

Figure S1. Bioinformatic analysis of DEMIs (a) The Heatmap of DEMIs in the low and high IMF groups; (b) GO enrichment of DEMIs.

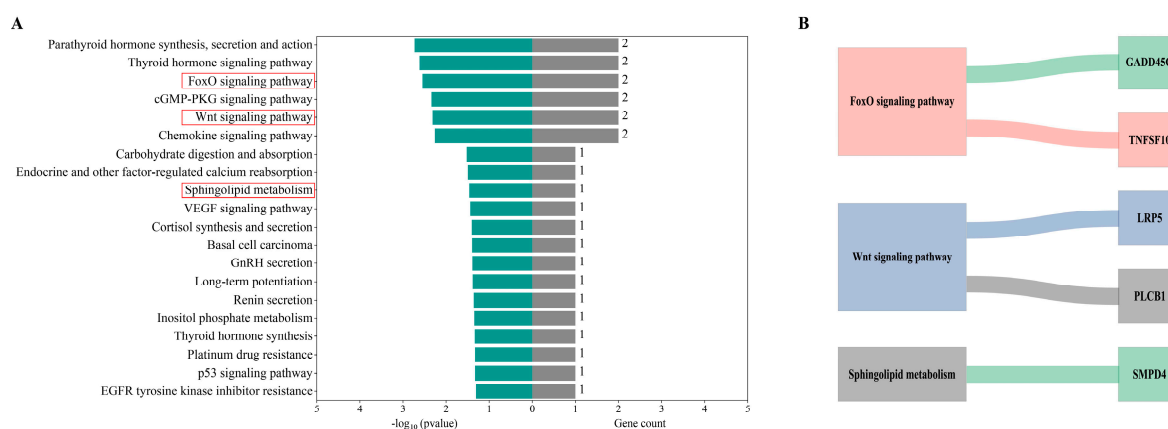


Figure S2. KEGG annotation analysis showing differentially expressed genes. (A) KEGG pathway annotation analysis. (B) Differentially expressed genes involved in fat associated pathways.