

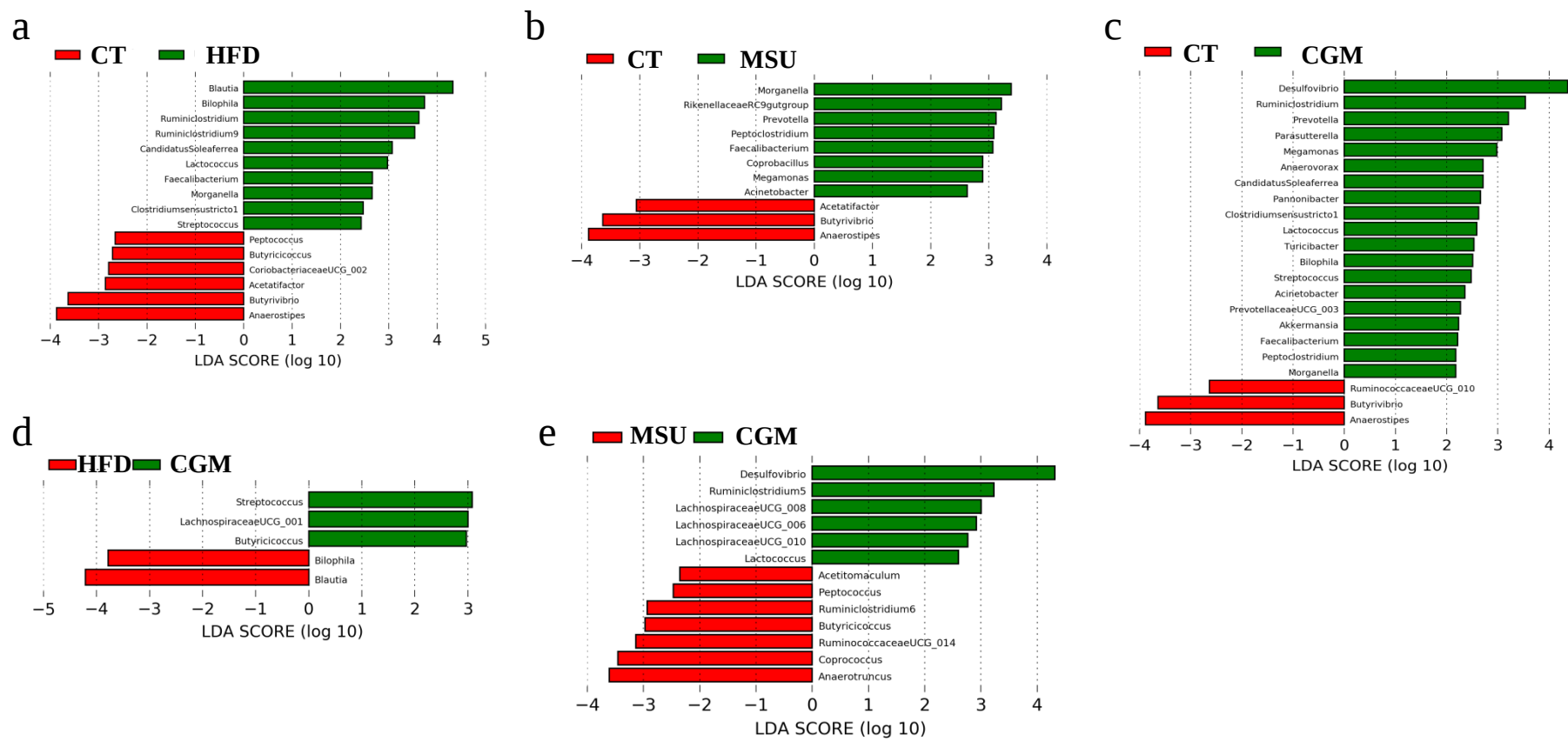


Fig. S1 Differentially expressed genera with the LDA scores >2.0 and adjusted p values <0.05 between CT and HFD (a), between CT and MSU (b), between CT and CGM (c), between HFD and MSU (d) and between CGM and CGM (e). Differences are represented by the color of over-represented genera. CT: control group; HFD: hyperuricaemia model; MSU: acute gout model; CGM: compound gout model.

PURINE METABOLISM

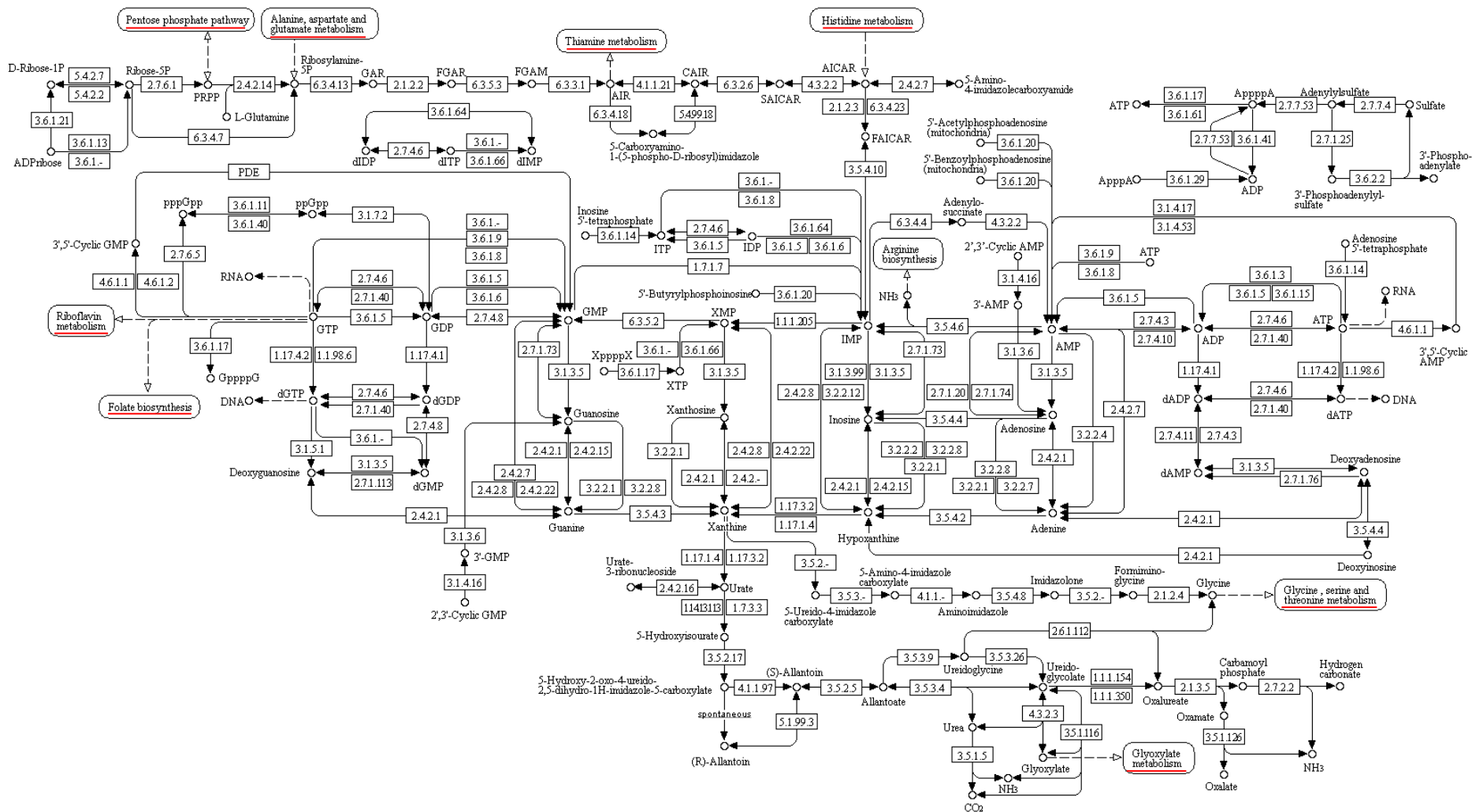


Fig. S2 Purine metabolism pathways in KEGG analysis