

acidaminococcus_intestini

FDR: 2.305e-02

Coefficient: -6.87e-02

N: 923

0.05
0.04
0.03
0.02
0.01
0.00

0.00

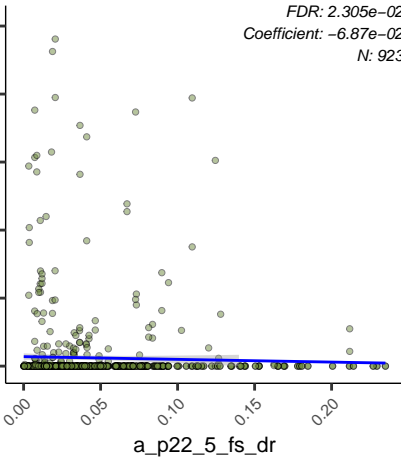
0.05

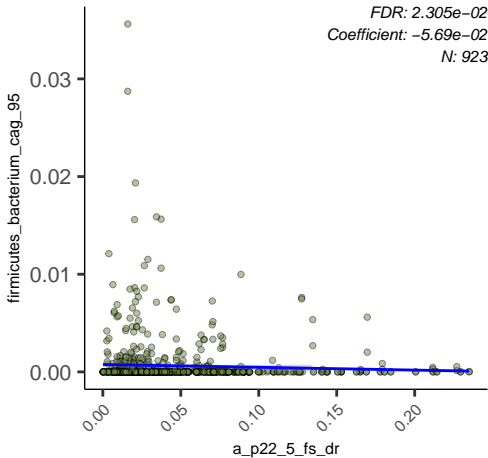
0.10

0.15

0.20

a_p22_5_fs_dr





butyrivibrio_sp_cag_318

FDR: 3.322e-02

Coefficient: -1.45e-01

N: 923

0.2

0.1

0.0

0.00

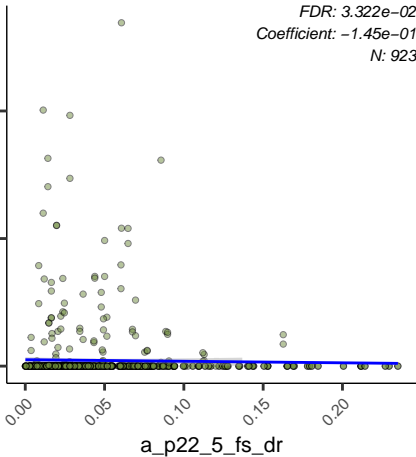
0.05

0.10

0.15

0.20

a_p22_5_fs_dr



parabacteroides_johnsonii

FDR: 6.849e-02
Coefficient: 6.72e-02
N: 923

0.06
0.04
0.02
0.00

0.00

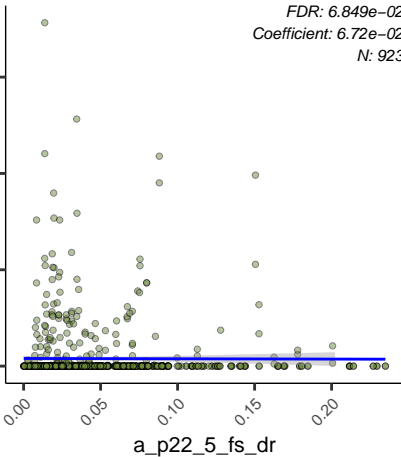
0.05

0.10

0.15

0.20

a_p22_5_fs_dr

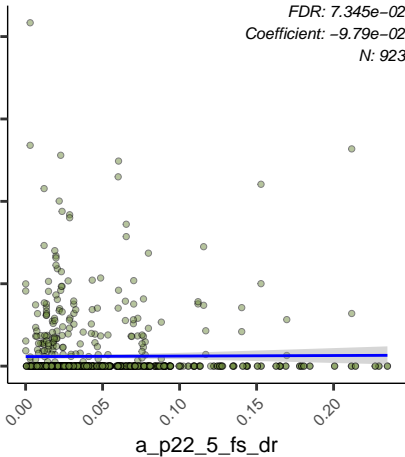


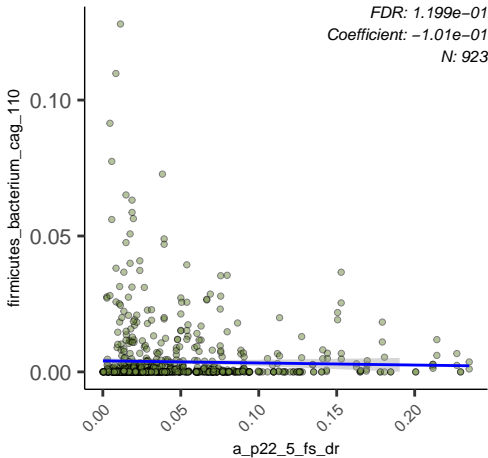
eubacterium_sp_cag_251

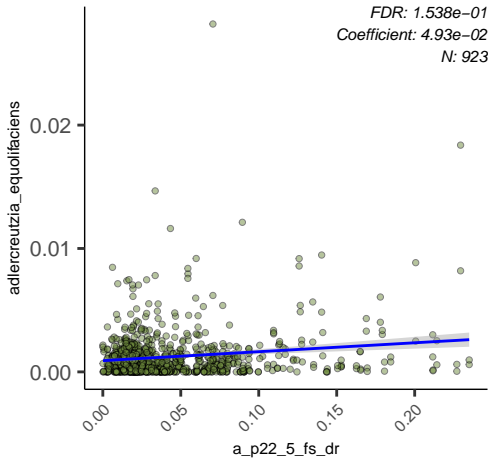
FDR: 7.345e-02

Coefficient: -9.79e-02

N: 923







eubacterium_sp_cag_180

FDR: 2.321e-01
Coefficient: 1.37e-01
N: 923

