

GCA.900066575

FDR: 1.185e-03
Coefficient: 5.58e-01
● *N: 101*

30

20

10

0

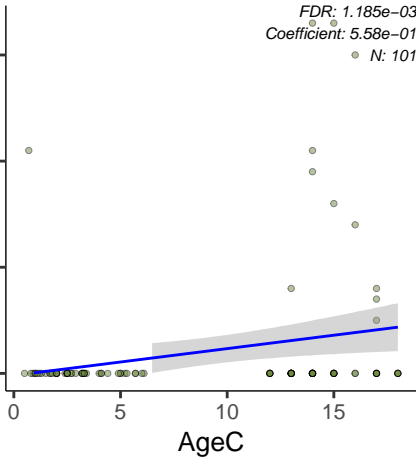
0

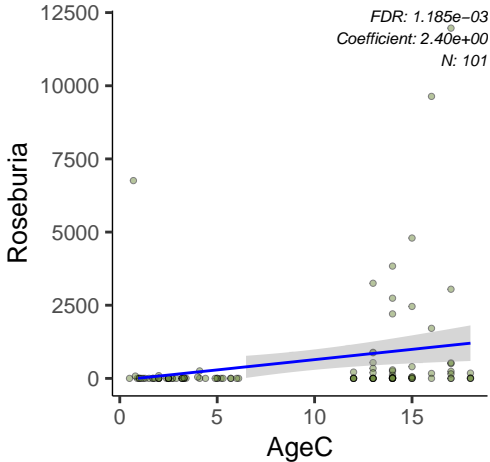
5

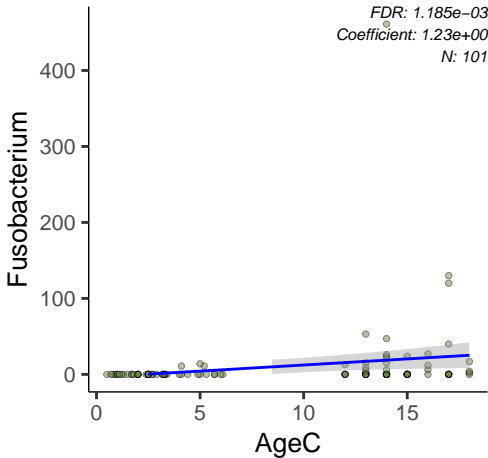
10

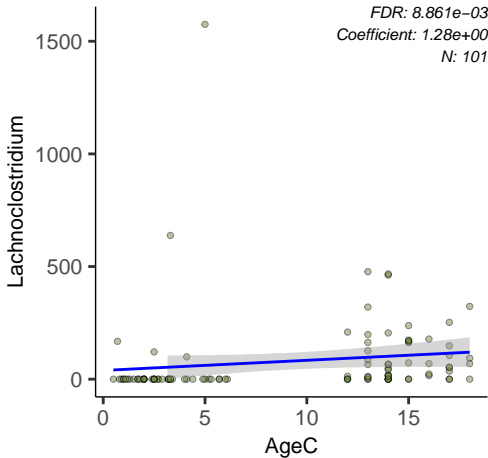
15

AgeC



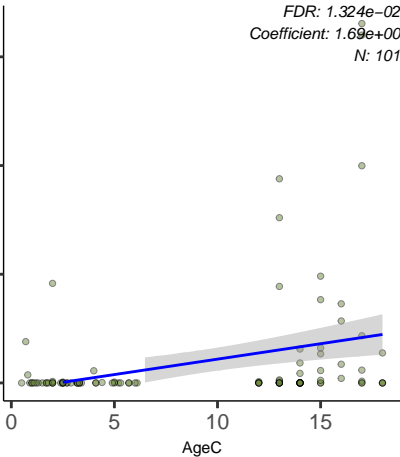


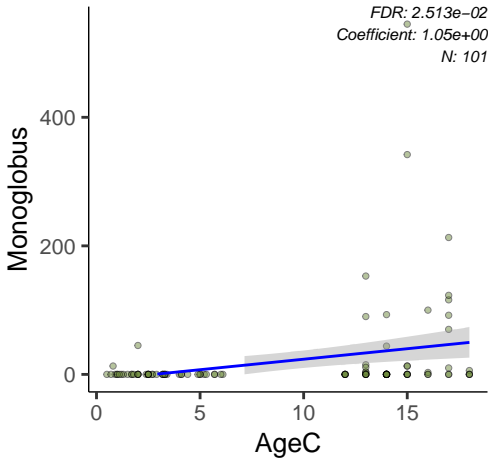




X.Eubacterium._coprostanoligenes_group

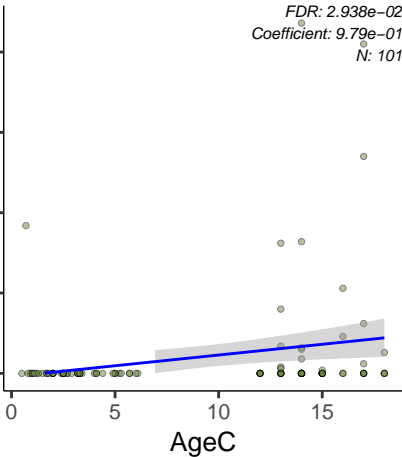
FDR: 1.324×10^{-2}
Coefficient: 1.69×10^0
N: 101

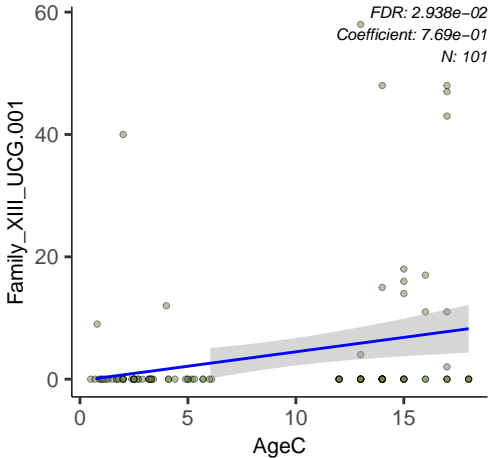


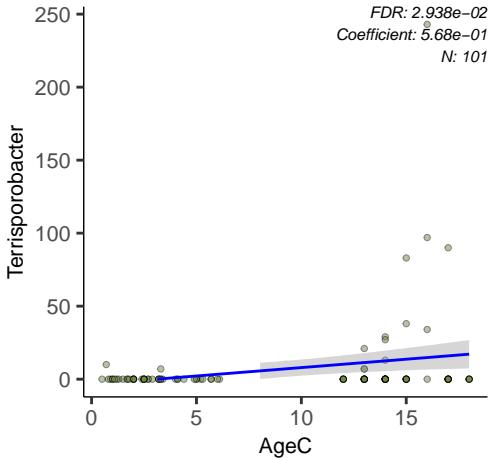


Lachnospira

FDR: 2.938e-02
Coefficient: 9.79e-01
N: 101







Desulfovibrio

FDR: 3.118e-02
Coefficient: 1.16e+00
N: 101

900

600

300

0

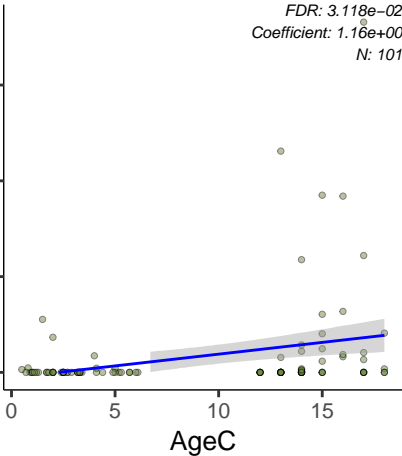
0

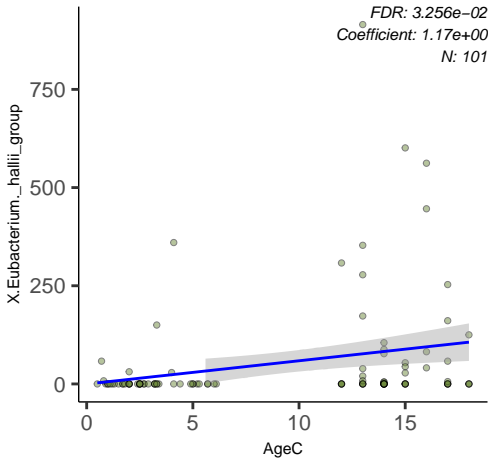
5

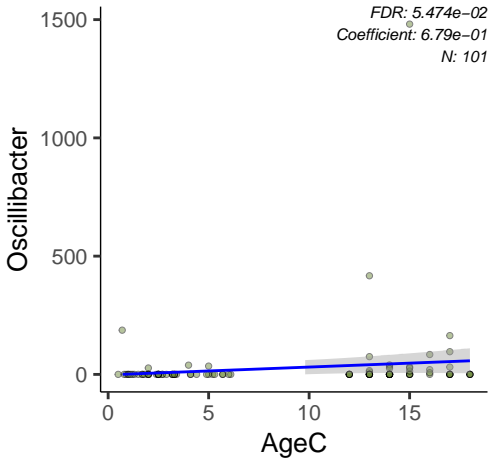
10

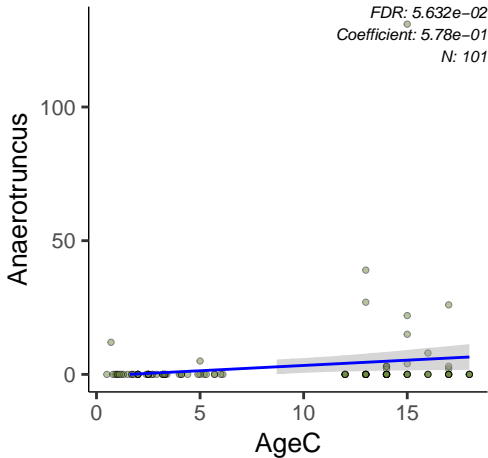
15

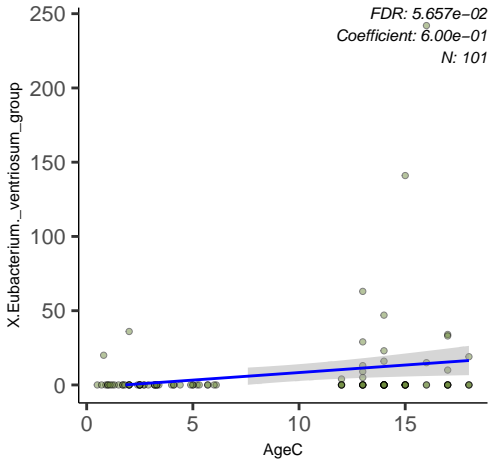
AgeC





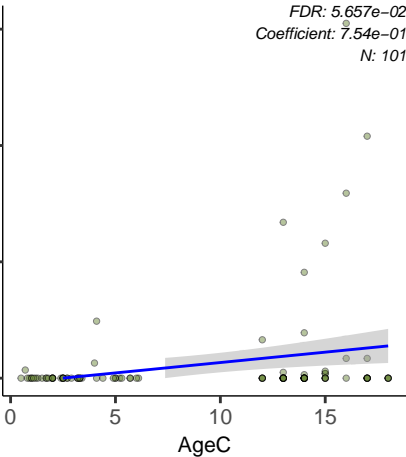


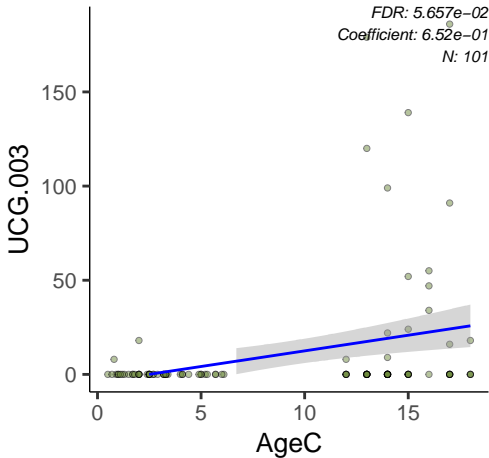


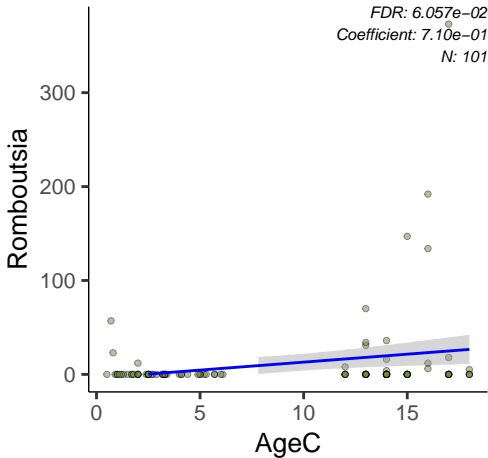


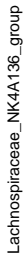
Lachnospiraceae_UCG.010

FDR: 5.657e-02
Coefficient: 7.54e-01
N: 101

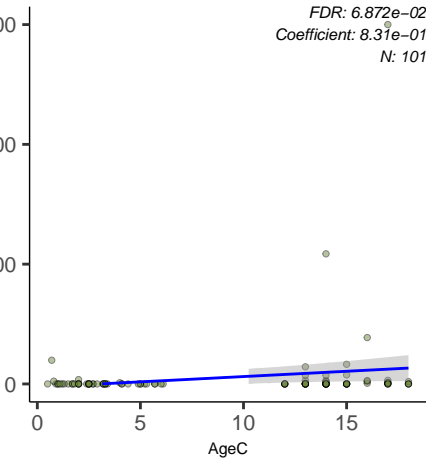


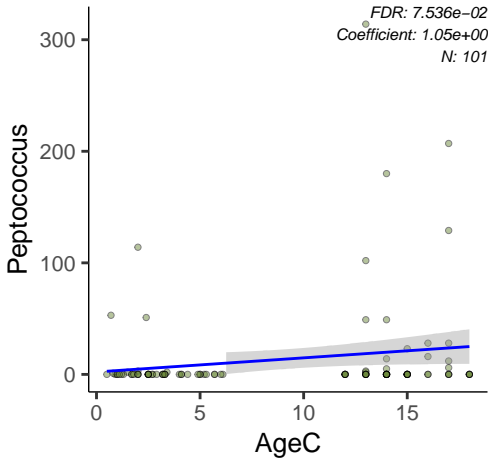


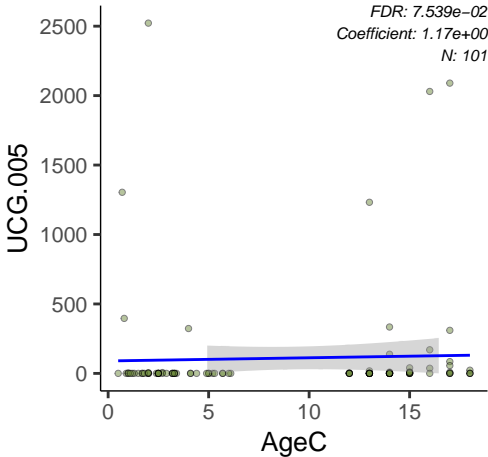




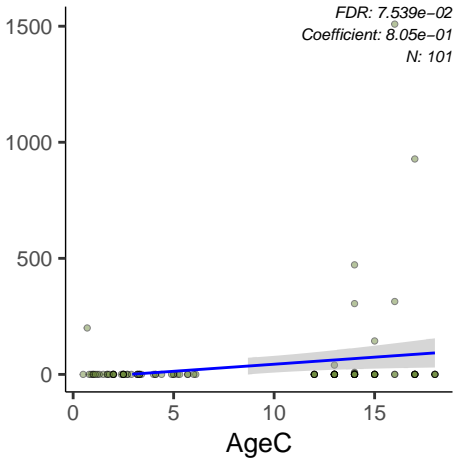
FDR: 6.872e-02
Coefficient: 8.31e-01
N: 101







Anaerovibrio



Turcibacter

