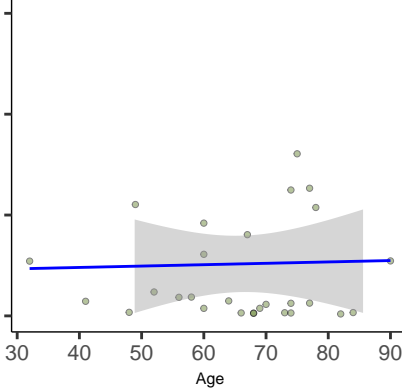
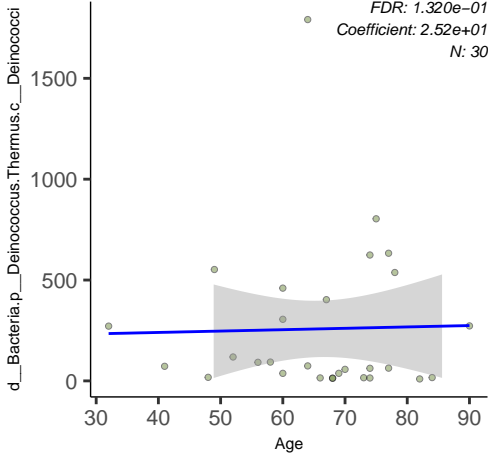
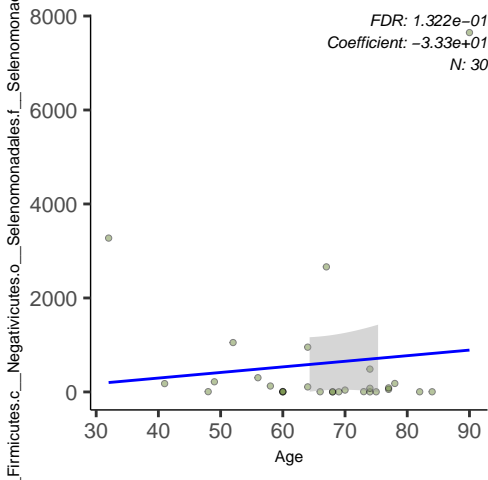


d_Bacteria.p_Deinococcus.Thermus

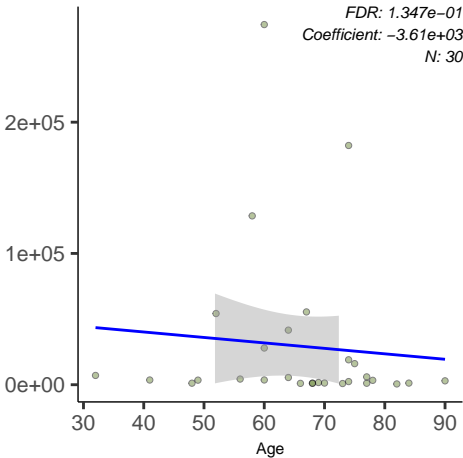
FDR: 1.320e-01
Coefficient: 2.52e+01
N: 30

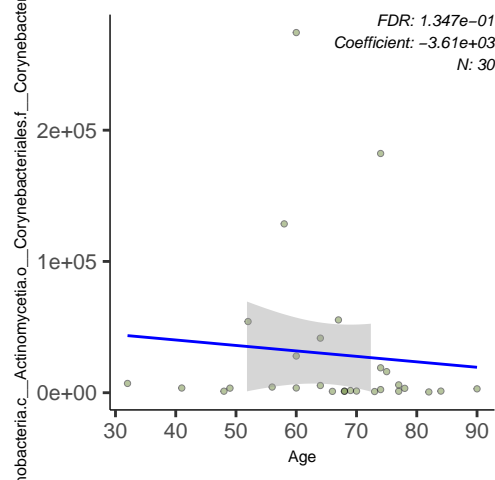


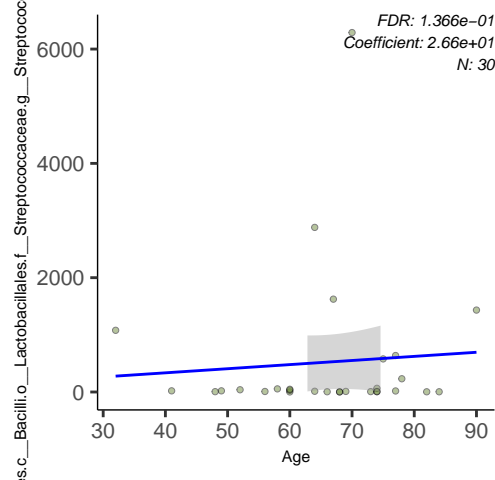


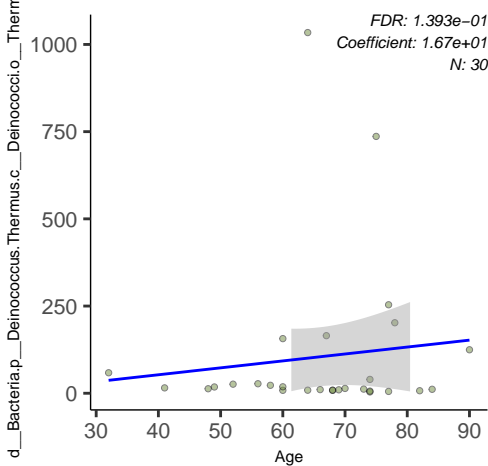


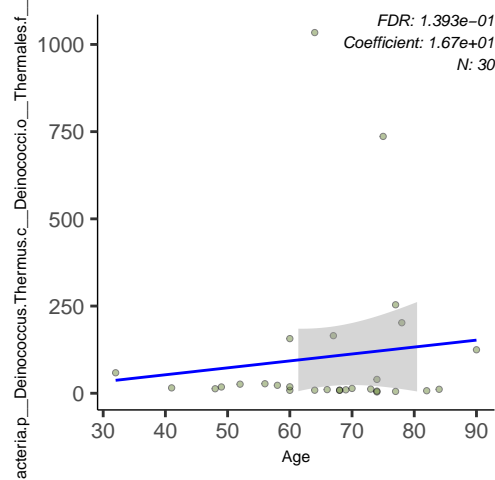
ia.p_Actinobacteria.c_Actinomycetia.o_Corynebacteriales.f_Co

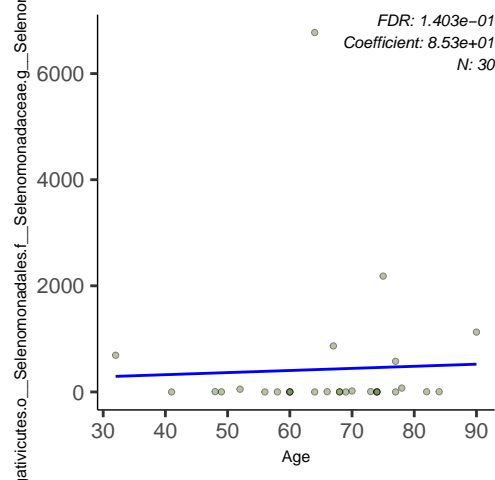


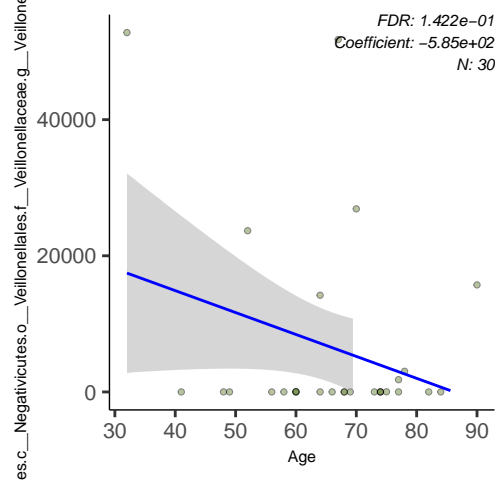


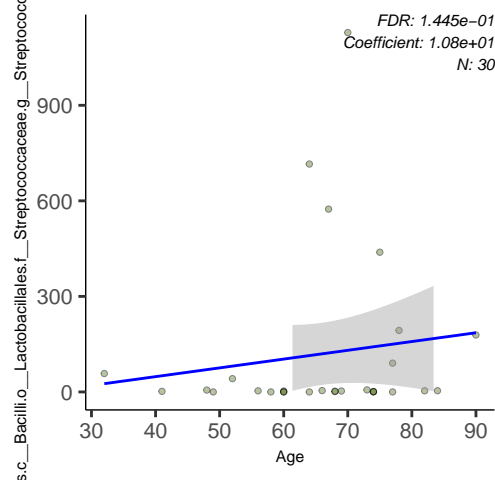


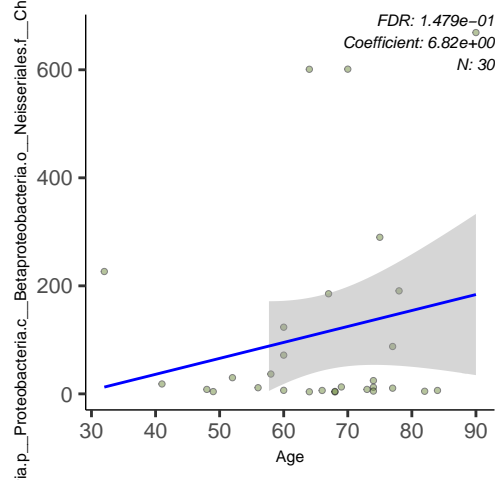


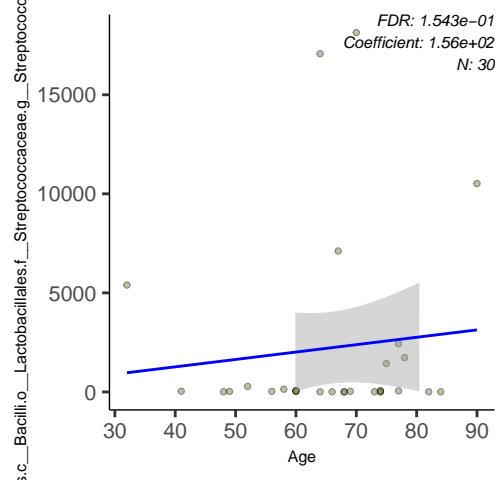


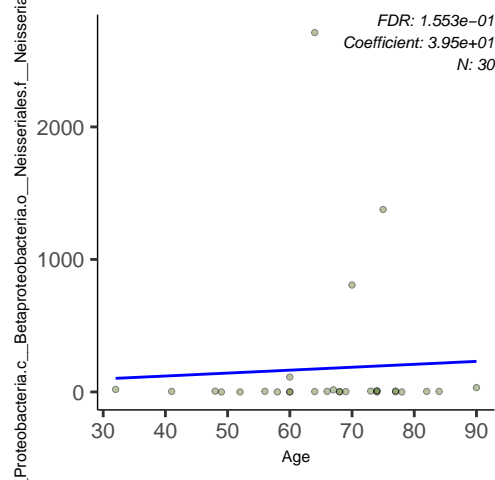


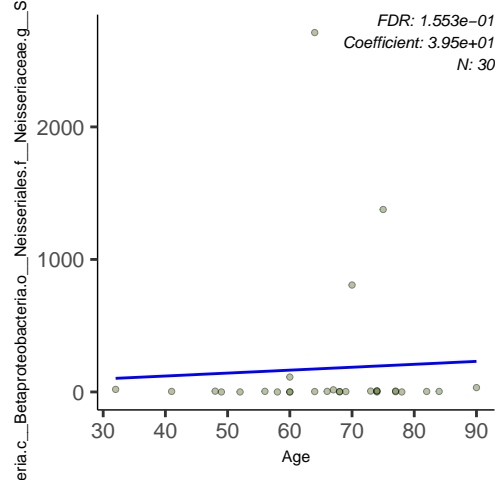


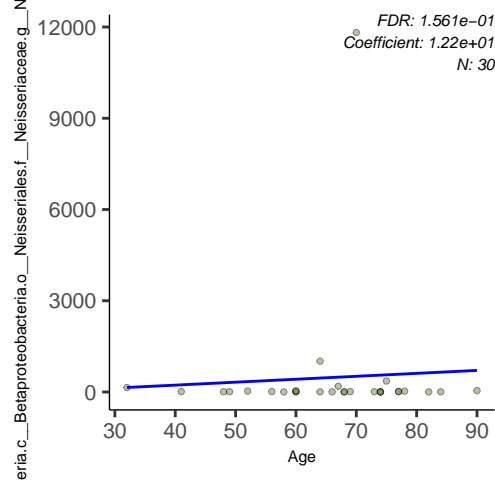


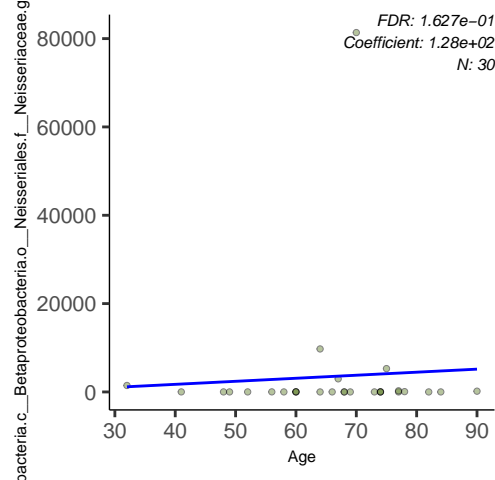


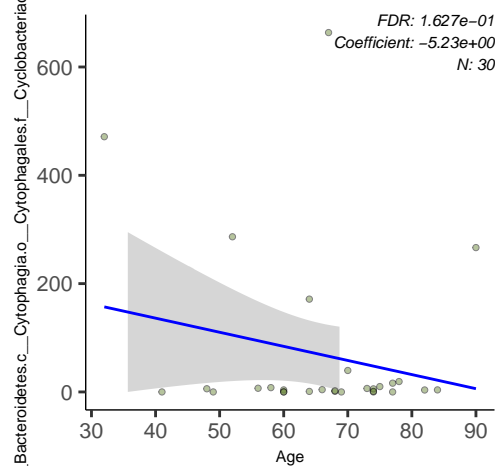


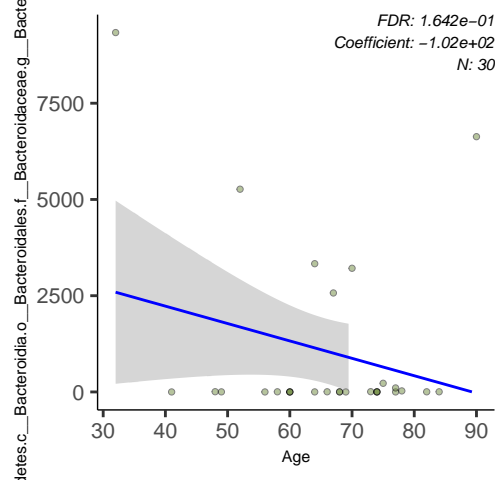


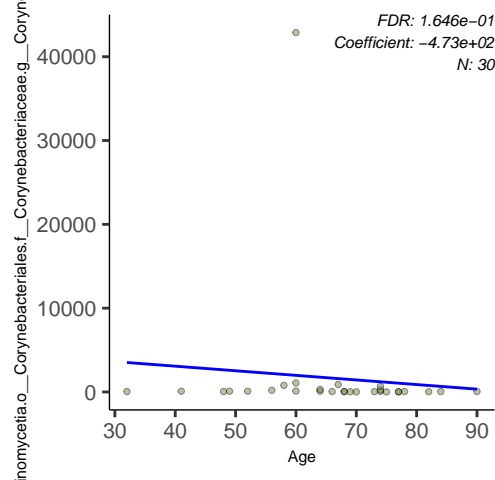


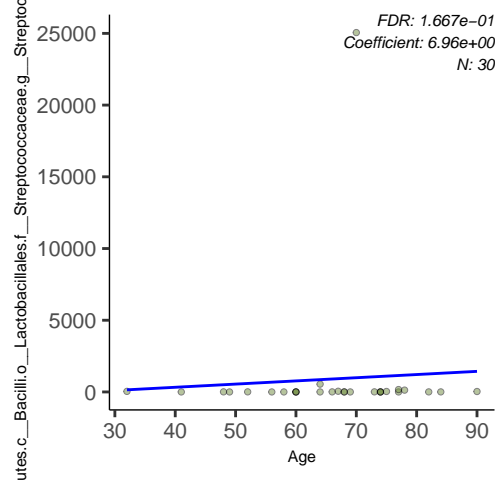


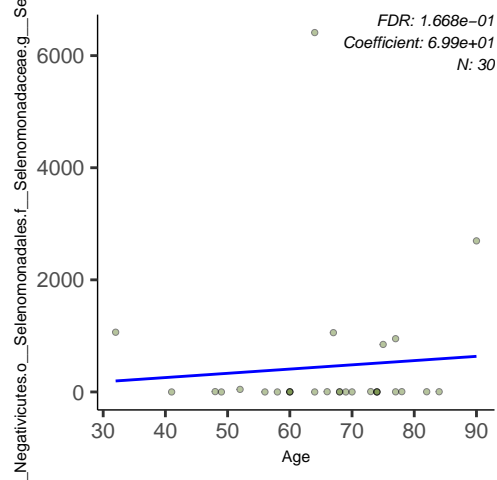


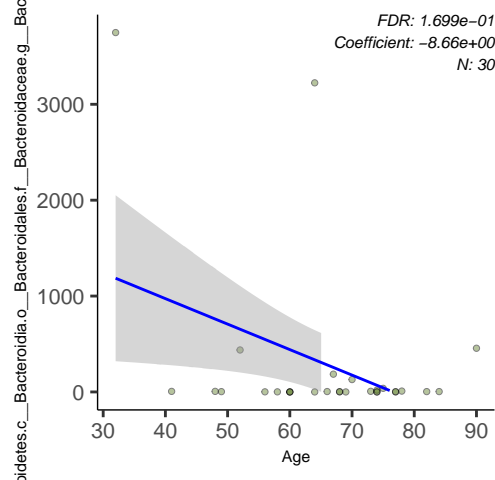


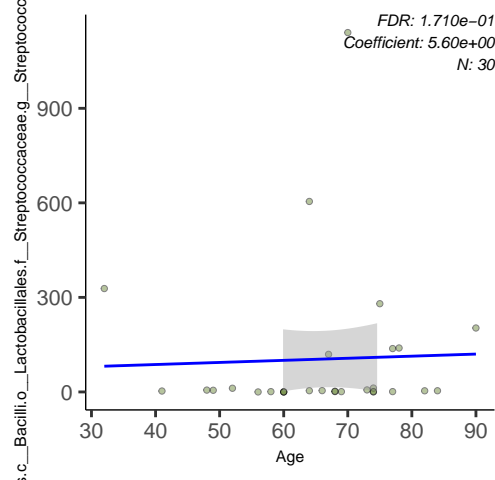


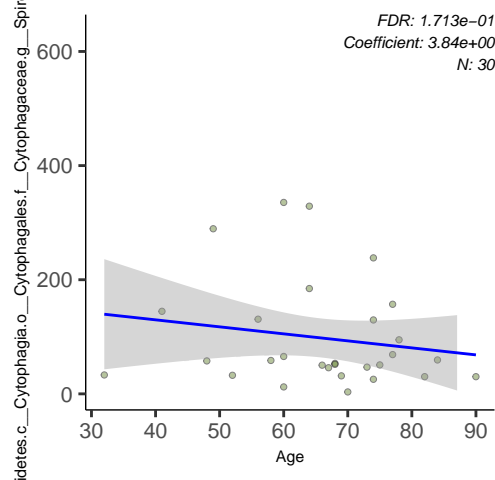


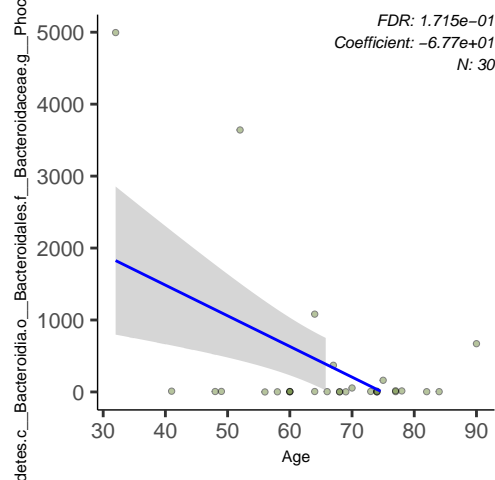


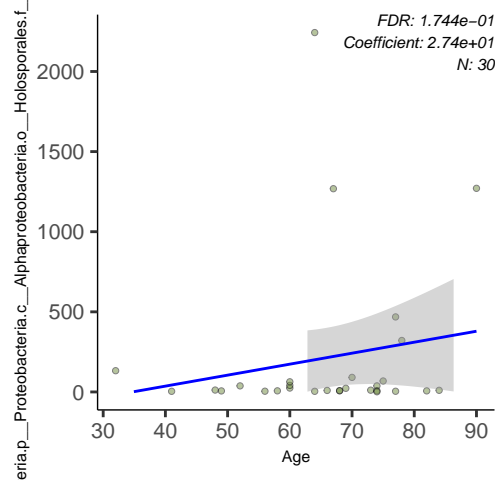


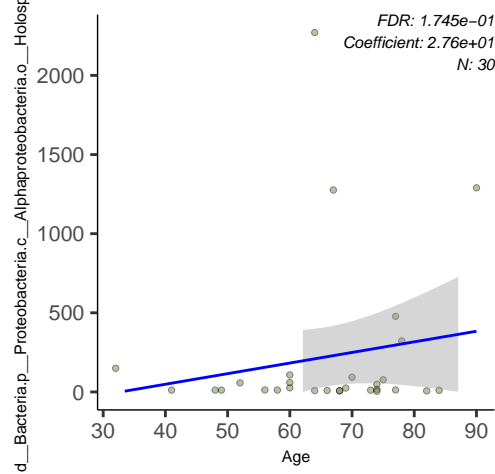


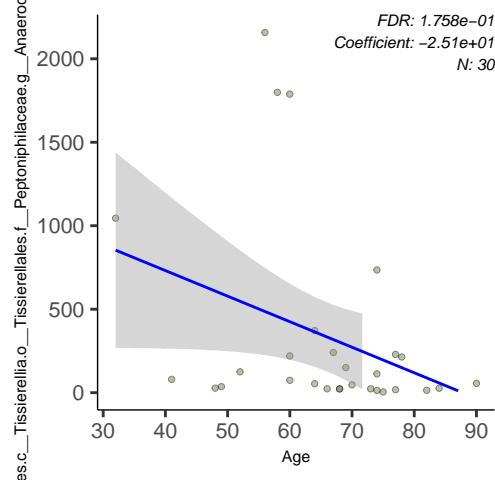


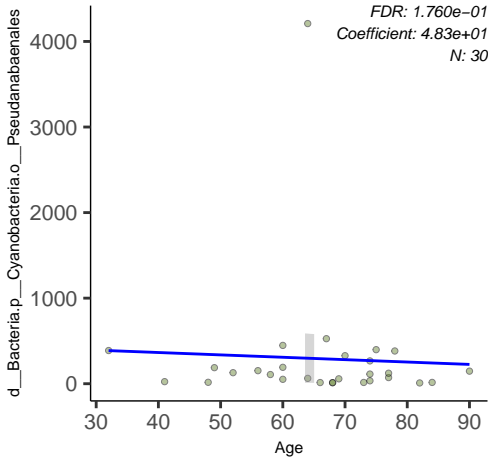


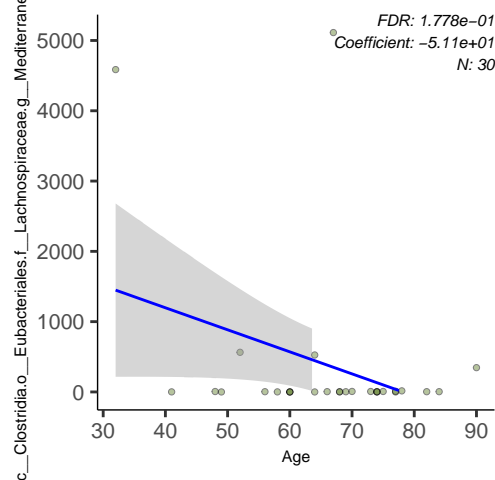


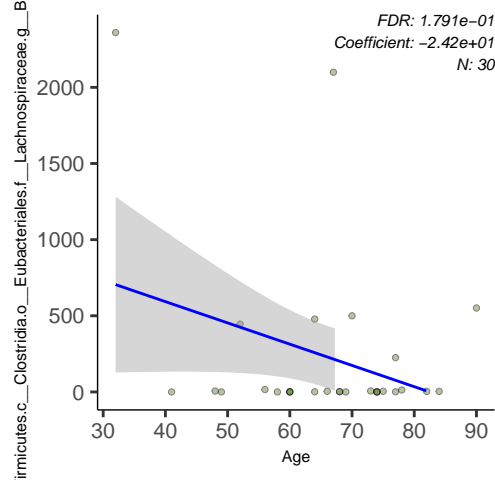


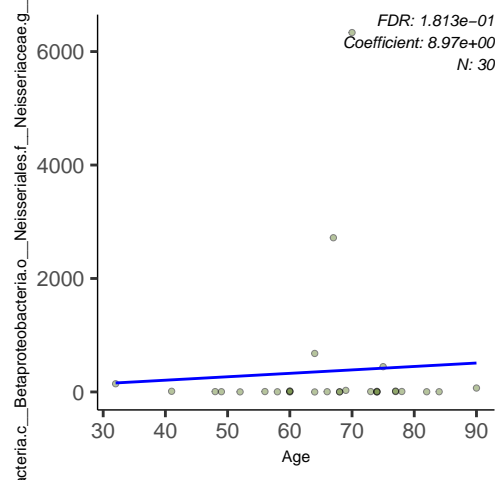


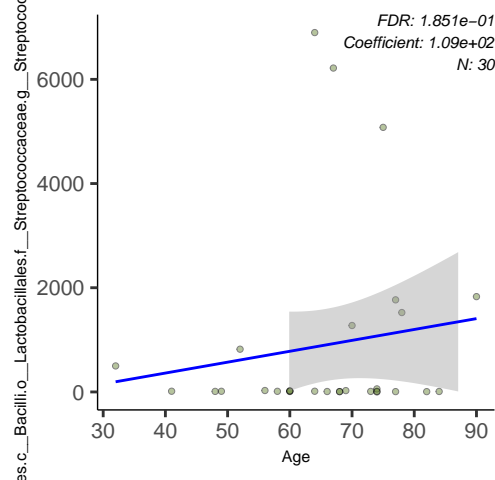


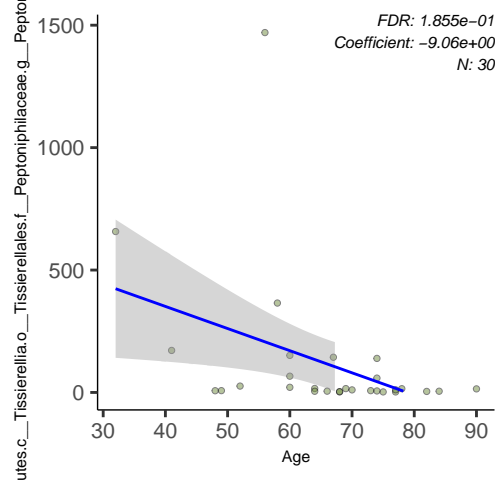


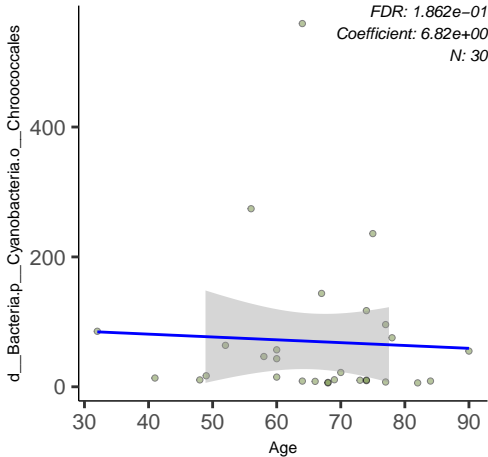


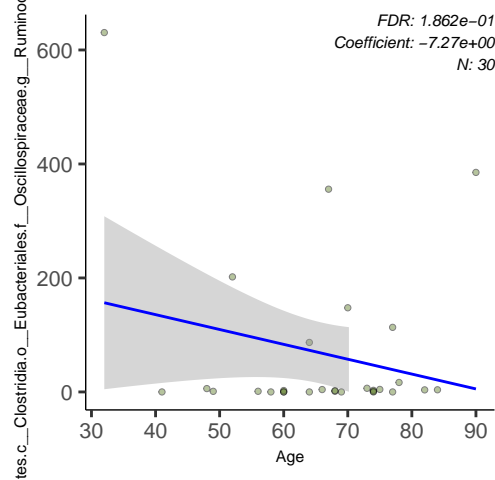


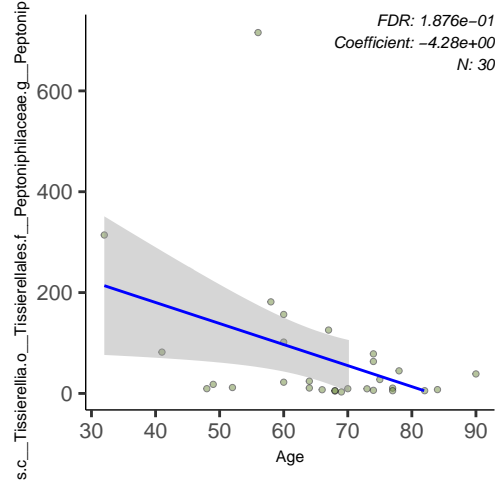


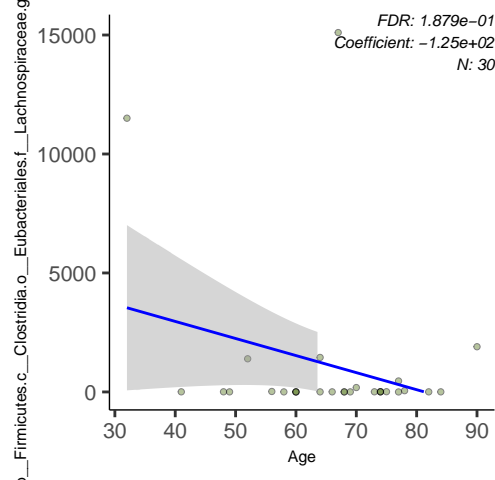


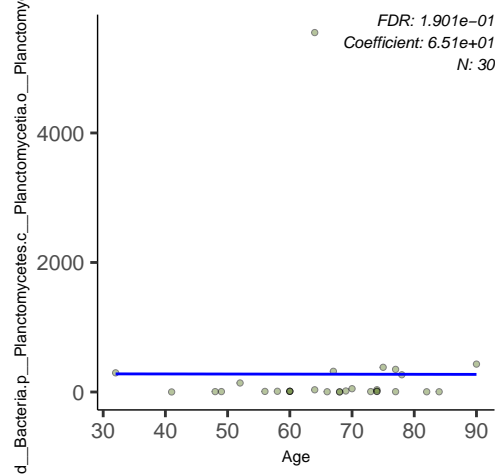


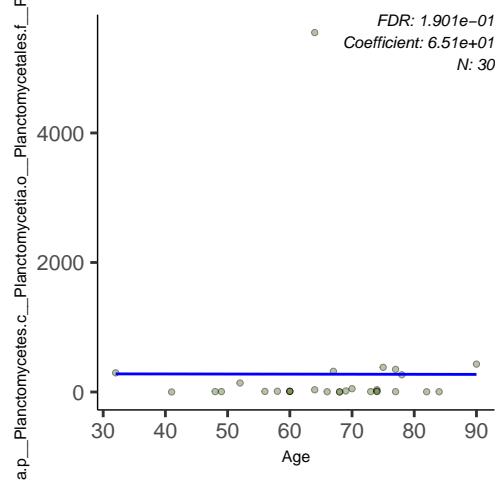


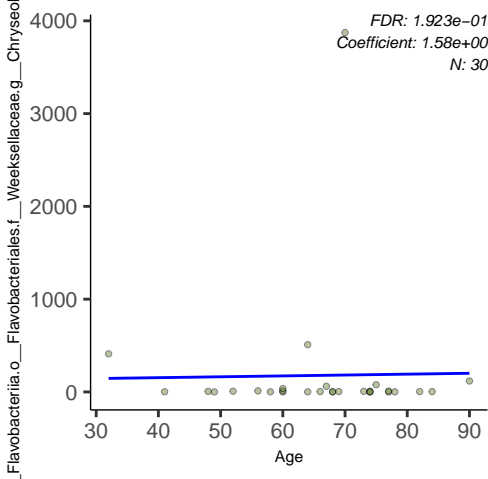


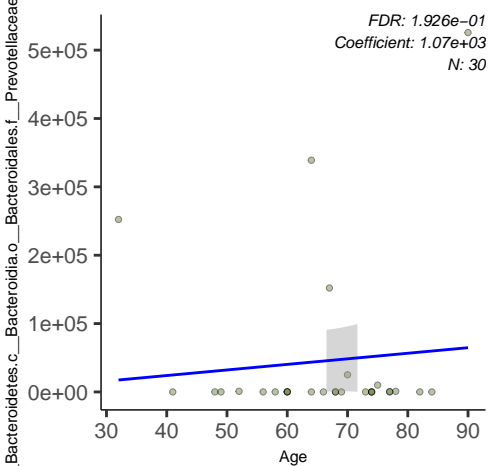


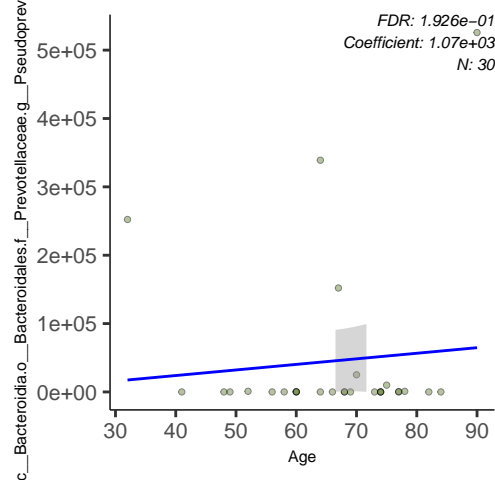


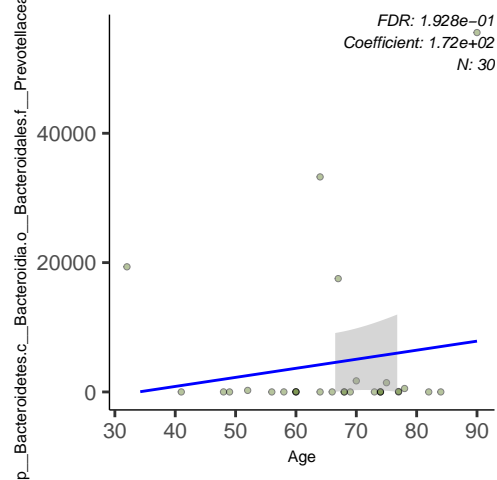


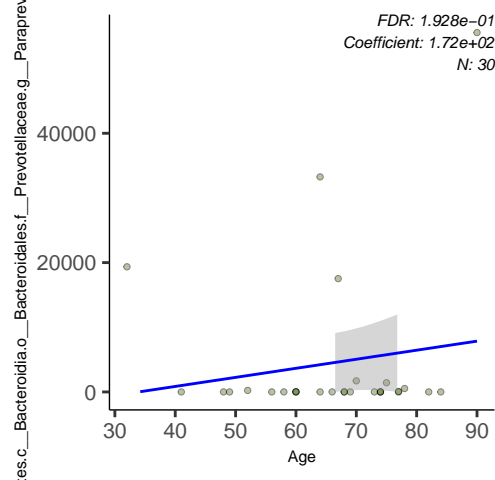


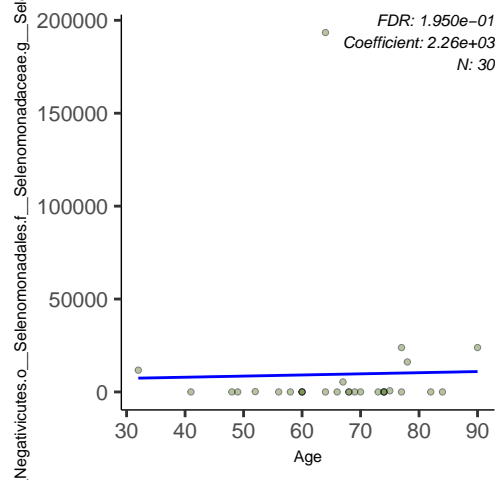


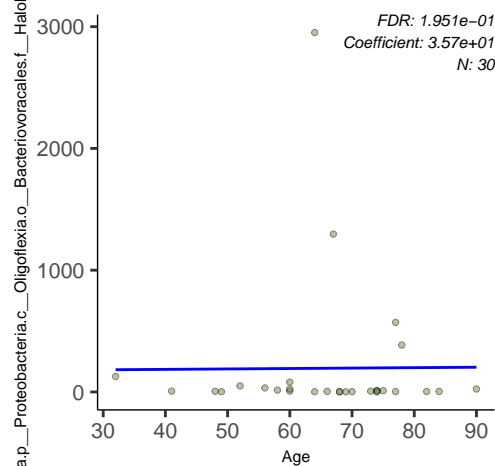


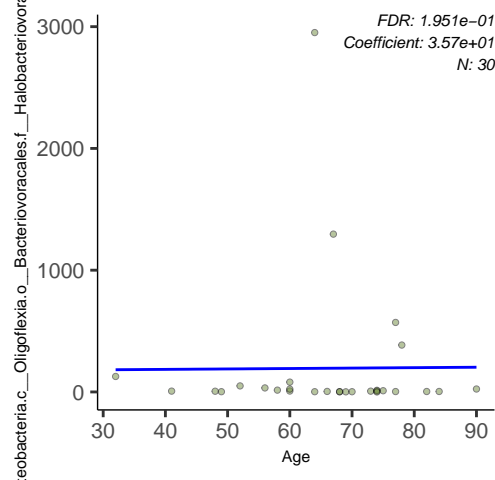


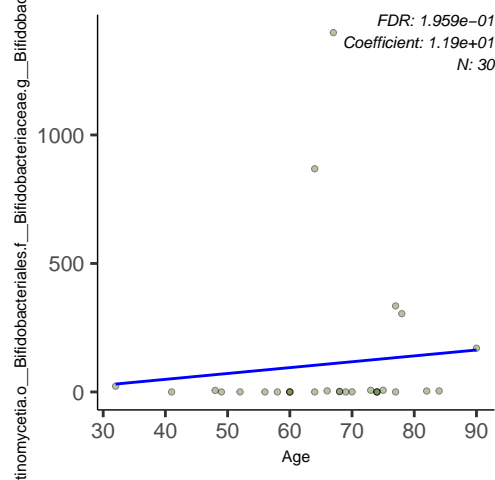


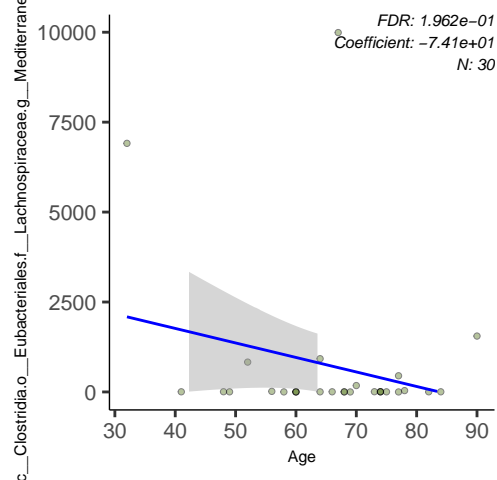


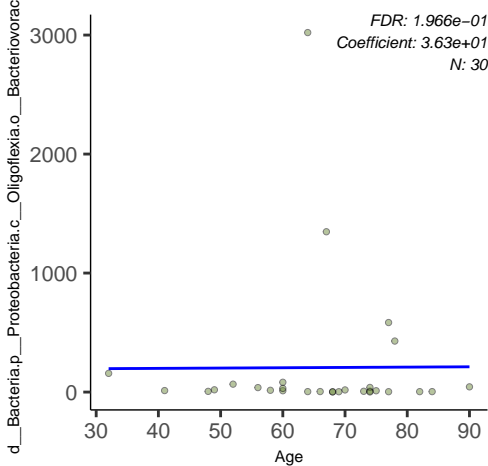




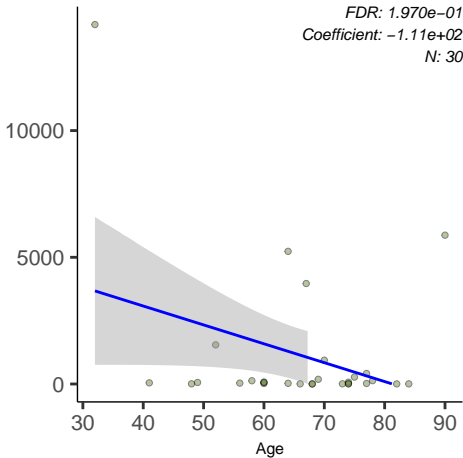


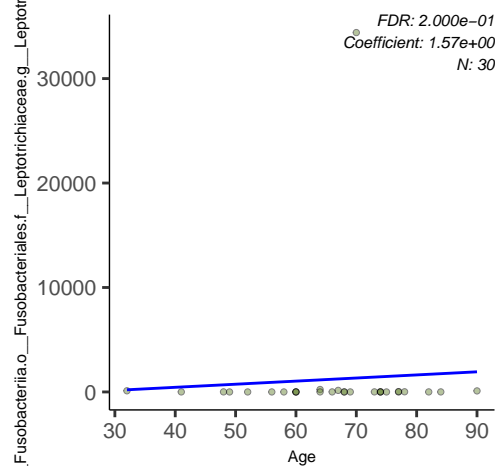


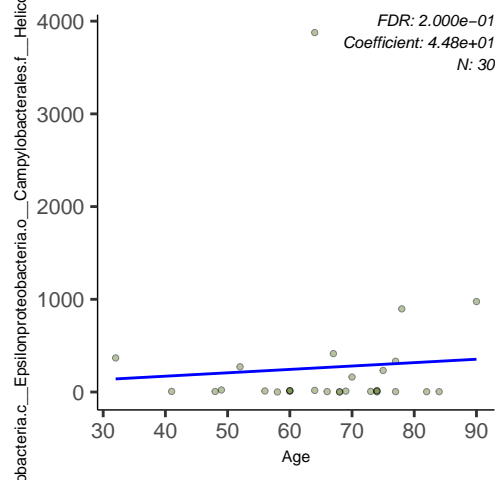


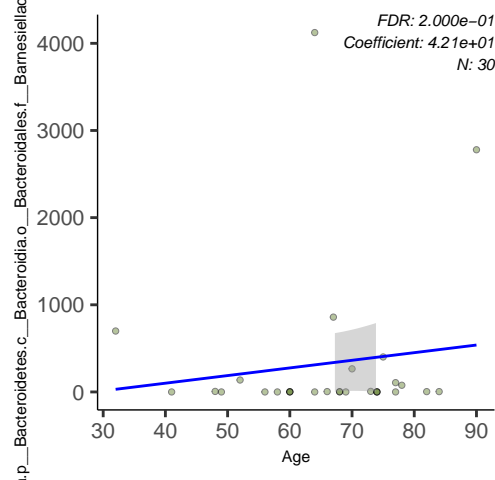


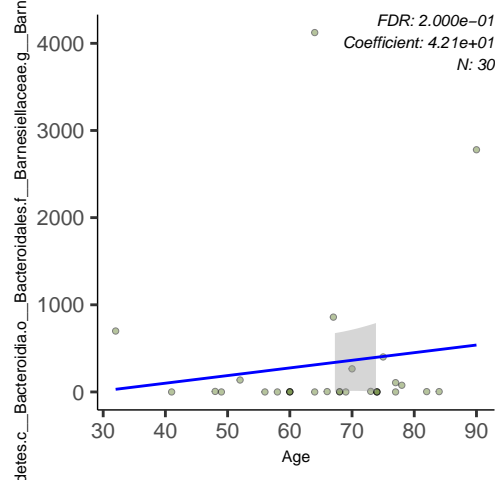
acteroidetes.c__Flavobacteriia.o__Flavobacteriales.f__Flavobacteri











Clostridia.o__Eubacteriales.f__Eubacteriales.Family.XIII..Incertae.S

FDR: 2.000e-01
Coefficient: 7.86e+01
N: 30

6000

4000

2000

0

30

40

50

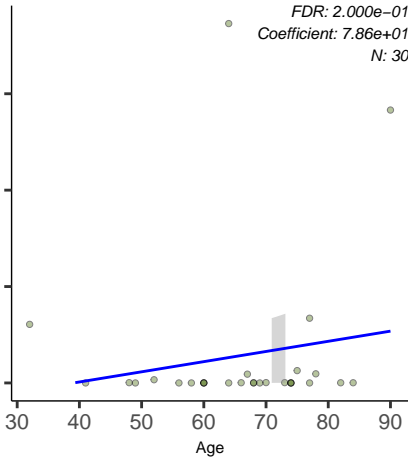
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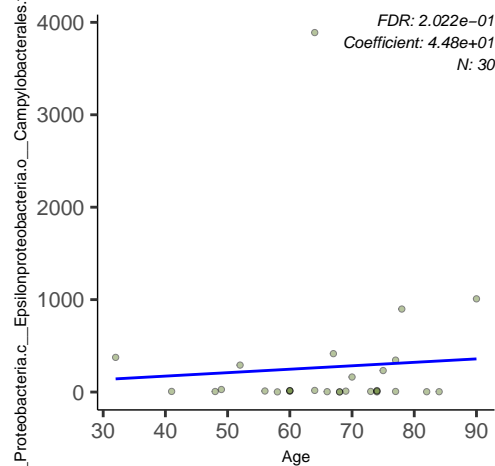
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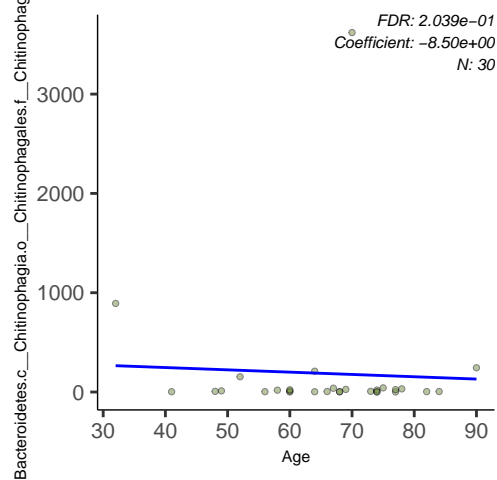
80

90

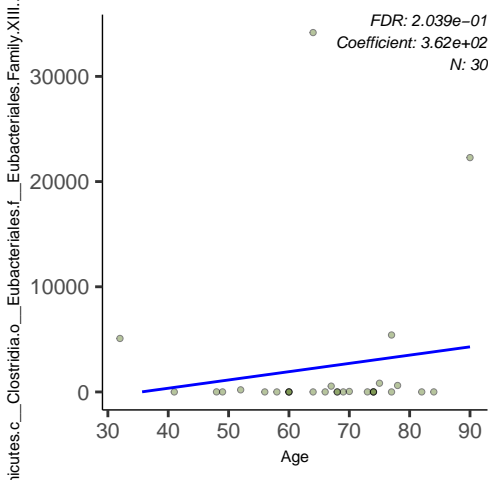
Age

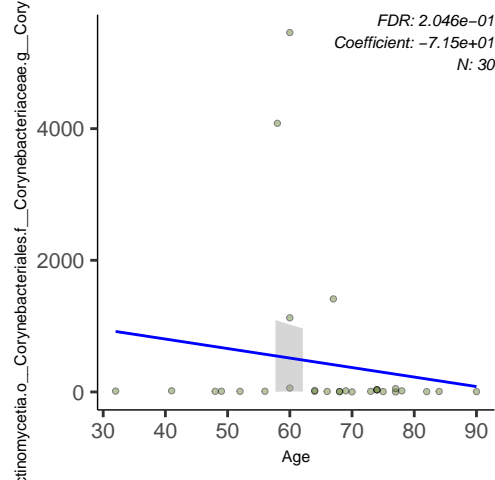


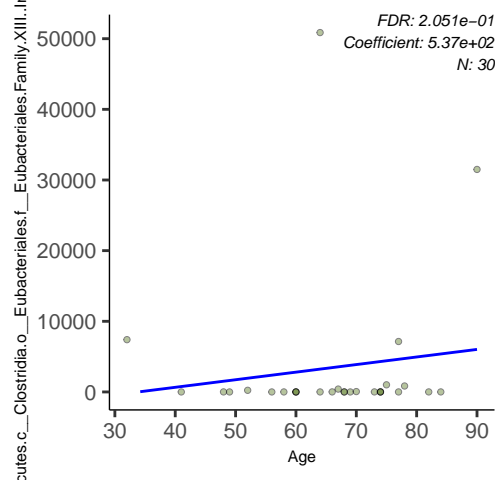


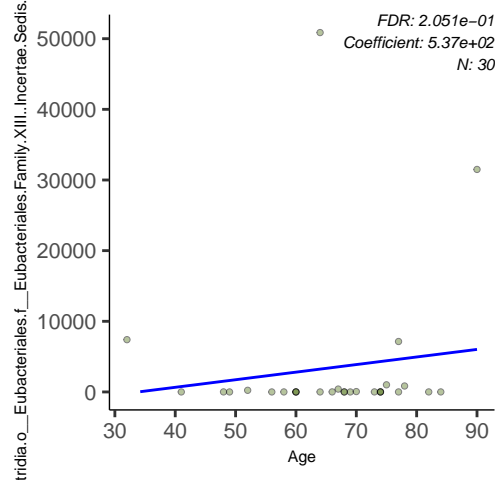


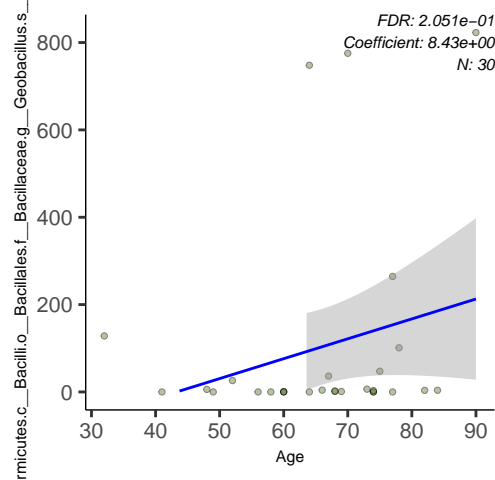
phylum: Bacteroidetes, class: Sphingobacteriia, order: Sphingobacteriales, family: Sphingobacteriaceae, genus: Sphingobacterium, species: Sphingobacterium

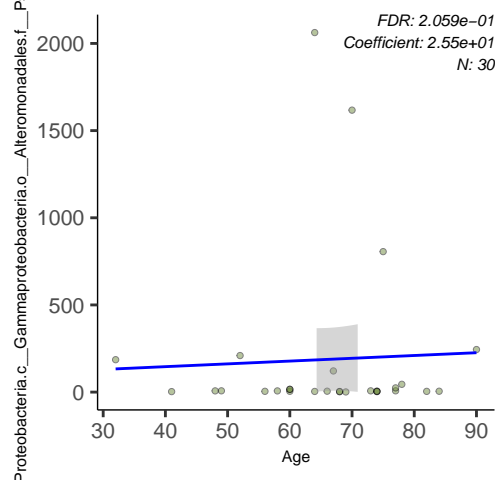


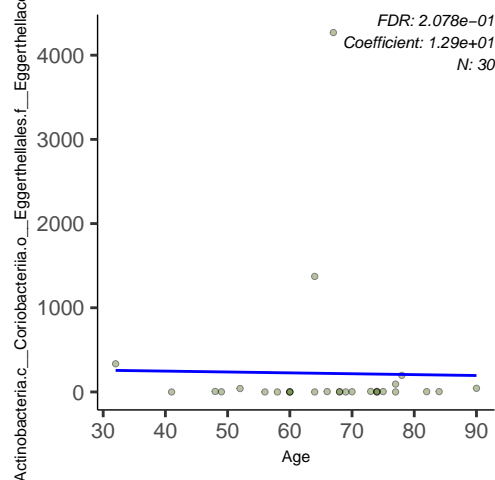


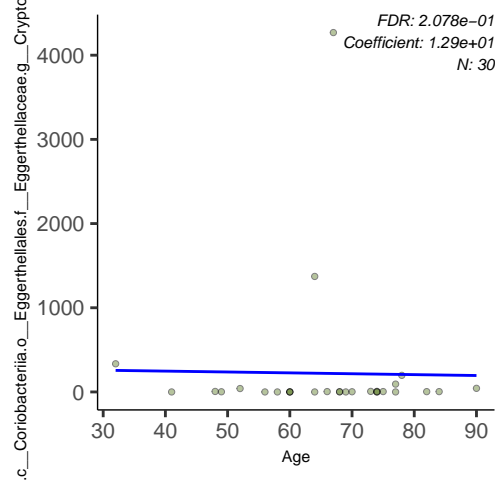


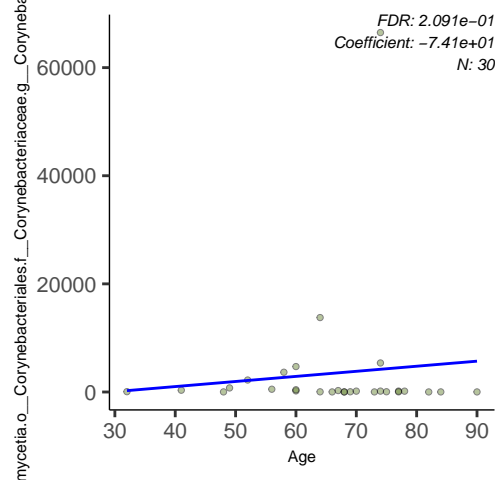


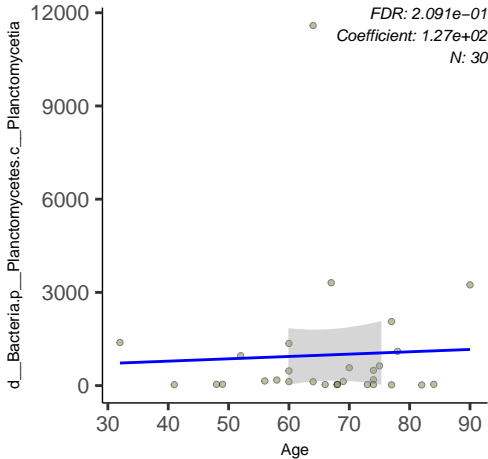


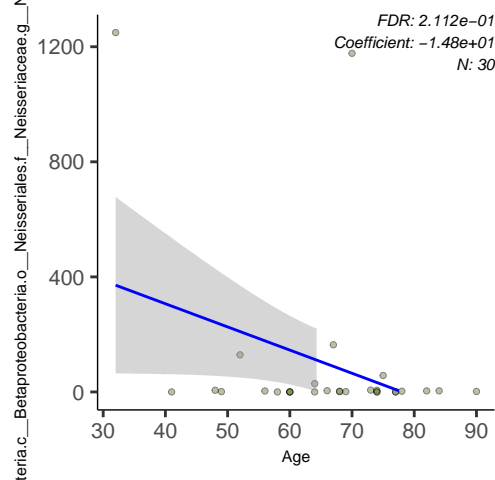










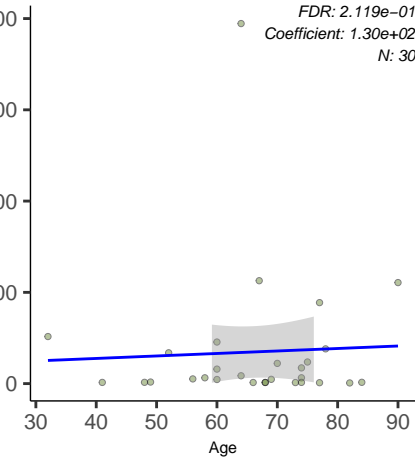


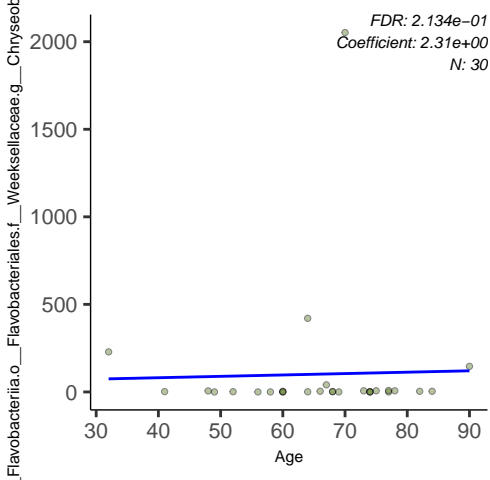
d_Bacteria.p_Planctomycetes

FDR: 2.119e-01

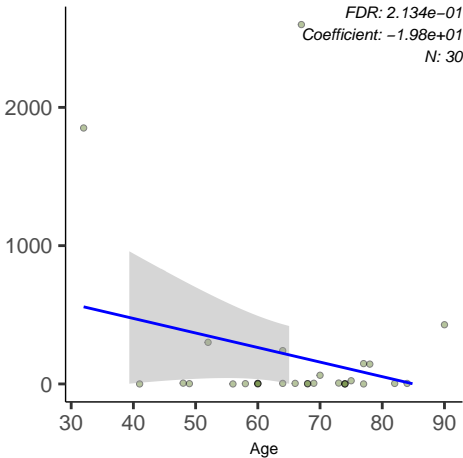
Coefficient: 1.30e+02

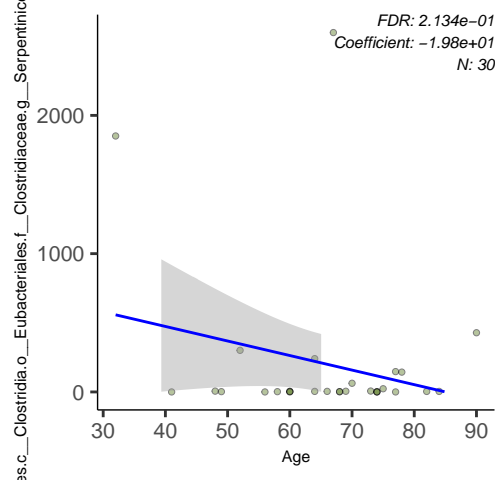
N: 30

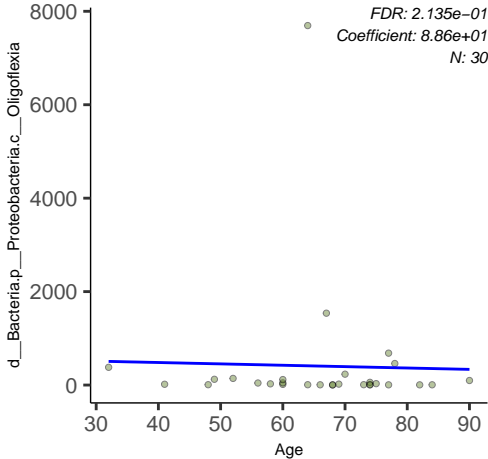


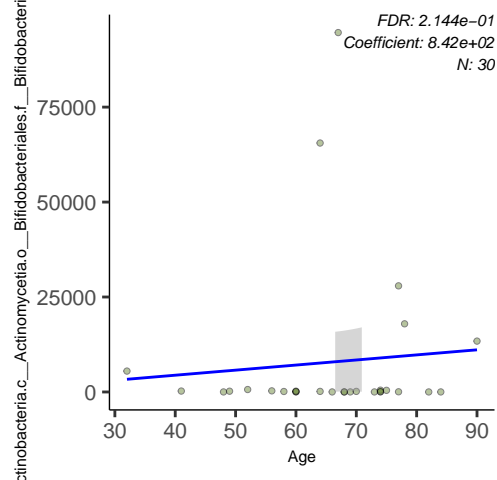


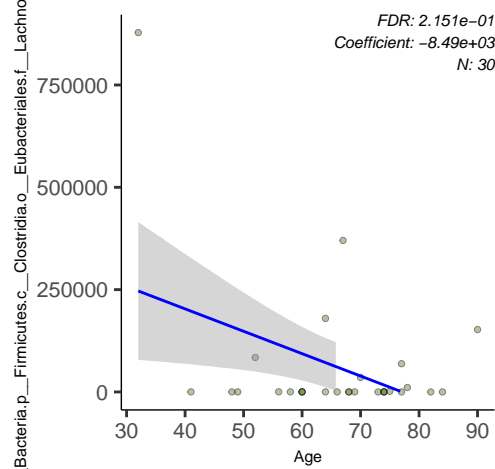
ia.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Clostridiaceae.g



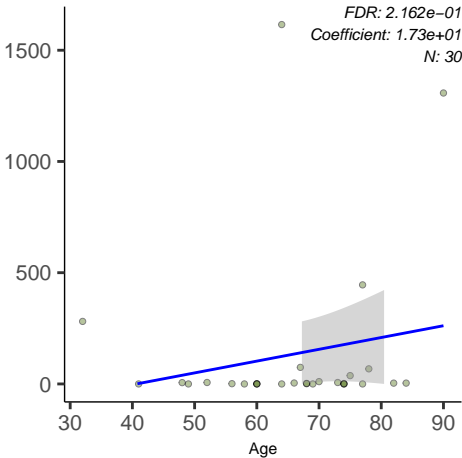


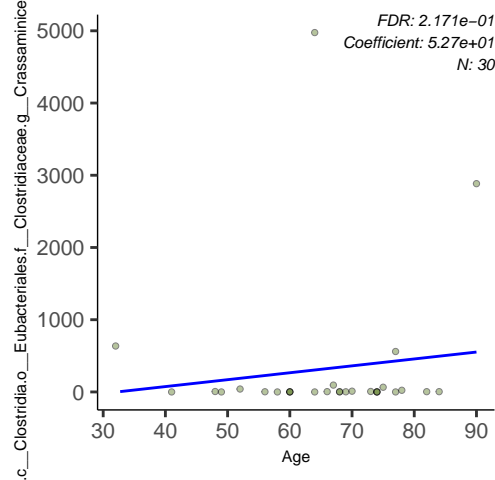


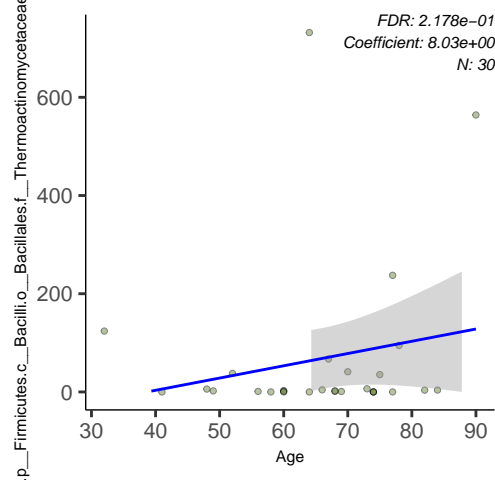


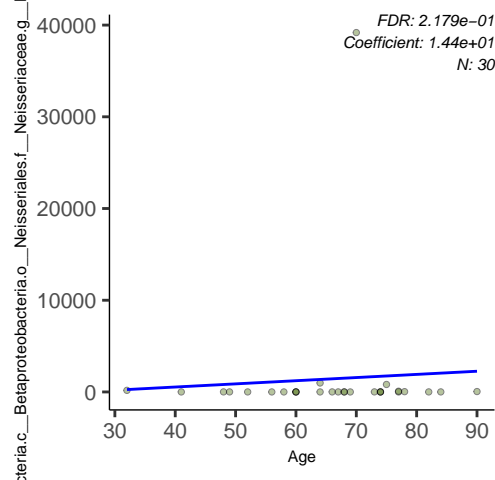


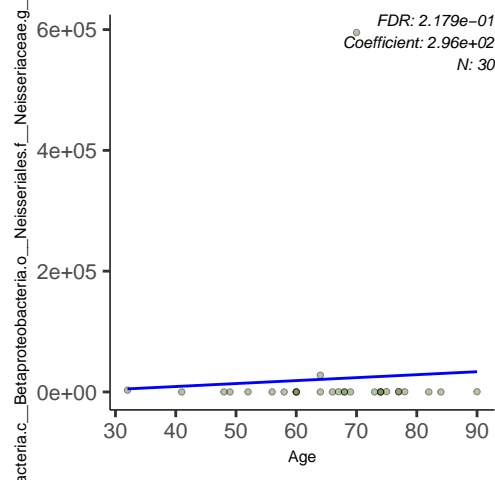
Clostridia.o__Eubacteriales.f__Eubacteriales.Family.XIII..Incertae.Se

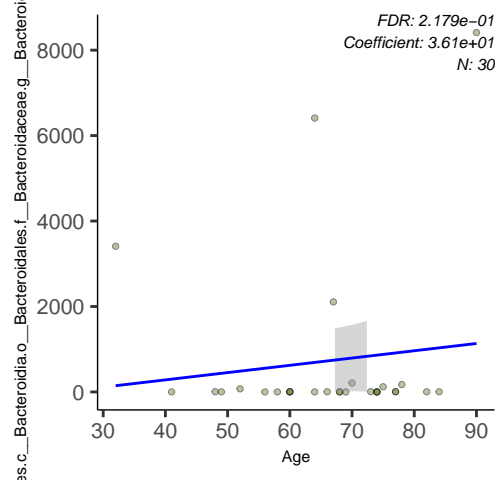


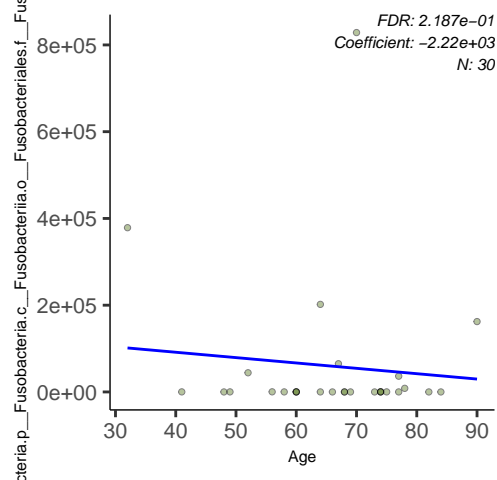


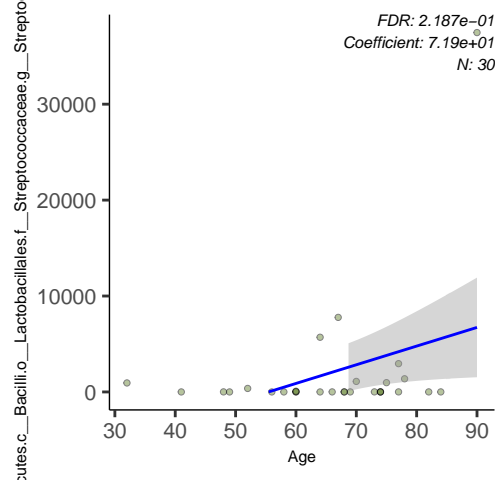


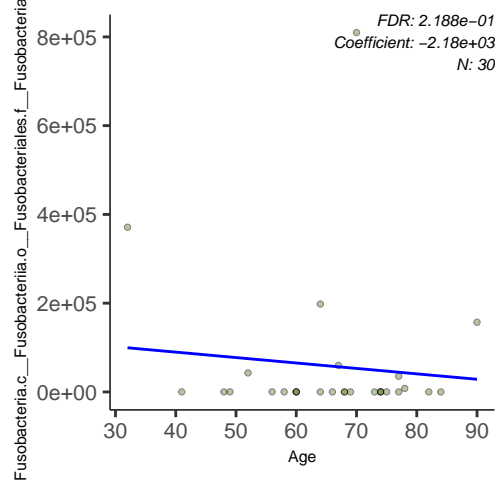


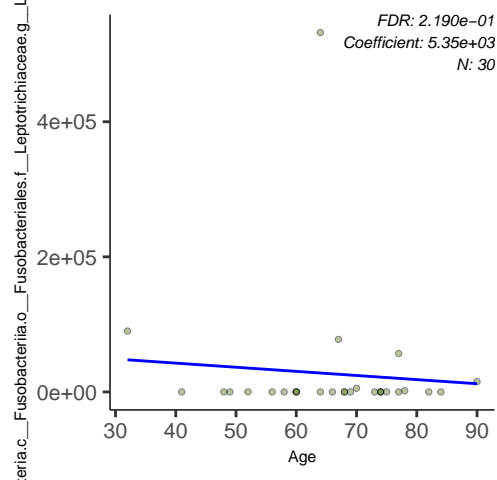


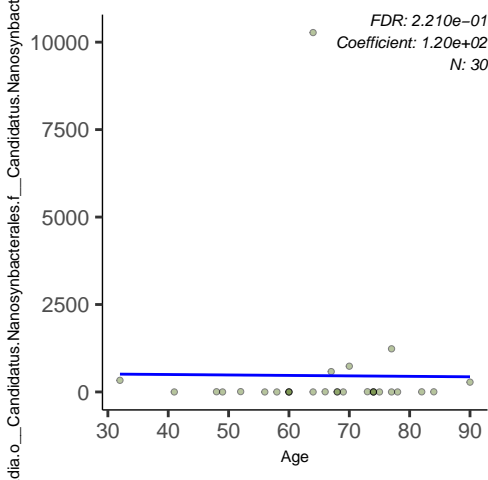


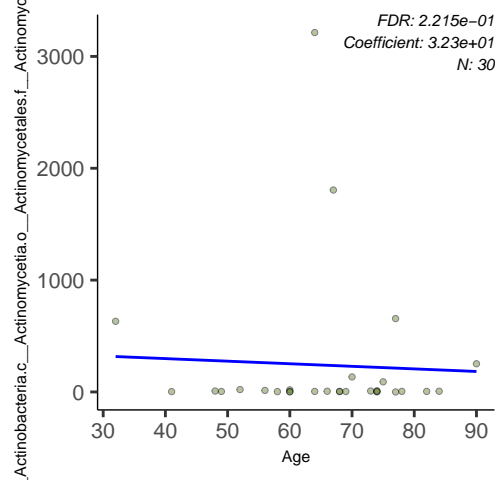




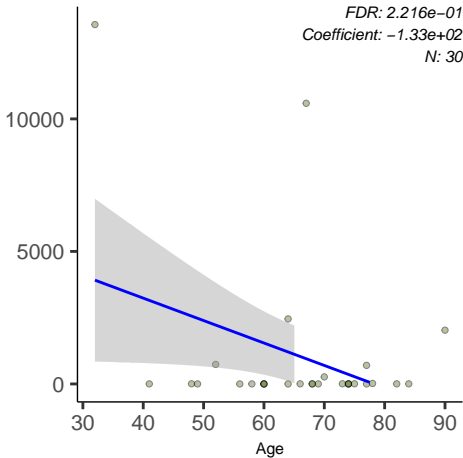


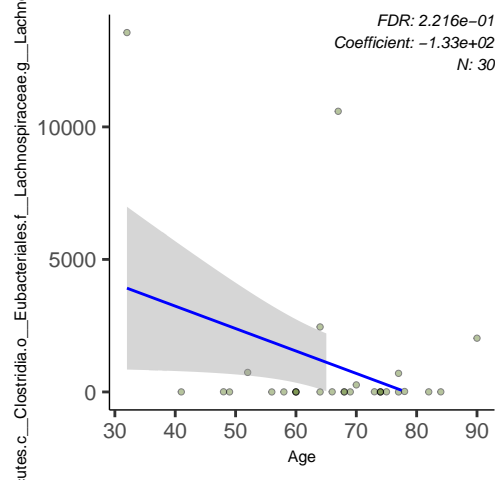


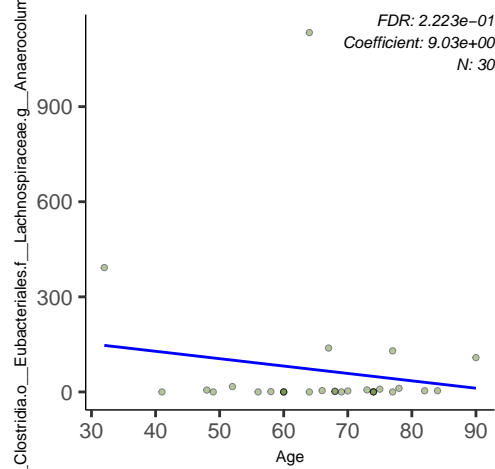


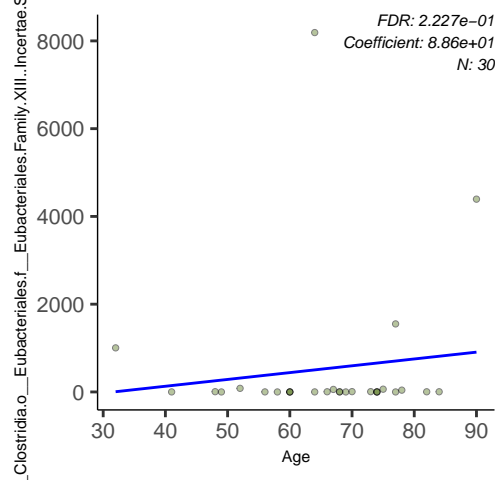


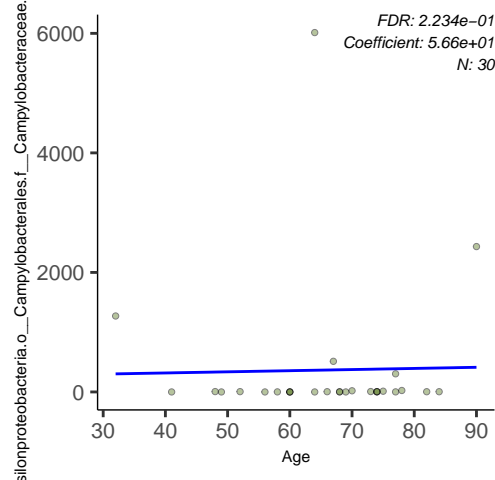
ia.p__Firmicutes.c__Clostridia.o__Eubacteriales.f__Lachnospiraceae

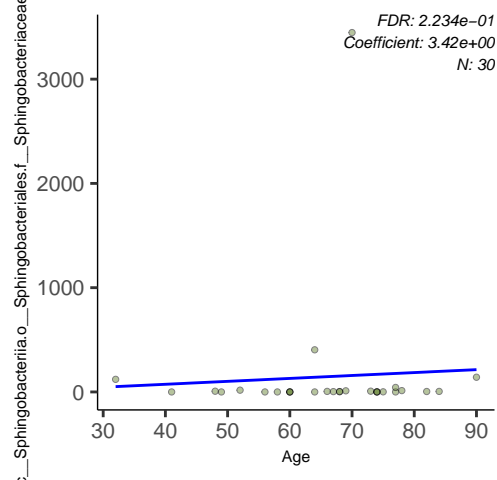






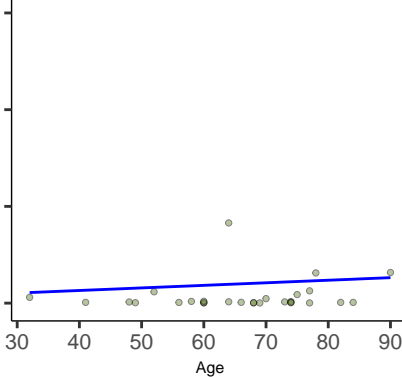


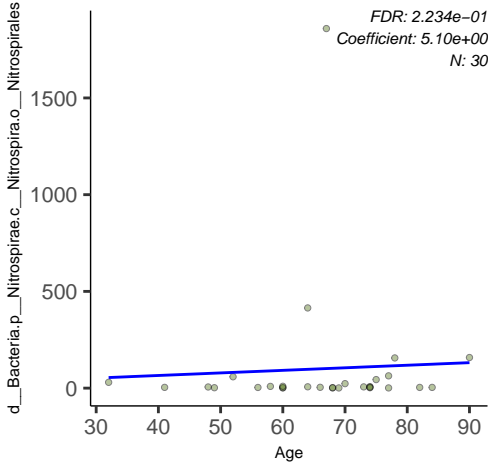


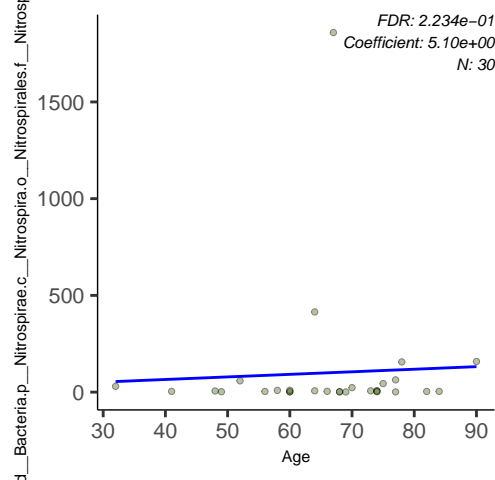


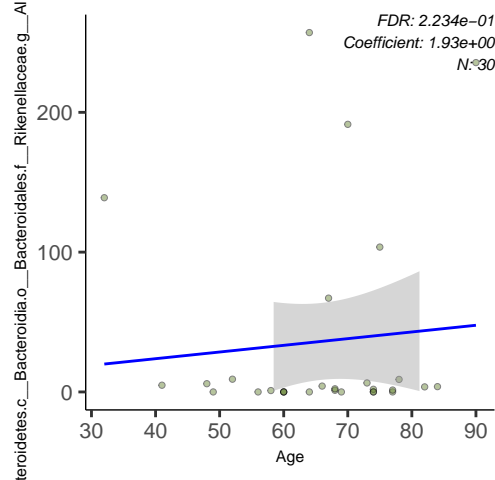
d__Bacteria.p__Nitrospirae.c__Nitrospira

FDR: 2.234e-01
Coefficient: 5.10e+00
N: 30







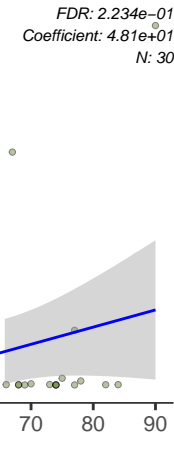


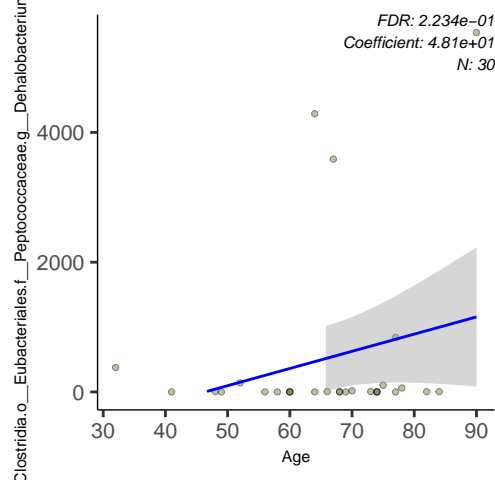
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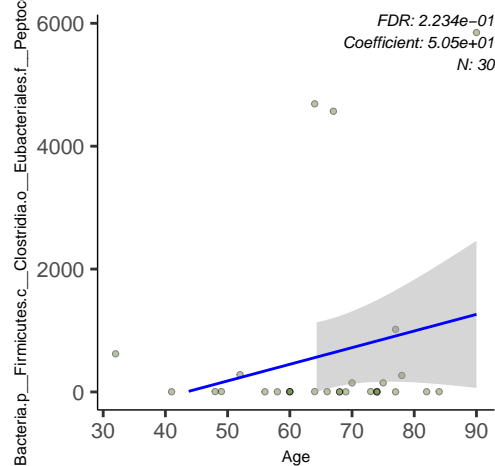
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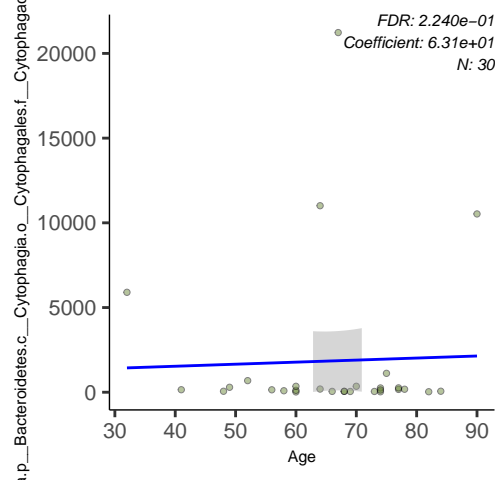
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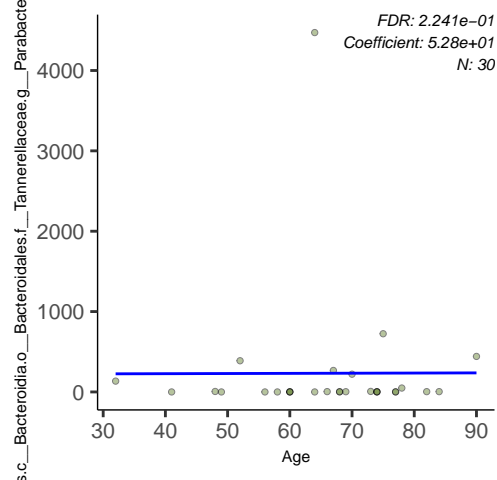
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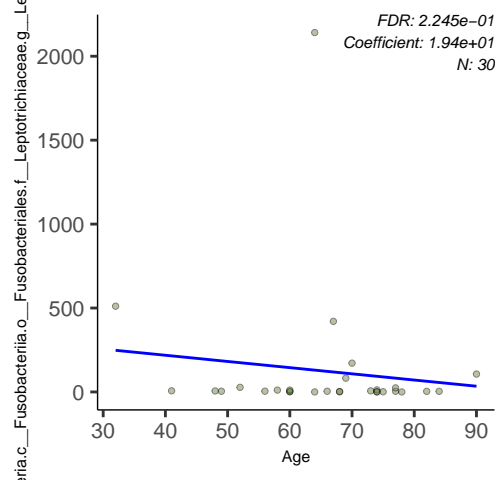


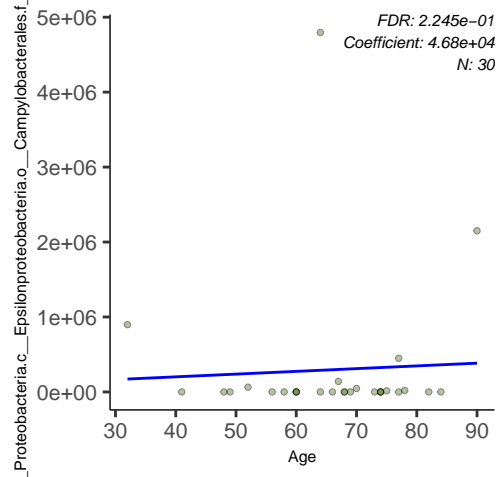


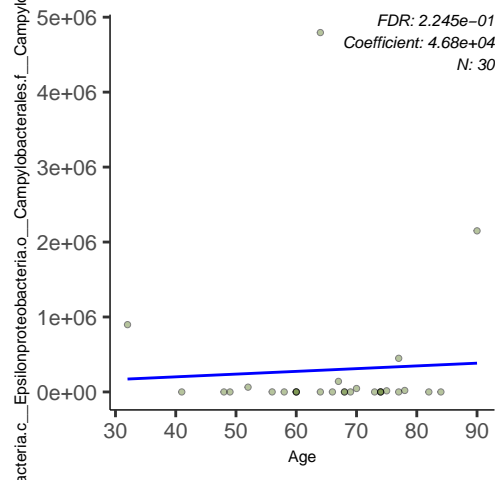


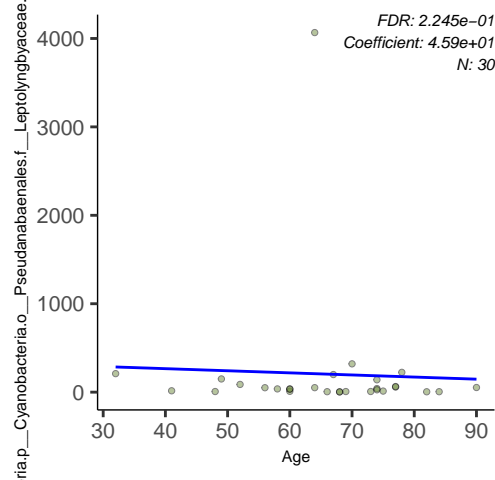


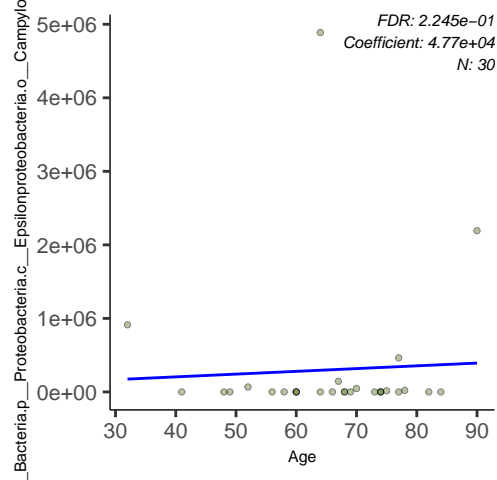


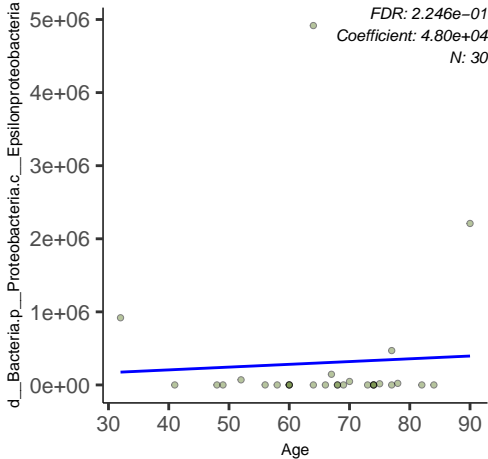






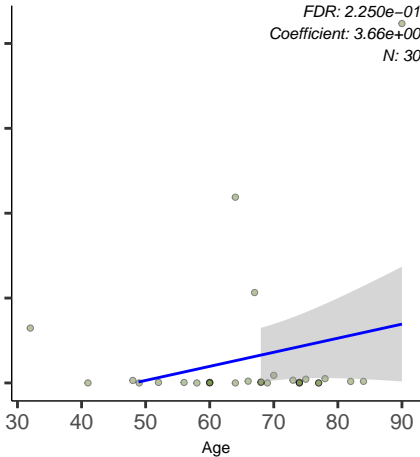


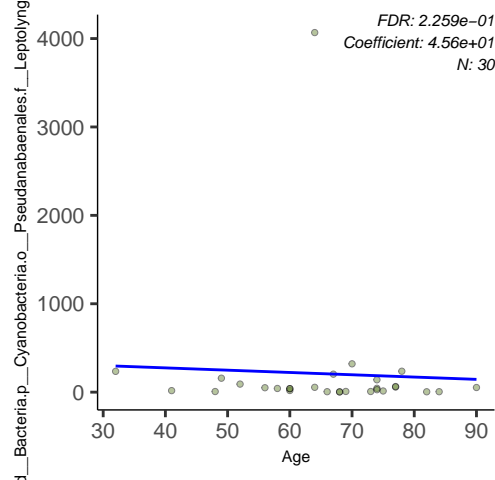


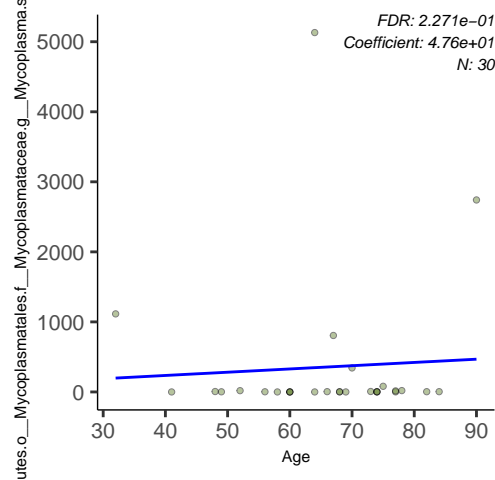


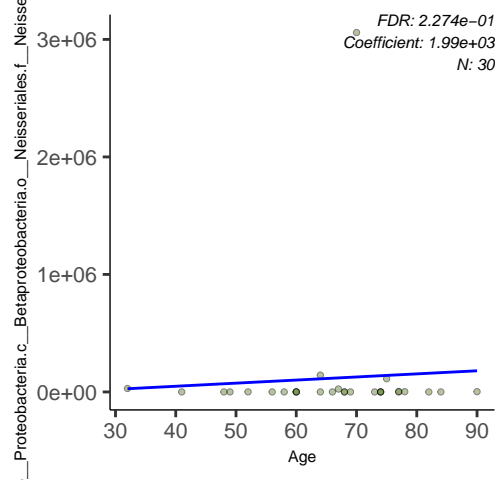
eroidetes.c__Bacteroidia.o__Bacteroidales.f__Rikenellaceae.g__Ali

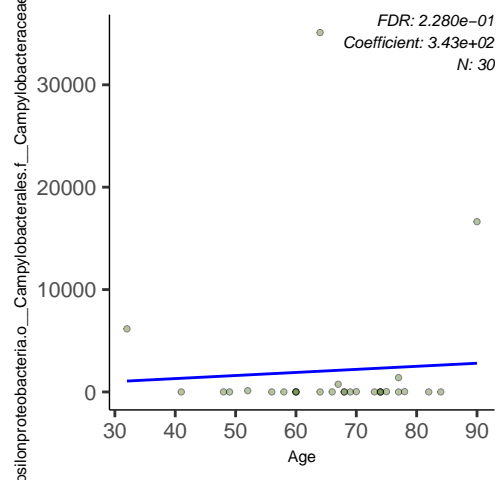
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Coefficient: 3.66e+00
N: 30

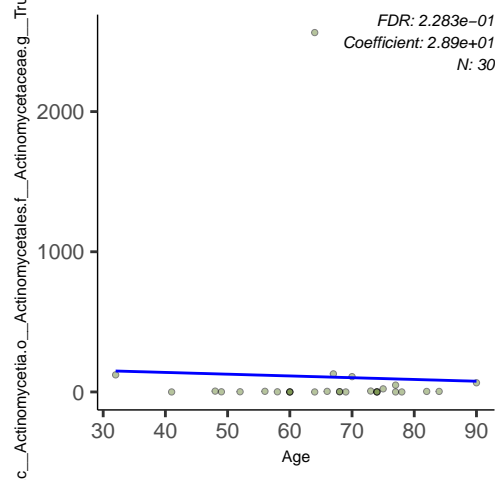


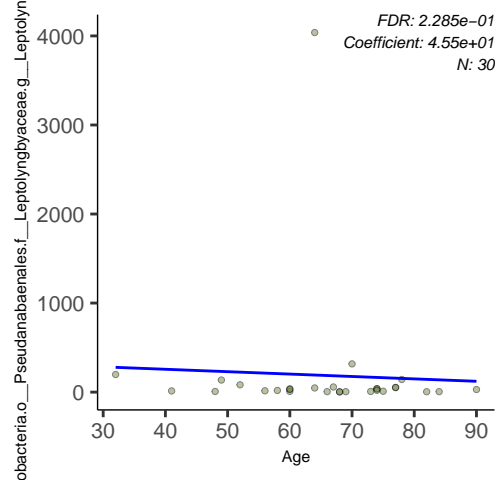


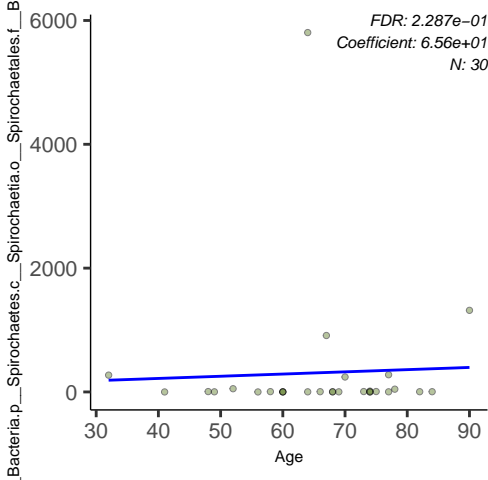




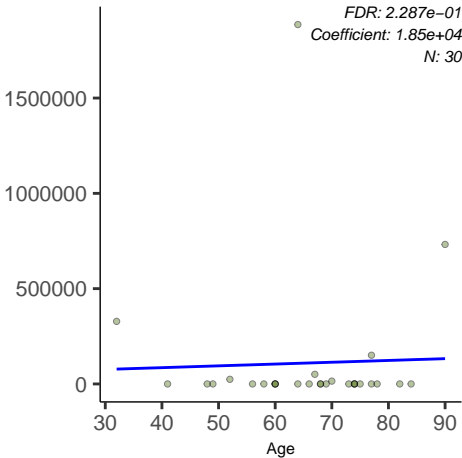


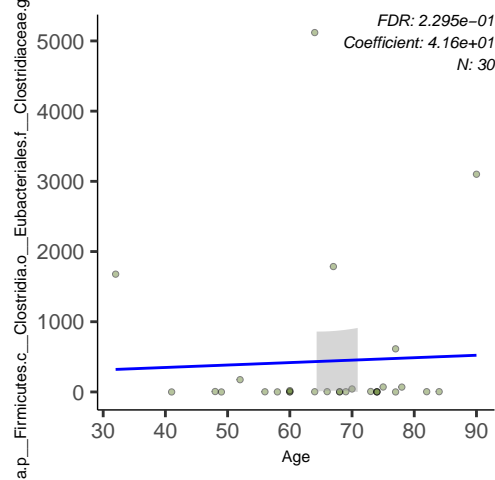


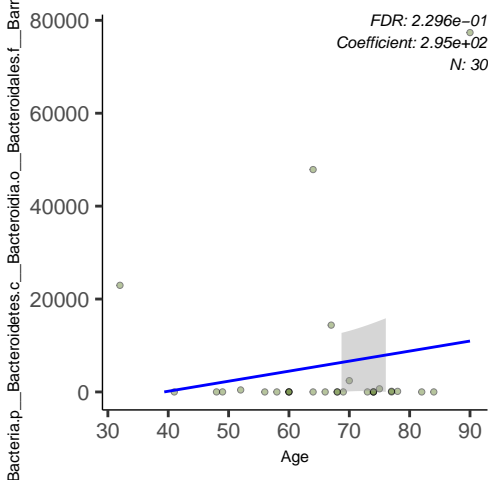


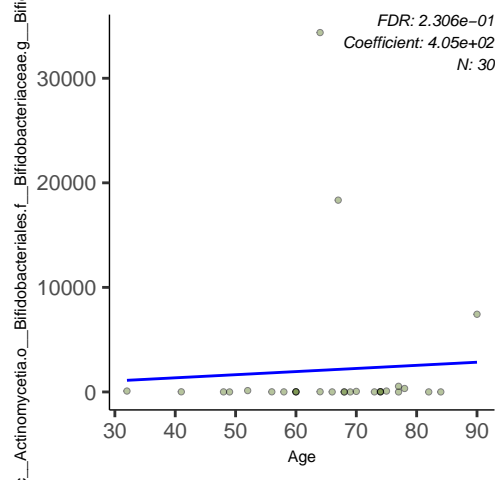


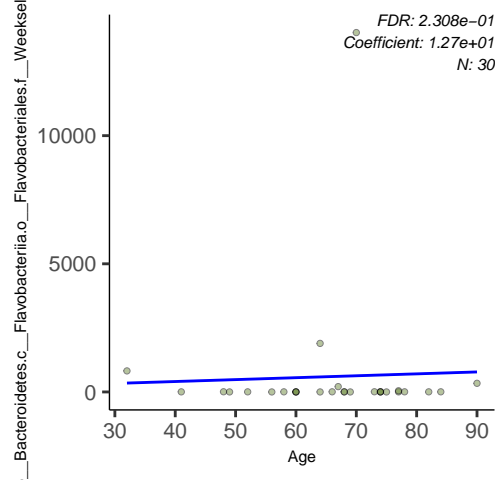
ilonproteobacteria.o__Campylobacteraceae











letes.c__Flavobacteriia.o__Flavobacteriales.f__Weeksellaceae.g__E

FDR: 2.308e-01
Coefficient: 1.27e+01
N: 30

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30

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50

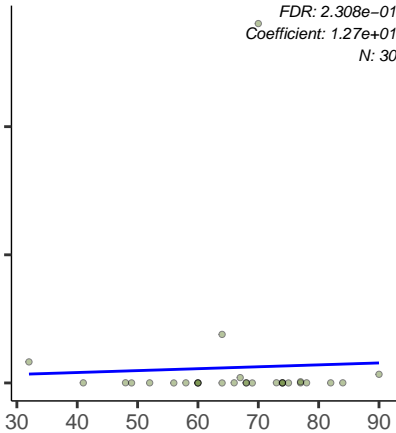
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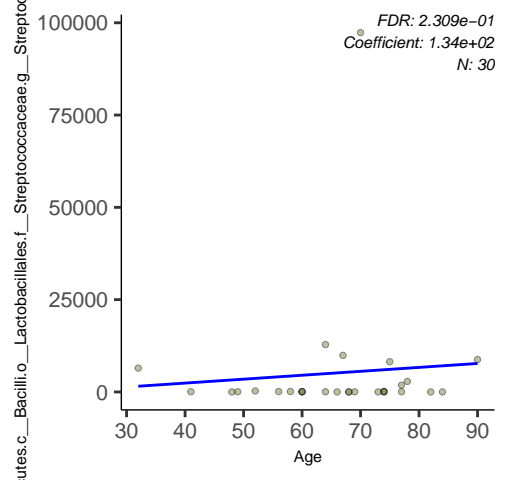
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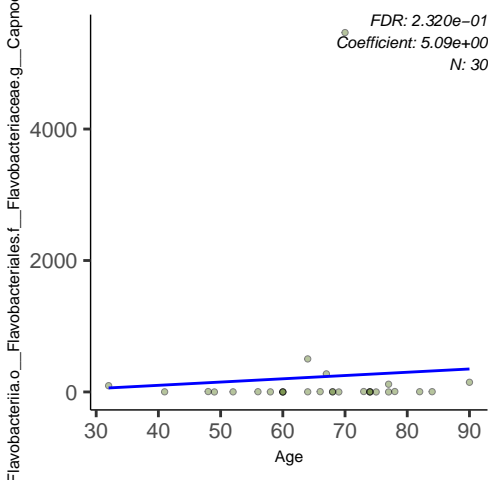
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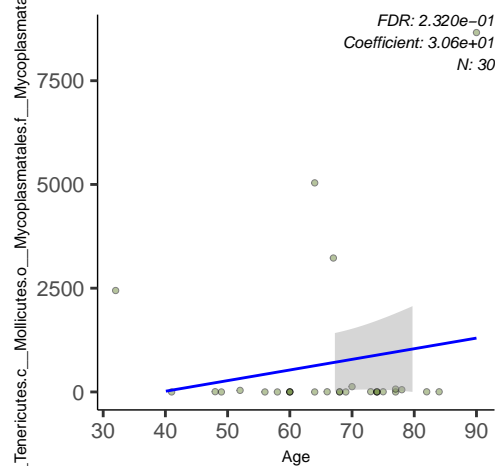
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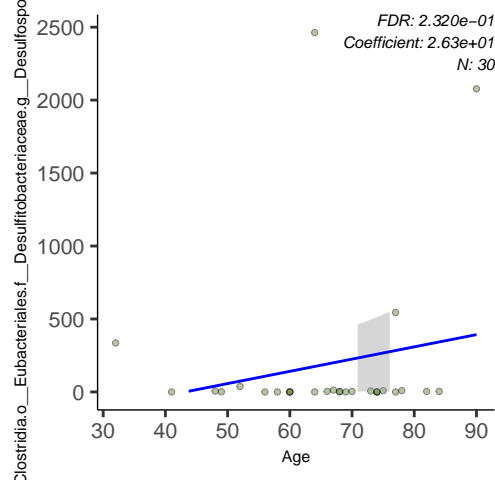
Age

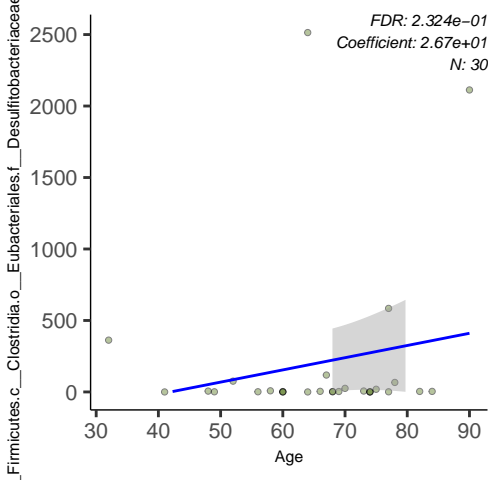


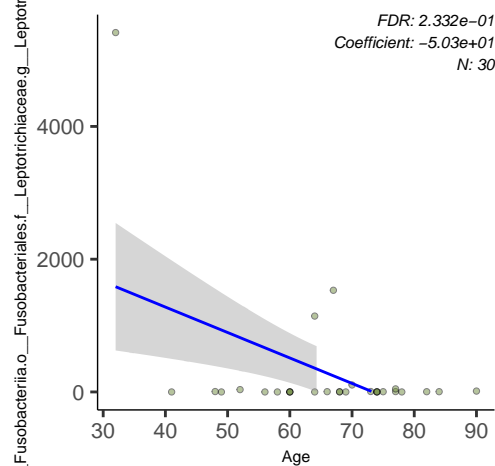


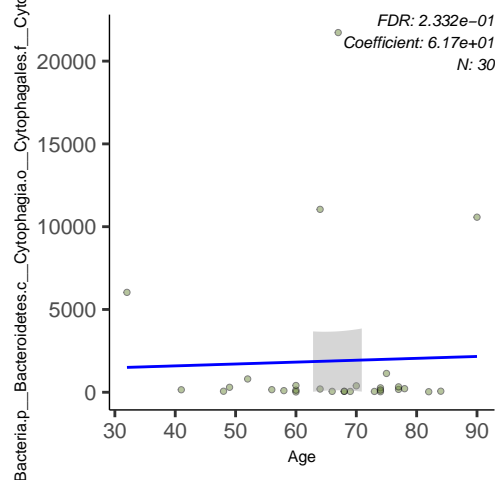


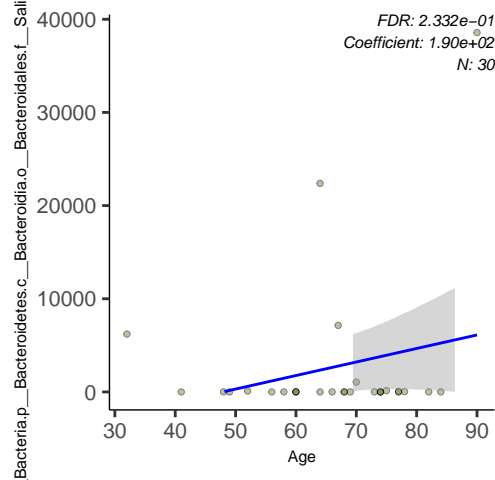


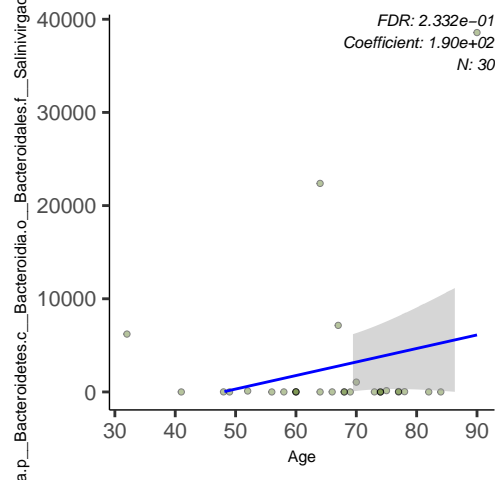


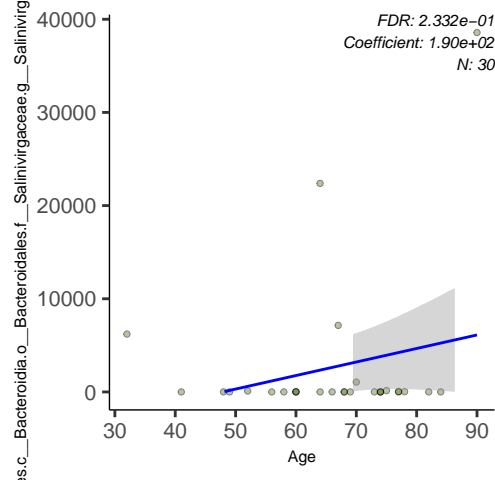


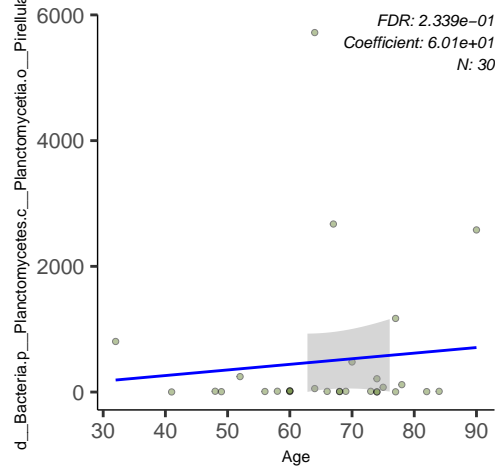


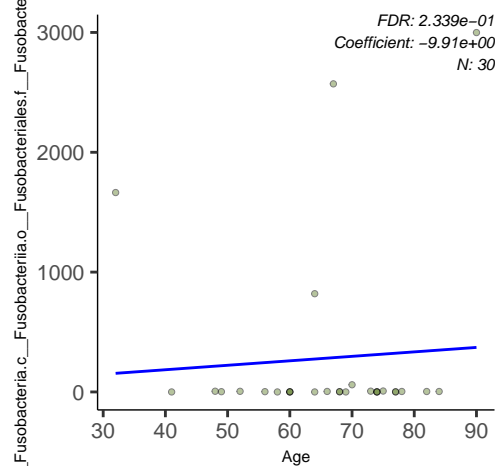


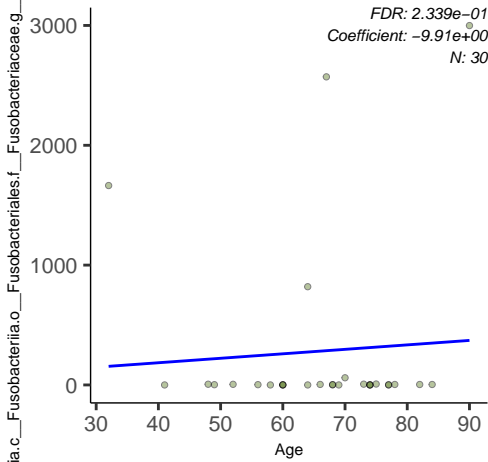


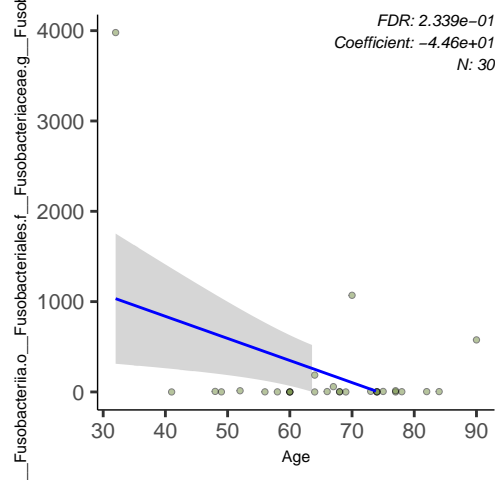


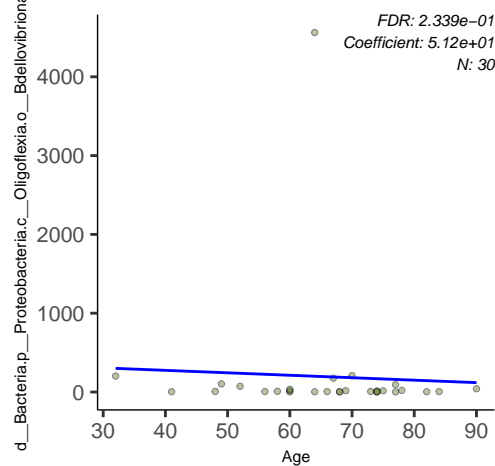


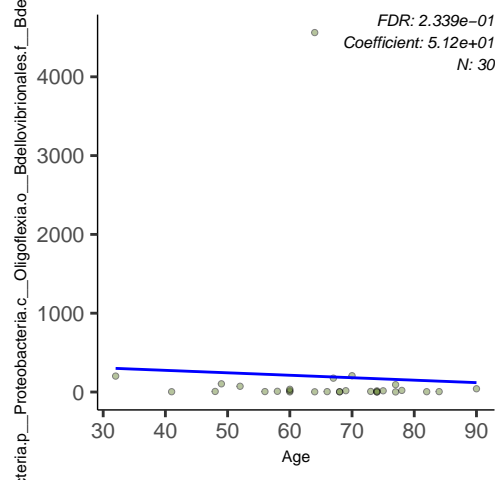


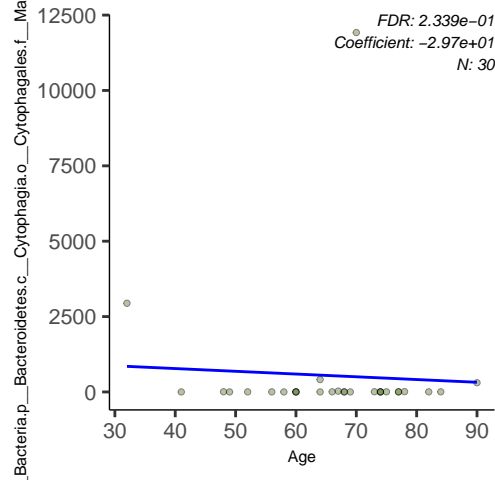


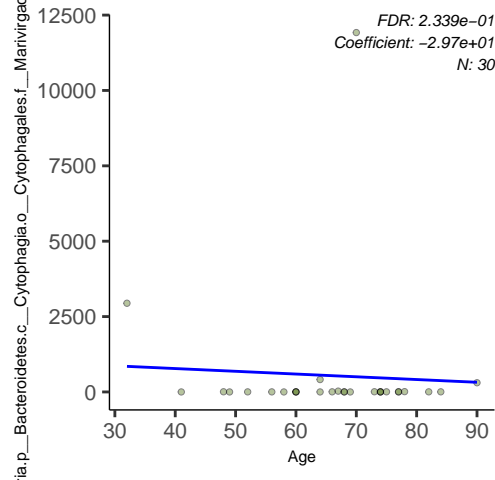


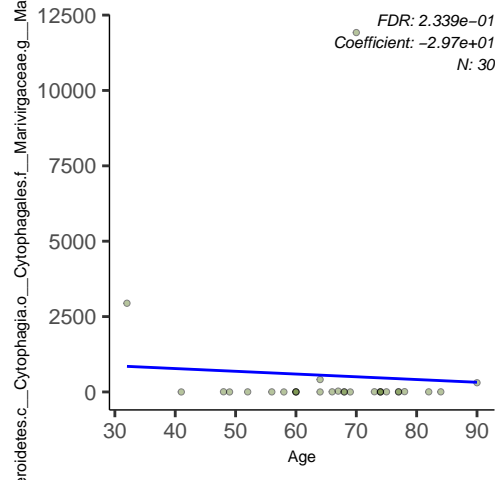


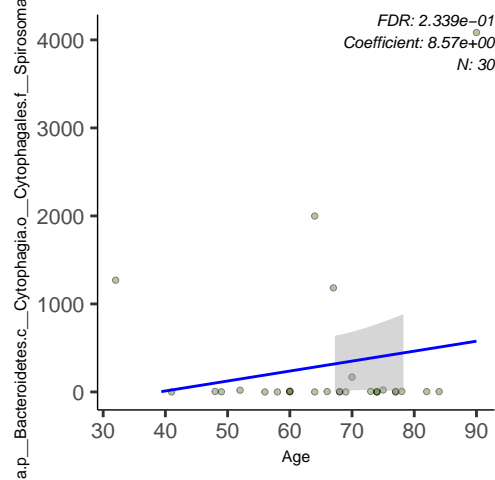


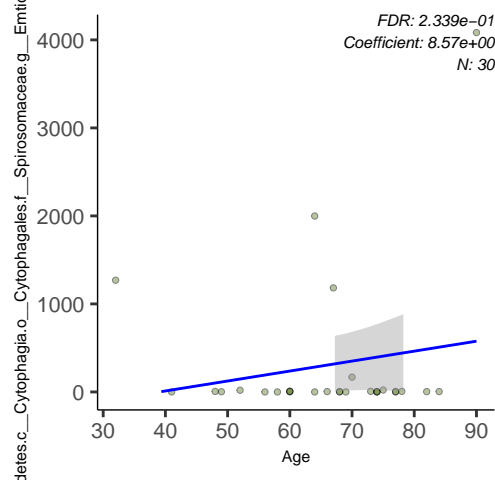


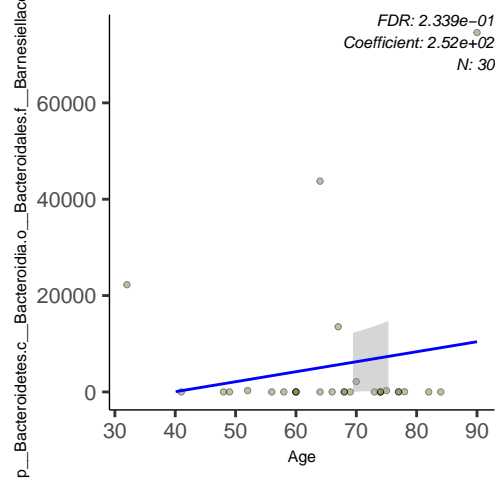


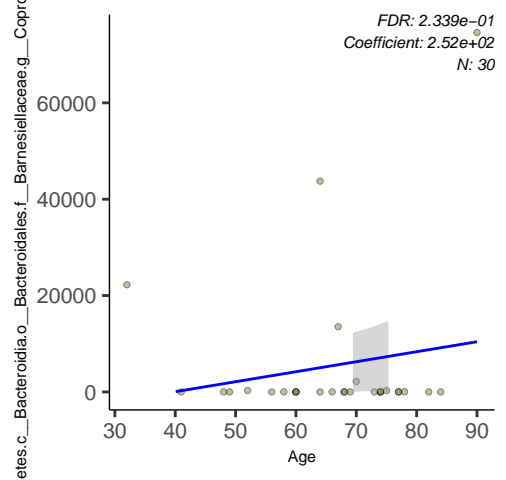


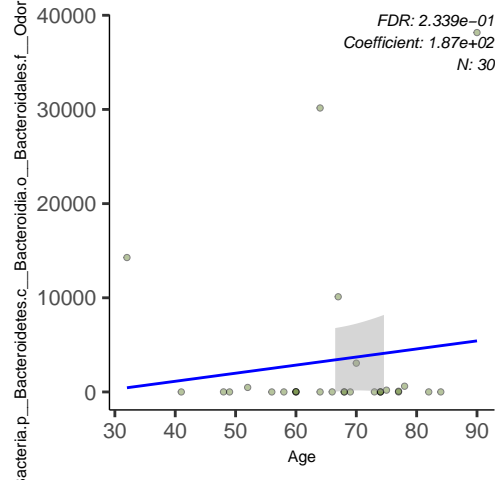


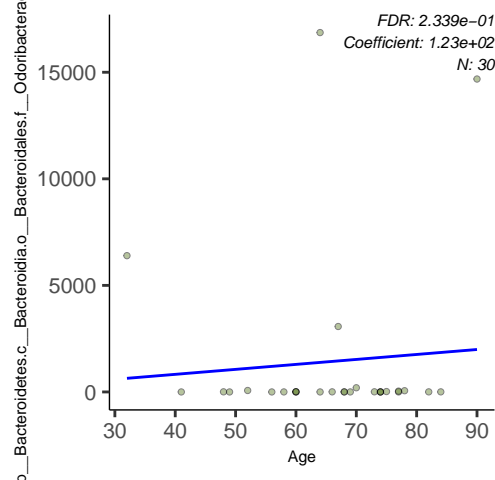


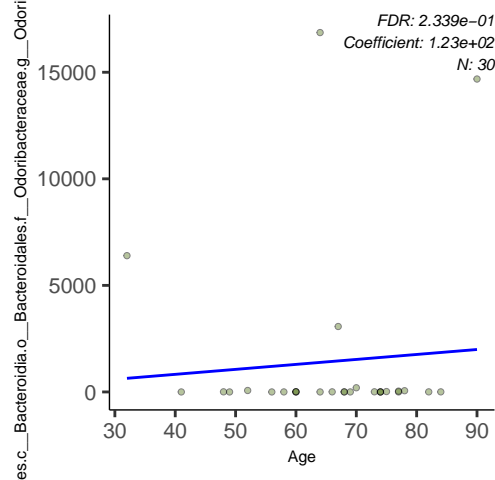


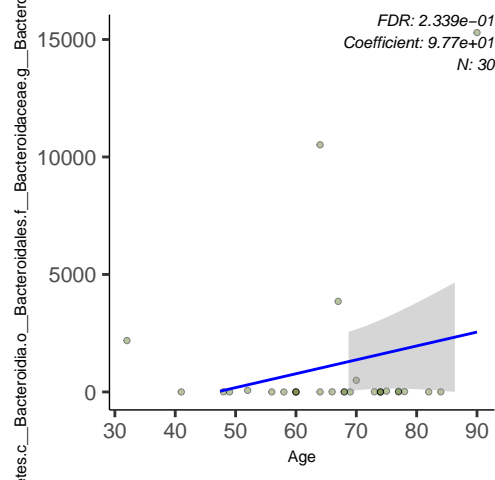


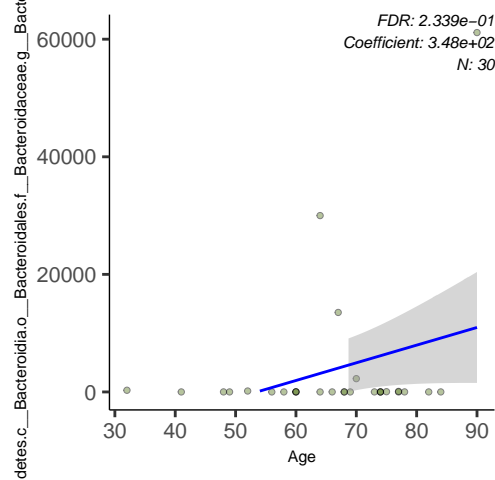






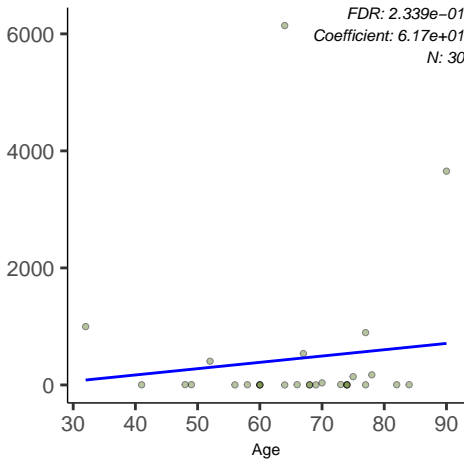


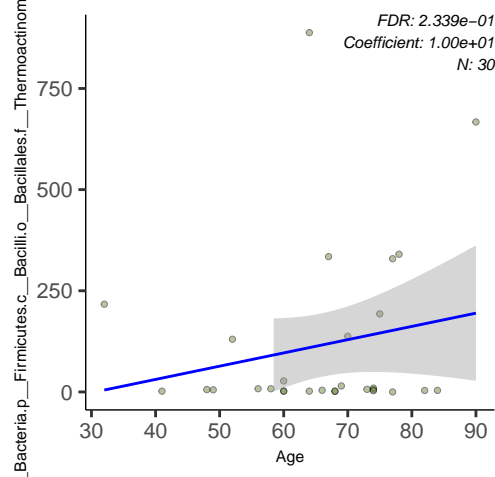


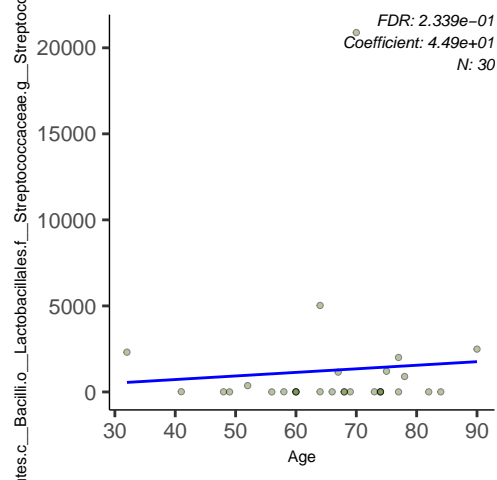


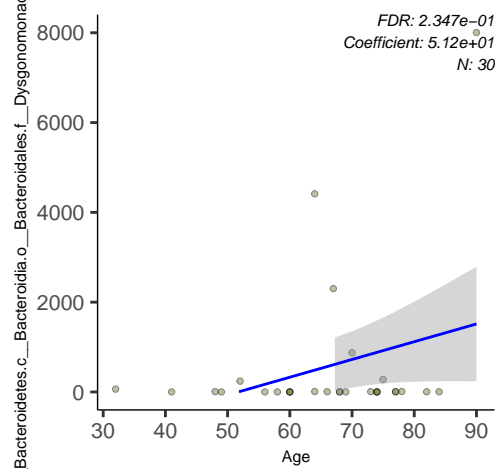
ria.p__Firmicutes.c__Clostridia.o__Eubacteriales.f__Eubacteriaceae

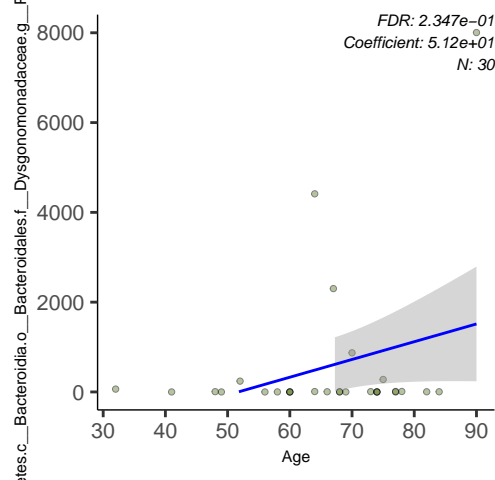
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Coefficient: 6.17e+01
N: 30

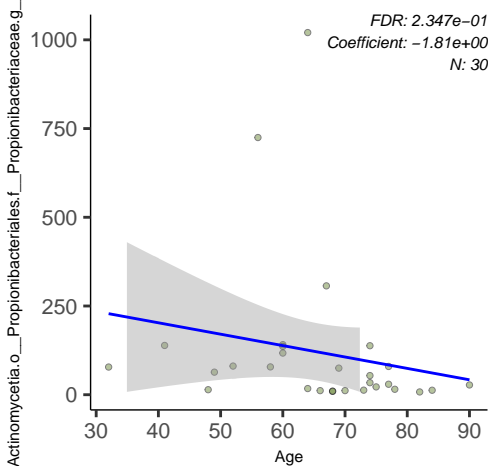


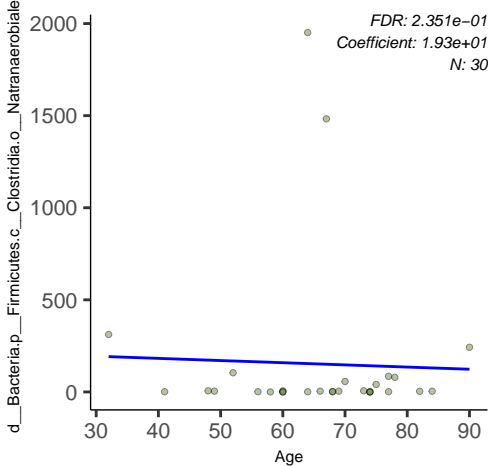


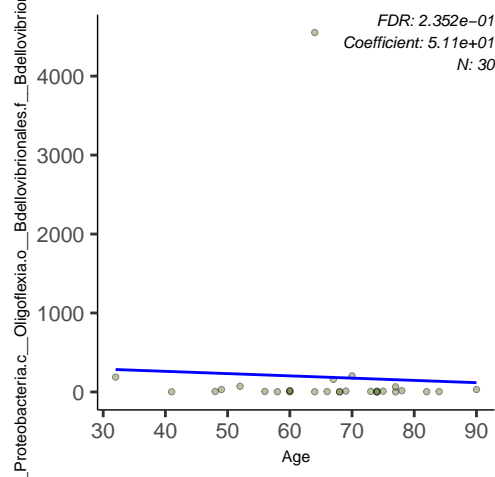


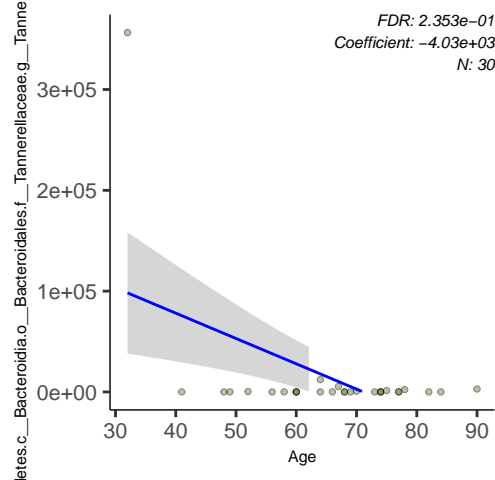


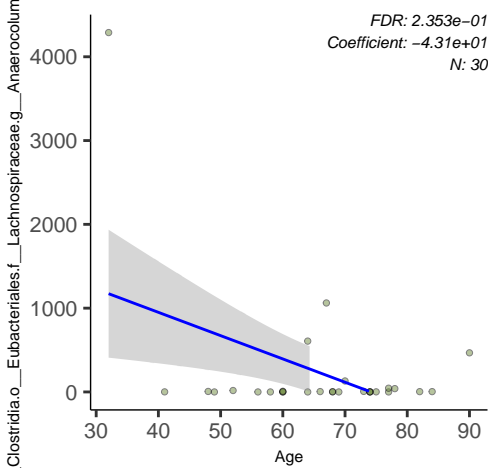


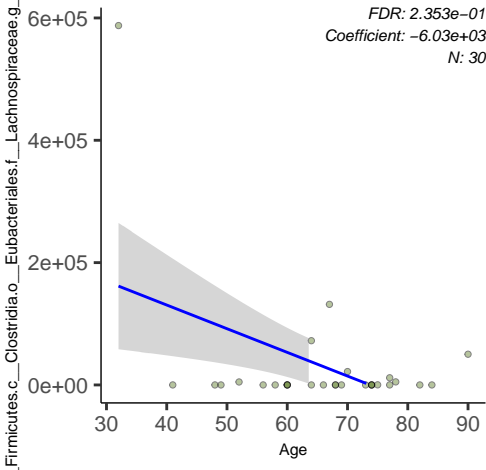


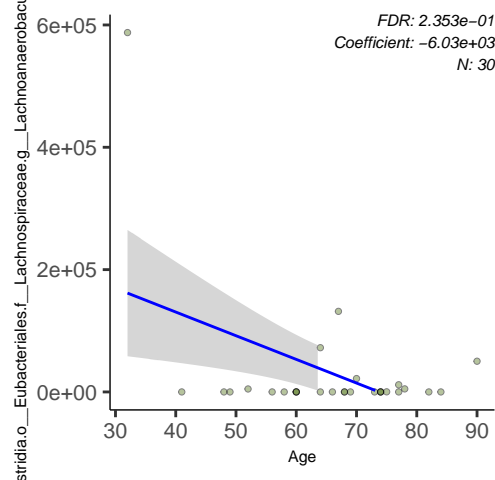


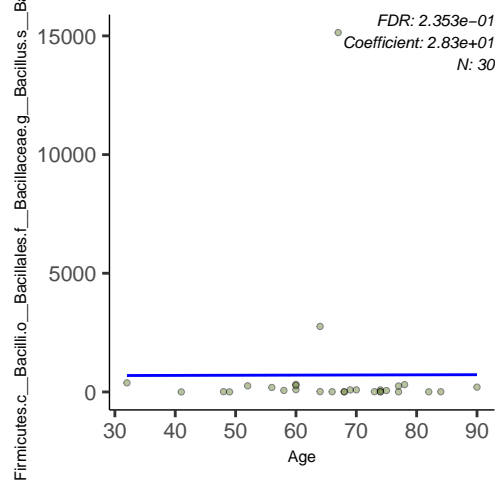


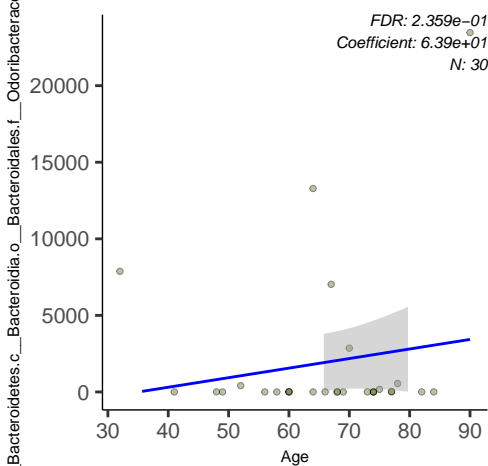


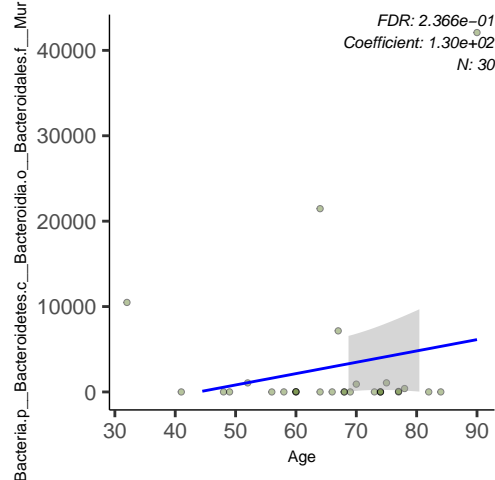


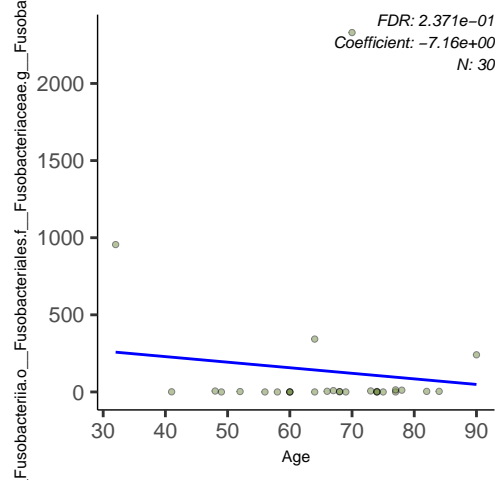


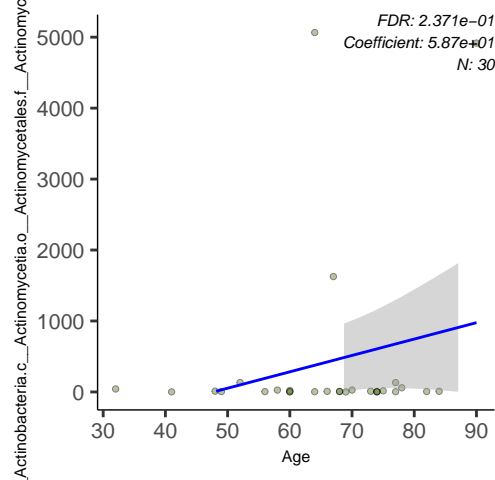


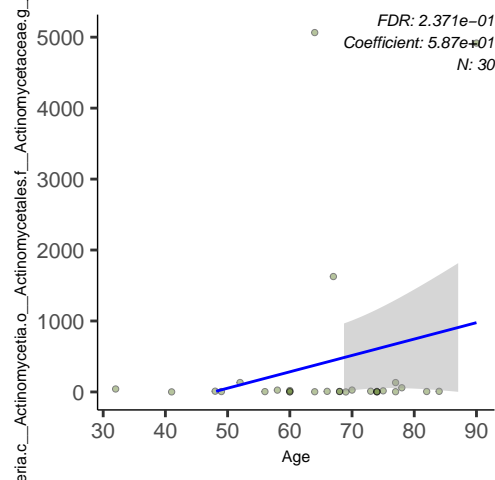


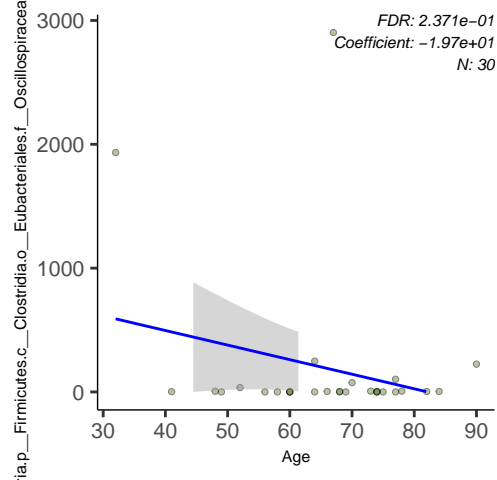


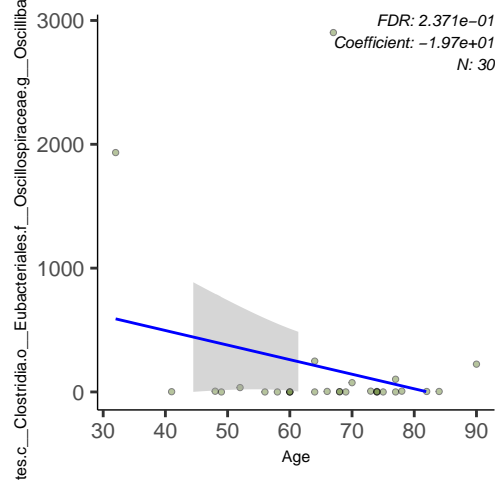


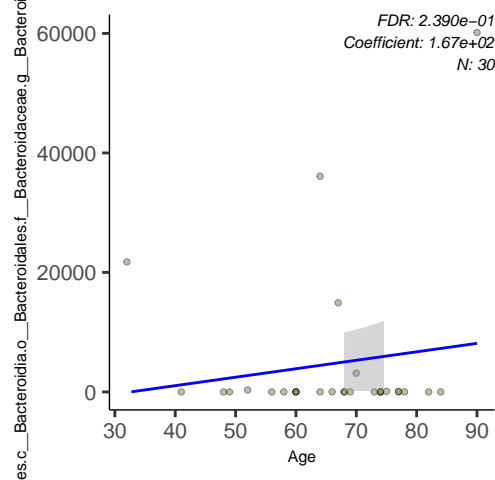


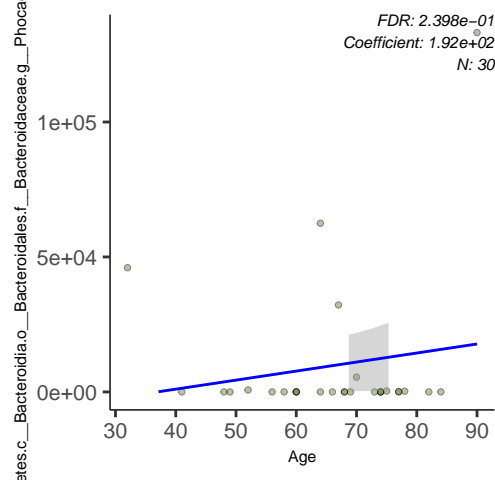


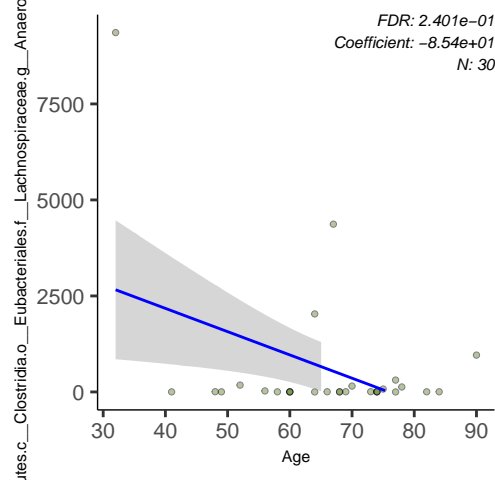


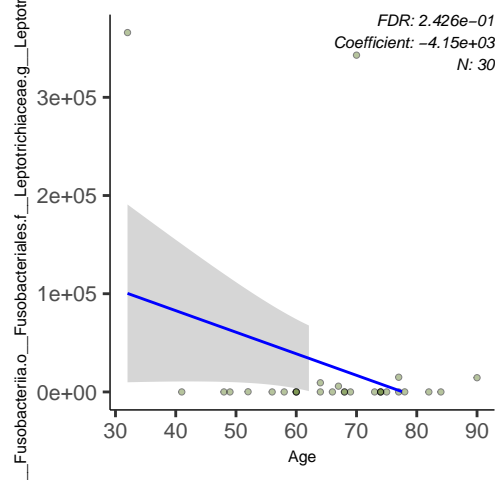


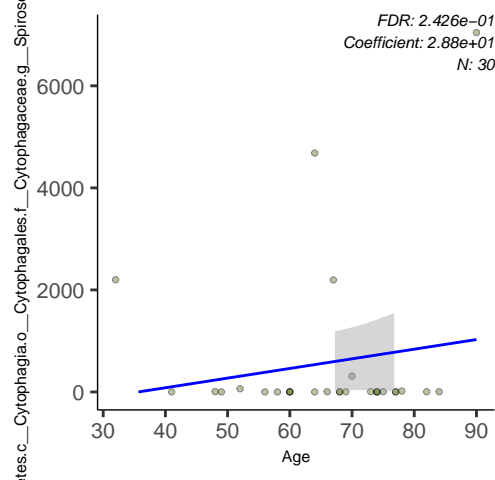


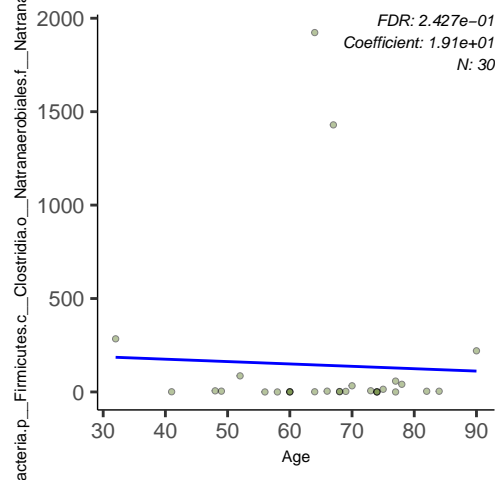


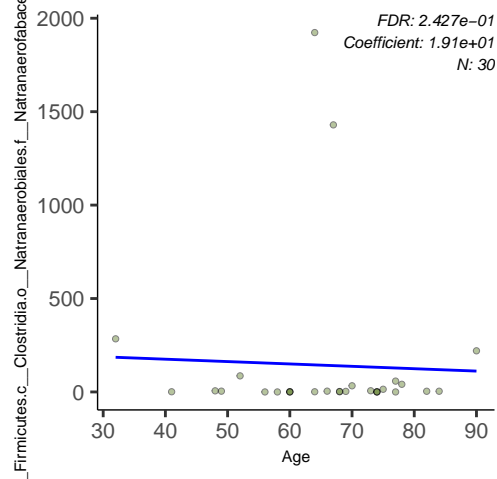


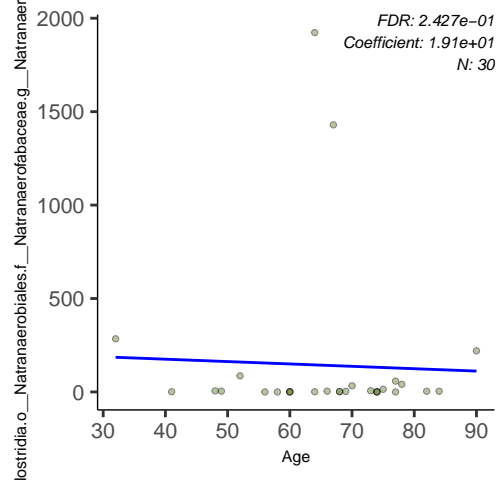




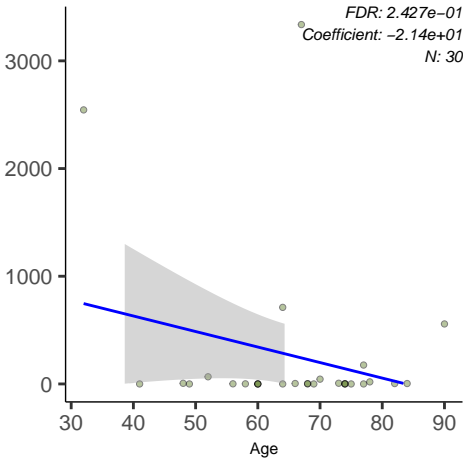


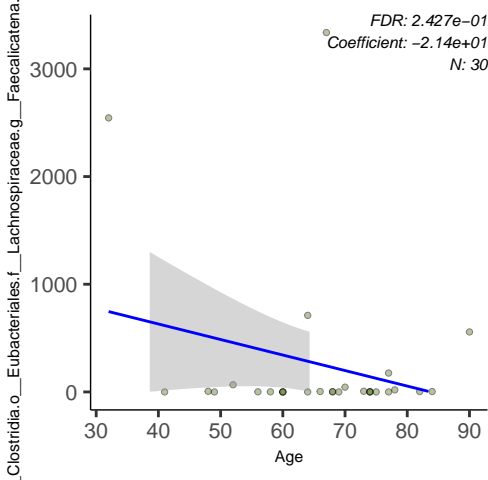


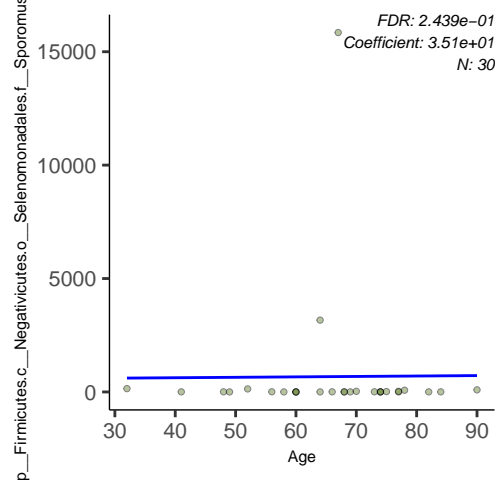


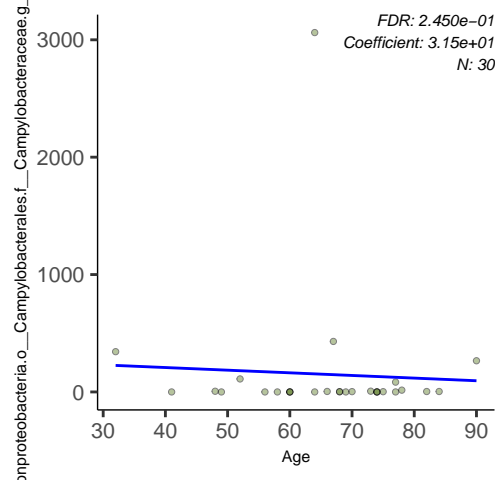


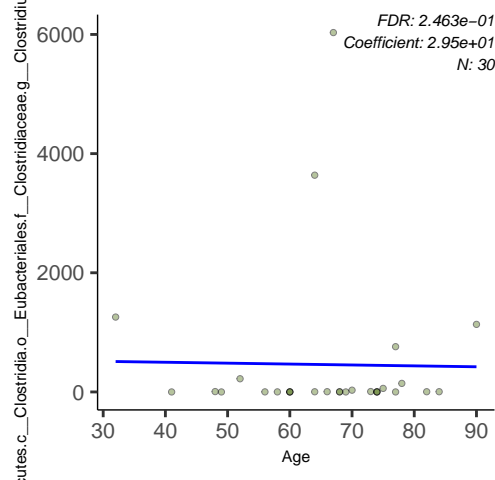
a.p_Firmicutes.c_Clostridia.o_Eubacteriales.f_Lachnospiraceae











a.p__Firmicutes.c__Clostridia.o__Eubacteriales.f__Lachnospiraceae

