

eubacterium_sp_cag_180

0.25
0.20
0.15
0.10
0.05
0.00

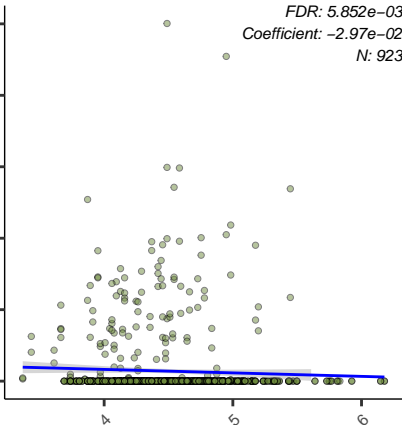
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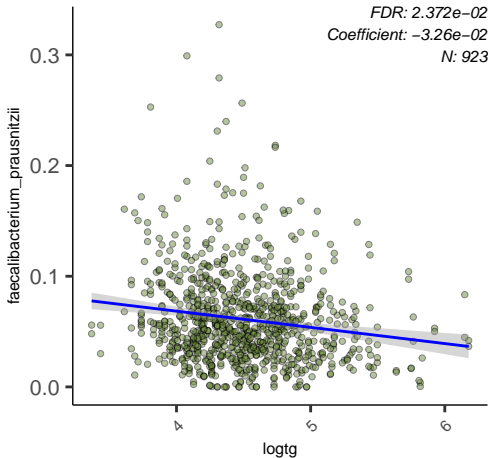
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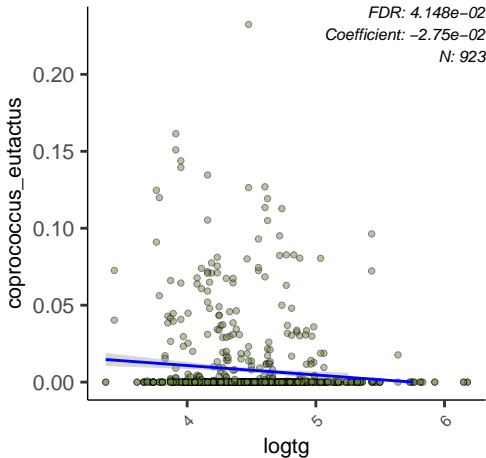
6

logtg

FDR: 5.852e-03
Coefficient: -2.97e-02
N: 923







clostridium_sp_cag_167

0.06

0.04

0.02

0.00

4

5

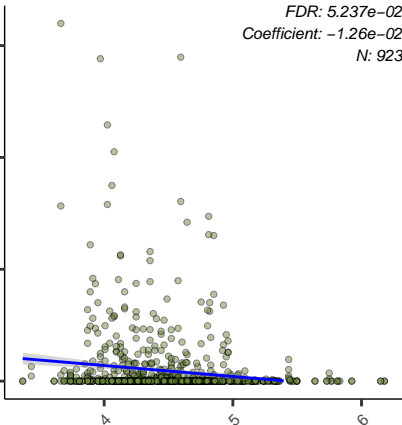
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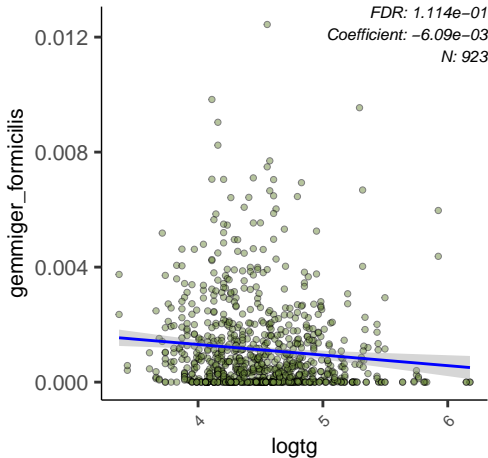
logtg

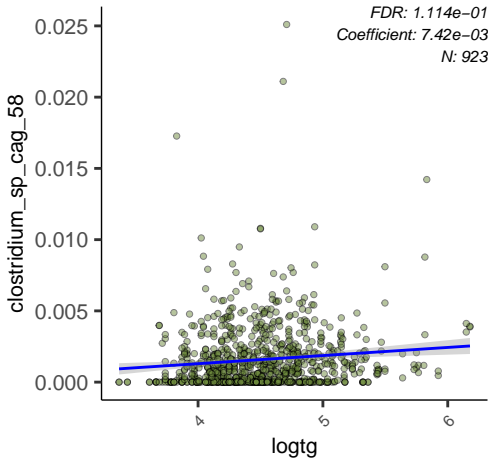
FDR: 5.237e-02

Coefficient: -1.26e-02

N: 923







bacteroides_xylanisolvens

0.100
0.075
0.050
0.025
0.000

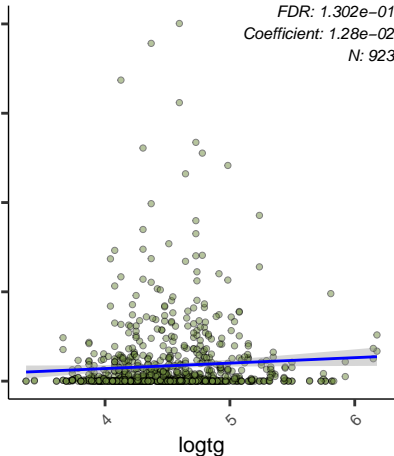
FDR: 1.302e-01
Coefficient: 1.28e-02
N: 923

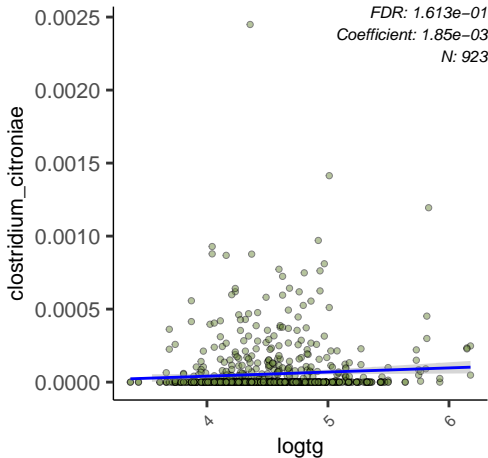
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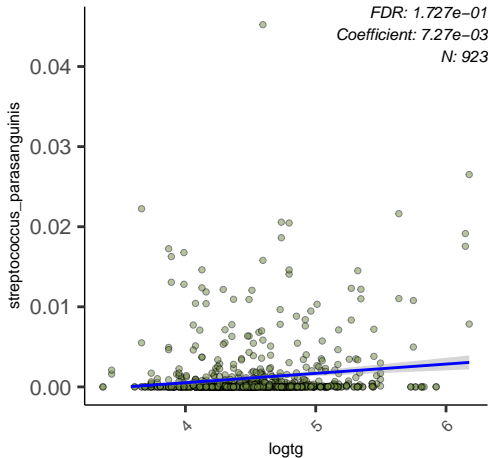
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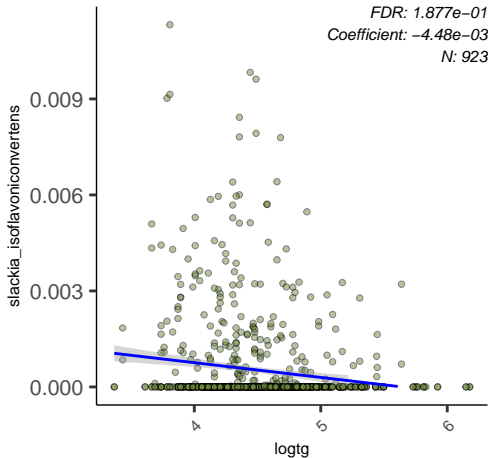
6

logtg









oscillibacter_sp_cag_241

0.06
0.04
0.02
0.00

4

5

6

logtg

FDR: 2.071e-01
Coefficient: -1.04e-02
N: 923

