

eubacterium\_sp\_cag\_251

*FDR: 3.618e-03*

*Coefficient: -5.85e-03*

*N: 923*

0.08  
0.06  
0.04  
0.02  
0.00

-1

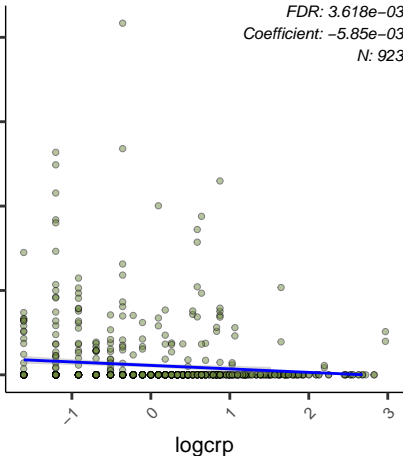
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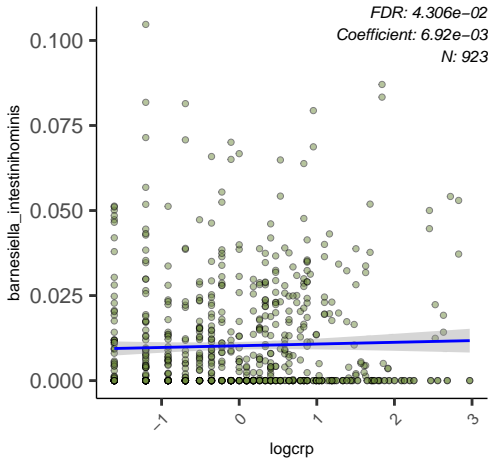
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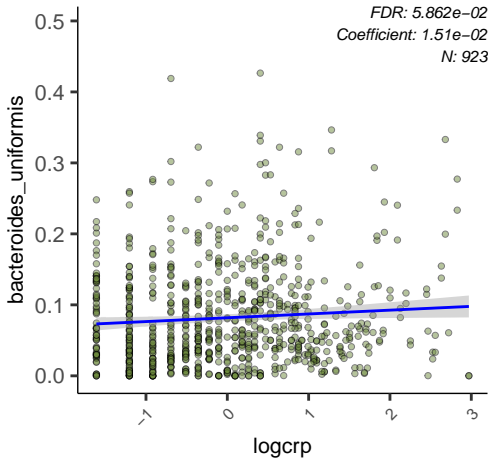
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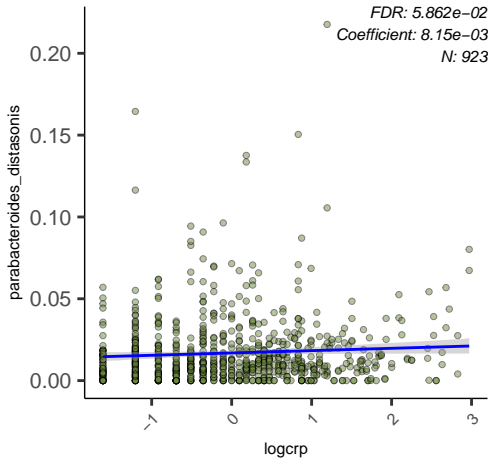
3

logcrp







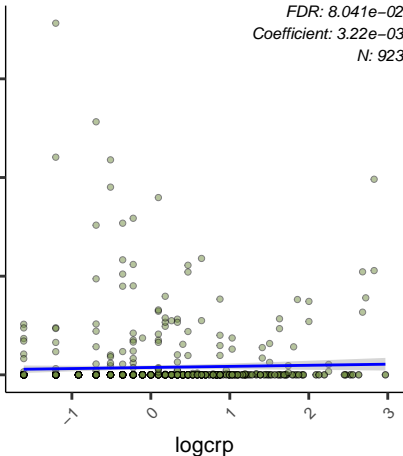


parabacteroides\_johnsonii

FDR: 8.041e-02  
Coefficient: 3.22e-03  
N: 923

0.06  
0.04  
0.02  
0.00

logcrp



butyrivibrio\_sp\_cag\_318

0.2  
0.1  
0.0

-1

0

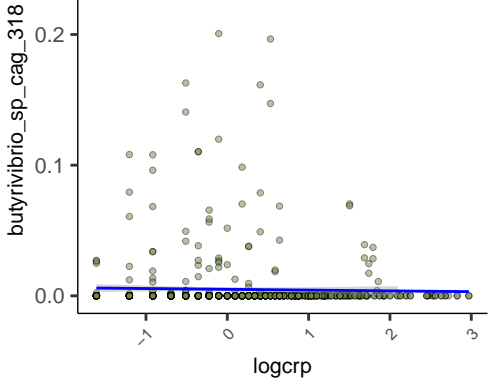
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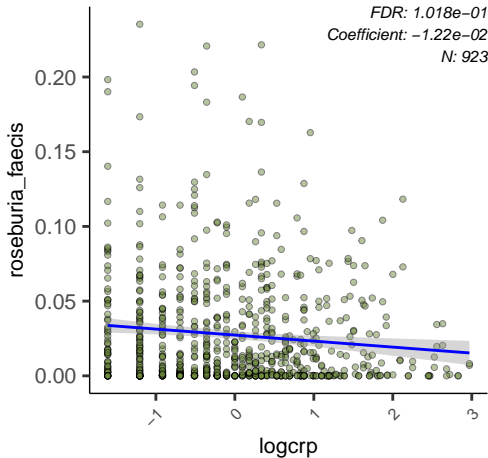
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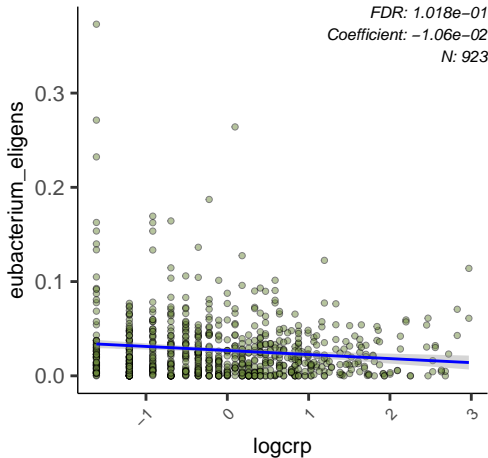
3

logcrp

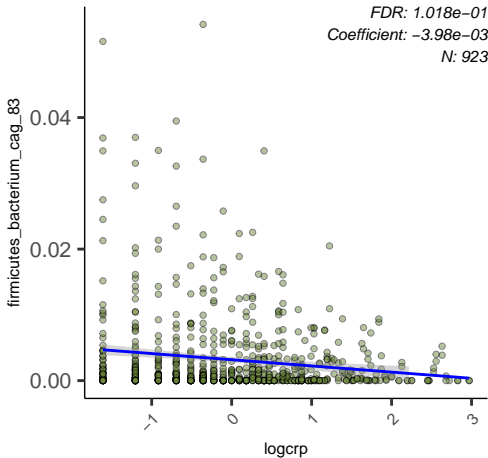
*FDR: 9.517e-02*  
*Coefficient: -6.18e-03*  
*N: 923*











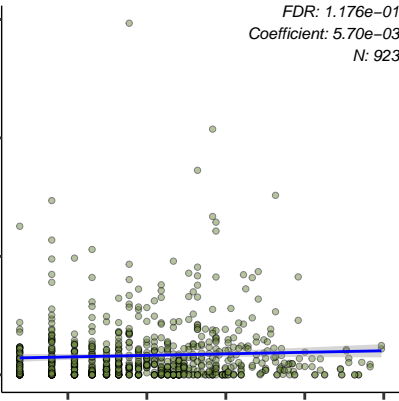
ruminococcus\_torques

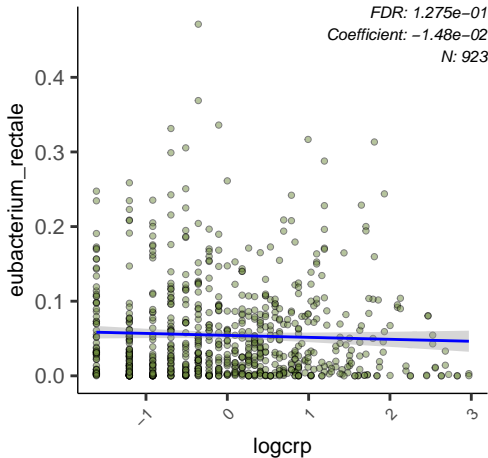
*FDR: 1.176e-01*  
*Coefficient: 5.70e-03*  
*N: 923*

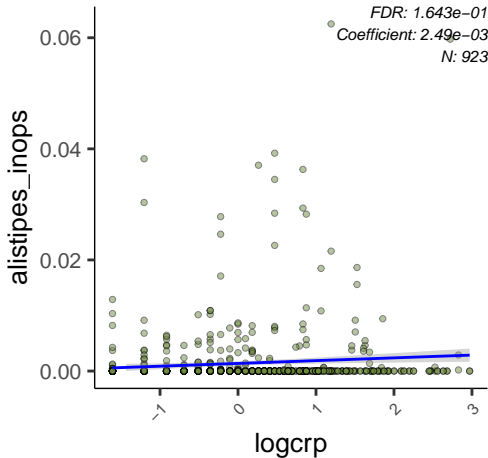
0.15  
0.10  
0.05  
0.00

logcrp

-1 0 1 2 3







bifidobacterium\_bifidum

FDR: 1.643e-01  
Coefficient: -2.73e-03  
N: 923

