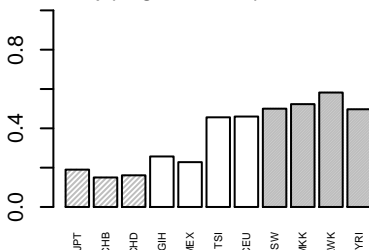
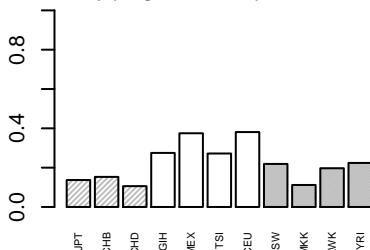


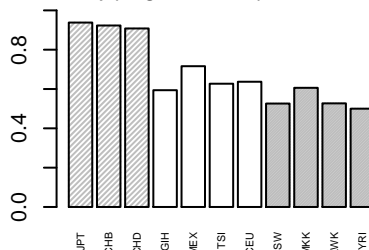
rs9939609_A_FTO(anc)
89 studies on 27 populations
p (vs. genomic SNPs)= 2.17e-02



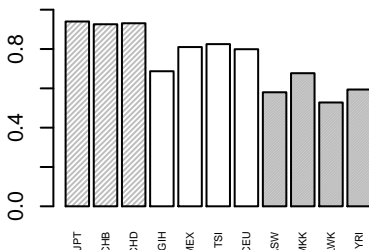
rs7498665_G_SH2B1(anc)
13 studies on 7 populations
p (vs. genomic SNPs)= 1.00e-01



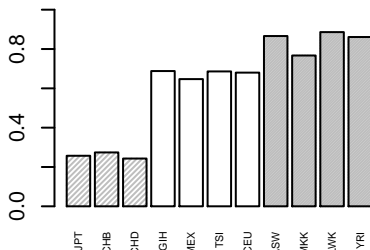
rs3101336_C_
8 studies on 7 populations
p (vs. genomic SNPs)= 5.62e-01



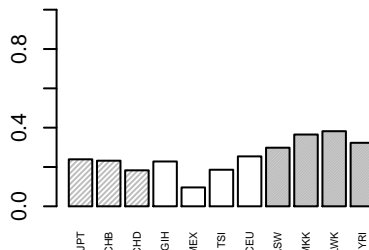
rs7647305_C_
6 studies on 7 populations
p (vs. genomic SNPs)= 6.24e-01



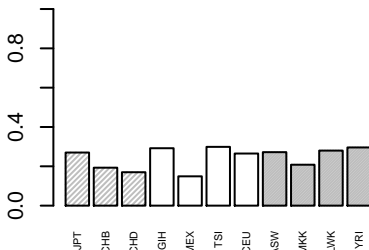
rs29941_G_
6 studies on 7 populations
p (vs. genomic SNPs)= 3.77e-03



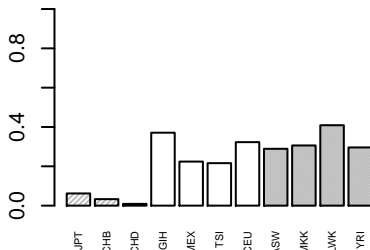
rs10913469_C_SEC16B(anc)
5 studies on 7 populations
p (vs. genomic SNPs)= 1.32e-01



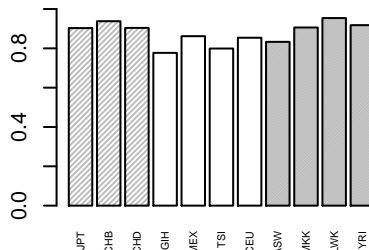
rs17782313_C_
30 studies on 6 populations
p (vs. genomic SNPs)= 1.60e-01



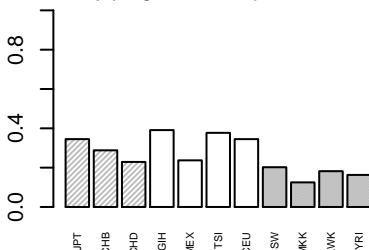
rs925946_T_
11 studies on 6 populations
p (vs. genomic SNPs)= 1.47e-02



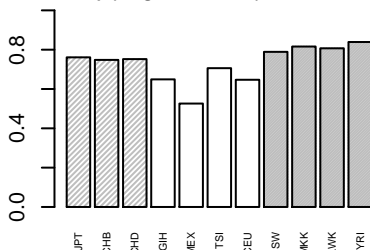
rs2867125_C_(anc)
7 studies on 6 populations
p (vs. genomic SNPs)= 1.53e-01



rs7138803_A_
7 studies on 6 populations
p (vs. genomic SNPs)= 2.80e-01



rs7190492_G_FTO(anc)
5 studies on 5 populations
p (vs. genomic SNPs)= 1.29e-01



rs6499640_A_FTO(anc)
5 studies on 5 populations
p (vs. genomic SNPs)= 7.09e-03

