

eubacterium_sp_cag_251

FDR: 1.880e-03

Coefficient: -5.95e-03

N: 775

0.08
0.06
0.04
0.02
0.00

logcrp

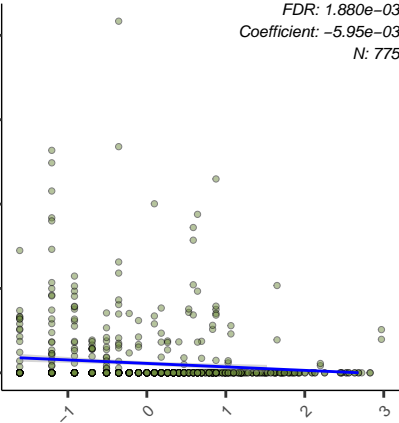
-1

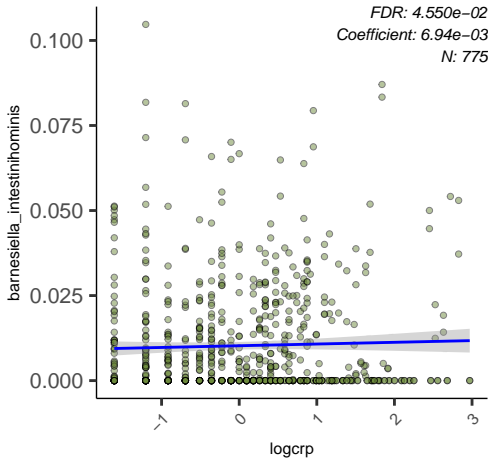
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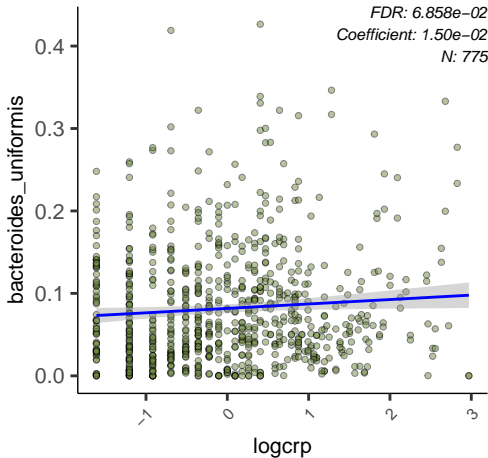
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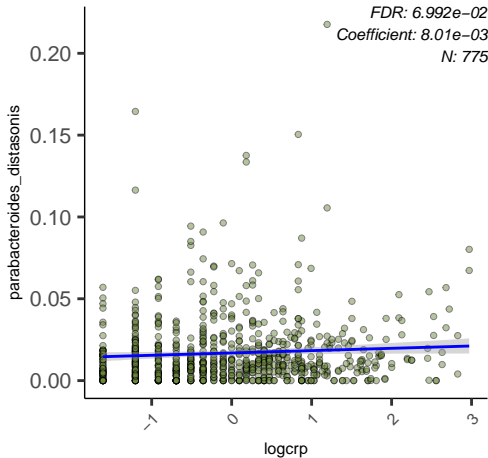
2

3









butyrivibrio_sp_cag_318

FDR: $7.016e-02$
Coefficient: $-6.47e-03$
N: 775

0.2

0.1

0.0

logcrp

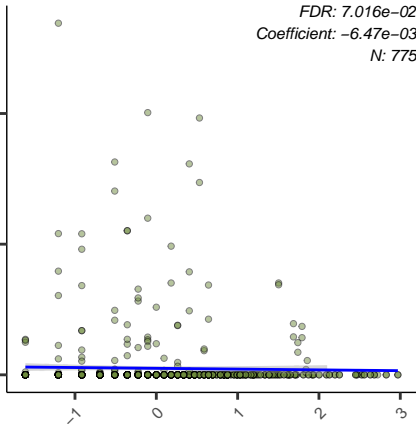
-1

0

1

2

3

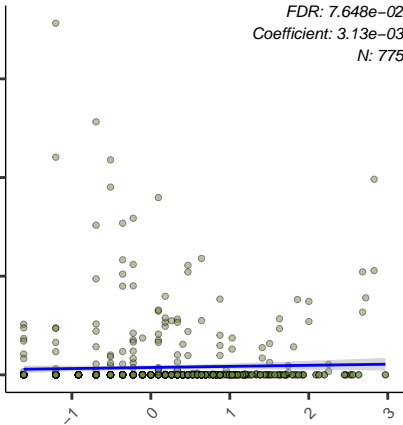


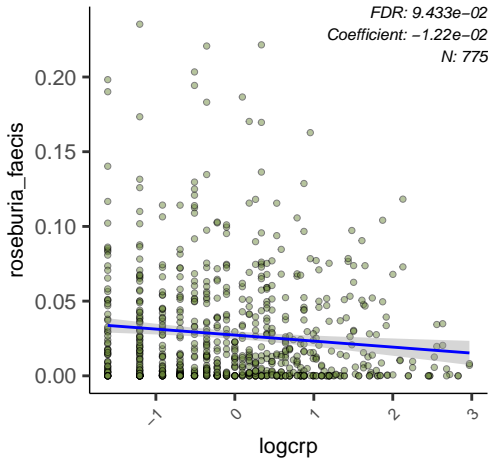
parabacteroides_johnsonii

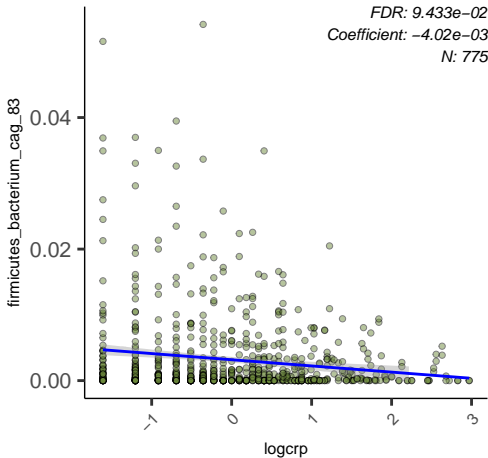
FDR: 7.648e-02
Coefficient: 3.13e-03
N: 775

0.06
0.04
0.02
0.00

logcrp





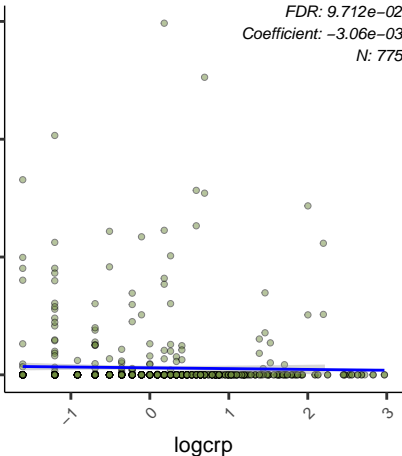


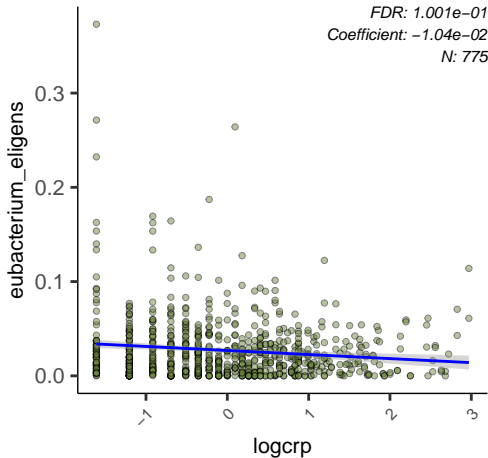
bifidobacterium_bifidum

0.06
0.04
0.02
0.00

FDR: 9.712e-02
Coefficient: -3.06e-03
N: 775

logcrp





ruminococcus_torques

FDR: 1.215e-01
Coefficient: 5.61e-03
N: 775

0.15
0.10
0.05
0.00

logcrp

