

clostridium_sp_cag_253

0.04
0.03
0.02
0.01
0.00

FDR: 3.088e-03
Coefficient: 6.80e-06
N: 923

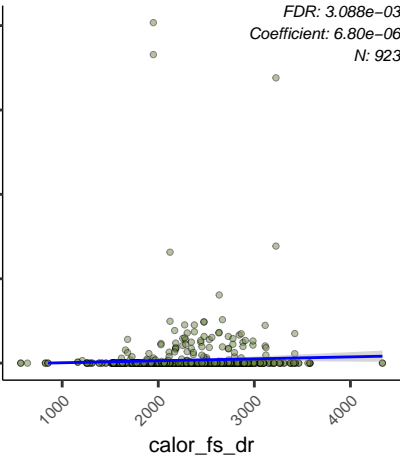
1000

2000

3000

4000

calor_fs_dr



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

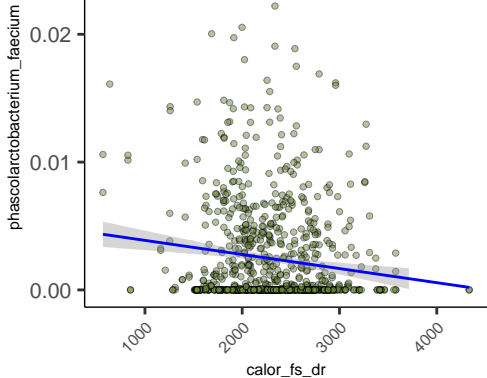
2000

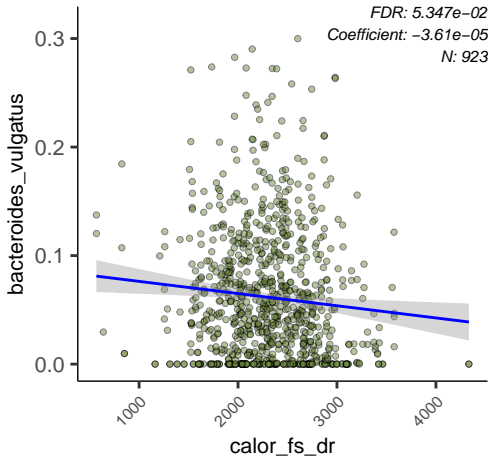
3000

4000

calor_fs_dr

FDR: 4.003e-02
Coefficient: -9.33e-06
N: 923





odoribacter_splanchnicus

FDR: 5.750e-02

Coefficient: -9.92e-06

N: 923

0.03

0.02

0.01

0.00

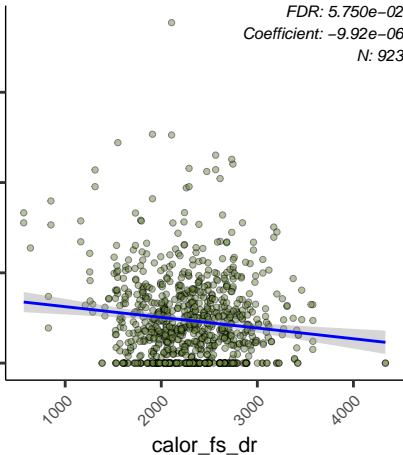
1000

2000

3000

4000

calor_fs_dr



ruminococcaceae_bacterium_d5

