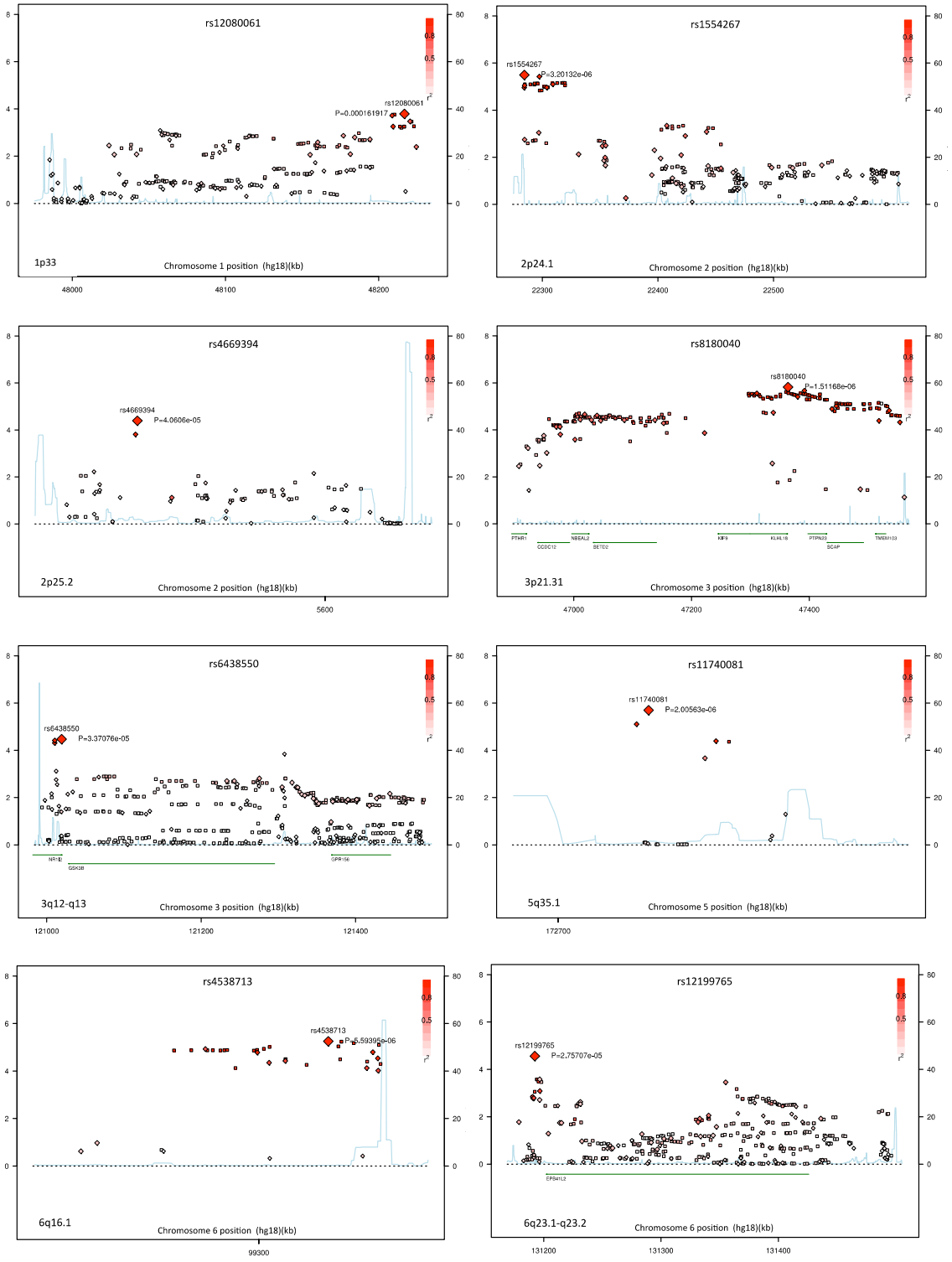
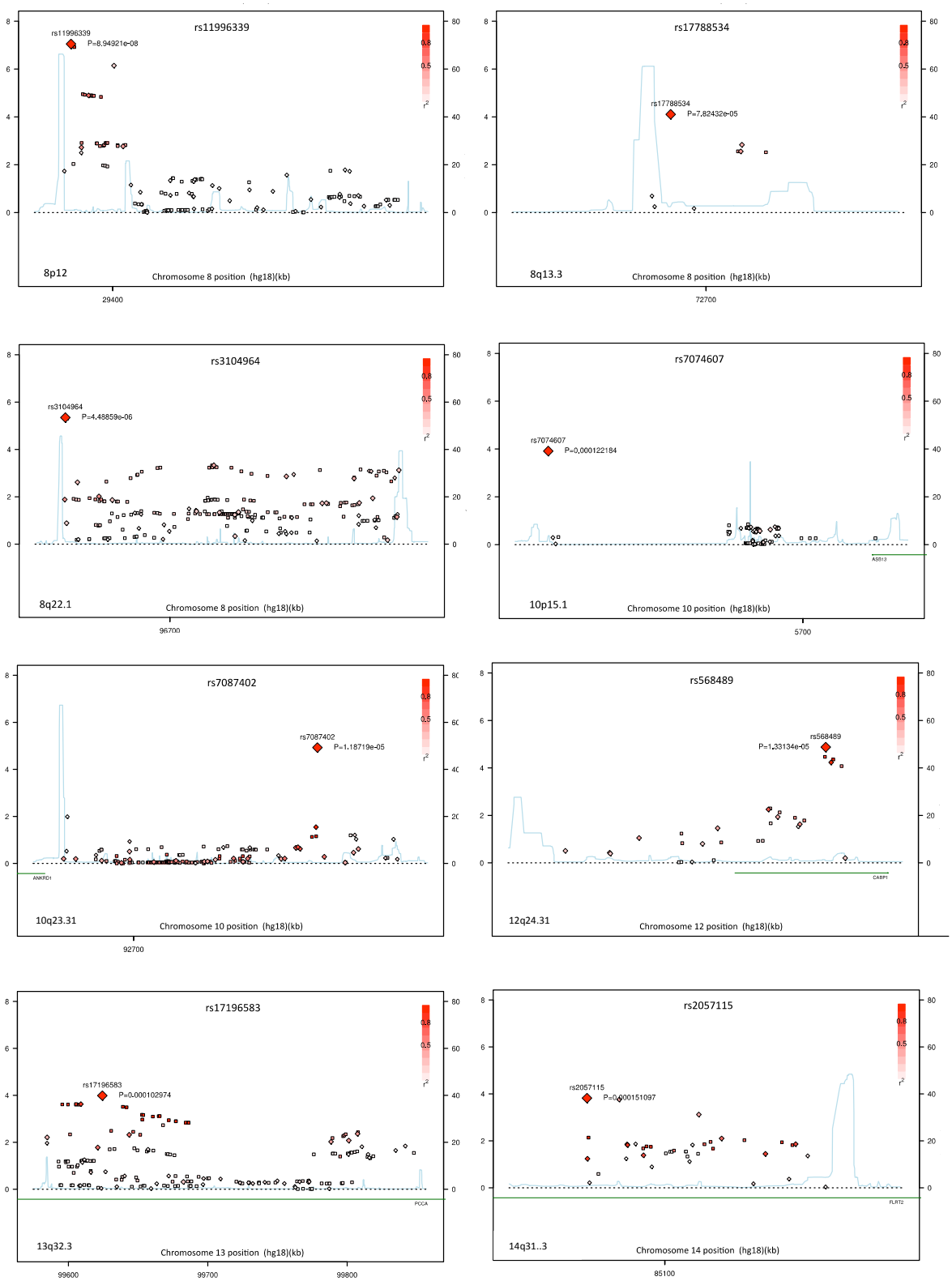
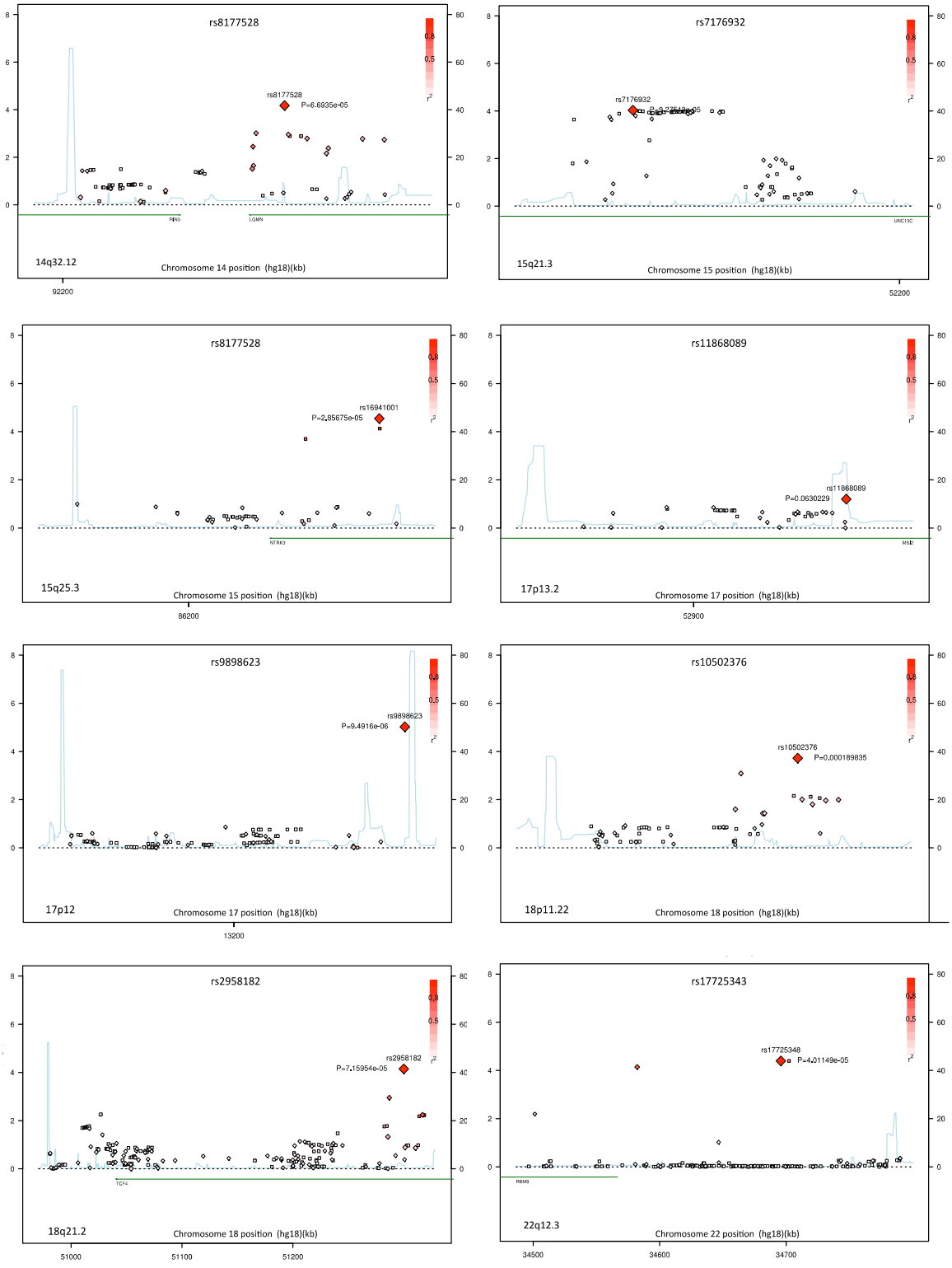
Supplementary Figure 2. Imputation plots for the 24 loci associated with CRC in EPICOLON. P-value plots for the imputed markers in the associated regions. Diamonds represent typed SNPs, squares depict imputed markers; the biggest diamond is the best-associated SNP in the region, irrespective of typed/imputed status; red grading represents LD relationships. X axis: Chromosome location; Y axis: observed (-logP); Z axis: Recombination rate (cM/Mb).



Supplementary Figure 2. (Continuation I).



Supplementary Figure 2. (Continuation II).