

Description of Additional Supplementary Files

Supplementary Data 1: Table listing top 40 SNPs within the PPARG locus with predicted the highest positive or negative Enformer effect on the *PPARG2* signal.

Supplementary Data 2: Table listing top 40 SNPs within the PPARG locus with predicted the highest positive or negative Enformer effect on enhancer accessibility.

Supplementary Data 3: Table listing fine-mapped phenotype association of top variants within the PPARG locus with predicted highest positive or negative Enformer effect on the PPARG2 signal/enhancer accessibility.

Supplementary Data 4: Table listing QTL colocalization of top variants within the PPARG locus with predicted highest positive or negative Enformer effect on the PPARG2 signal/enhancer accessibility.

