

clostridium\_sp\_cag\_253

0.04  
0.03  
0.02  
0.01  
0.00

*FDR: 3.001e-03*  
*Coefficient: 6.72e-06*  
*N: 923*

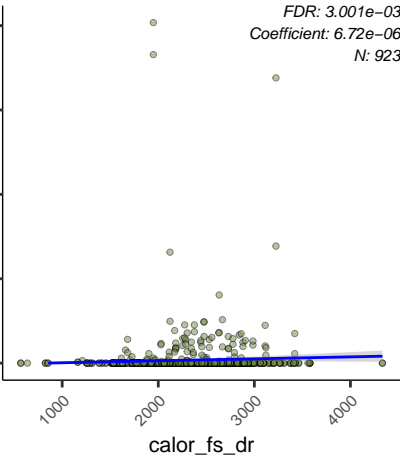
1000

2000

3000

4000

calor\_fs\_dr



phascolarctobacterium\_faecium

0.02

0.01

0.00

1000

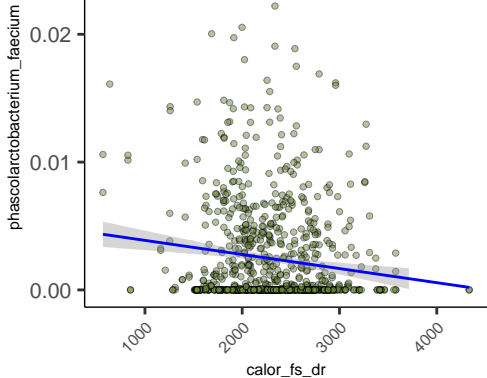
2000

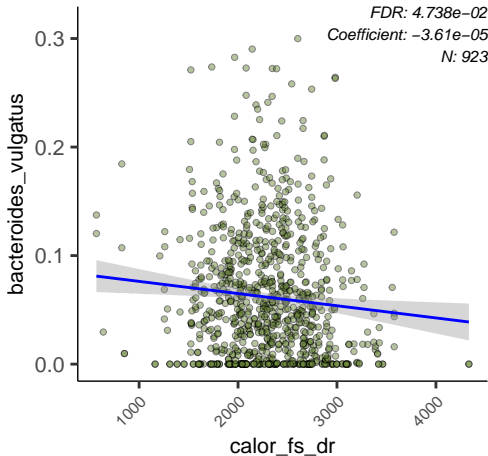
3000

4000

calor\_fs\_dr

*FDR: 3.372e-02*  
*Coefficient: -9.34e-06*  
*N: 923*





odoribacter\_splanchnicus

*FDR: 5.120e-02*

*Coefficient: -9.96e-06*

*N: 923*

0.03

0.02

0.01

0.00

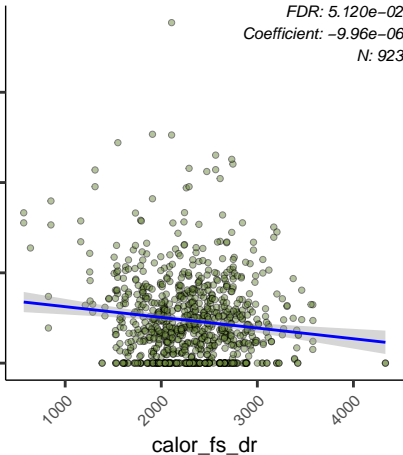
1000

2000

3000

4000

calor\_fs\_dr



ruminococcaceae\_bacterium\_d5

