

eubacterium\_sp\_cag\_180

0.25  
0.20  
0.15  
0.10  
0.05  
0.00

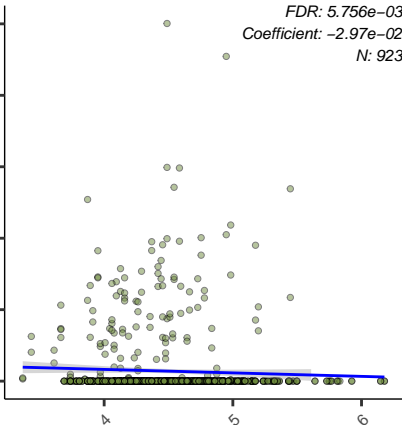
4

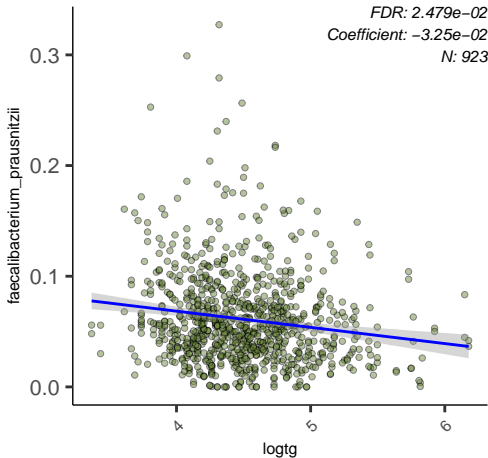
5

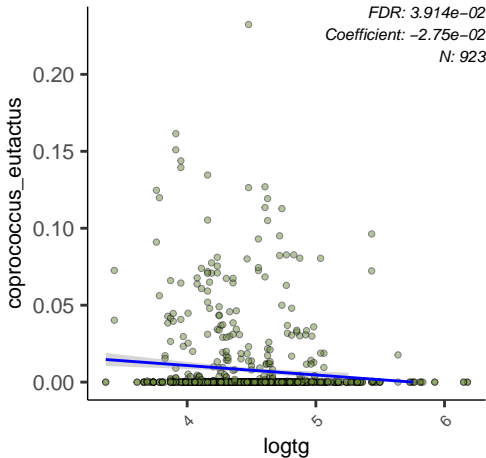
6

logtg

*FDR: 5.756e-03*  
*Coefficient: -2.97e-02*  
*N: 923*







clostridium\_sp\_cag\_167

0.06

0.04

0.02

0.00

4

5

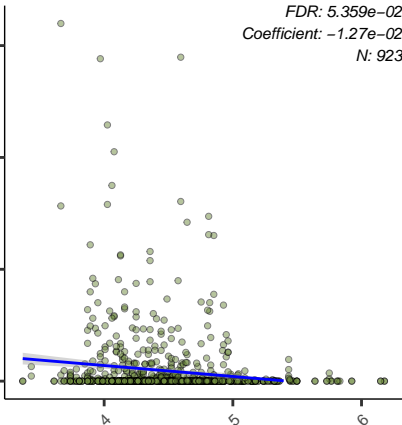
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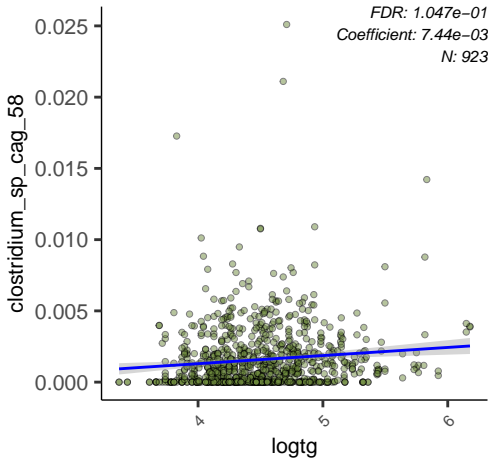
logtg

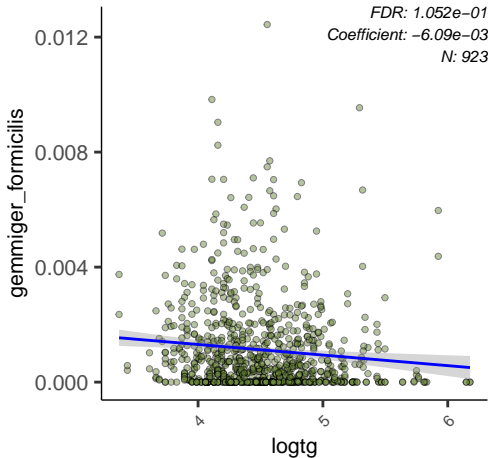
*FDR: 5.359e-02*

*Coefficient: -1.27e-02*

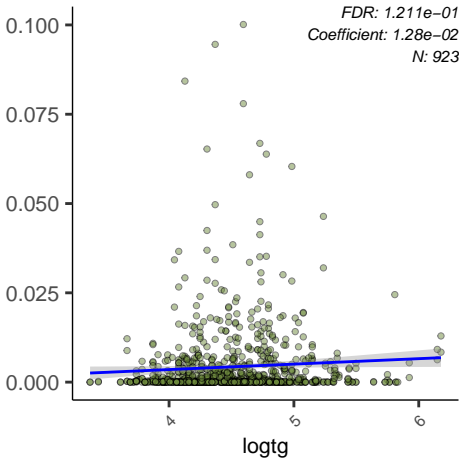
*N: 923*

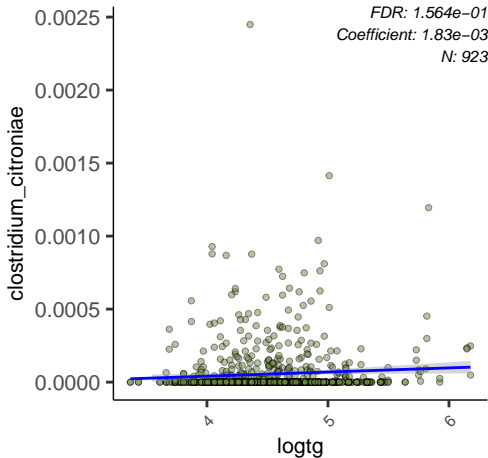




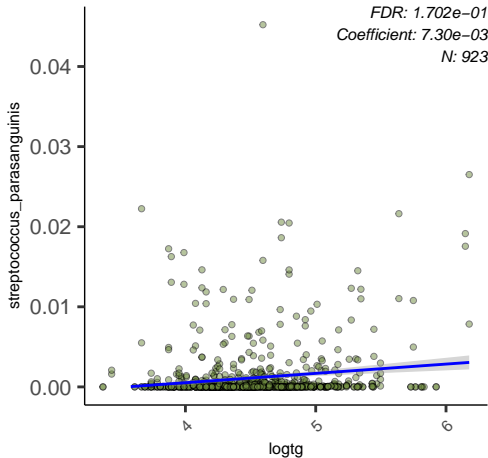


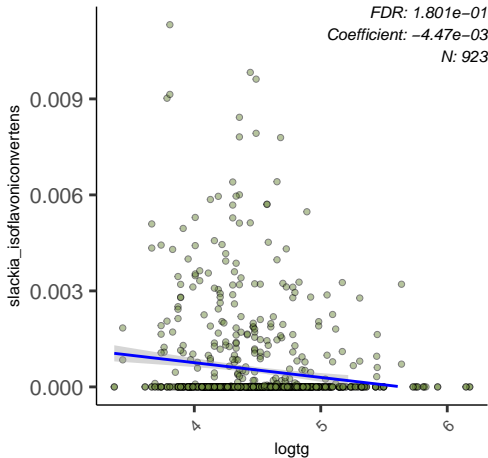
bacteroides\_xylanisolvens











oscillibacter\_sp\_cag\_241

0.06

0.04

0.02

0.00

4

5

6

logtg

*FDR: 1.976e-01*

*Coefficient: -1.03e-02*

*N: 923*

