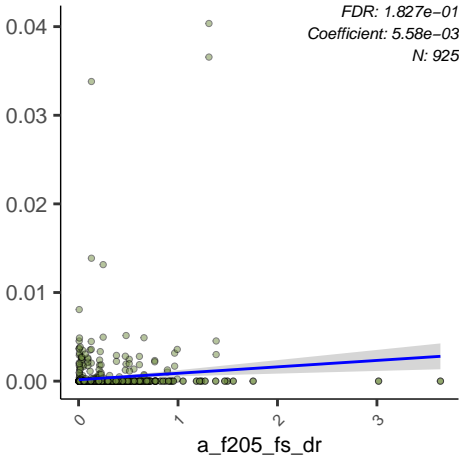


clostridium_sp_cag_253



eubacterium_sp_cag_251

0.08

0.06

0.04

0.02

0.00

0

1

2

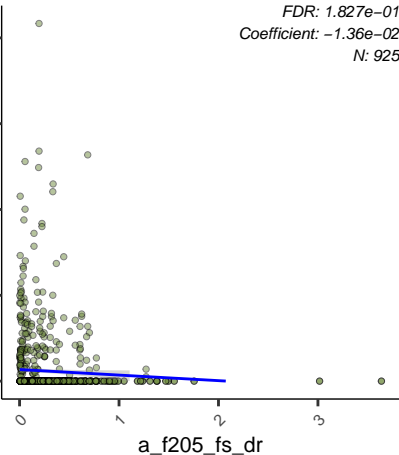
3

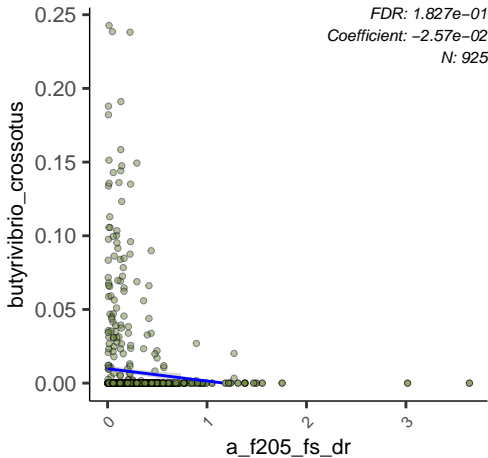
a_f205_fs_dr

FDR: 1.827e-01

Coefficient: -1.36e-02

N: 925





oscillibacter_sp_cag_241

FDR: $1.827e-01$
Coefficient: $1.27e-02$
N: 925

0.06
0.04
0.02
0.00

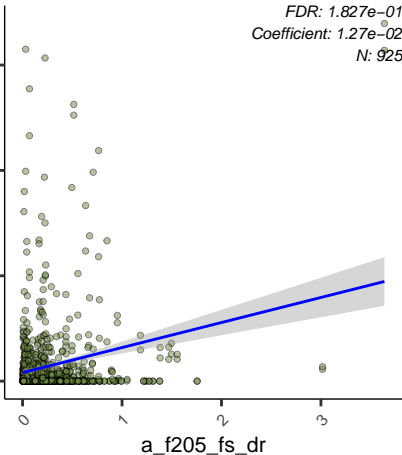
0

1

2

3

a_f205_fs_dr



odoribacter_splanchnicus

0.03
0.02
0.01
0.00

0

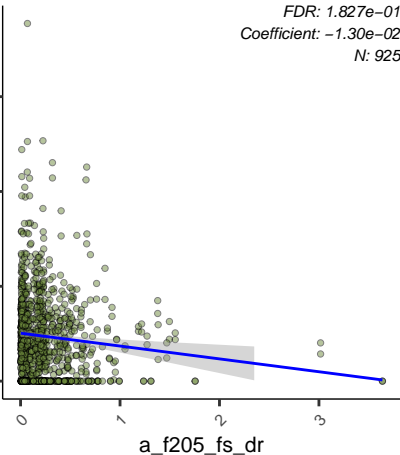
1

2

3

a_f205_fs_dr

FDR: 1.827e-01
Coefficient: -1.30e-02
N: 925



enterorhabdus_caecimuris

FDR: 1.827e-01
Coefficient: 1.93e-03
N: 925

