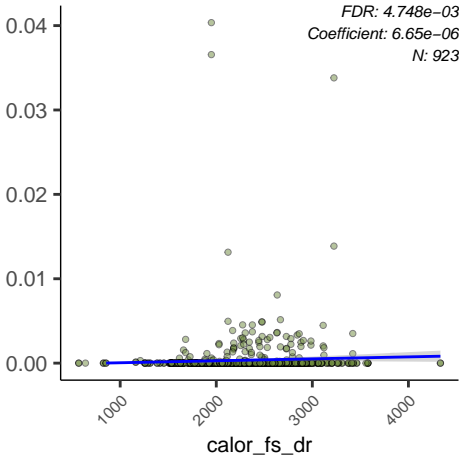


clostridium_sp_cag_253



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

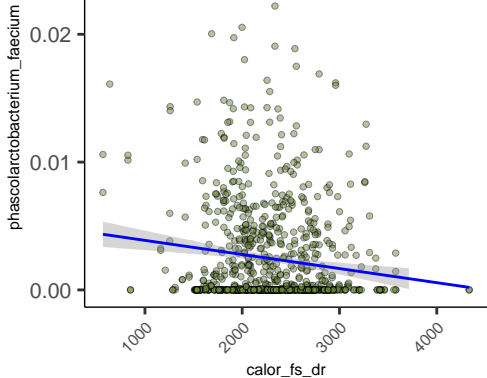
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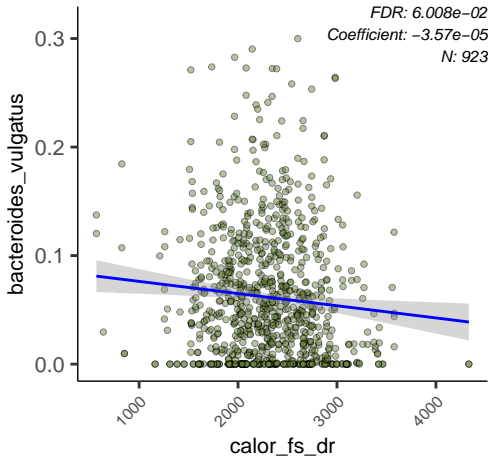
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4000

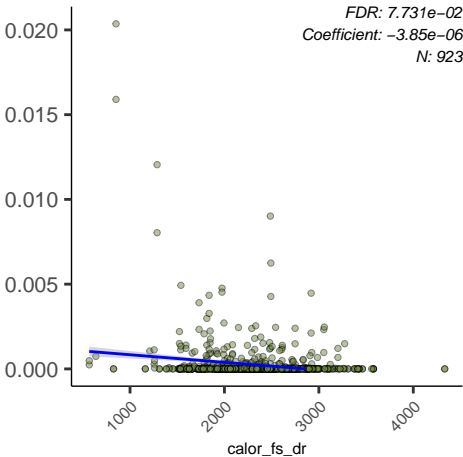
calor_fs_dr

FDR: 4.178e-02
Coefficient: -9.30e-06
N: 923





ruminococcaceae_bacterium_d5



odoribacter_splanchnicus

FDR: 8.593e-02

Coefficient: -9.47e-06

N: 923

0.03

0.02

0.01

0.00

1000

2000

3000

4000

calor_fs_dr

