

Collinsella

FDR: 1.160e-02
Coefficient: 7.20e-01
N: 114

0

5

10

15

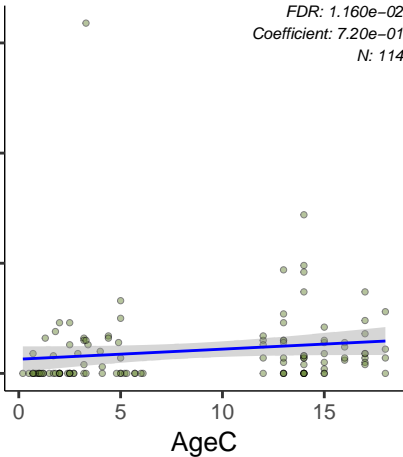
AgeC

150

100

50

0

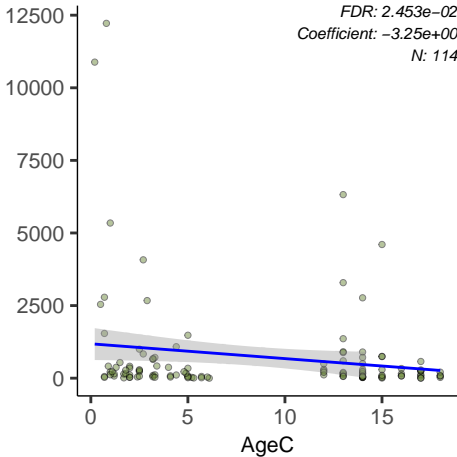


Escherichia.Shigella

FDR: 2.453e-02

Coefficient: -3.25e+00

N: 114



Mitsuokella

FDR: 4.483e-02
Coefficient: 7.86e-01
N: 114

2000

1000

0

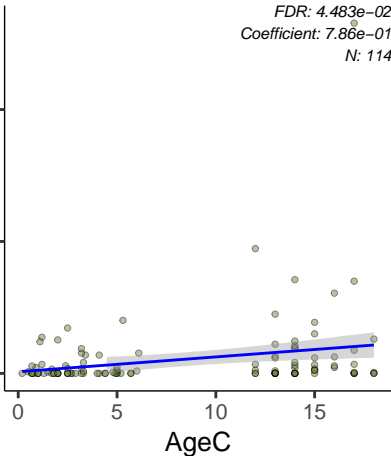
0

5

10

15

AgeC



Bifidobacterium

30000
20000
10000
0

0

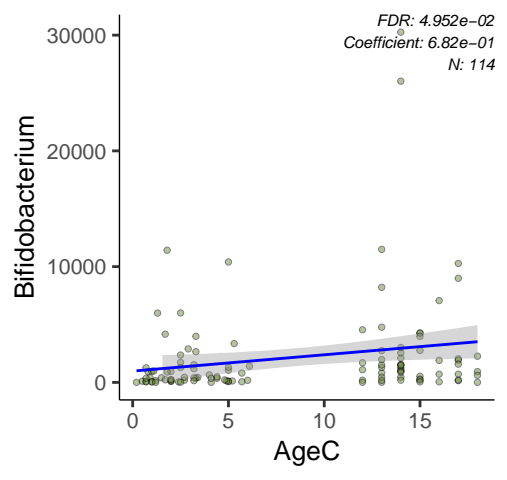
5

10

15

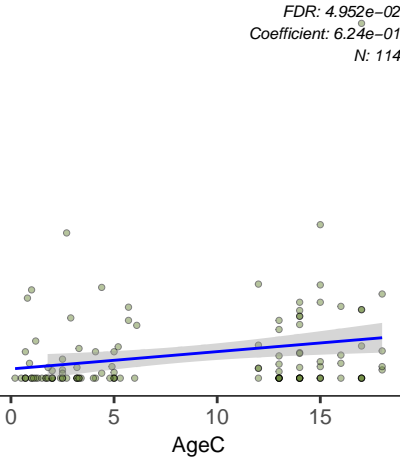
AgeC

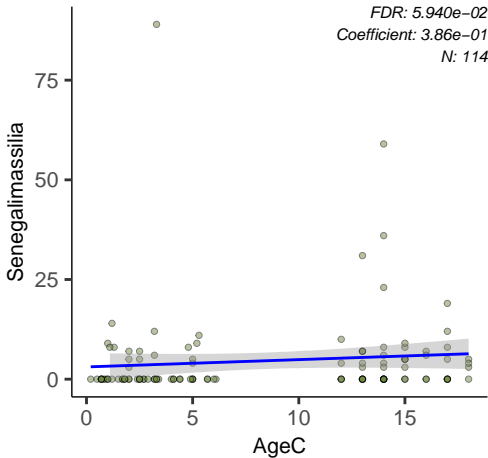
FDR: 4.952e-02
Coefficient: 6.82e-01
N: 114

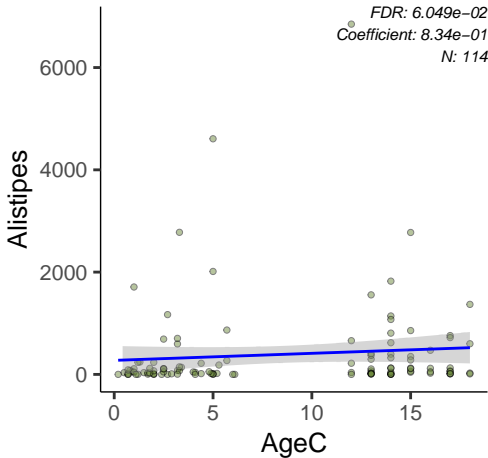


Lachnospiraceae_UCG.001

FDR: 4.952e-02
Coefficient: 6.24e-01
N: 114







Succinivibrio

FDR: 6.049e-02
Coefficient: 6.60e-01
N: 114

1000

500

0

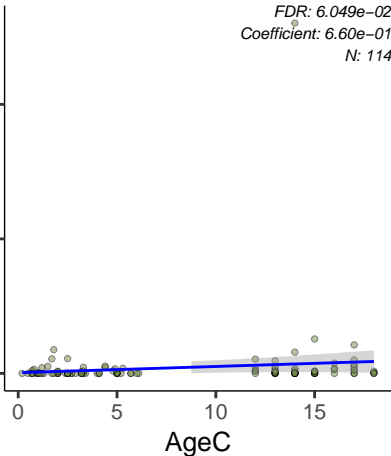
0

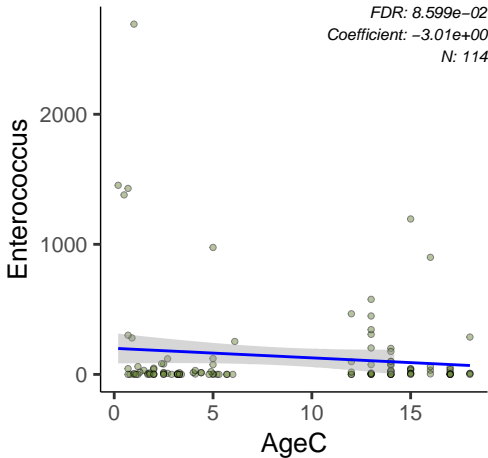
5

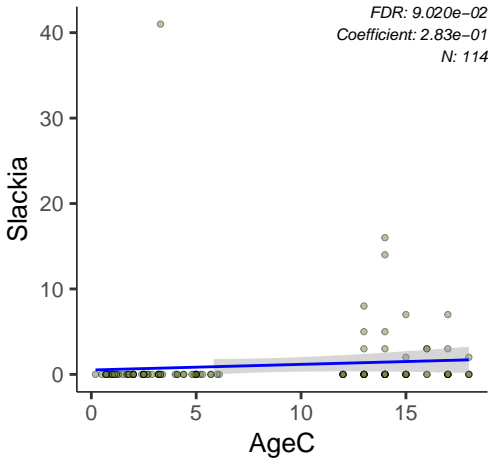
10

15

AgeC

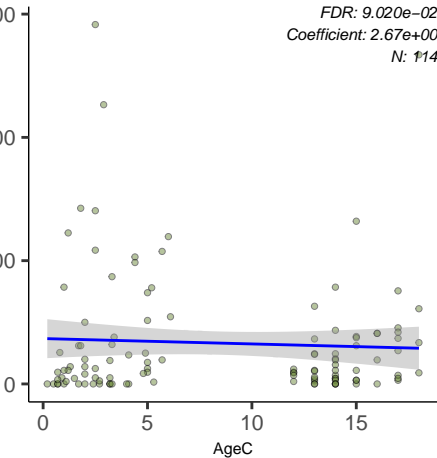






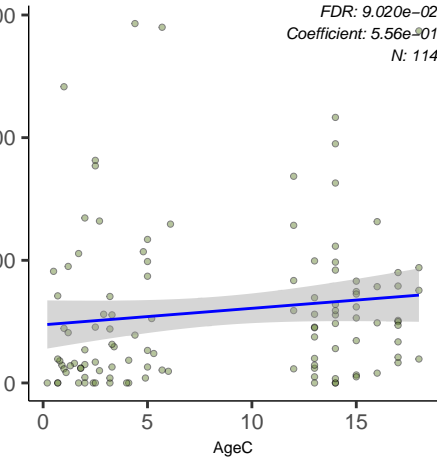
X.Eubacterium._ventriosum_group

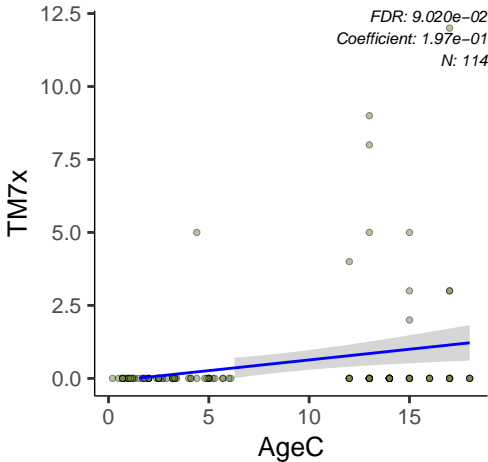
FDR: 9.020e-02
Coefficient: 2.67e+00
N: 114



Lachnospiraceae_NK4A136_group

FDR: 9.020e-02
Coefficient: 5.56e-01
N: 114





Peptoniphilus

FDR: 9.063e-02
Coefficient: 3.10e-01
N: 114

20
10
0

0

5

10

15

AgeC

