

clostridium_sp_cag_253

0.04
0.03
0.02
0.01
0.00

FDR: 2.842e-03
Coefficient: 6.99e-06
N: 925

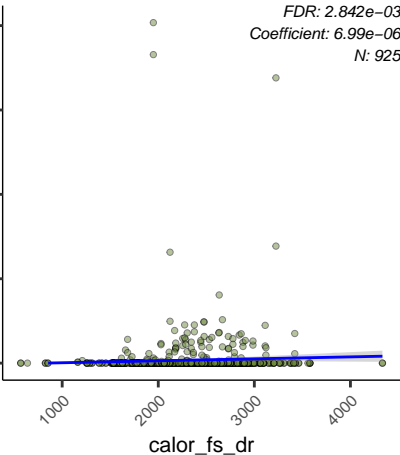
1000

2000

3000

4000

calor_fs_dr



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

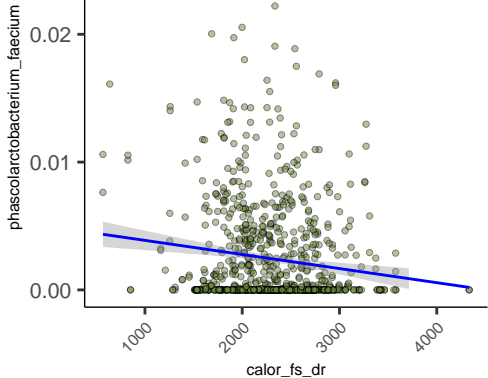
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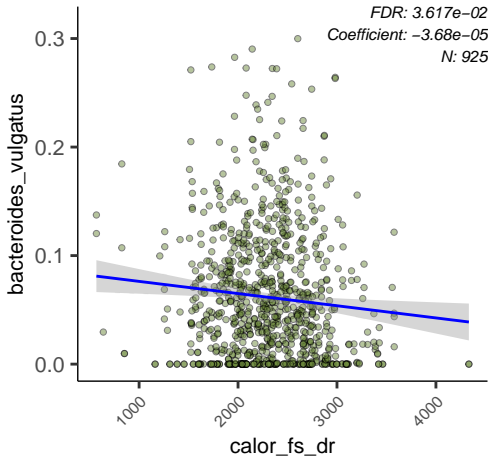
3000

4000

calor_fs_dr

FDR: 2.177e-02
Coefficient: -9.73e-06
N: 925





odoribacter_splanchnicus

FDR: 3.617e-02

Coefficient: -1.05e-05

N: 925

0.03

0.02

0.01

0.00

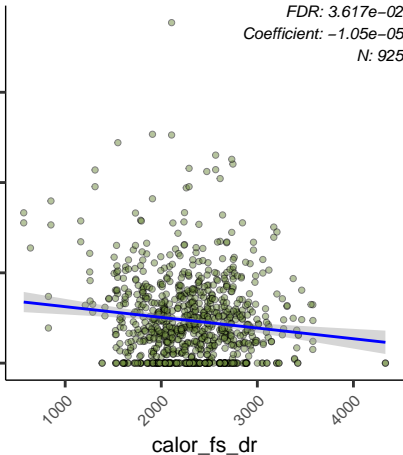
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2000

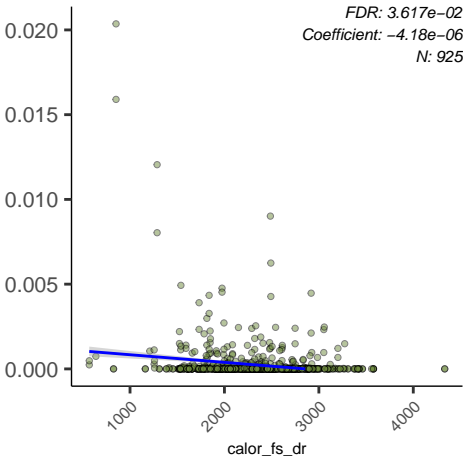
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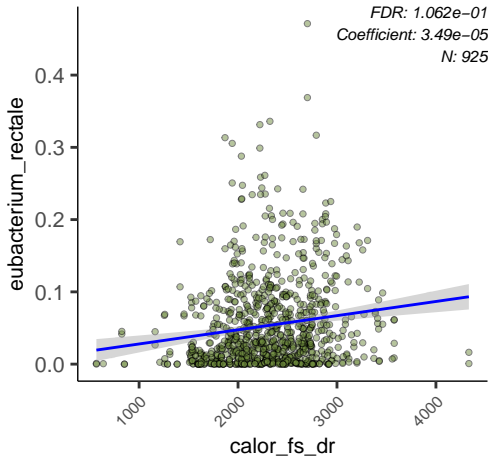
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calor_fs_dr



ruminococcaceae_bacterium_d5





eubacterium_siraeum

FDR: 2.051e-01
Coefficient: -2.86e-05
N: 925

0.3
0.2
0.1
0.0

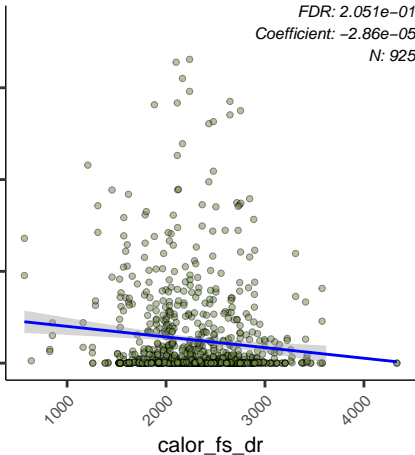
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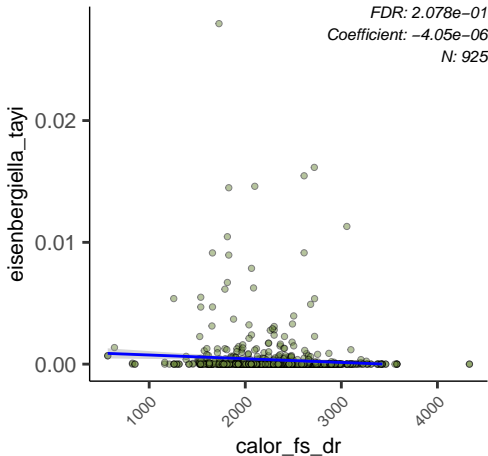
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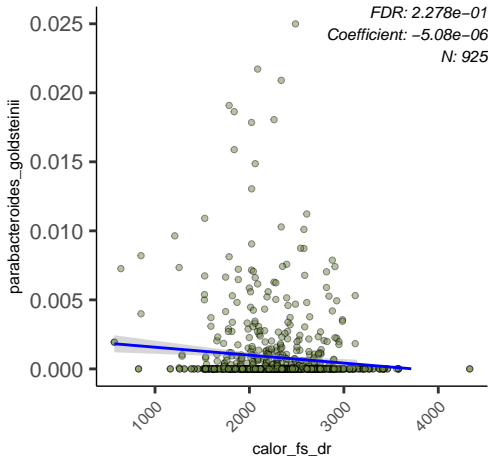
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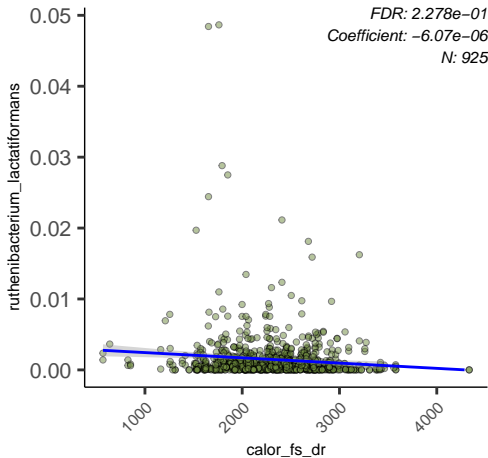
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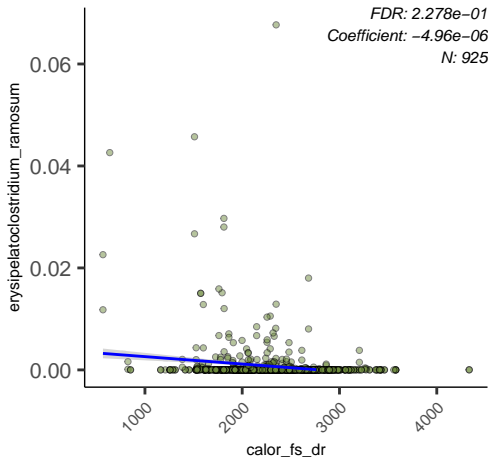
calor_fs_dr











clostridium_sp_cag_58

0.025
0.020
0.015
0.010
0.005
0.000

FDR: 2.322e-01
Coefficient: -5.58e-06
N: 925

1000

2000

3000

4000

calor_fs_dr

