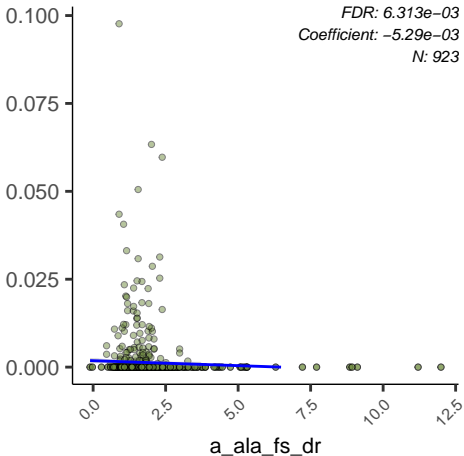
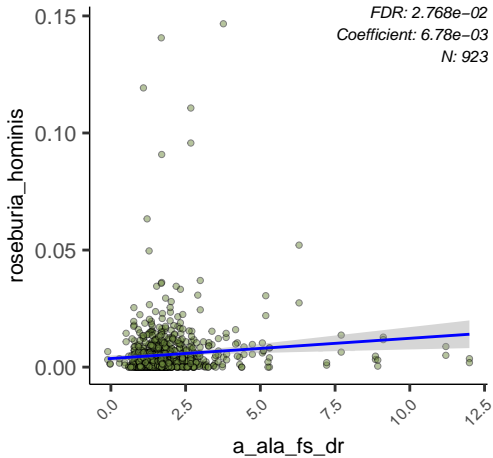
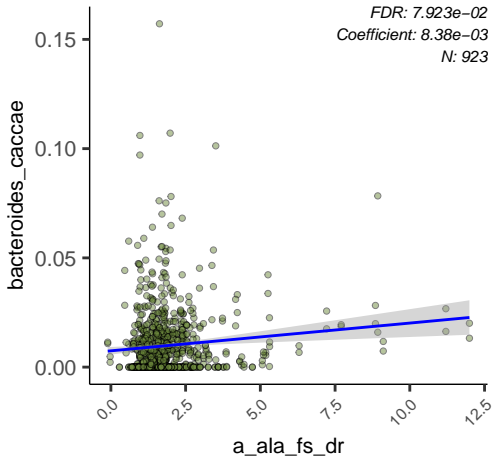
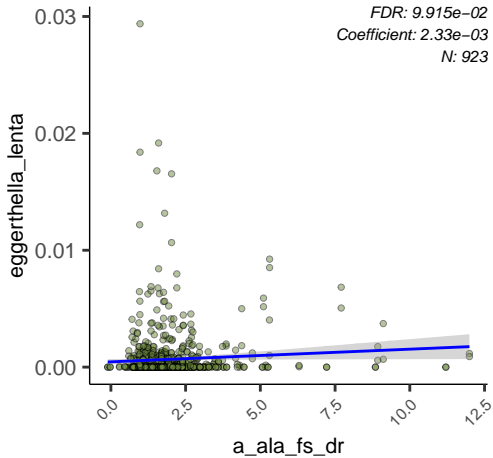


bifidobacterium_bifidum



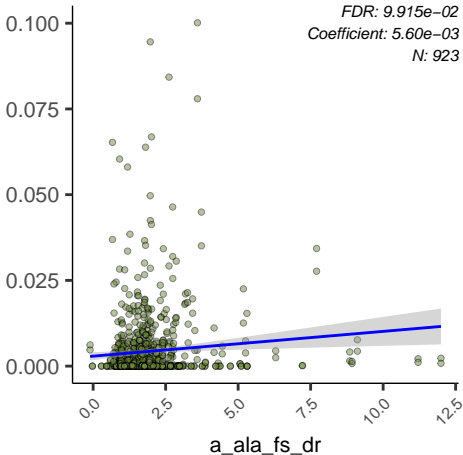


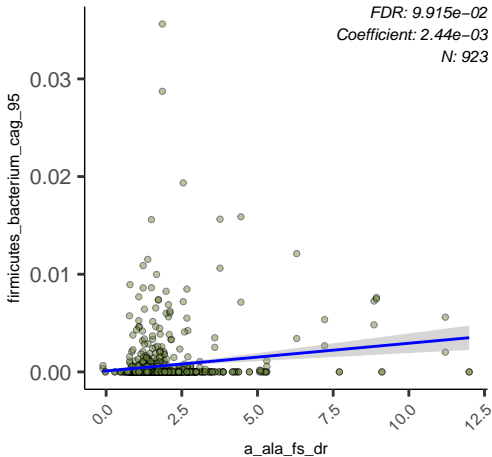




bacteroides_xylanisolvens

FDR: 9.915e-02
Coefficient: 5.60e-03
N: 923

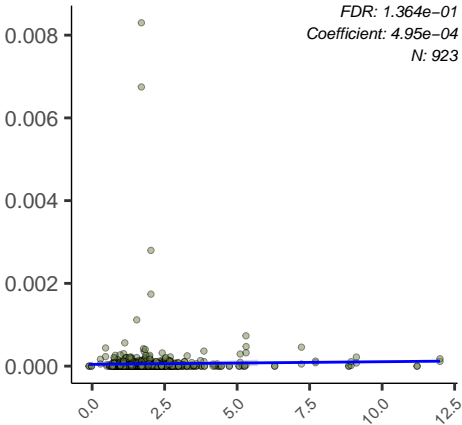




enterorhabdus_caecimuris

FDR: 1.364e-01
Coefficient: 4.95e-04
N: 923

a_ala_fs_dr



proteobacteria_bacterium_cag_139

FDR: 1.862e-01
Coefficient: 2.39e-03
N: 923

0.03
0.02
0.01
0.00

0.0

2.5

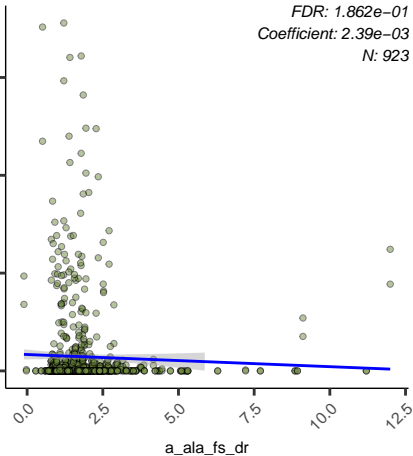
5.0

7.5

10.0

12.5

a_ala_fs_dr



erysipelatoclostridium_amosum

FDR: 2.427e-01
Coefficient: -2.52e-03
N: 923

0.06
0.04
0.02
0.00

0.0

2.5

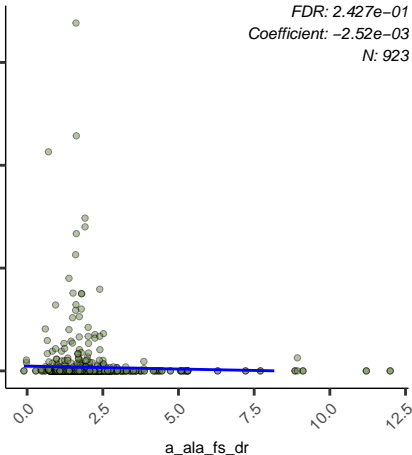
5.0

7.5

10.0

12.5

a_ala_fs_dr



roseburia_sp_cag_303

FDR: 2.457e-01

Coefficient: 2.94e-03

N: 923

0.0

2.5

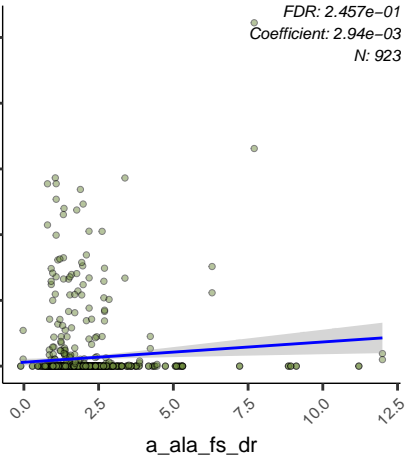
5.0

7.5

10.0

12.5

a_ala_fs_dr



lachnospira_pectinoschiza

FDR: 2.457e-01

Coefficient: -6.80e-03

N: 923

a_ala_fs_dr

