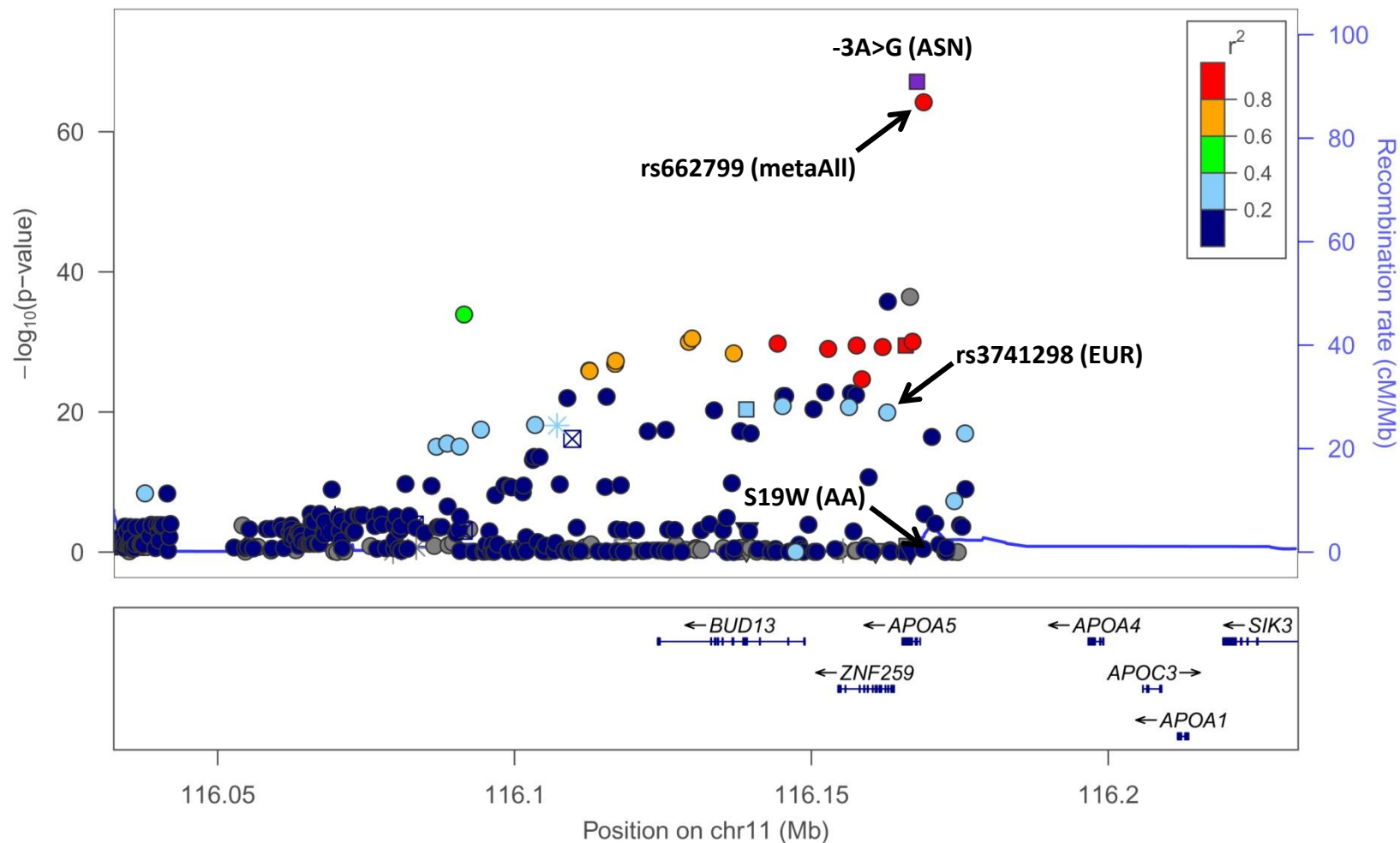
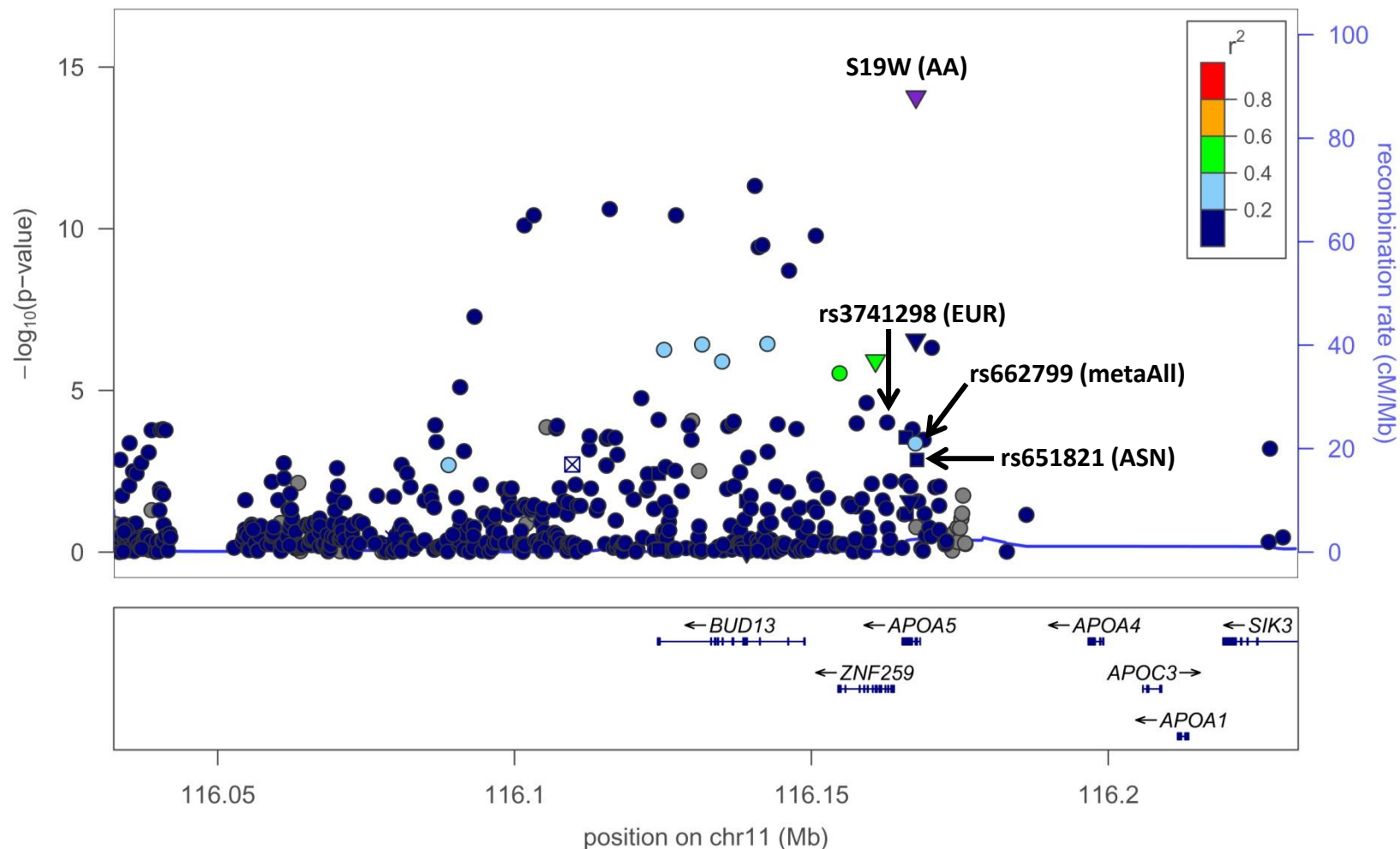


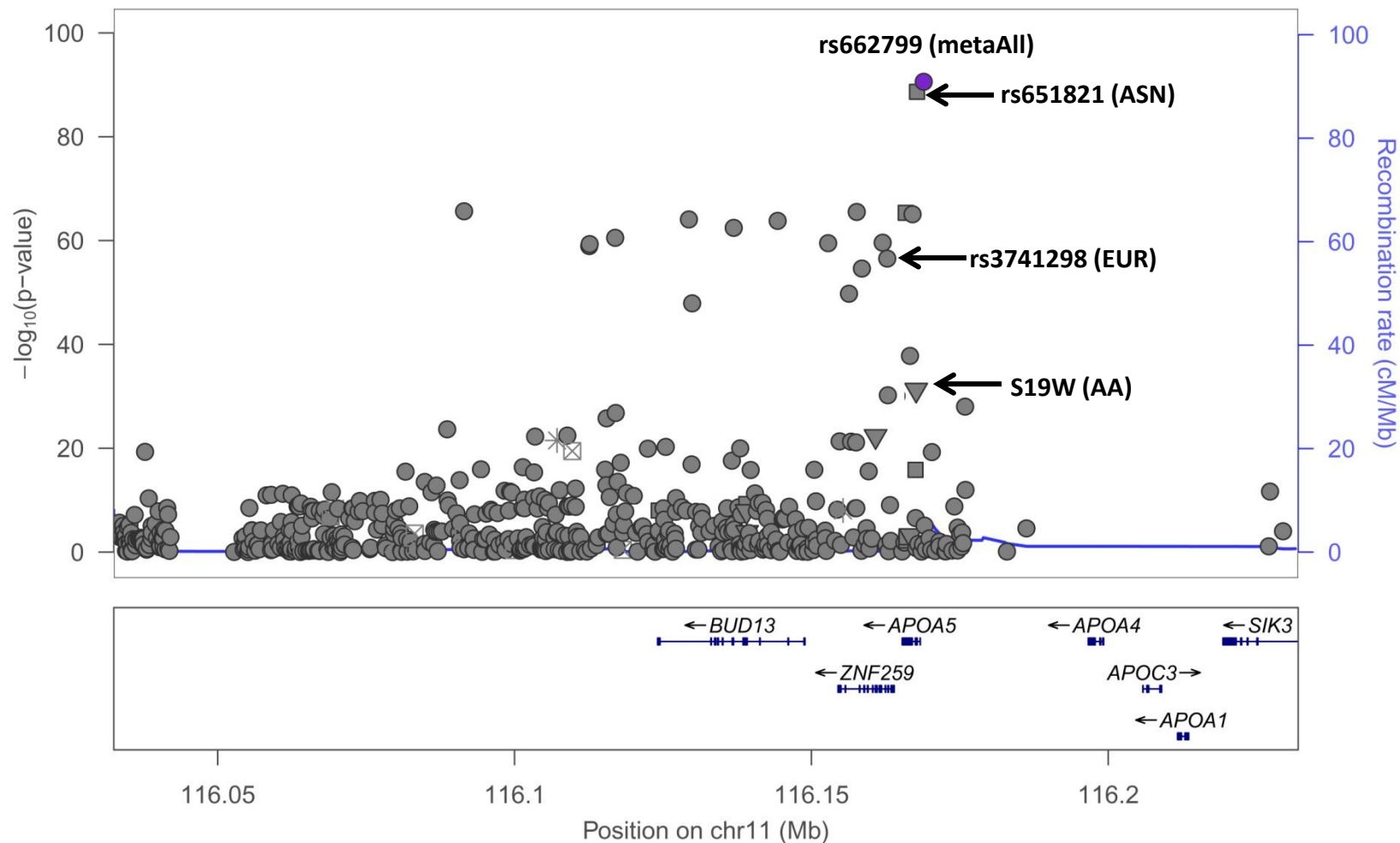
FigureS6A. Association at TG locus *APOA5* in Europeans. Index SNP rs3741298 is the variant showing the strongest evidence of association in Europeans. The SNPs with the smallest P values in East Asians (-3A>G), African Americans (S19W) and the trans-ethnic meta-analysis (rs662799) are also indicated.



FigureS6B. Association at TG locus *APOA5* in East Asians. Index SNP -3A>G is the variant showing the strongest evidence of association in East Asians. The SNPs with the smallest P values in Europeans (rs3741298), African Americans (S19W) and the trans-ethnic meta-analysis (rs662799) are also indicated.



FigureS6C. Association at TG locus *APOA5* in African Americans. Index SNP S19W is the variant showing the strongest evidence of association in African Americans. The SNPs with the smallest P values in Europeans (rs3741298), East Asians (-3A>G) and the trans-ethnic meta-analysis (rs662799) are also indicated.



FigureS6D. Association at TG locus *APOA5* in the trans-ethnic meta-analysis. Index SNP rs662799 is the variant showing the strongest evidence of association in trans-ethnic meta-analysis. The SNPs with the smallest P values in Europeans (rs3741298), East Asians (-3A>G) and African Americans (S19W) are also indicated.