

clostridium\_sp\_cag\_253

0.04  
0.03  
0.02  
0.01  
0.00

*FDR: 2.420e-03*  
*Coefficient: 7.06e-06*  
*N: 925*

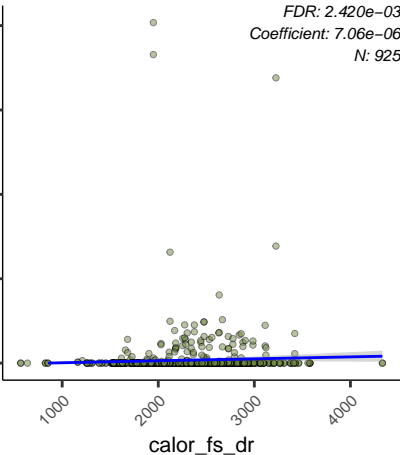
1000

2000

3000

4000

calor\_fs\_dr



phascolarctobacterium\_faecium

0.02

0.01

0.00

1000

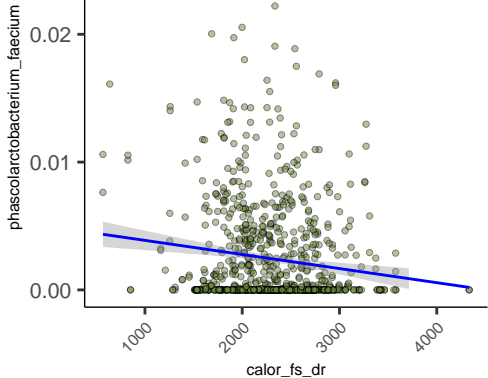
2000

3000

4000

calor\_fs\_dr

*FDR: 2.494e-02*  
*Coefficient: -9.67e-06*  
*N: 925*



odoribacter\_splanchnicus

*FDR: 4.147e-02*

*Coefficient: -1.06e-05*

*N: 925*

0.03

0.02

0.01

0.00

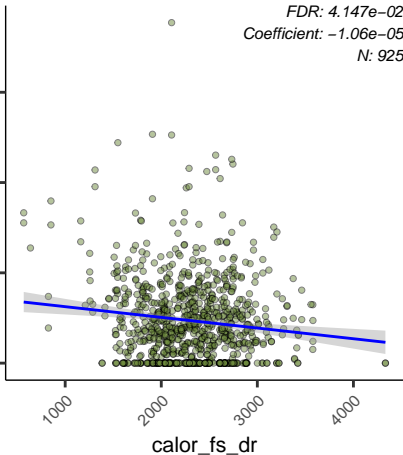
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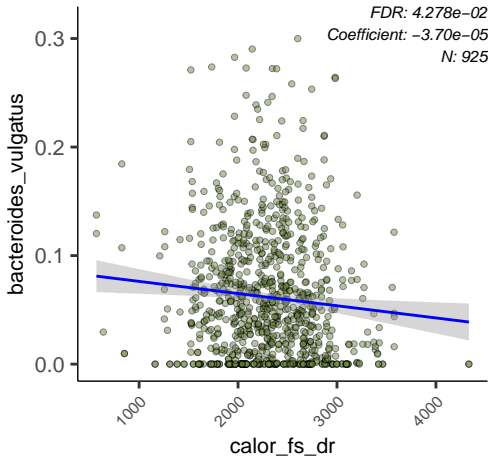
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3000

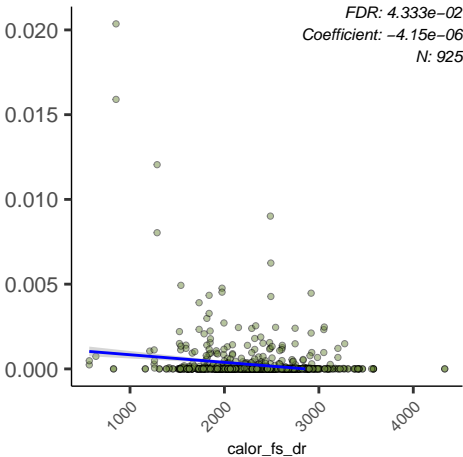
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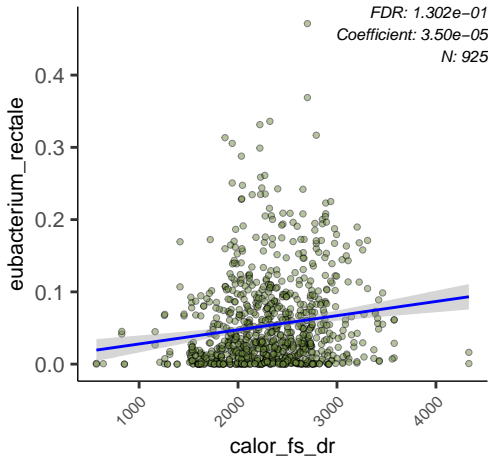
calor\_fs\_dr

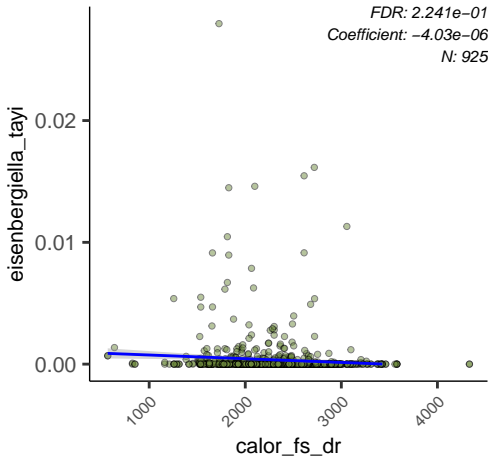




ruminococcaceae\_bacterium\_d5







eubacterium\_siraeum

*FDR: 2.241e-01*  
*Coefficient: -2.82e-05*  
*N: 925*

0.3  
0.2  
0.1  
0.0

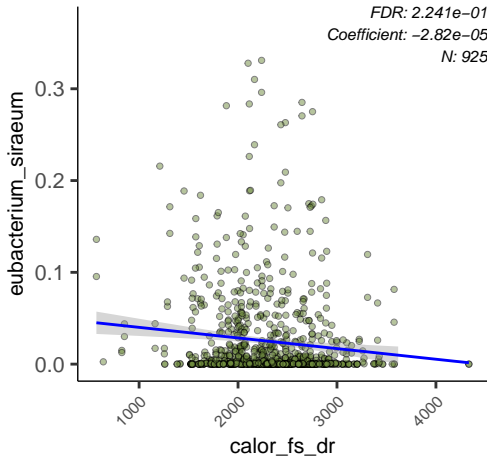
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2000

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calor\_fs\_dr





clostridium\_sp\_cag\_58

0.025  
0.020  
0.015  
0.010  
0.005  
0.000

*FDR: 2.443e-01*  
*Coefficient: -5.64e-06*  
*N: 925*

1000

2000

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4000

calor\_fs\_dr

