

acidaminococcus_intestini

FDR: 2.076e-02

Coefficient: -6.59e-02

N: 925

0.05
0.04
0.03
0.02
0.01
0.00

0.00

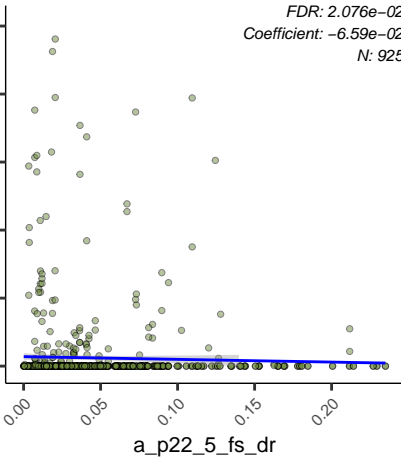
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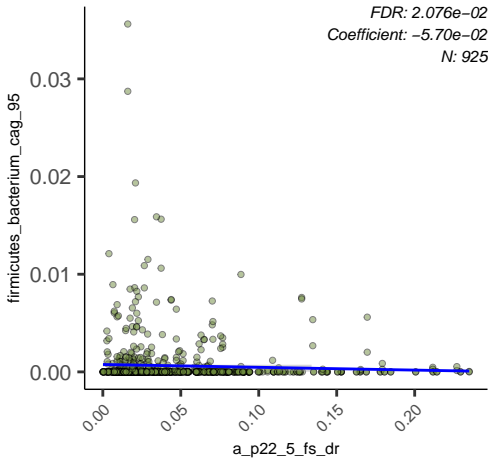
0.10

0.15

0.20

a_p22_5_fs_dr





butyrivibrio_sp_cag_318

FDR: 2.567e-02

Coefficient: -1.42e-01

N: 925

0.2

0.1

0.0

0.00

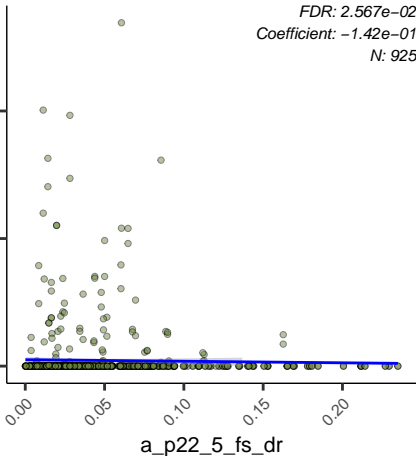
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0.10

0.15

0.20

a_p22_5_fs_dr

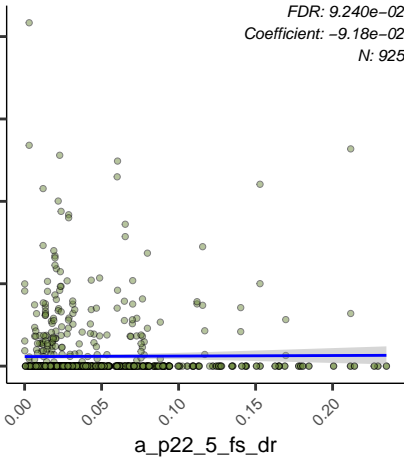


eubacterium_sp_cag_251

FDR: $9.240e-02$

Coefficient: $-9.18e-02$

N: 925



a_p22_5_fs_dr

parabacteroides_johnsonii

FDR: 9.240e-02
Coefficient: 6.39e-02
N: 925

0.06
0.04
0.02
0.00

0.00

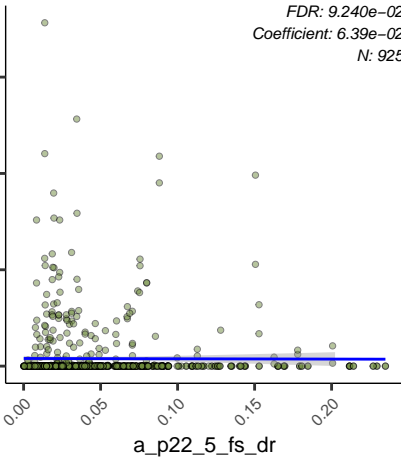
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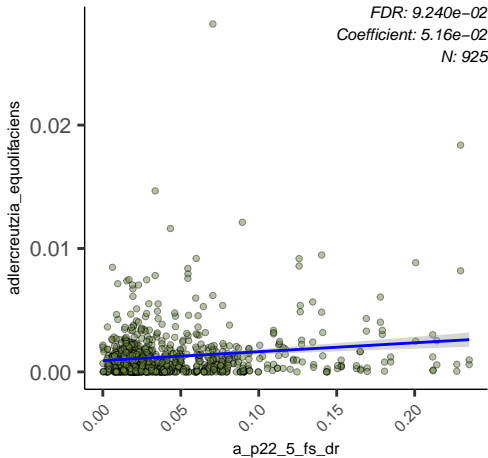
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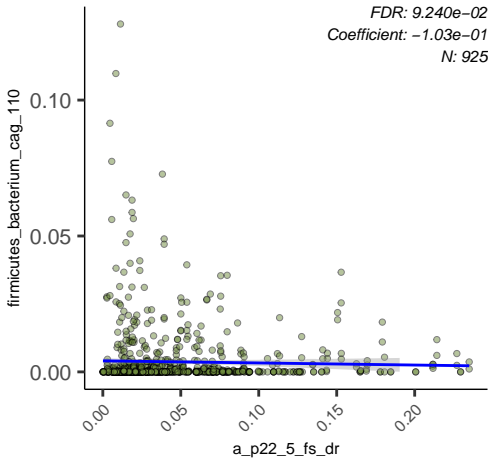
0.15

0.20

a_p22_5_fs_dr

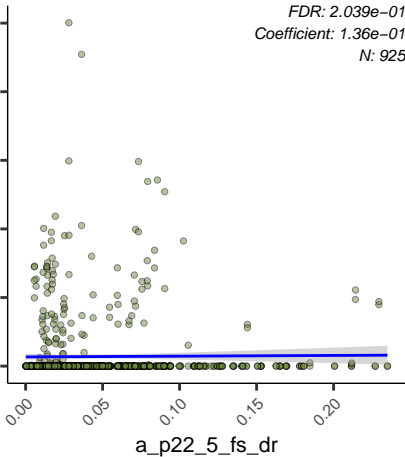






eubacterium_sp_cag_180

FDR: 2.039e-01
Coefficient: 1.36e-01
N: 925



enterorhabdus_caecimuris

FDR: 2.118e-01
Coefficient: 1.32e-02
N: 925

