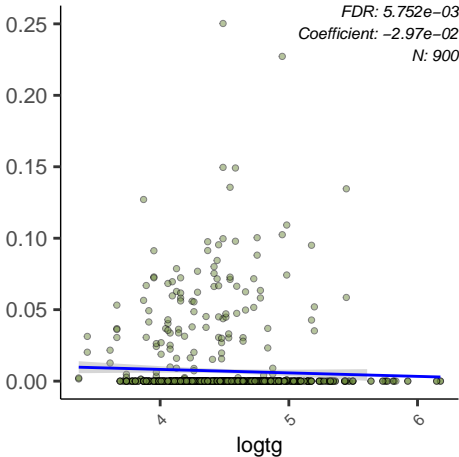
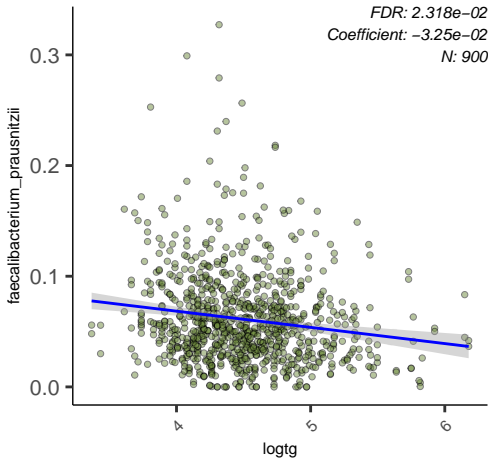
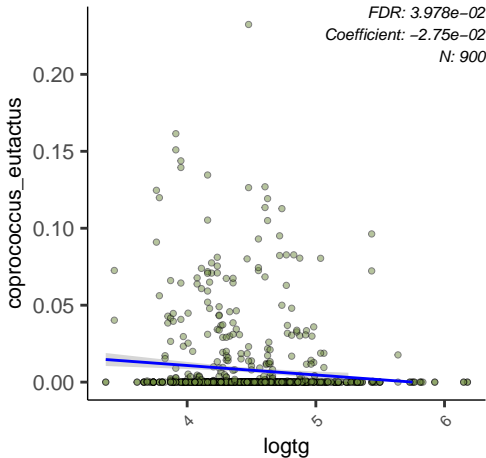


eubacterium_sp_cag_180







clostridium_sp_cag_167

0.06

0.04

0.02

0.00

4

5

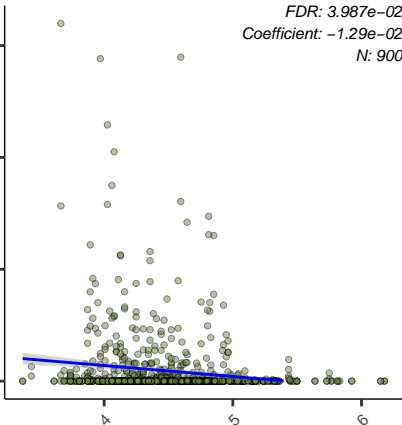
6

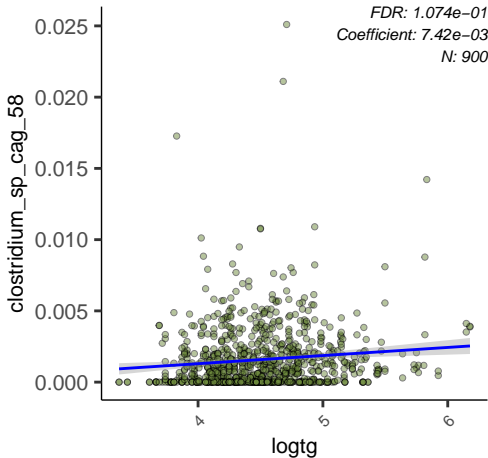
logtg

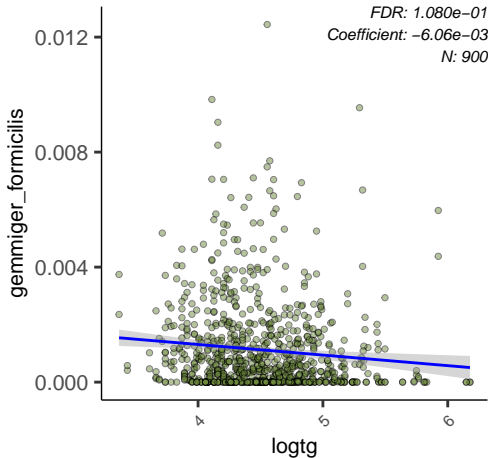
FDR: $3.987e-02$

Coefficient: $-1.29e-02$

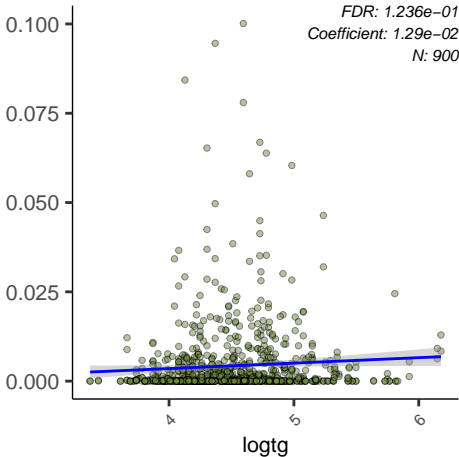
N: 900

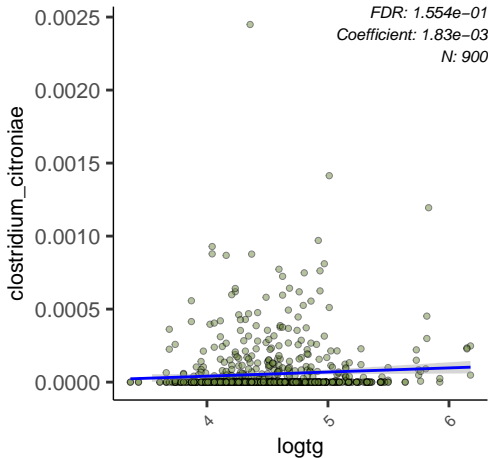


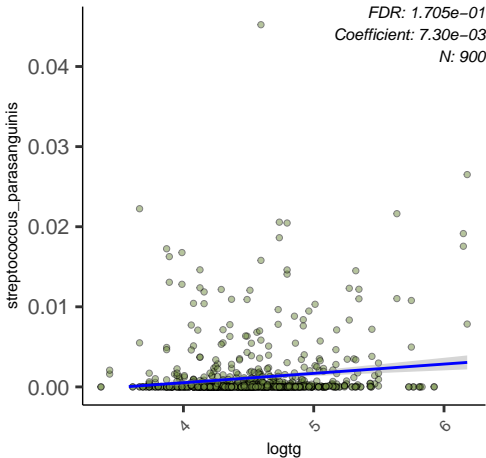


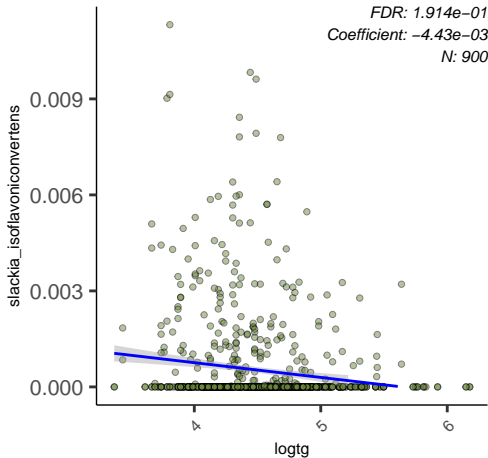


bacteroides_xylanisolvens









oscillibacter_sp_cag_241

0.06

0.04

0.02

0.00

4

5

6

logtg

FDR: 1.941e-01

Coefficient: -1.04e-02

N: 900

