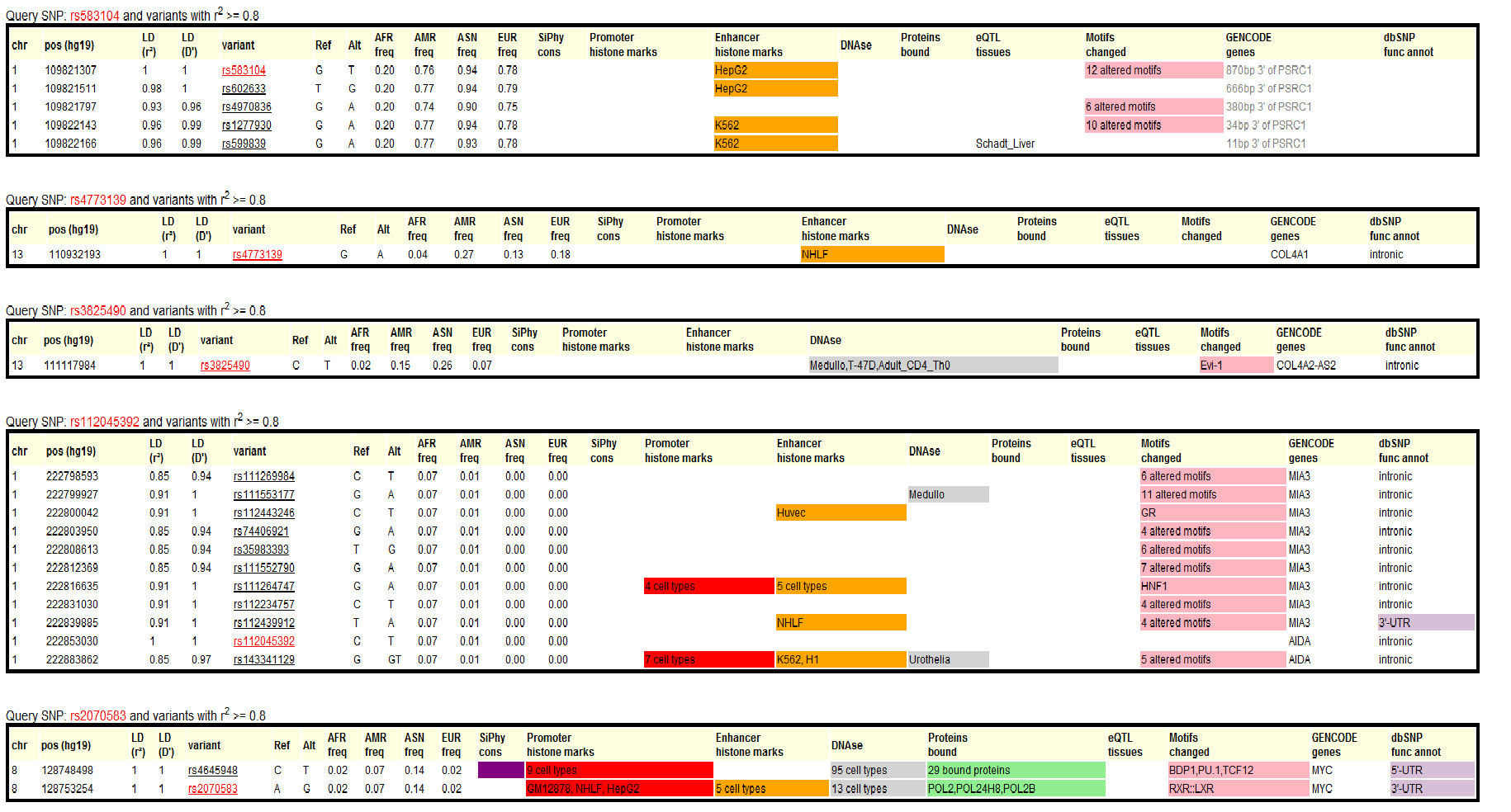
**Table S3.** Bioinformatic functional annotation of SNPs in high LD (r2>0.8 in AFR) identified in validated and new loci for incident coronary heart disease. Data obtained from HaploReg at www.broadinstitute.org/mammals/haploreg/haploreg.php



ed using R v2.15.1 (survival package), under an additive model.