**S5 Fig. Unconditional plots of the five lipid and lipoprotein subclass-associated loci.** Each circle represents a single variant. The color is based on LD ($r^2$) between each variant and the reference variant (purple diamond); X-axis, genomic (GRCh37/hg19) position in Mb; Left y-axis, p-value of variant-trait association in $-\log_{10}$; Right y-axis, local estimates of genomic recombination rate in cM/Mb, represented by blue vertical lines.