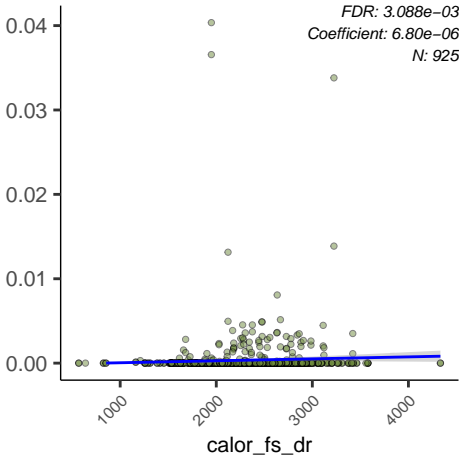


clostridium_sp_cag_253



phascolarctobacterium_faecium

0.02

0.01

0.00

1000

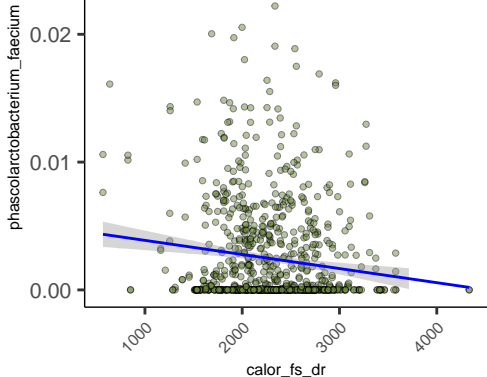
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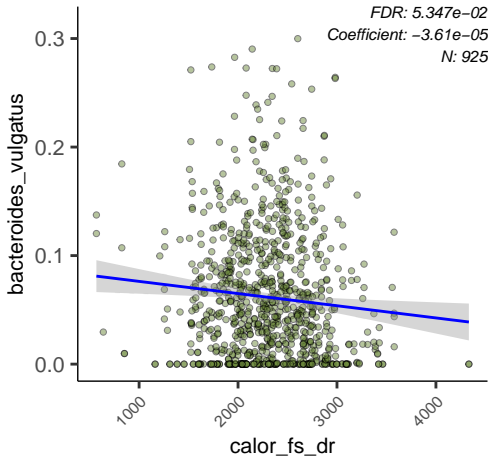
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4000

calor_fs_dr

FDR: 4.003e-02
Coefficient: -9.33e-06
N: 925





odoribacter_splanchnicus

FDR: 5.750e-02

Coefficient: -9.92e-06

N: 925

0.03

0.02

0.01

0.00

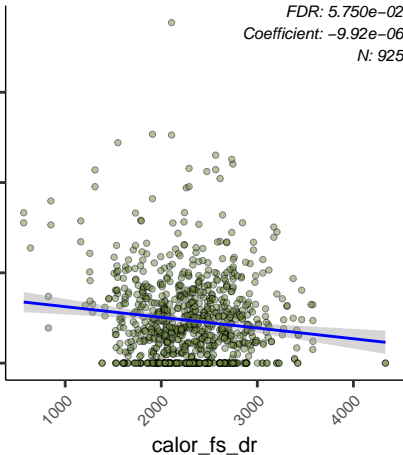
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2000

3000

4000

calor_fs_dr



ruminococcaceae_bacterium_d5

