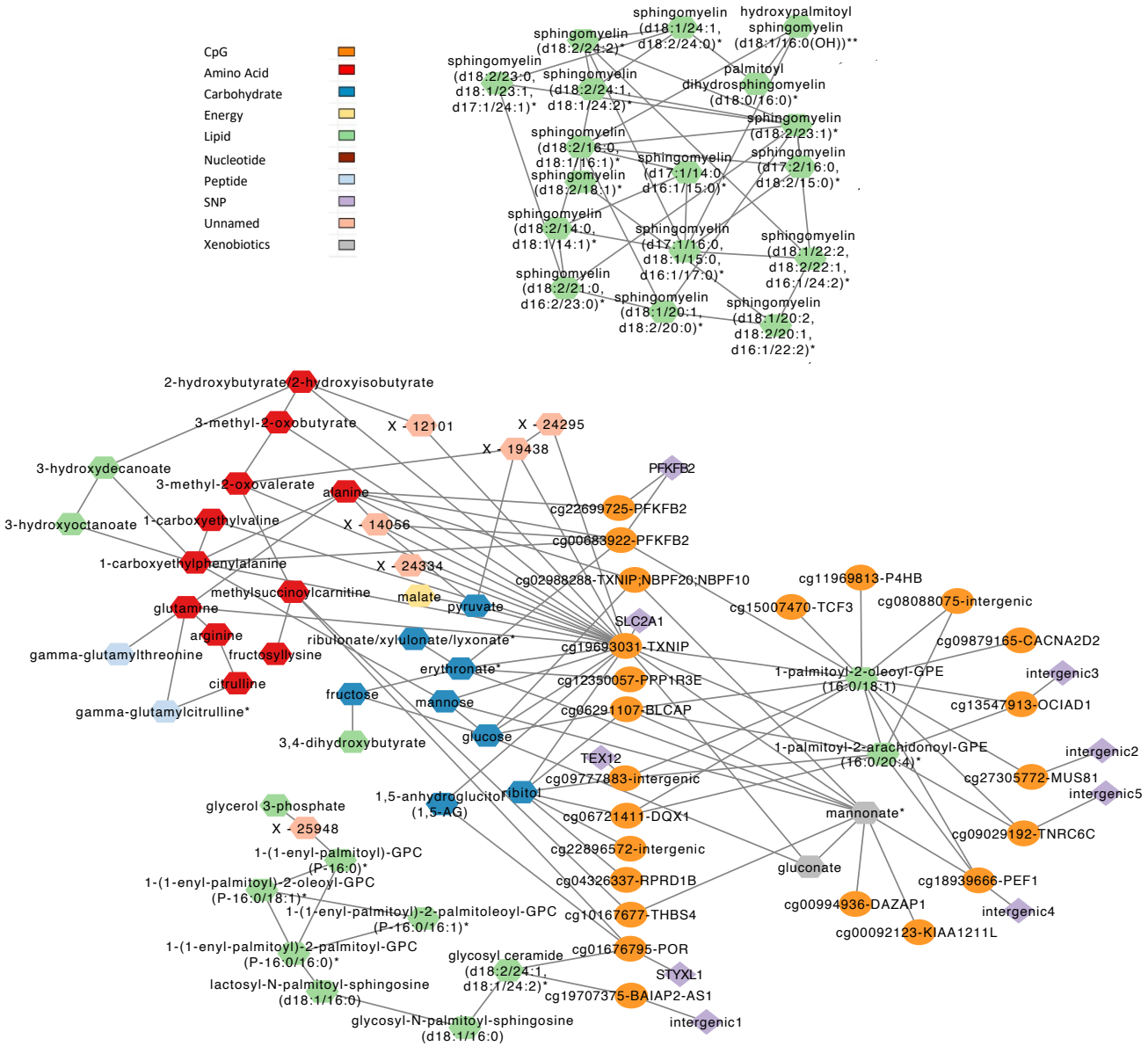

Figure S1: Full multi-omics network of T2D methylated genes associations with metabolites, and their meQTLs integrated with metabolic networks. Oval nodes are CpGs, diamonds are SNPs, hexagons are metabolites.