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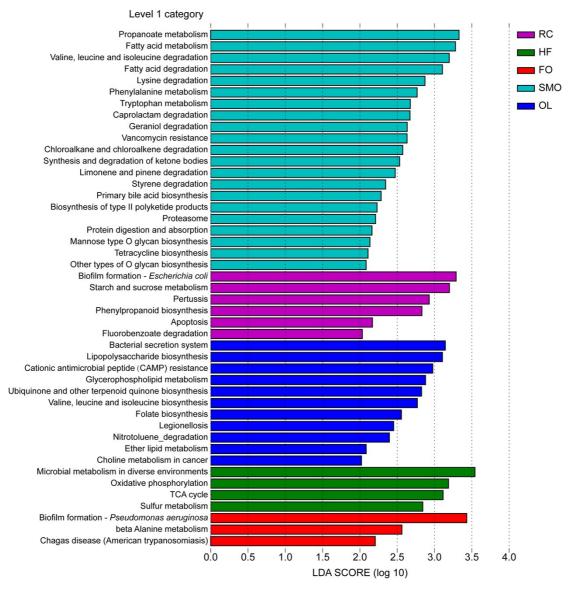


Fig. S4 LDA analysis revealing the differentially enriched bacterial community functional pathways predicted by Tax4Fun2 among the five groups.