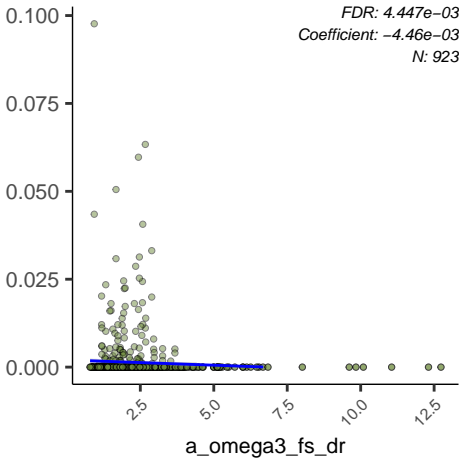
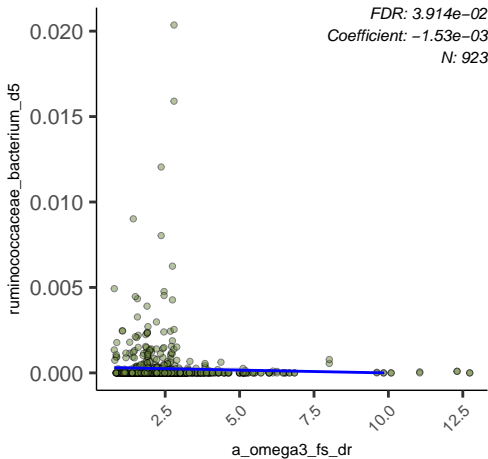
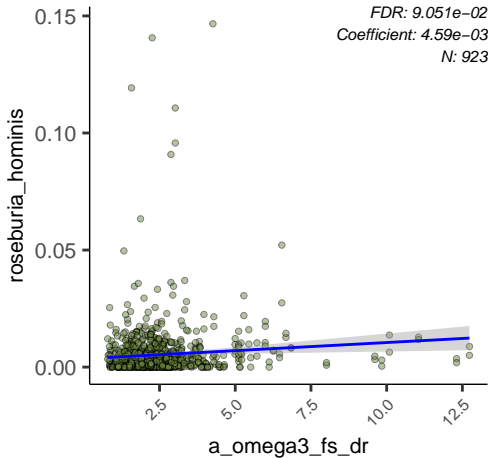
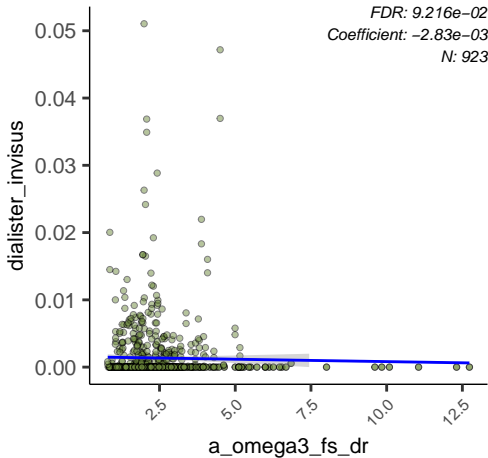


bifidobacterium_bifidum









bacteroides_xylanisolvens

FDR: 1.047e-01
Coefficient: 4.22e-03
N: 923

0.100
0.075
0.050
0.025
0.000

2.5

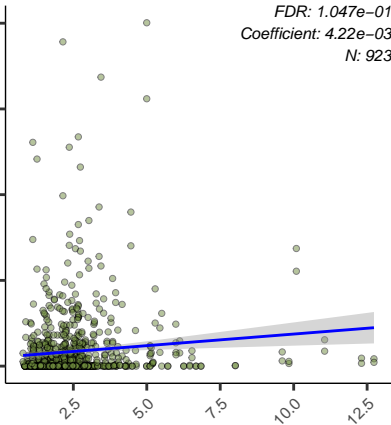
5.0

7.5

10.0

12.5

a_omega3_fs_dr



clostridium_sp_cag_253

0.04
0.03
0.02
0.01
0.00

FDR: 1.202e-01
Coefficient: 1.56e-03
N: 923

2.5

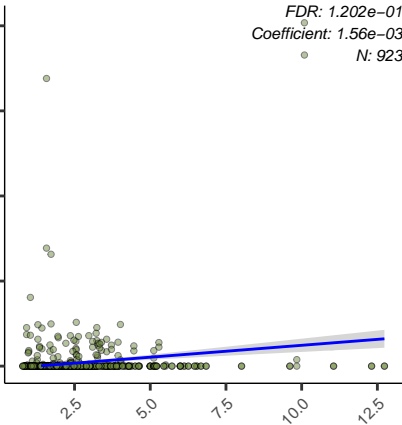
5.0

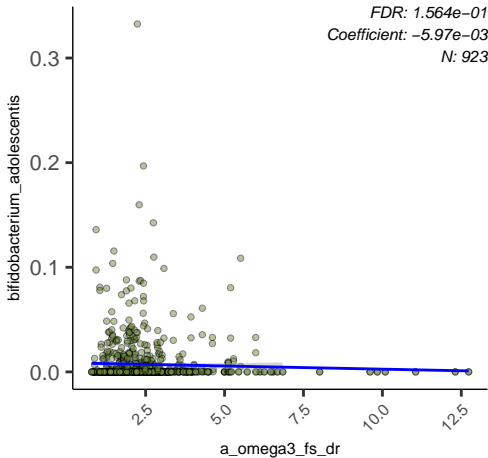
7.5

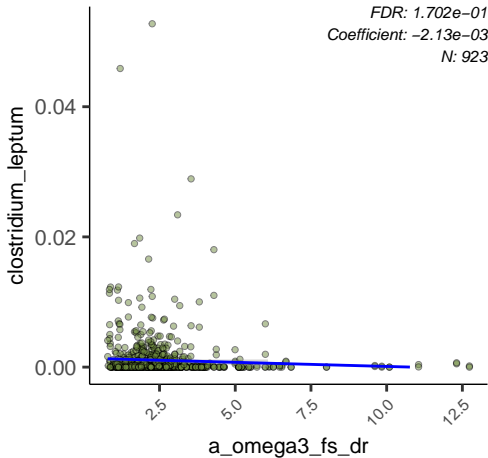
10.0

12.5

a_omega3_fs_dr

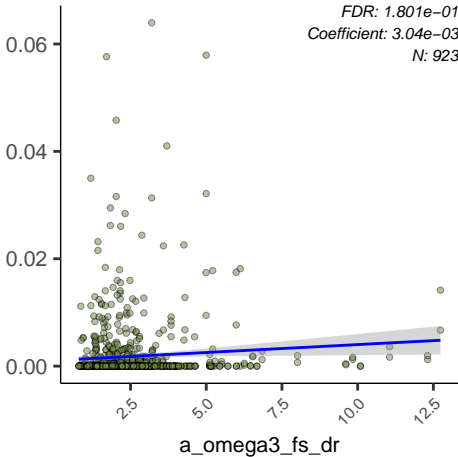






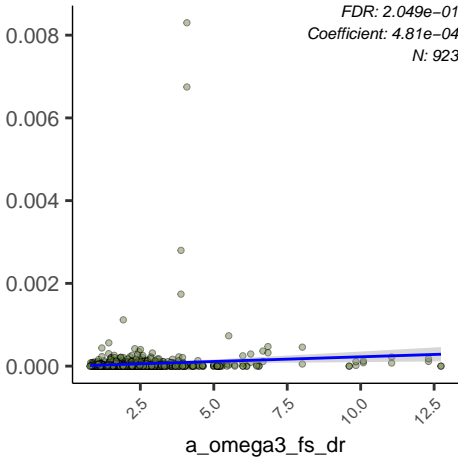
clostridium_sp_cag_167

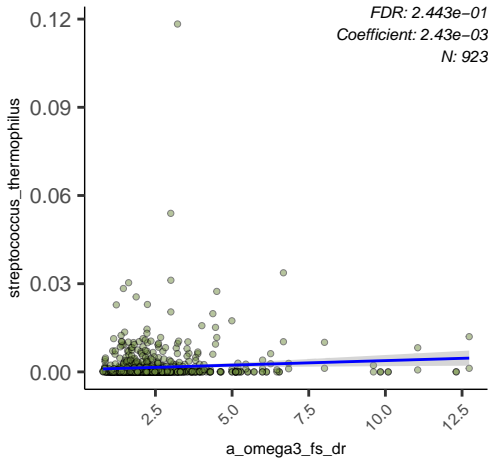
FDR: 1.801e-01
Coefficient: 3.04e-03
N: 923



enterorhabdus_caecimuris

FDR: 2.049e-01
Coefficient: 4.81e-04
N: 923





ruminococcus_callidus

0.03
0.02
0.01
0.00

2.5

5.0

7.5

10.0

12.5

a_omega3_fs_dr

FDR: 2.456e-01
Coefficient: -1.14e-03
N: 923

