

erysipelatoclostridium\_ramosum

*FDR: 1.934e-01*  
*Coefficient: 1.23e-02*  
*N: 923*

0.06

0.04

0.02

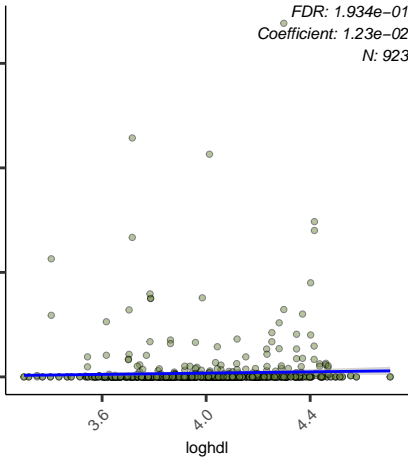
0.00

3.6

4.0

4.4

loghdl



eubacterium\_sp\_cag\_180

0.25  
0.20  
0.15  
0.10  
0.05  
0.00

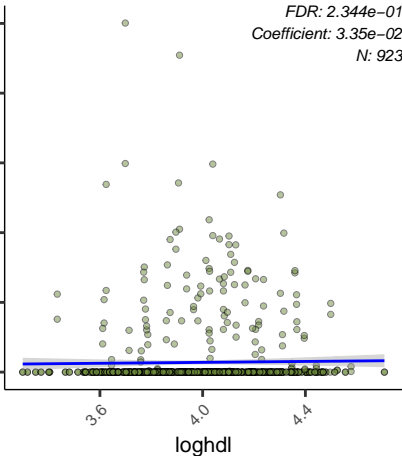
*FDR: 2.344e-01*  
*Coefficient: 3.35e-02*  
*N: 923*

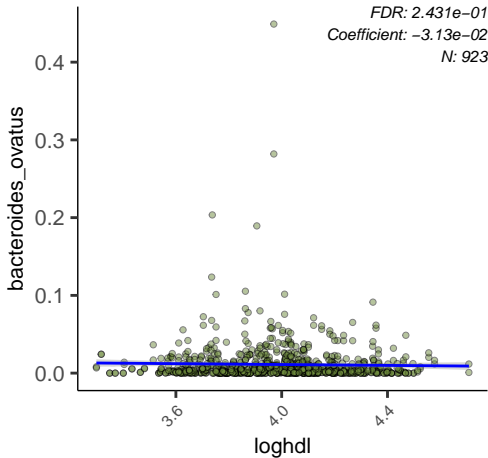
3.6

4.0

4.4

loghdi





enterorhabdus\_caecimuris

*FDR: 2.431e-01*  
*Coefficient: 2.90e-03*  
*N: 923*

0.008

0.006

0.004

0.002

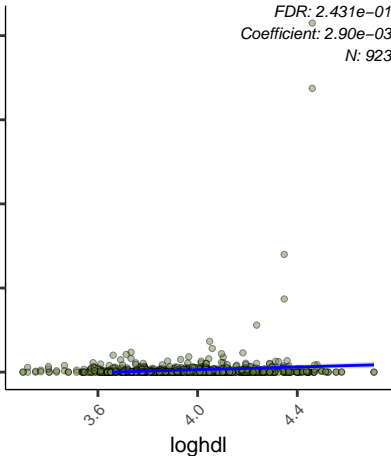
0.000

3.6

4.0

4.4

loghdl



bacteroides\_xylanisolvens

