

eubacterium_sp_cag_251

FDR: 1.880e-03

Coefficient: -5.95e-03

N: 923

0.08
0.06
0.04
0.02
0.00

logcrp

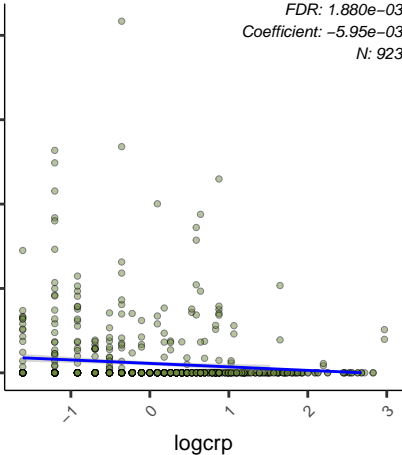
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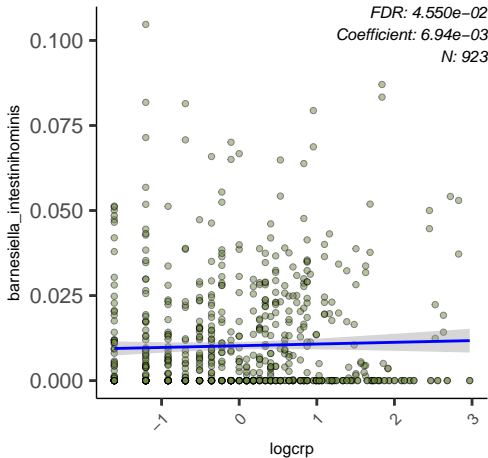
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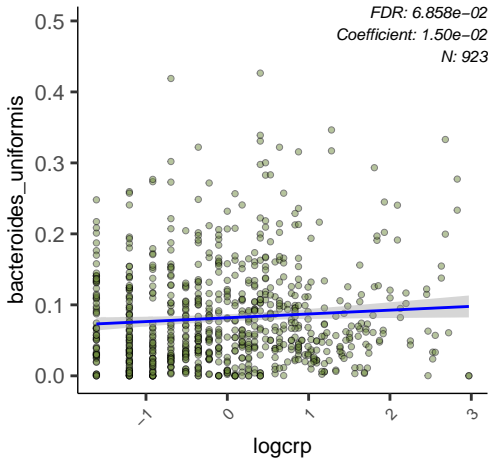
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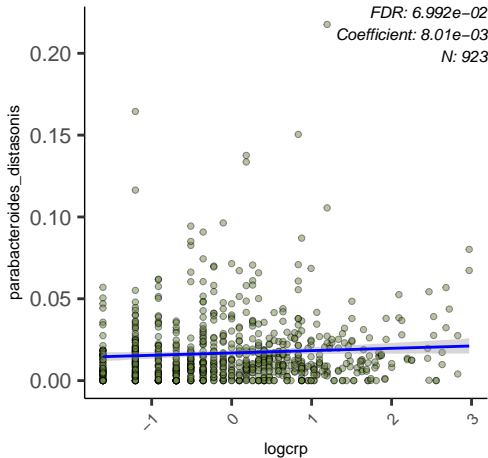
2

3









butyrivibrio_sp_cag_318

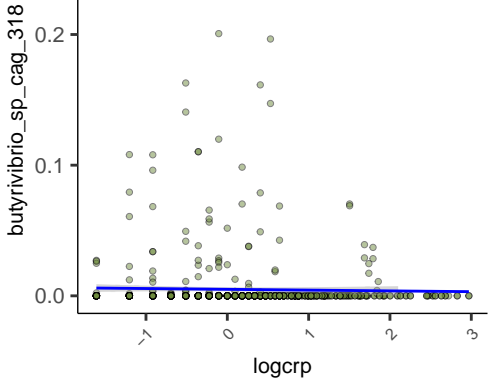
0.2

0.1

0.0

logcrp

FDR: 7.016e-02
Coefficient: -6.47e-03
N: 923



parabacteroides_johnsonii

FDR: 7.648e-02
Coefficient: 3.13e-03
N: 923

0.06
0.04
0.02
0.00

logcrp

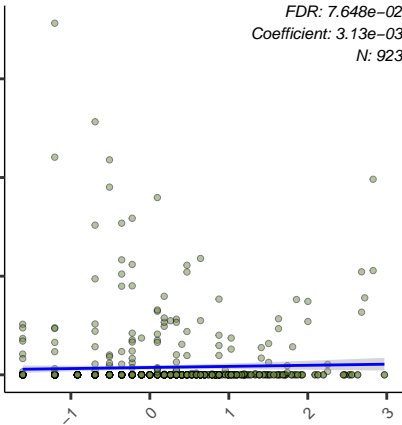
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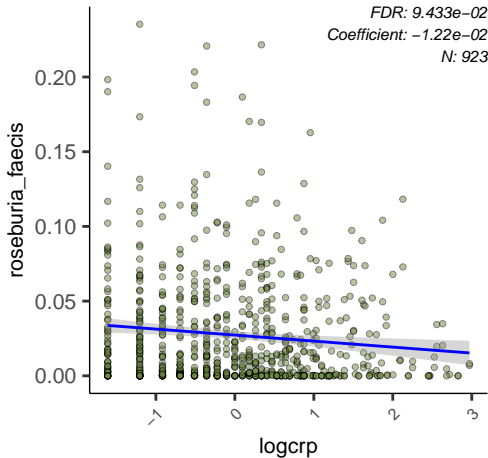
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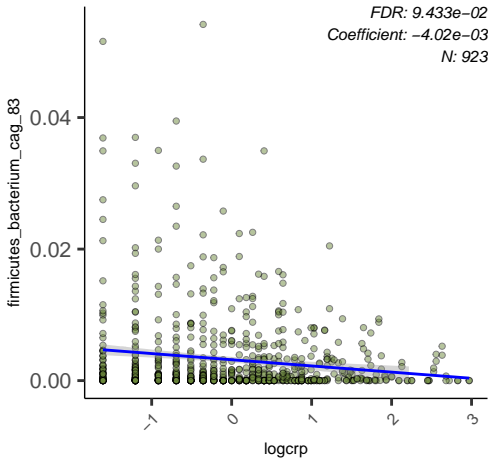
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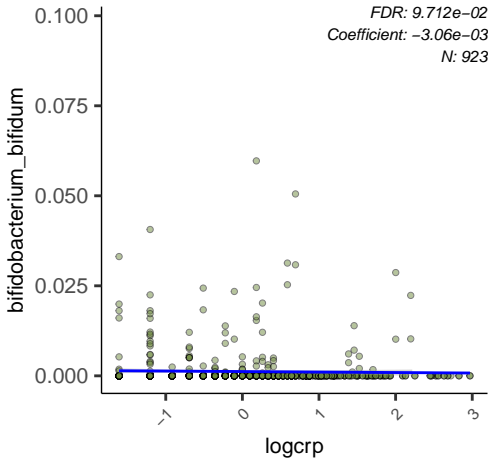
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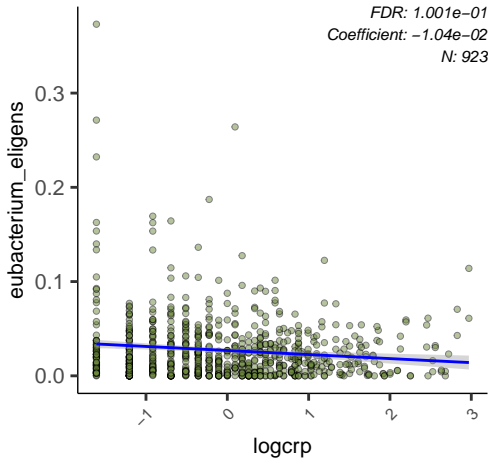
3











ruminococcus_torques

FDR: 1.215e-01
Coefficient: 5.61e-03
N: 923

0.15
0.10
0.05
0.00

logcrp

