

odoribacter_splanchnicus

FDR: 6.668e-02

Coefficient: -1.29e-02

N: 923

0.03
0.02
0.01
0.00

0

1

2

3

a_f205_fs_dr

0

1

2

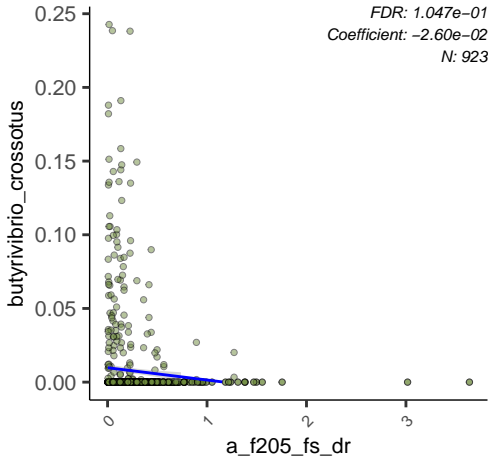
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0

1

2

3



eubacterium_sp_cag_251

0.08

0.06

0.04

0.02

0.00

0

1

2

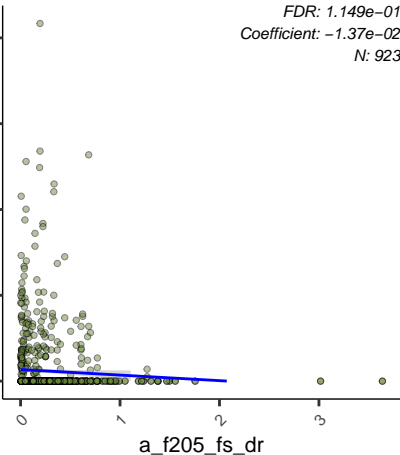
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a_f205_fs_dr

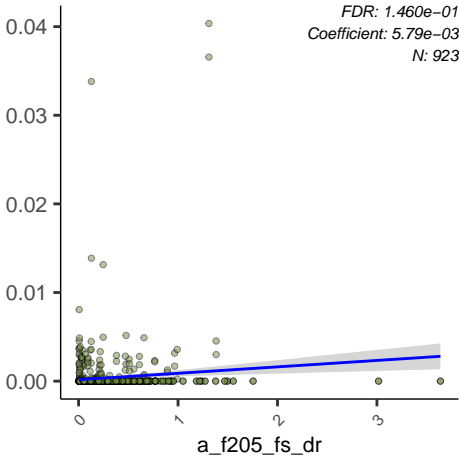
FDR: 1.149e-01

Coefficient: -1.37e-02

N: 923



clostridium_sp_cag_253



oscillibacter_sp_cag_241

FDR: $1.523e-01$
Coefficient: $1.27e-02$
N: 923

0.06
0.04
0.02
0.00

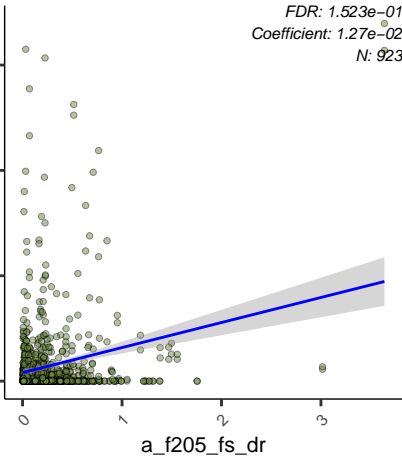
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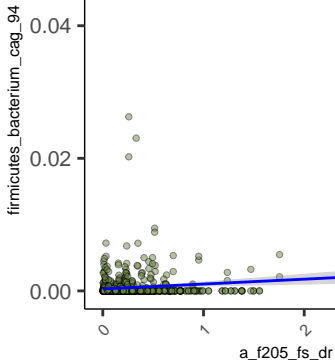
2

3

a_f205_fs_dr



FDR: 1.523e-01
Coefficient: 5.71e-03
N: 923



enterorhabdus_caecimuris

FDR: 1.902e-01
Coefficient: 1.90e-03
N: 923

