

clostridium_sp_cag_253

0.04
0.03
0.02
0.01
0.00

FDR: 1.983e-03
Coefficient: 6.97e-06
N: 923

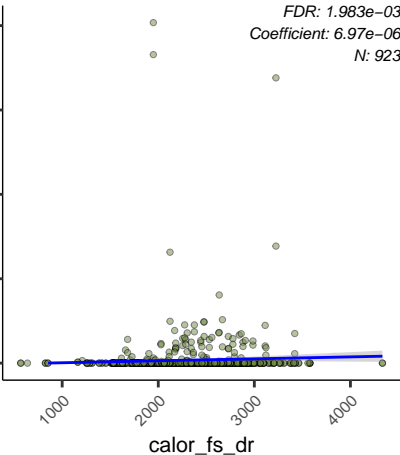
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2000

3000

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calor_fs_dr



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

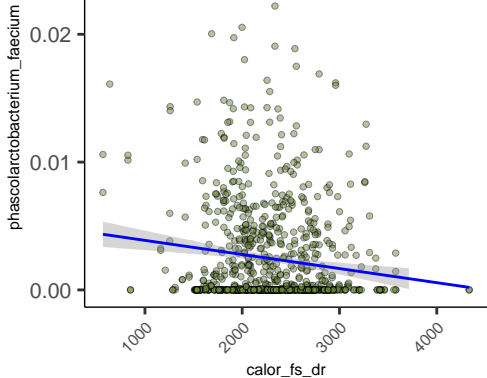
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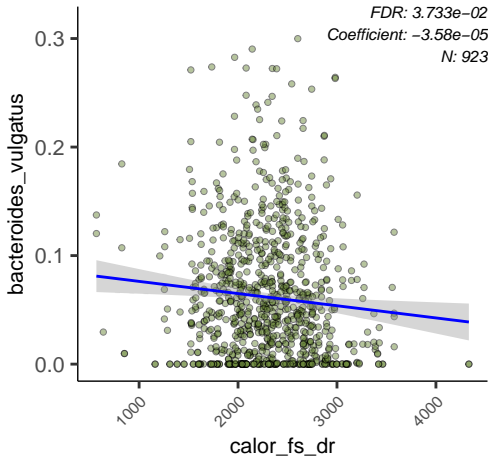
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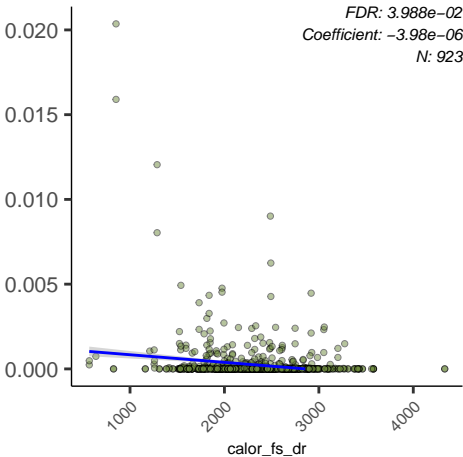
calor_fs_dr

FDR: 2.772e-02
Coefficient: -9.36e-06
N: 923





ruminococcaceae_bacterium_d5



odoribacter_splanchnicus

FDR: 4.268e-02

Coefficient: -9.77e-06

N: 923

0.03

0.02

0.01

0.00

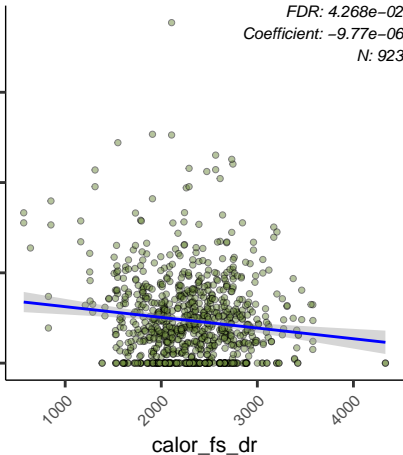
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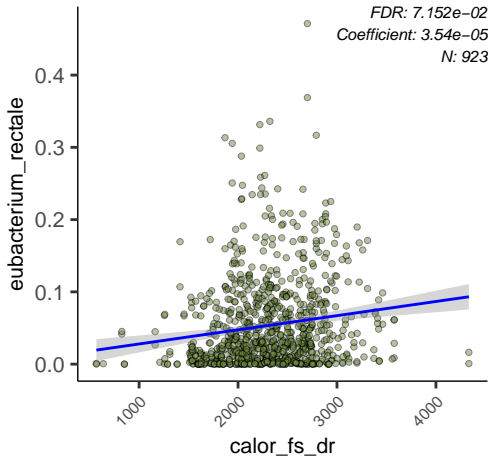
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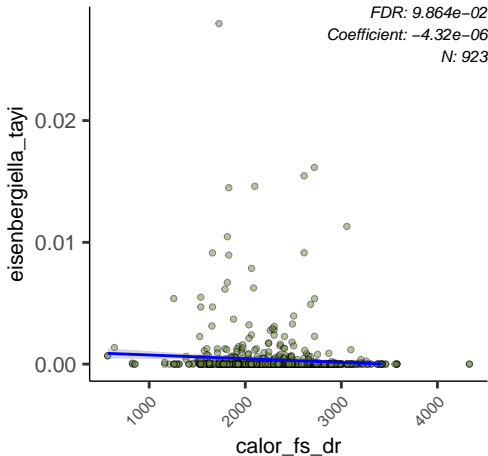
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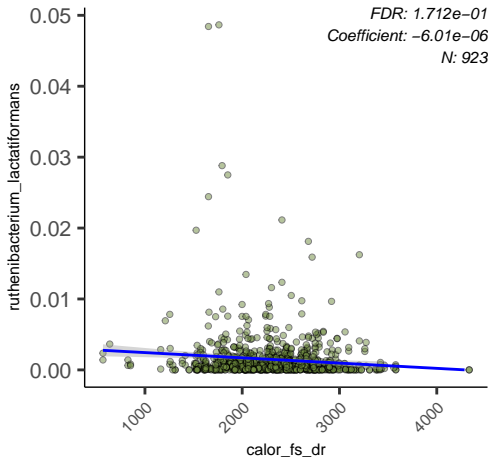
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calor_fs_dr









clostridium_sp_cag_58

0.025
0.020
0.015
0.010
0.005
0.000

FDR: 2.158e-01
Coefficient: -5.39e-06
N: 923

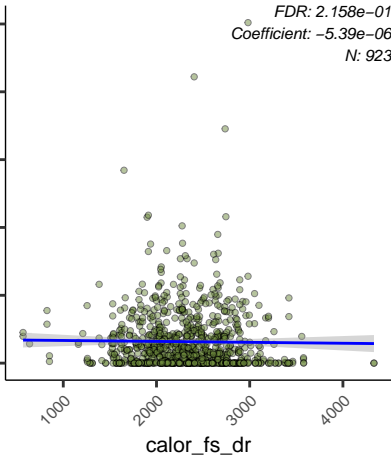
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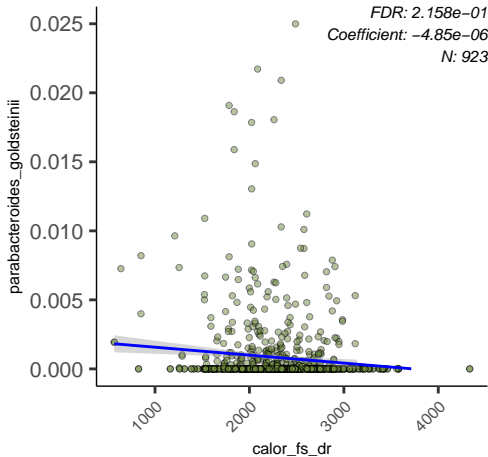
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calor_fs_dr





catenibacterium_mitsuokai

FDR: 2.322e-01
Coefficient: 3.23e-06
N: 923

0.020
0.015
0.010
0.005
0.000

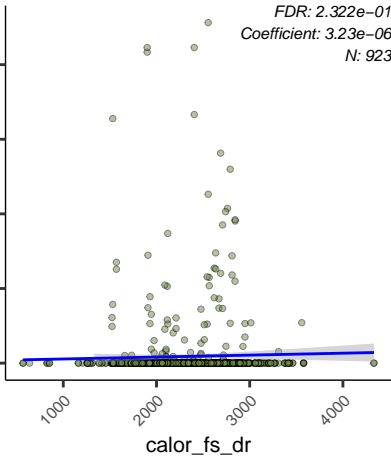
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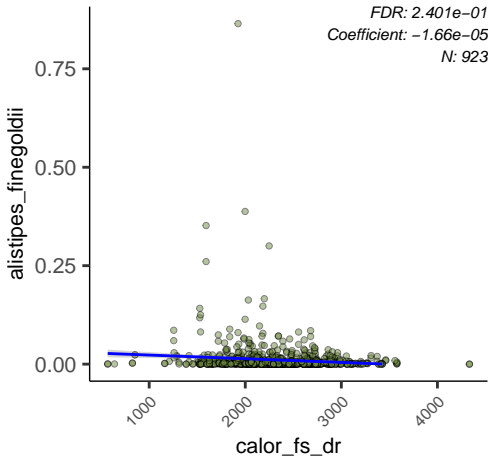
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calor_fs_dr





anaerotruncus_colihominis

FDR: 2.401e-01

Coefficient: -1.84e-06

N: 923

0.006

0.004

0.002

0.000

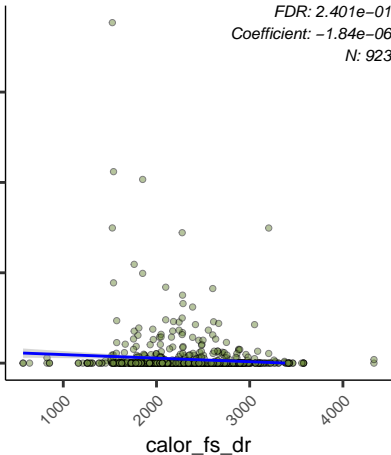
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2000

3000

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calor_fs_dr



agathobaculum_butyriciproducens

FDR: 2.401e-01

Coefficient: 6.13e-06

N: 923

0.03

0.02

0.01

0.00

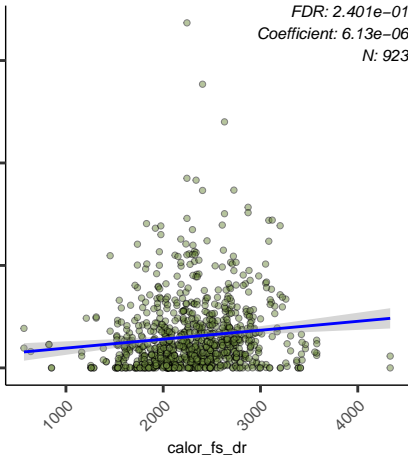
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2000

3000

4000

calor_fs_dr



bacteroides_salysiae

FDR: 2.454e-01

Coefficient: -6.48e-06

N: 923

0.04

0.03

0.02

0.01

0.00

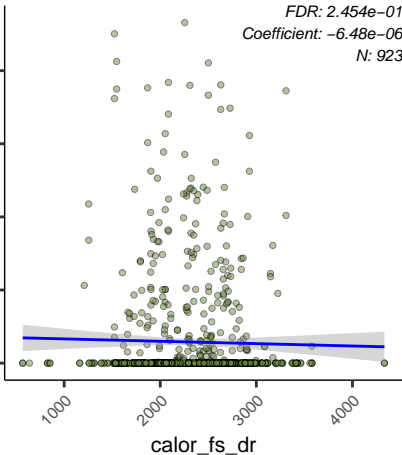
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2000

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4000

calor_fs_dr



eubacterium_siraeum

FDR: 2.454e-01
Coefficient: -2.48e-05
N: 923

0.3
0.2
0.1
0.0

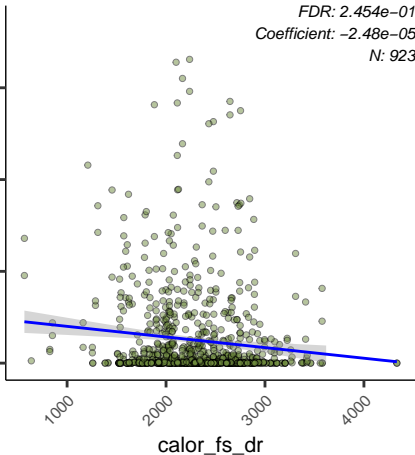
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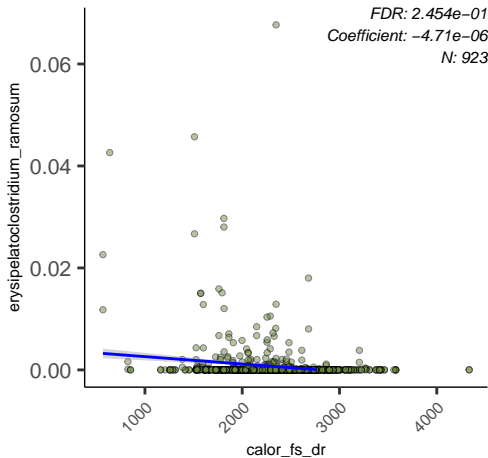
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calor_fs_dr





proteobacteria_bacterium_cag_139

FDR: 2.454e-01
Coefficient: 4.43e-06
N: 923

0.03
0.02
0.01
0.00

1000

2000

3000

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calor_fs_dr

