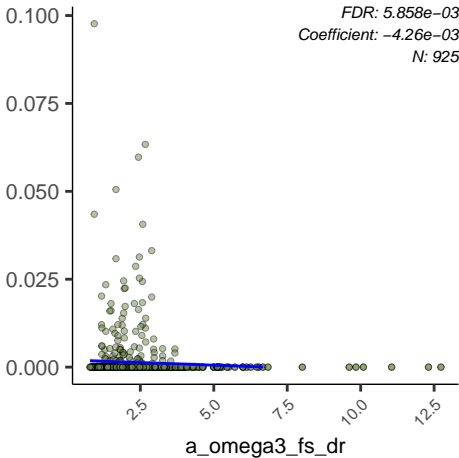
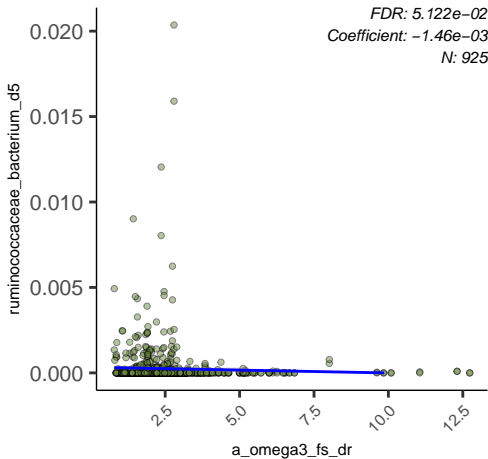
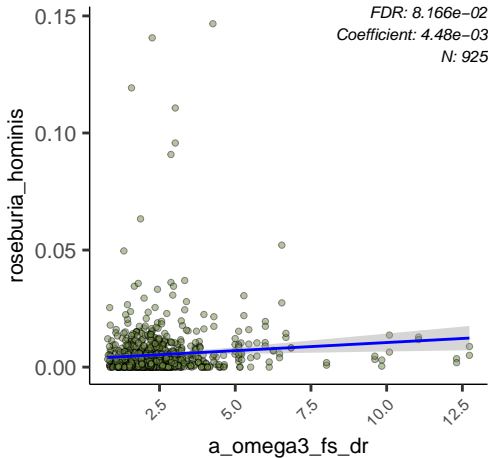
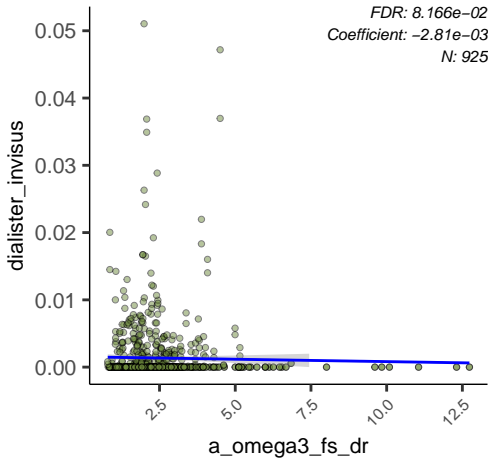


bifidobacterium\_bifidum



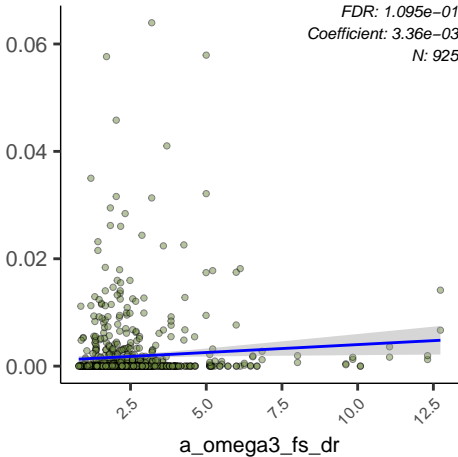






clostridium\_sp\_cag\_167

FDR: 1.095e-01  
Coefficient: 3.36e-03  
N: 925



bacteroides\_xylanisolvens

FDR: 1.095e-01  
Coefficient: 4.10e-03  
N: 925

0.100  
0.075  
0.050  
0.025  
0.000

2.5

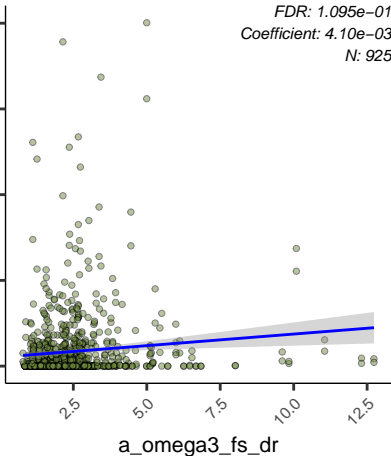
5.0

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: 1.467e-01  
Coefficient: 1.51e-03  
N: 925

2.5

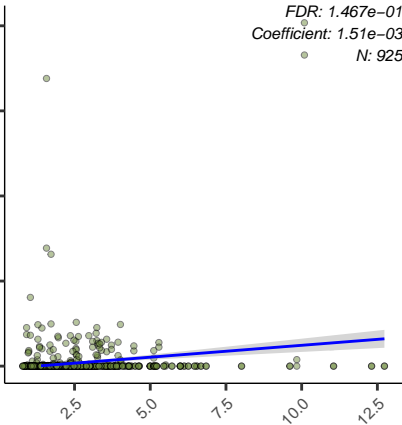
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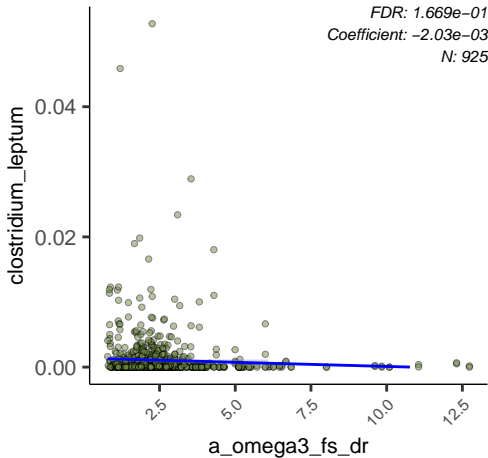
7.5

10.0

12.5

a\_omega3\_fs\_dr







enterorhabdus\_caecimuris

*FDR: 1.669e-01*  
*Coefficient: 4.95e-04*  
*N: 925*

