

SUPPORTING INFORMATION FOR

Gut microbiota-derived metabolite signature in suckling and weaned piglets

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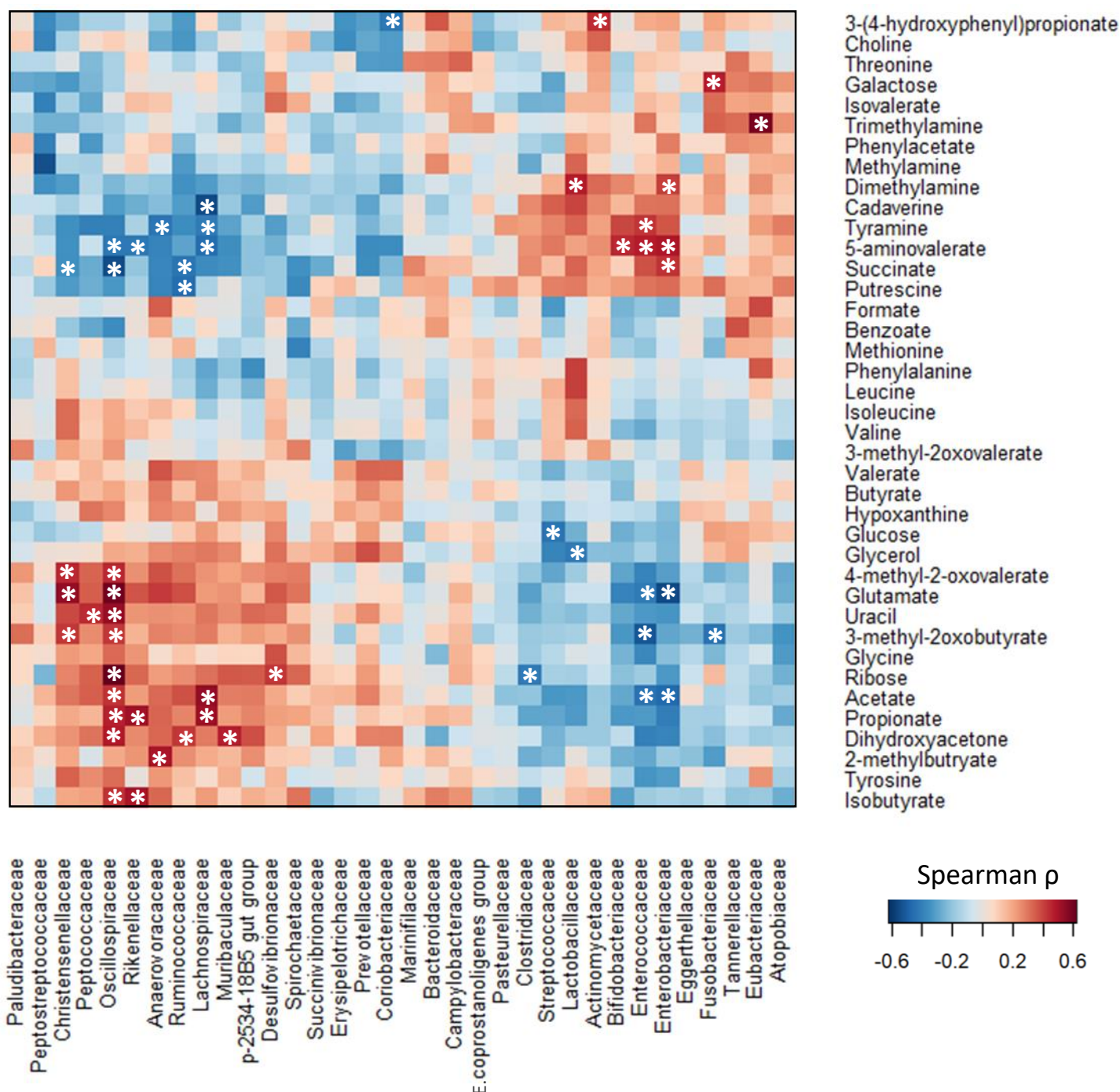


Figure S1: Correlation analysis. The full dataset (n=56 piglet fecal samples) was used for correlation analysis. The heatmap represents the Spearman correlation coefficient between the relative concentrations of metabolites (rows) and the relative abundances of bacterial families (columns). Rows and columns were clustered with the average method. Negative correlations are represented in blue and positive correlations in red. *: adjusted P-value < 0.05.