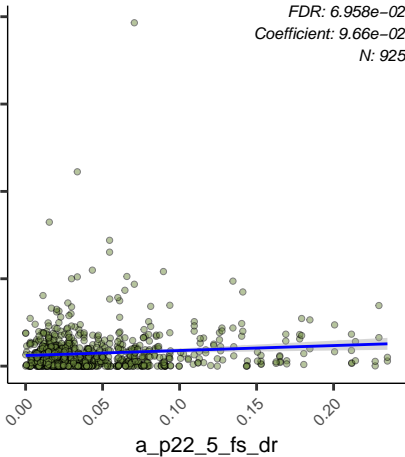
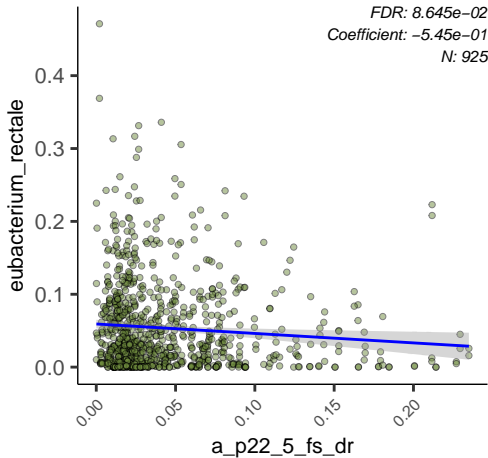


asaccharobacter_celatus

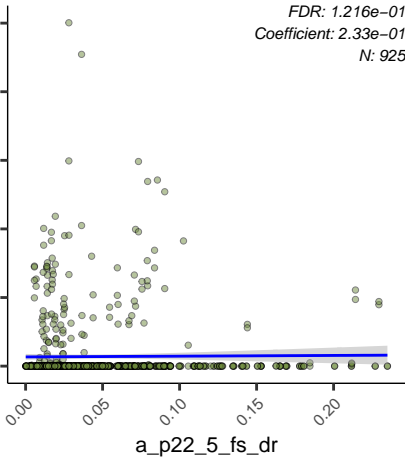
FDR: 6.958e-02
Coefficient: 9.66e-02
N: 925





eubacterium_sp_cag_180

FDR: 1.216e-01
Coefficient: 2.33e-01
N: 925



enterorhabdus_caecimuris

FDR: 1.216e-01
Coefficient: 2.32e-02
N: 925

0.008
0.006
0.004
0.002
0.000

0.00

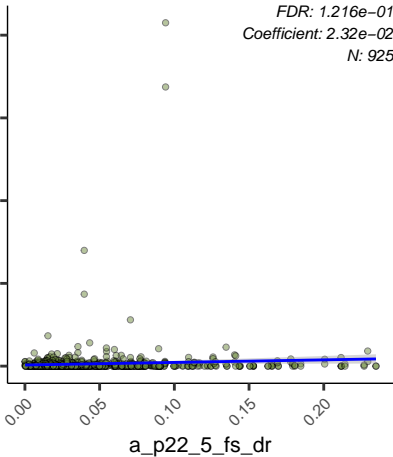
0.05

0.10

0.15

0.20

a_p22_5_fs_dr



bacteroides_salyersiae

FDR: 1.484e-01
Coefficient: 9.44e-02
N: 925

0.04
0.03
0.02
0.01
0.00

0.00

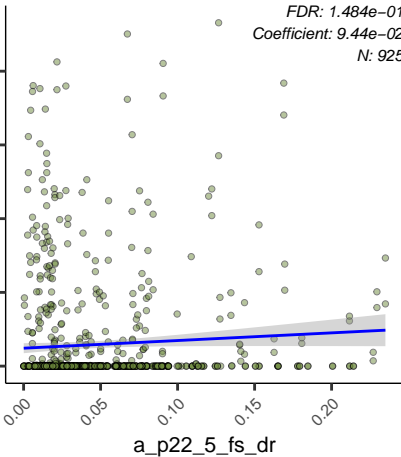
0.05

0.10

0.15

0.20

a_p22_5_fs_dr



holdemanella_biformis

FDR: 1.484e-01
Coefficient: 5.65e-02
N: 925

0.015
0.010
0.005
0.000

0.00

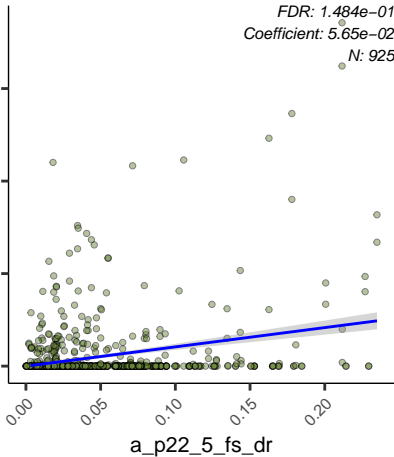
0.05

0.10

0.15

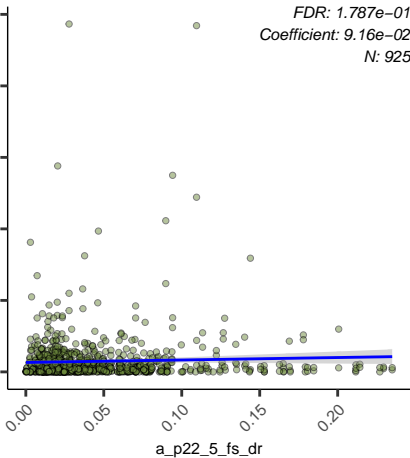
0.20

a_p22_5_fs_dr



ruthenibacterium_lactatiformans

FDR: 1.787e-01
Coefficient: 9.16e-02
N: 925



acidaminococcus_intestini

FDR: 2.167e-01

Coefficient: -6.98e-02

N: 925

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

