

clostridium_sp_cag_253

0.04
0.03
0.02
0.01
0.00

FDR: 4.004e-03
Coefficient: 6.71e-06
N: 923

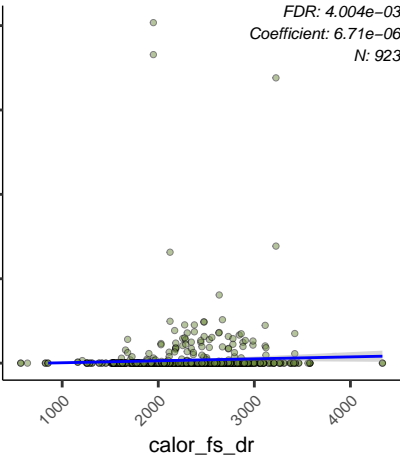
1000

2000

3000

4000

calor_fs_dr



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

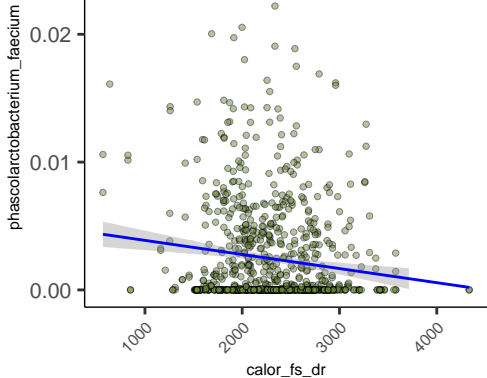
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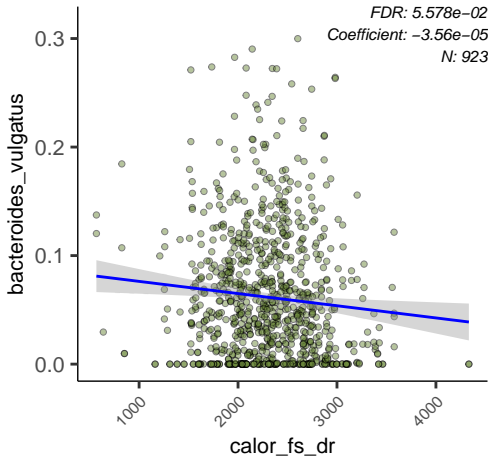
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4000

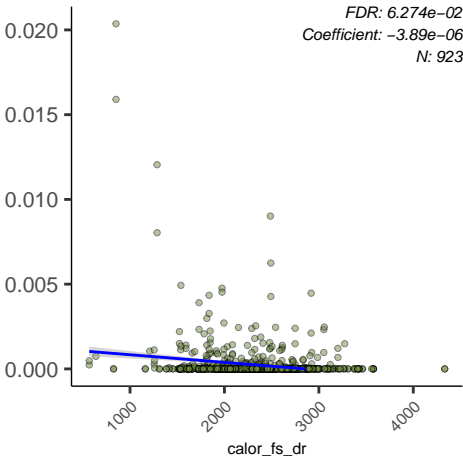
calor_fs_dr

FDR: 4.214e-02
Coefficient: -9.29e-06
N: 923





ruminococcaceae_bacterium_d5



odoribacter_splanchnicus

FDR: 6.577e-02

Coefficient: -9.64e-06

N: 923

0.03

0.02

0.01

0.00

1000

2000

3000

4000

calor_fs_dr

