

eubacterium\_sp\_cag\_251

*FDR: 1.880e-03*

*Coefficient: -5.95e-03*

*N: 923*

0.08  
0.06  
0.04  
0.02  
0.00

logcrp

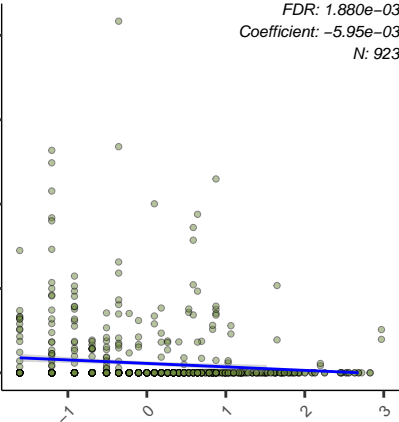
-1

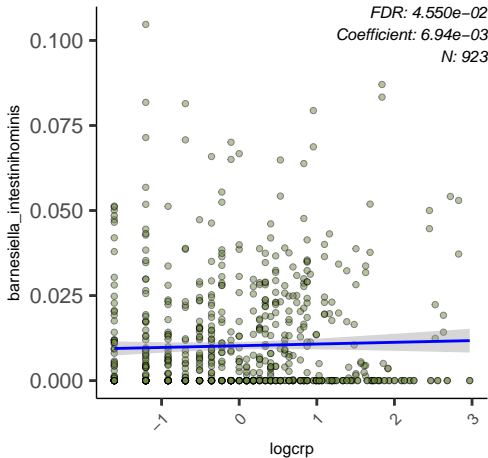
0

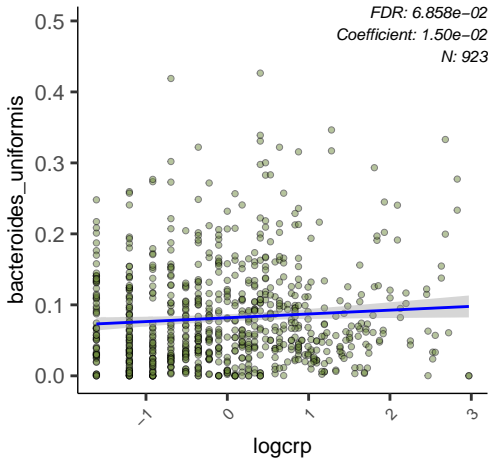
1

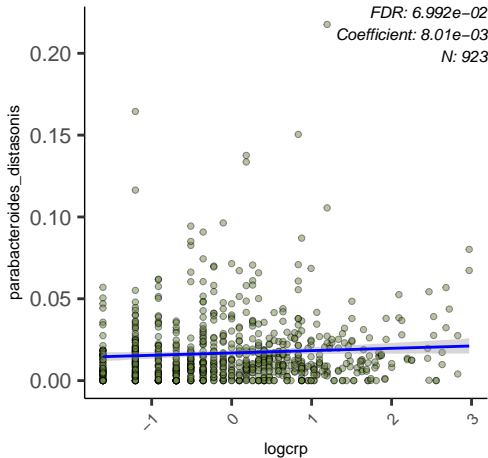
2

3









butyrivibrio\_sp\_cag\_318

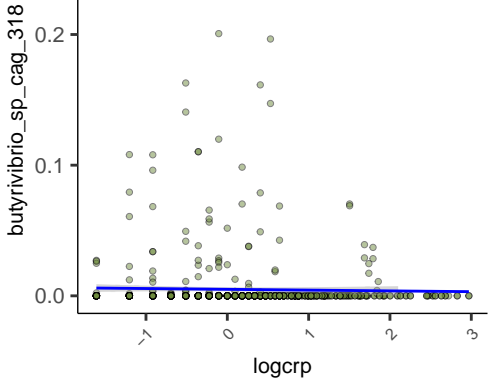
0.2

0.1

0.0

logcrp

*FDR: 7.016e-02*  
*Coefficient: -6.47e-03*  
*N: 923*



parabacteroides\_johnsonii

*FDR: 7.648e-02*  
*Coefficient: 3.13e-03*  
*N: 923*

0.06  
0.04  
0.02  
0.00

logcrp

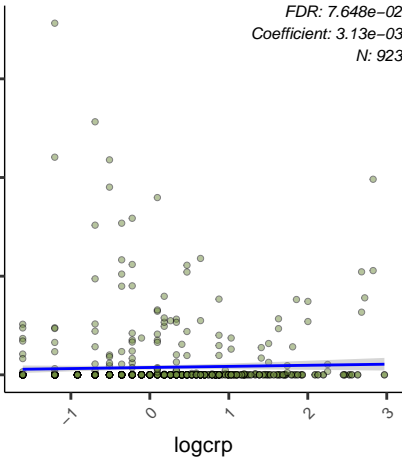
-1

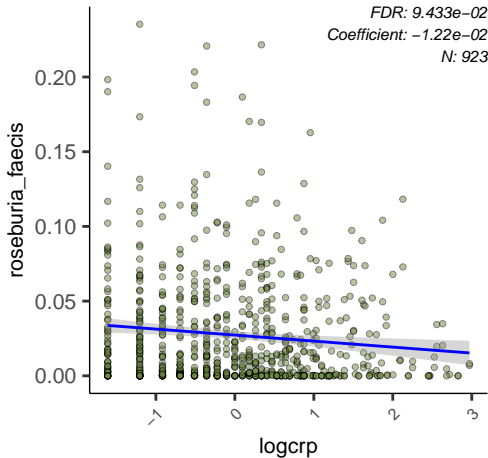
0

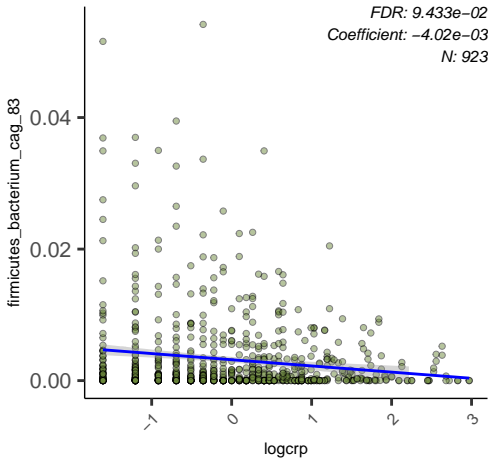
1

2

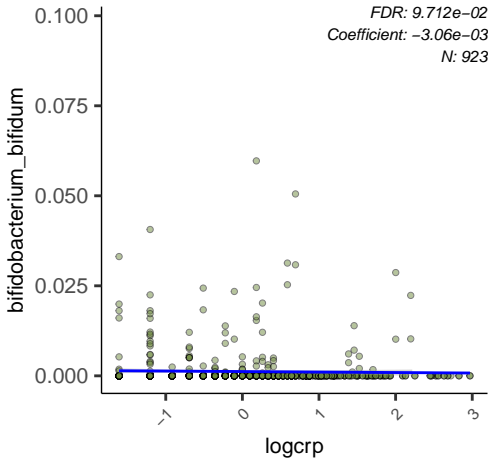
3

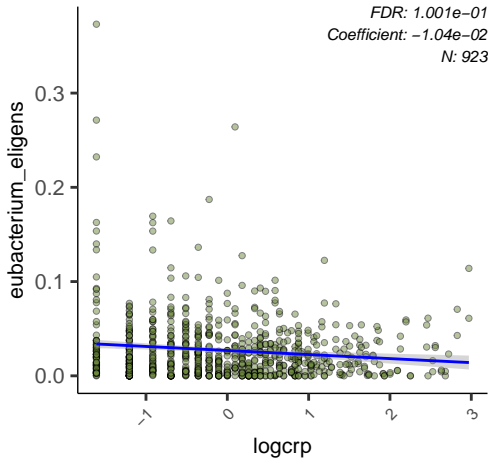












ruminococcus\_torques

*FDR: 1.215e-01*  
*Coefficient: 5.61e-03*  
*N: 923*

0.15  
0.10  
0.05  
0.00

logcrp

