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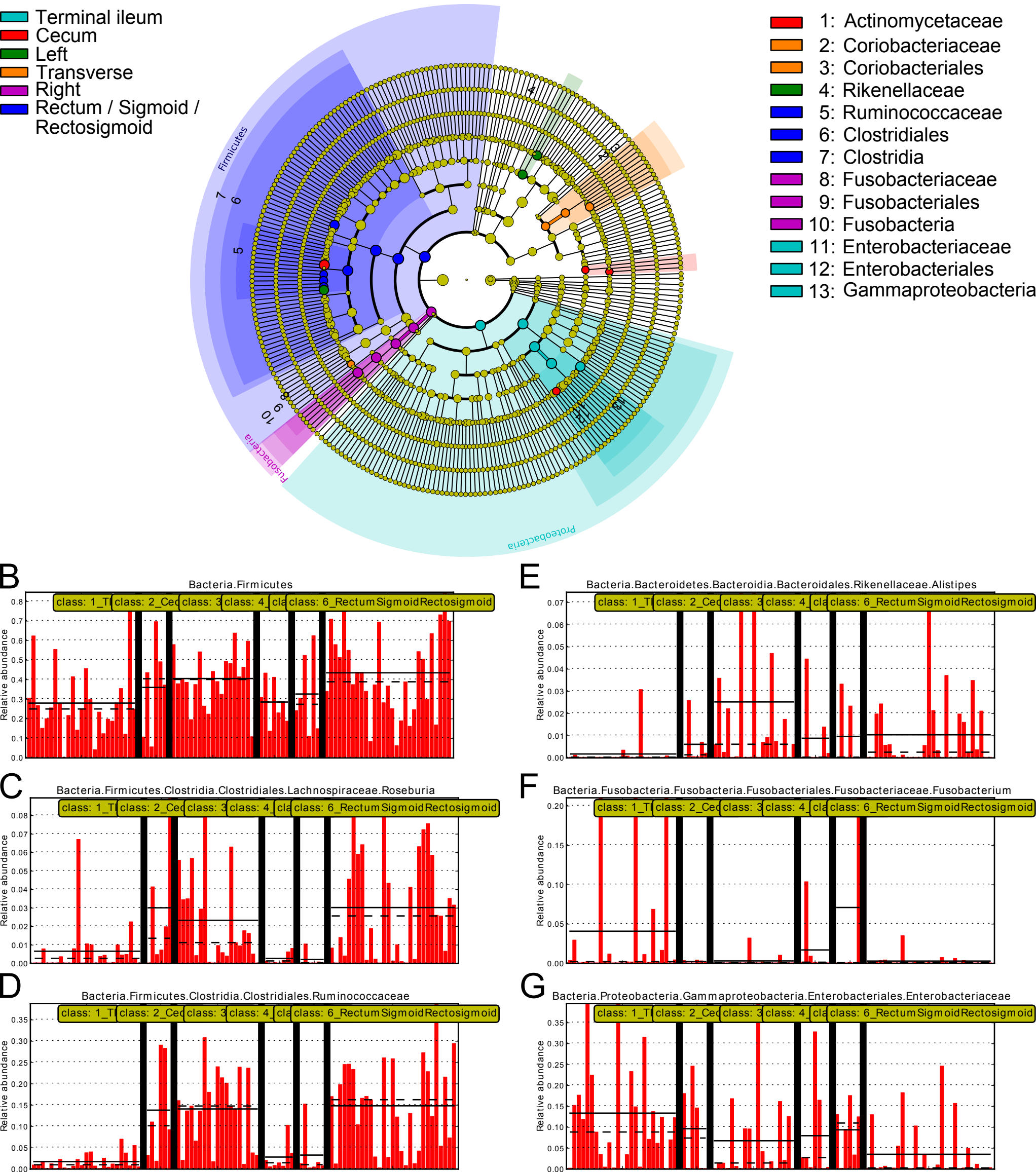
Univariate analysis for associations between gut biogeography and microbial taxa