

eubacterium_sp_cag_180

0.25
0.20
0.15
0.10
0.05
0.00

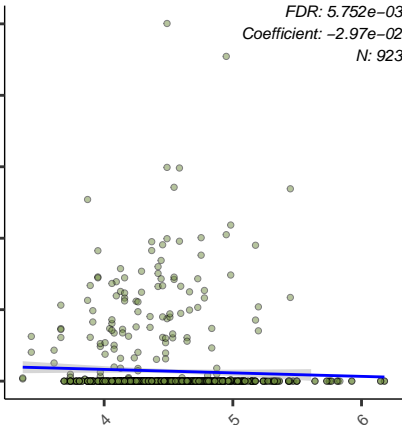
4

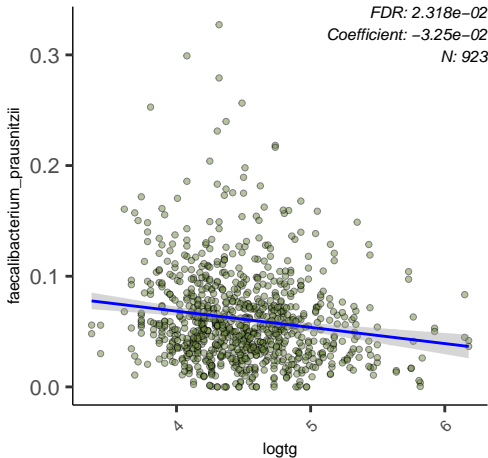
5

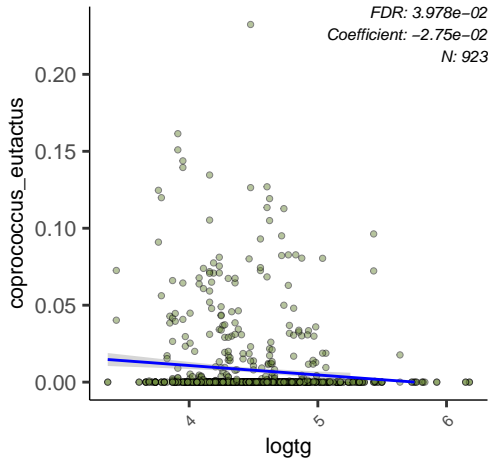
6

logtg

FDR: 5.752e-03
Coefficient: -2.97e-02
N: 923







clostridium_sp_cag_167

0.06
0.04
0.02
0.00

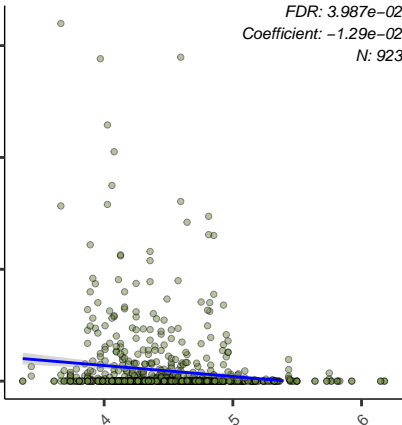
4

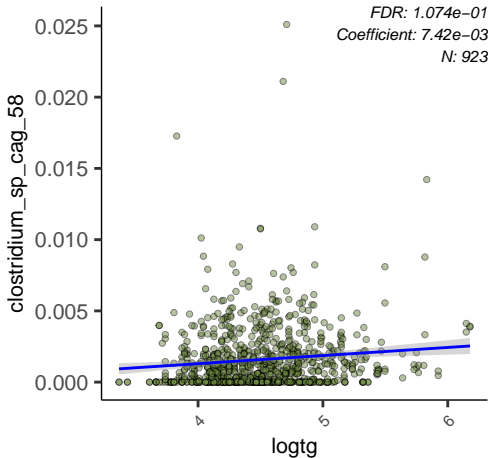
5

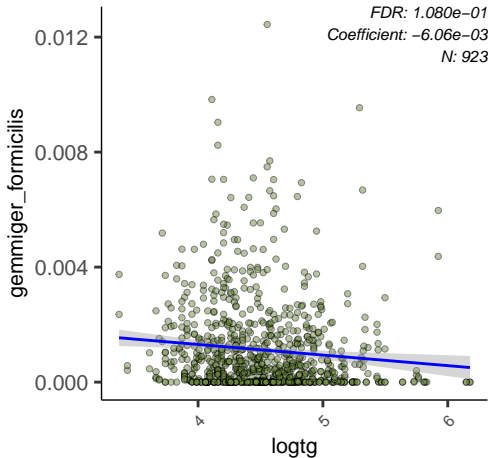
6

logtg

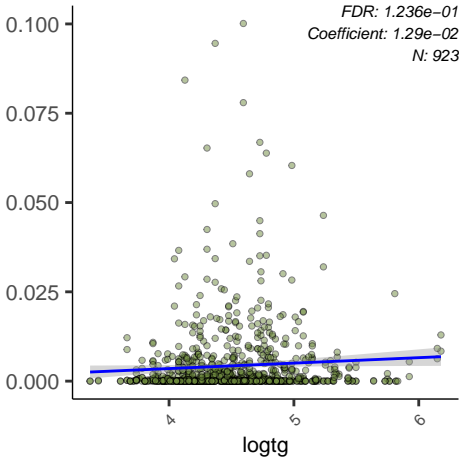
FDR: 3.987e-02
Coefficient: -1.29e-02
N: 923

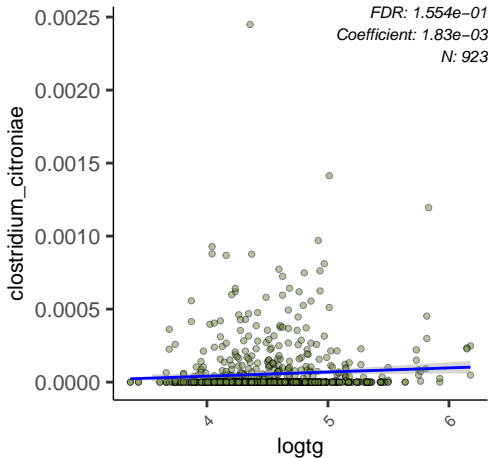


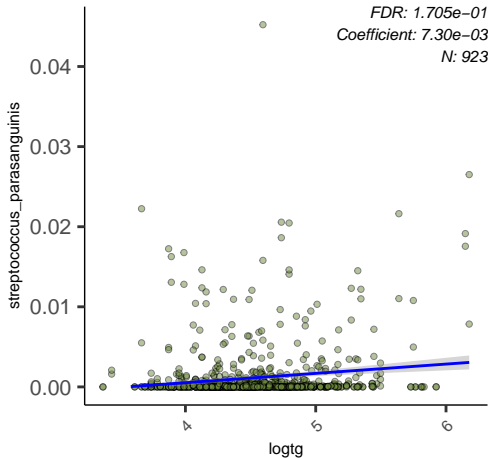


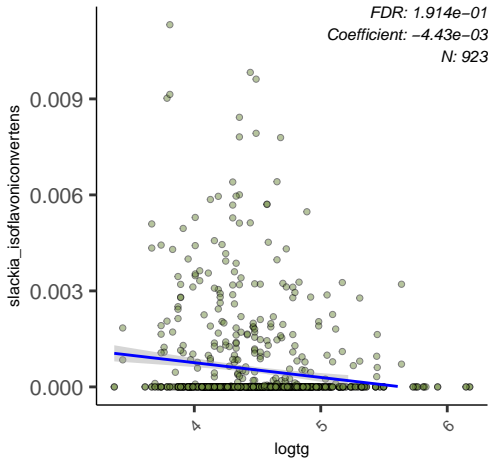


bacteroides_xylanisolvens









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: 923

