

Bifidobacterium

1.00
0.75
0.50
0.25
0.00

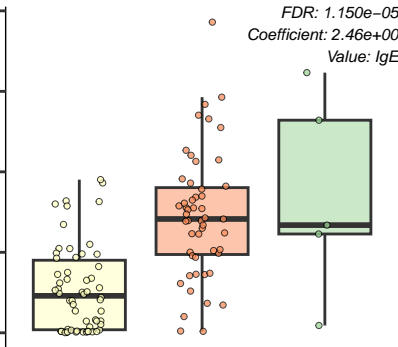
N (n=58)

IgE (n=51)

nE (n=5)

group

FDR: 1.150e-05
Coefficient: 2.46e+00
Value: IgE



Parasutterella

FDR: 1.150×10^{-5}
Coefficient: 2.24×10^0
Value: nE

0.00100

0.00075

0.00050

0.00025

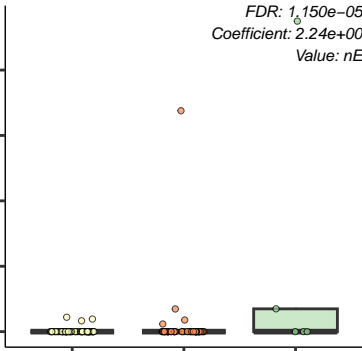
0.00000

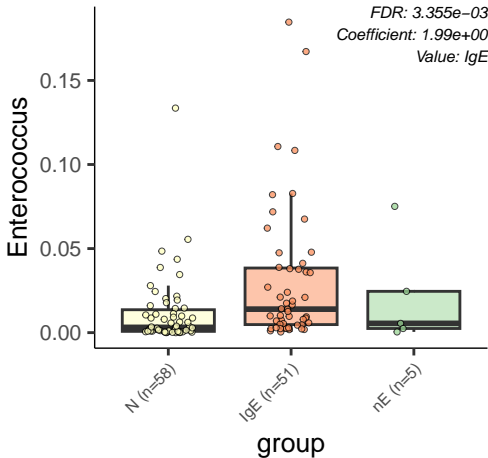
N (n=58)

IgE (n=51)

nE (n=5)

group





Altererythrobacter

FDR: $6.418e-03$
Coefficient: $6.24e-01$
Value: nE

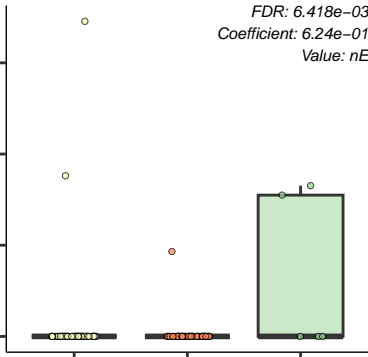
0.00015
0.00010
0.00005
0.00000

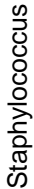
N (n=58)

IgE (n=51)

nE (n=5)

group





FDR: 1.828e-02

Coefficient: 1.52e+00

Value: IgE

0.004

0.002

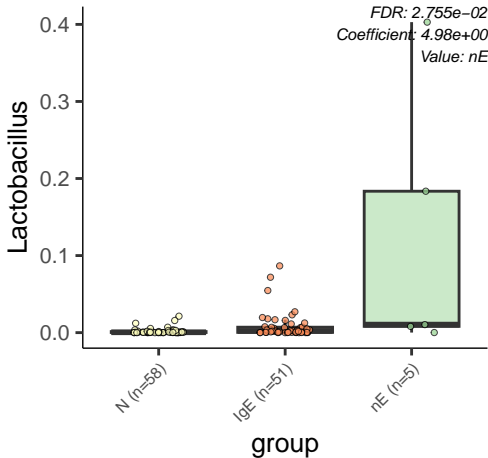
0.000

N (*n*=58)

IgE (n=51)

nE (n=5)

group



Pseudomonas

FDR: $2.755e-02$
Coefficient: $1.02e+00$
Value: IgE

0.03

0.02

0.01

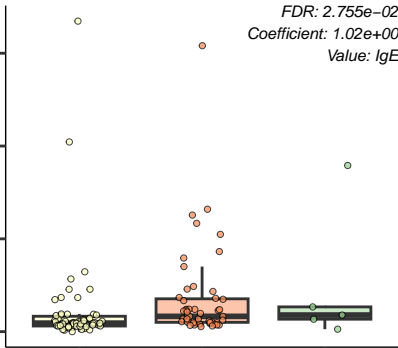
0.00

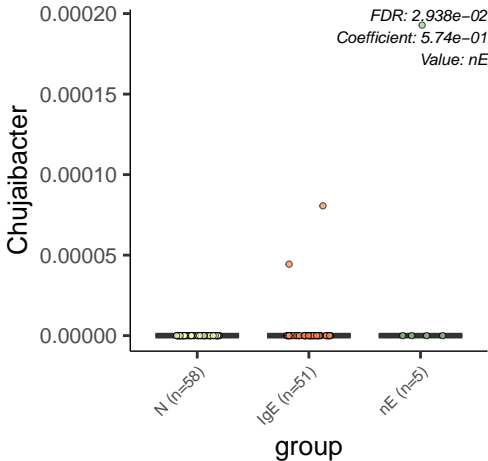
N (n=58)

IgE (n=51)

nE (n=5)

group





Escherichia.Shigella

1.00
0.75
0.50
0.25
0.00

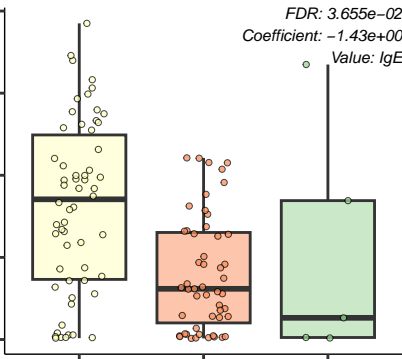
N (n=58)

IgE (n=51)

nE (n=5)

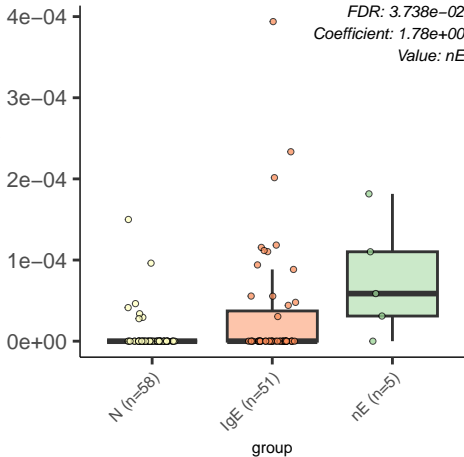
group

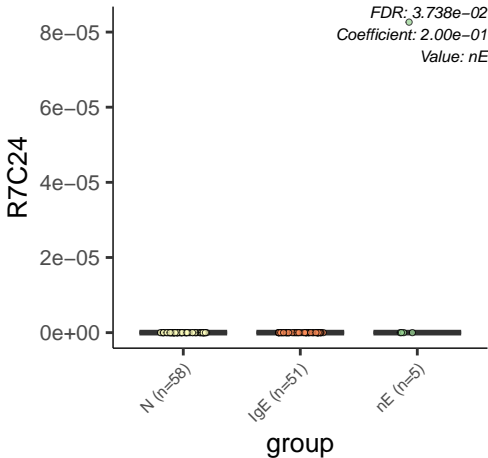
FDR: 3.655e-02
Coefficient: -1.43e+00
Value: IgE



Rikenellaceae_RC9_gut_group

FDR: $3.738e-02$
Coefficient: $1.78e+00$
Value: nE





Burkholderia.Caballeronia.Paraburkholderia

FDR: 3.776e-02
Coefficient: 2.00e-01
Value: nE

6e-05

4e-05

2e-05

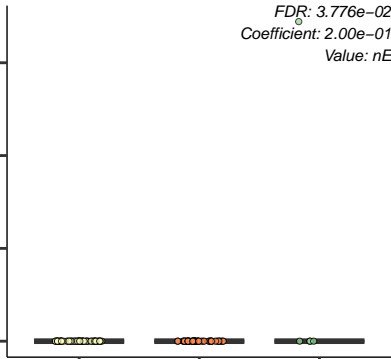
0e+00

N (n=58)

IgE (n=51)

nE (n=5)

group



Pontibacter

FDR: 3.776e-02
Coefficient: 2.00e-01
Value: nE

6e-05

4e-05

2e-05

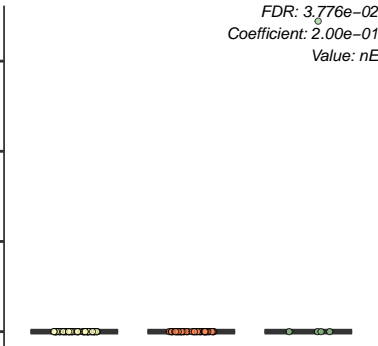
0e+00

N (n=58)

IgE (n=51)

nE (n=5)

group



Caulobacter

6e-05

4e-05

2e-05

0e+00

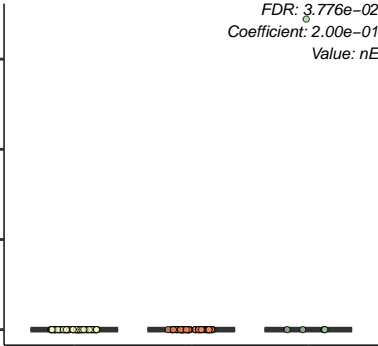
N (n=58)

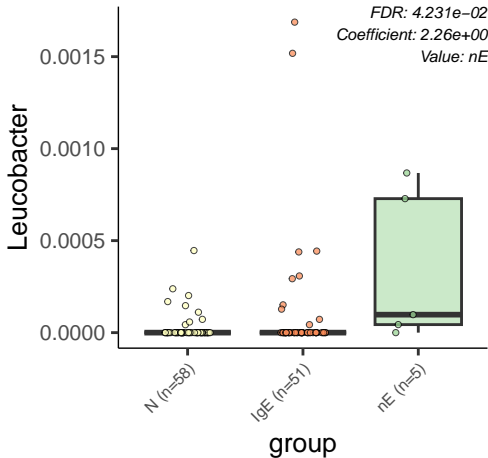
IgE (n=51)

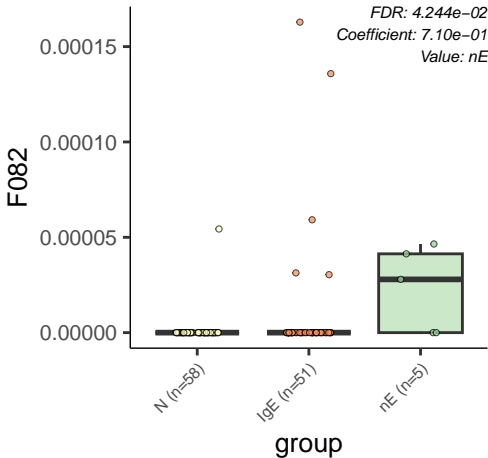
nE (n=5)

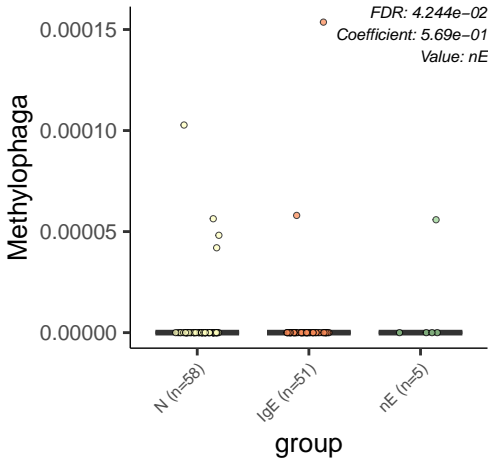
group

FDR: 3.776e-02
Coefficient: 2.00e-01
Value: nE

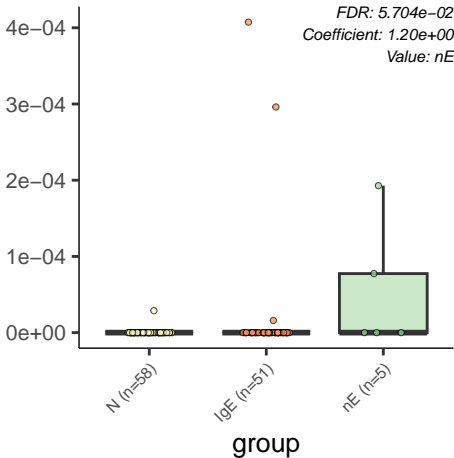


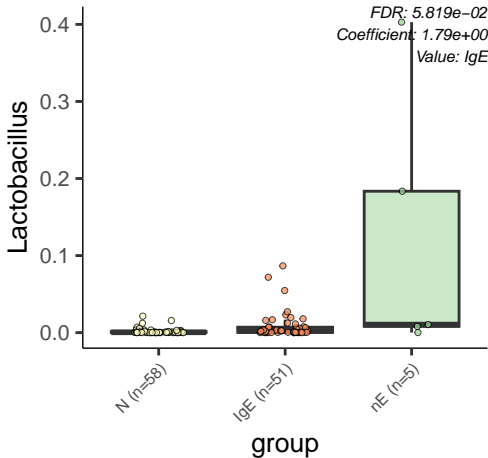


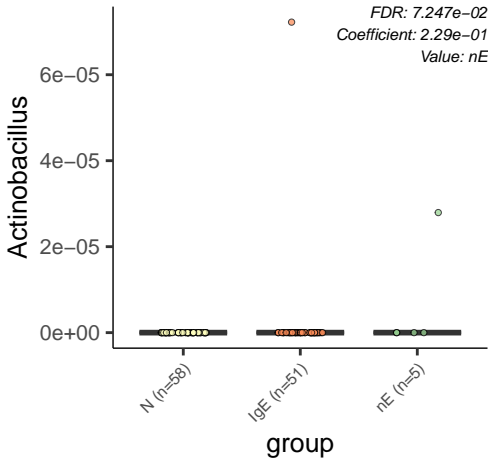


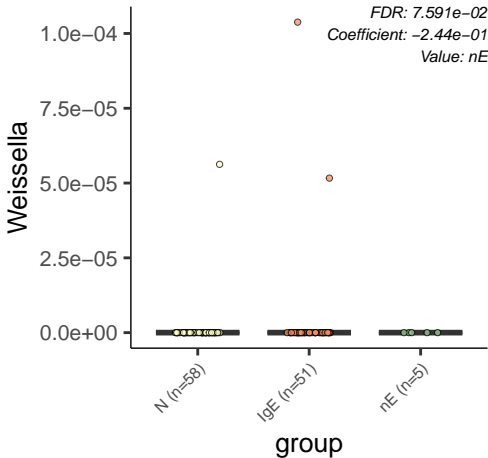


Leptotrichia









Methylobacterium.Methylorubrum

2e-04

1e-04

0e+00

N (n=58)

IgE (n=51)

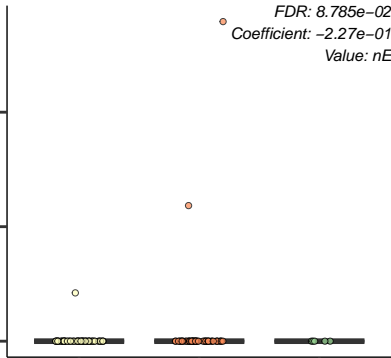
nE (n=5)

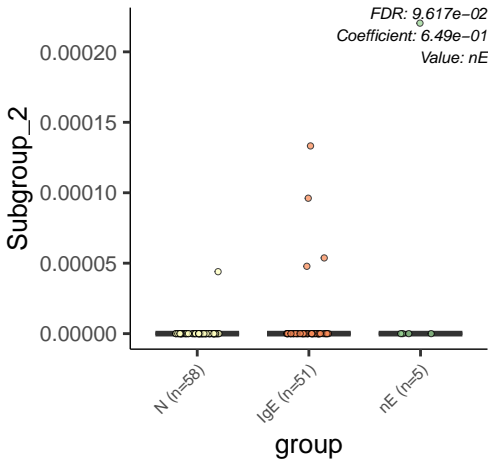
group

FDR: 8.785e-02

Coefficient: -2.27e-01

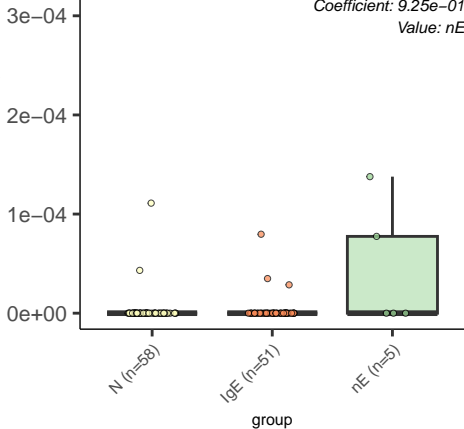
Value: nE

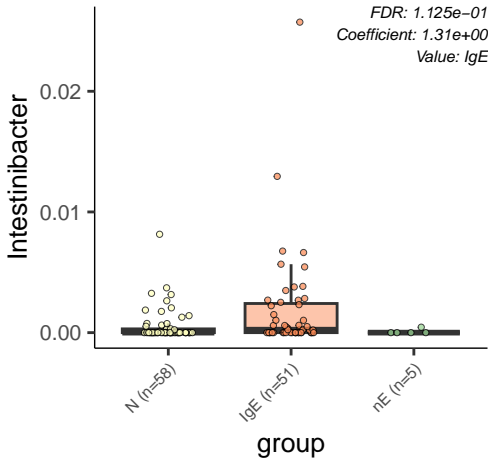




X.Eubacterium._siraеum_group

FDR: 1.026e-01
Coefficient: 9.25e-01
Value: nE





Rahnella1

FDR: 1.125e-01
Coefficient: 2.94e+00
Value: nE

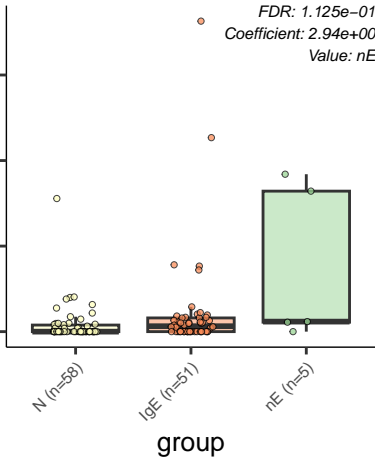
0.0075
0.0050
0.0025
0.0000

N (n=58)

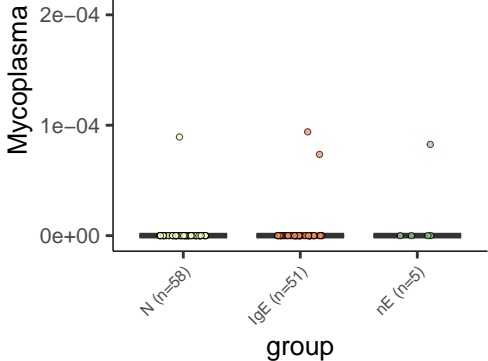
IgE (n=51)

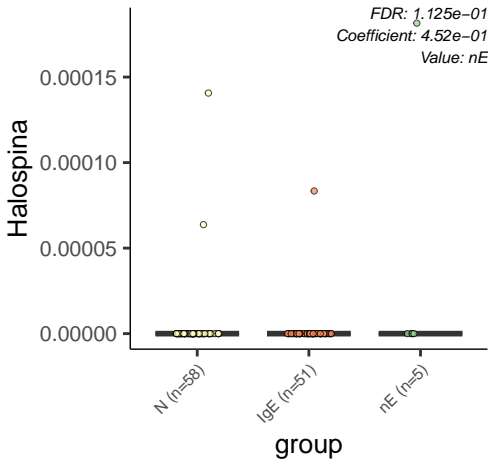
nE (n=5)

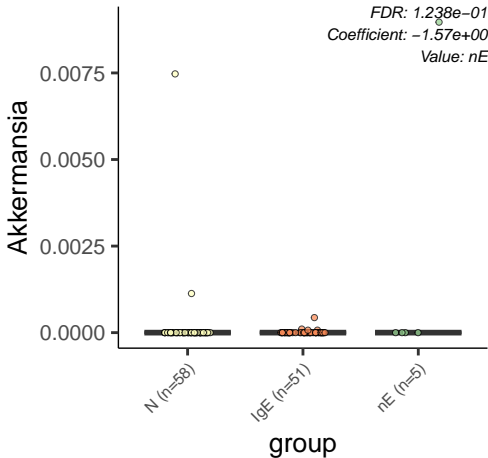
group



FDR: 1.125e-01
Coefficient: 2.72e-01
Value: nE







Spingobacterium

FDR: 1.238e-01
Coefficient: 4.47e-01
Value: IgE

3e-04

2e-04

1e-04

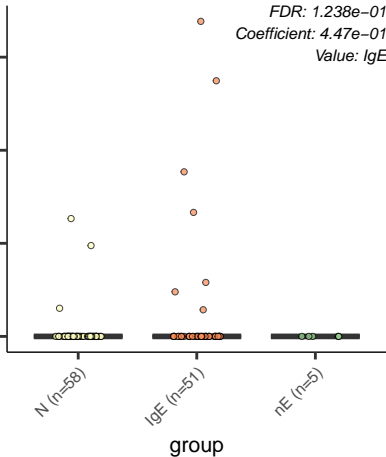
0e+00

N (n=58)

IgE (n=51)

nE (n=5)

group



Salmonella

0.00025
0.00020
0.00015
0.00010
0.00005
0.00000

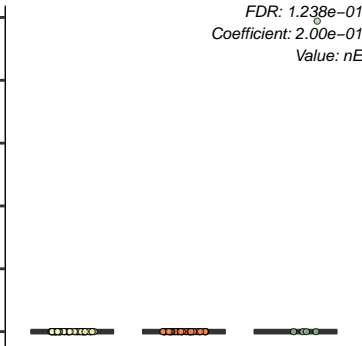
FDR: 1.238e-01
Coefficient: 2.00e-01
Value: nE

N (n=58)

IgE (n=51)

nE (n=5)

group



Corynebacterium

FDR: 1.311e-01
Coefficient: 4.49e-01
Value: IgE

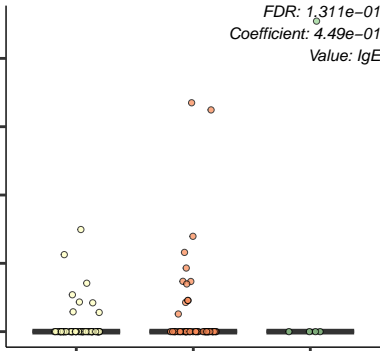
4e-04
3e-04
2e-04
1e-04
0e+00

N (n=58)

IgE (n=51)

nE (n=5)

group



Streptococcus

FDR: 1.362e-01
Coefficient: 1.39e+00
Value: IgE

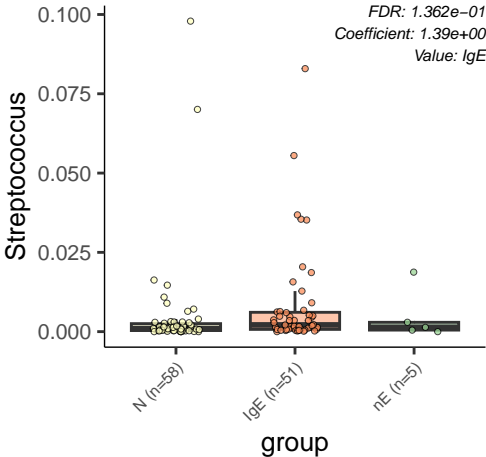
0.100
0.075
0.050
0.025
0.000

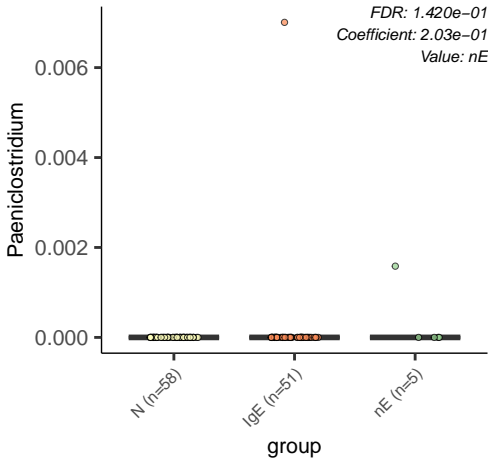
N (n=58)

IgE (n=51)

nE (n=5)

group





Tsukamurella

1e-04
5e-05
0e+00

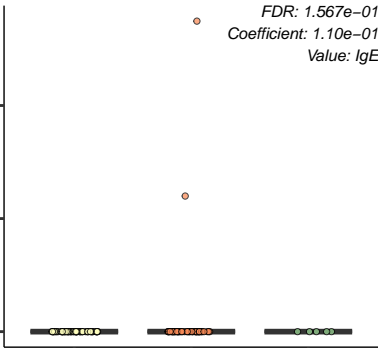
N (n=58)

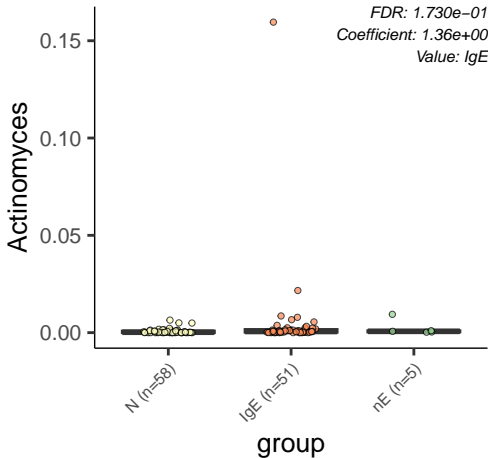
IgE (n=51)

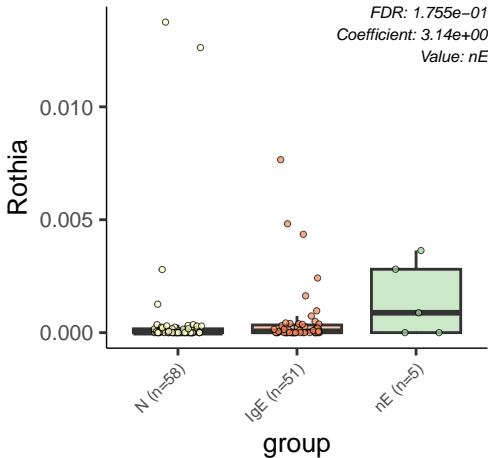
nE (n=5)

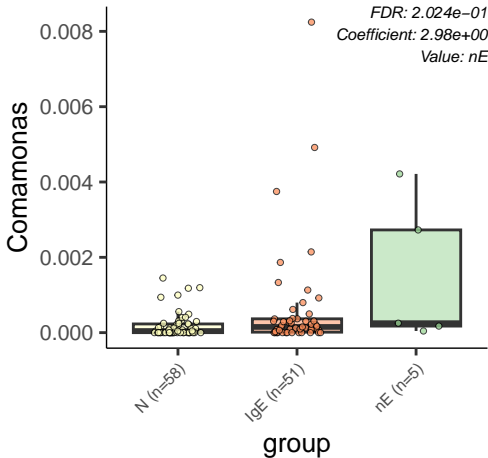
group

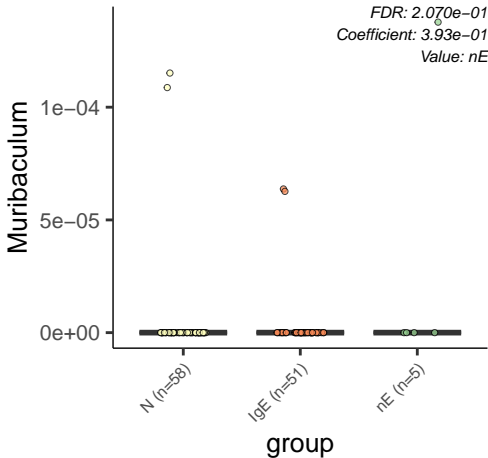
FDR: 1.567e-01
Coefficient: 1.10e-01
Value: IgE

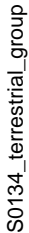












FDR: 2.236e-01

Coefficient: 1.19e-01

Value: IgE

0.00020

0.00015

0.00010

0.00005

0.00000

N (*n*=58)

IgE (n=51)

nE (n=5)

group

Granulicatella

FDR: 2.303e-01
Coefficient: 1.96e+00
Value: nE

0.002

0.001

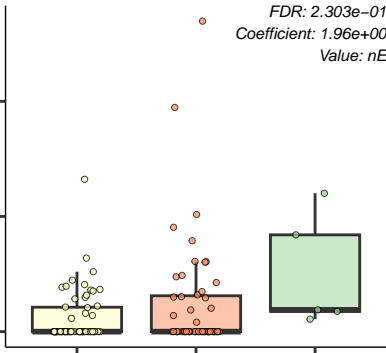
0.000

N (n=58)

IgE (n=51)

nE (n=5)

group



Rikenellaceae_RC9_gut_group

FDR: 2.407e-01
Coefficient: 5.20e-01
Value: IgE

