

Gordonibacter

FDR: 1.520e-08

Coefficient: -1.70e+00

Value: NO

200

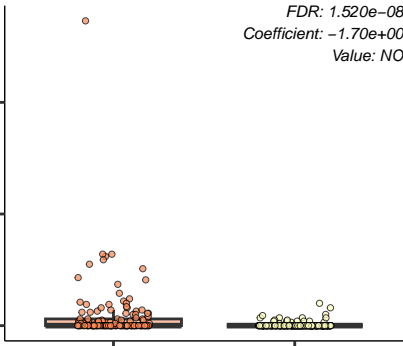
100

0

CZ (n=137)

NO (n=274)

Country



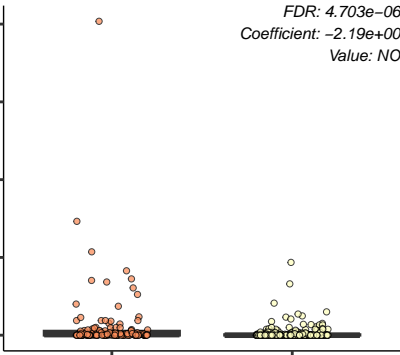
X.Clostridium._innocuum_group

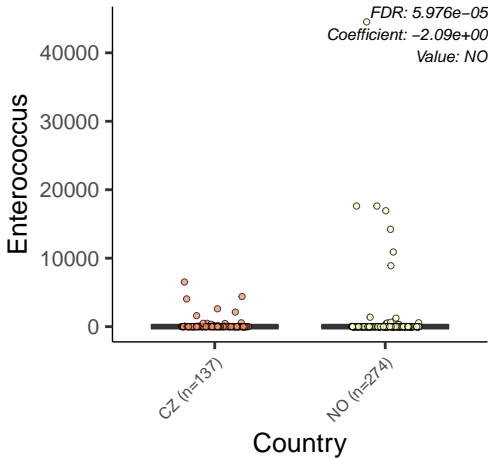
FDR: 4.703e-06
Coefficient: -2.19e+00
Value: NO

CZ (n=137)

NO (n=274)

Country





f__Peptococcaceae.g__uncultured

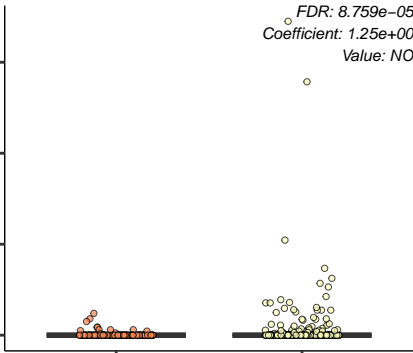
FDR: 8.759×10^{-5}
Coefficient: 1.25×10^0
Value: NO

600
400
200
0

CZ (n=137)

NO (n=274)

Country



Mogibacterium

FDR: $1.003e-04$

Coefficient: $-1.13e+00$

Value: NO

200

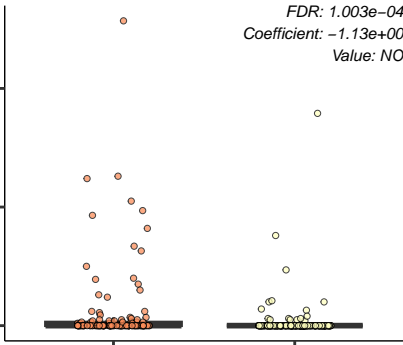
100

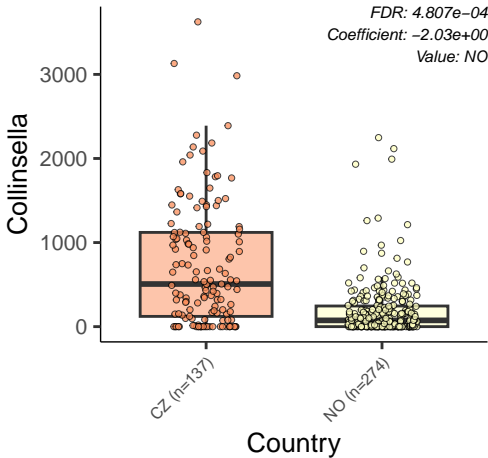
0

CZ (n=137)

NO (n=274)

Country





Eggerthella

1000

500

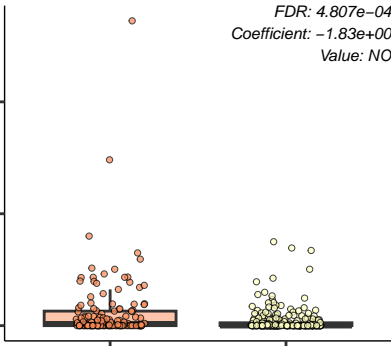
0

CZ (n=137)

NO (n=274)

Country

FDR: 4.807e-04
Coefficient: -1.83e+00
Value: NO



Faecalibacterium

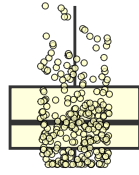
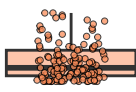
30000
20000
10000
0

CZ (n=137)

NO (n=274)

Country

FDR: 1.752e-03
Coefficient: 1.59e+00
Value: NO



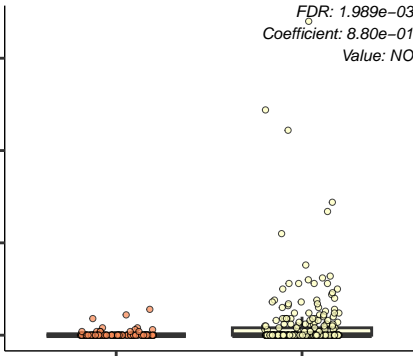
X.Eubacterium._xylanophilum_group

FDR: 1.989e-03
Coefficient: 8.80e-01
Value: NO

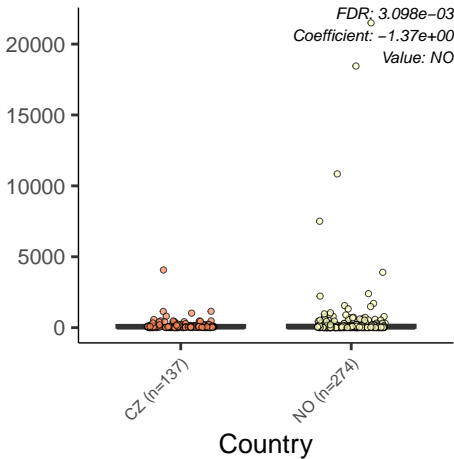
CZ (n=137)

NO (n=274)

Country



Streptococcus



Peptostreptococcus

200

100

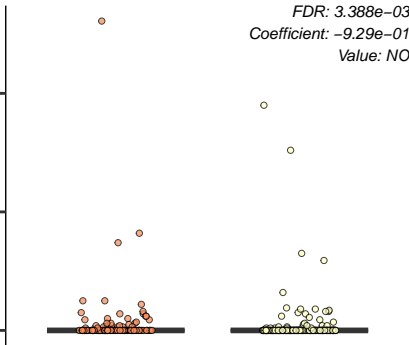
0

CZ (n=137)

NO (n=274)

Country

FDR: $3.388e-03$
Coefficient: $-9.29e-01$
Value: NO



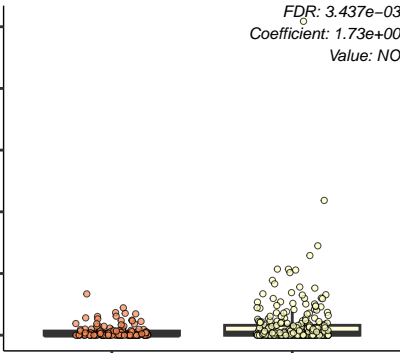
Christensenellaceae_R.7_group

FDR: 3.437e-03
Coefficient: 1.73e+00
Value: NO

CZ (n=137)

NO (n=274)

Country



Pseudomonas

FDR: 5.170e-03
Coefficient: 1.08e+00
Value: NO

6000

4000

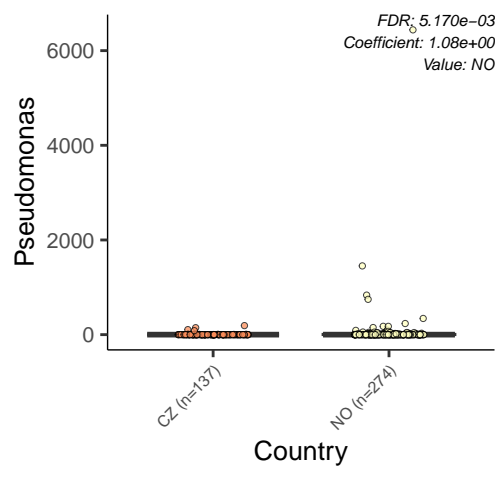
2000

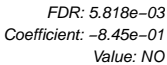
0

CZ (n=137)

NO (n=274)

Country





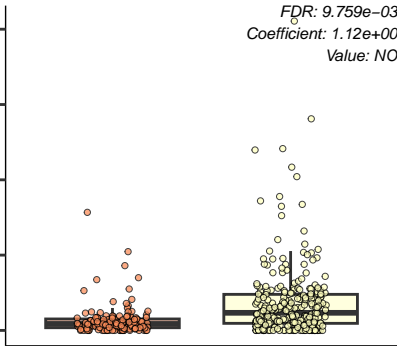
Lachnospiraceae_UCG.004

FDR: $9.759e-03$
Coefficient: $1.12e+00$
Value: NO

CZ (n=137)

NO (n=274)

Country



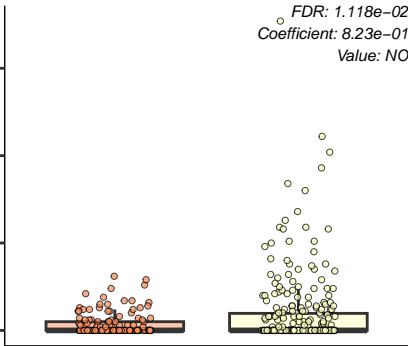
Family_XIII_UCG.001

FDR: 1.118e-02
Coefficient: 8.23e-01
Value: NO

CZ (n=137)

NO (n=274)

Country



X.Eubacterium._siraenum_group

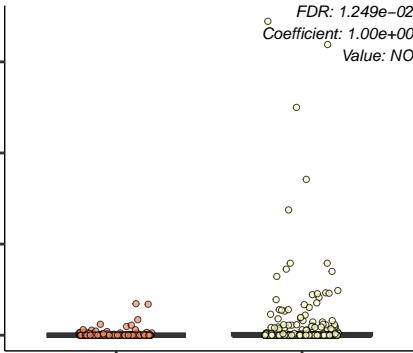
FDR: 1.249e-02
Coefficient: 1.00e+00
Value: NO

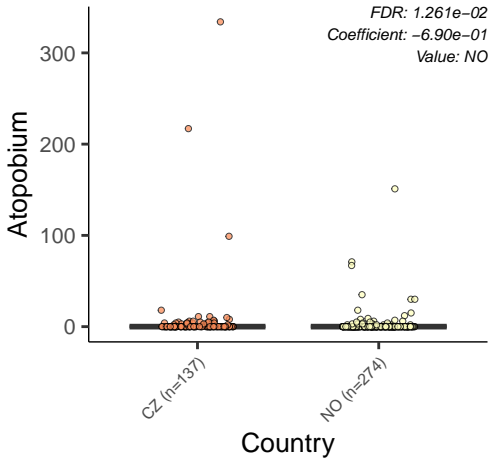
600
400
200
0

CZ (n=137)

NO (n=274)

Country





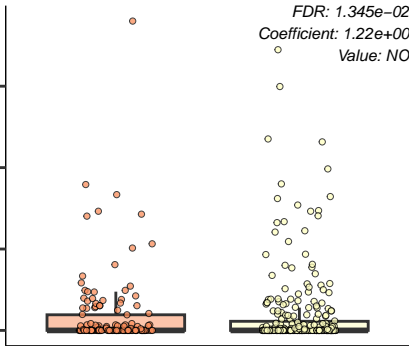
Clostridia_UCG.014

FDR: 1.345e-02
Coefficient: 1.22e+00
Value: NO

CZ (n=137)

NO (n=274)

Country



Turicibacter

10000

5000

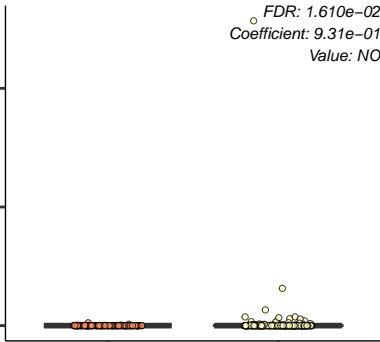
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CZ (n=137)

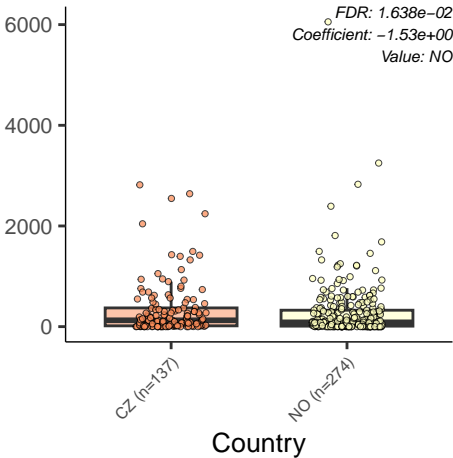
NO (n=274)

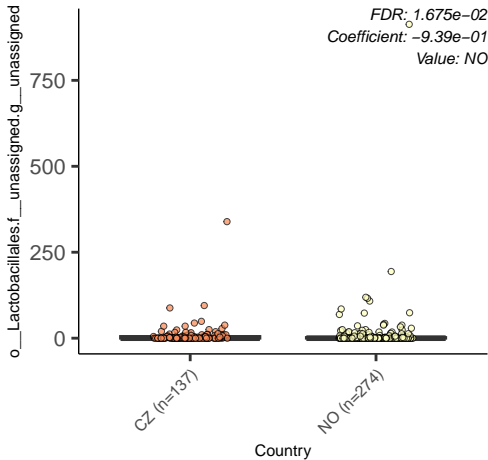
Country

FDR: 1.610e-02
Coefficient: 9.31e-01
Value: NO

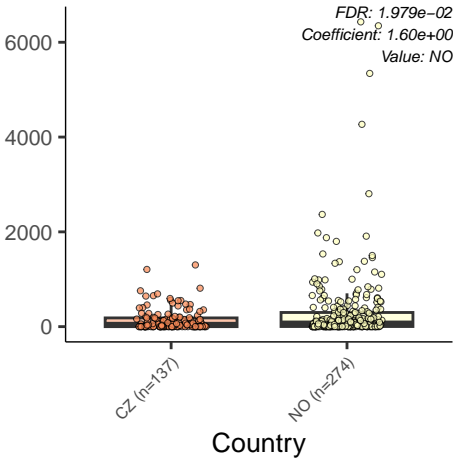


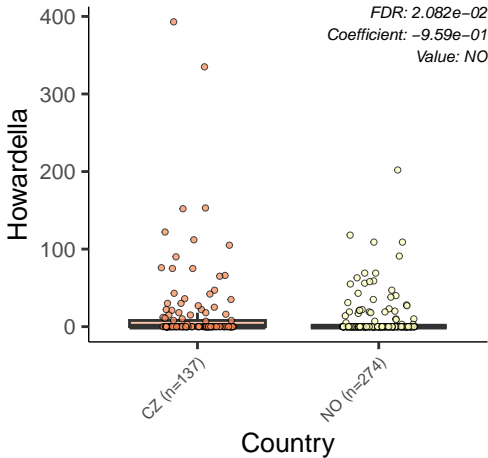
Bifidobacterium



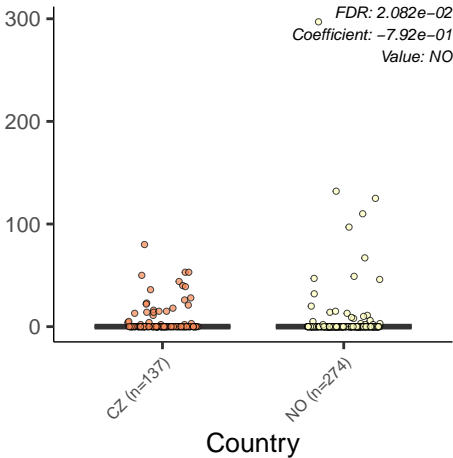


Agathobacter



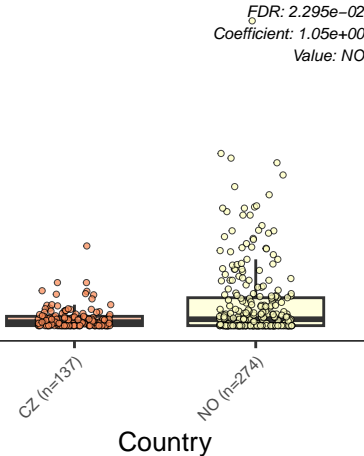


Megasphaera



GCA.900066575

FDR: 2.295e-02
Coefficient: 1.05e+00
Value: NO



Escherichia.Shigella

FDR: 2.377e-02
Coefficient: 2.07e+00
Value: NO

40000

30000

20000

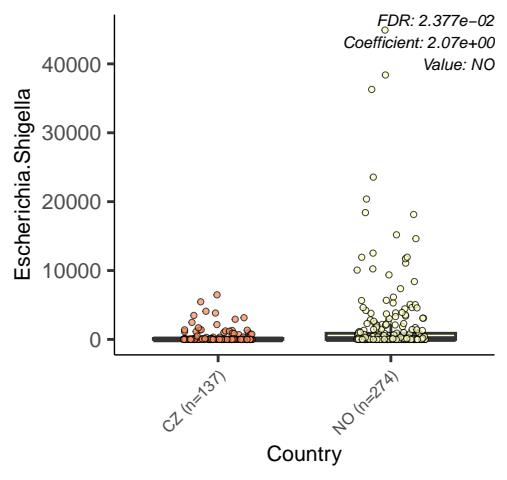
10000

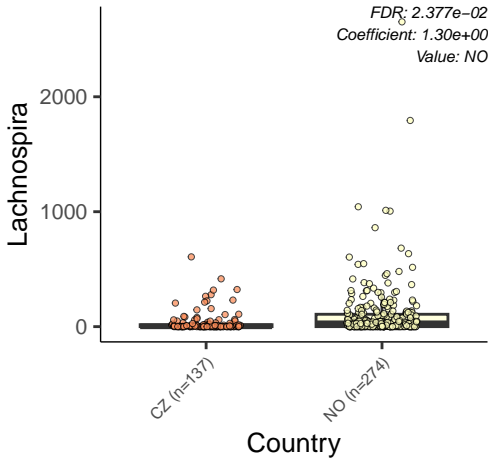
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CZ (n=137)

NO (n=274)

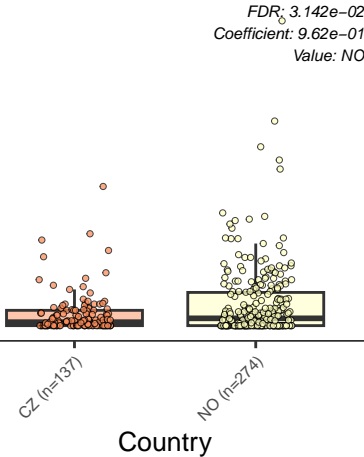
Country





Marvinbryantia

FDR: $3.142e-02$
Coefficient: $9.62e-01$
Value: NO



f_Ruminococcaceae.g__uncultured

FDR: $3.162e-02$
Coefficient: $8.85e-01$
Value: NO

150

100

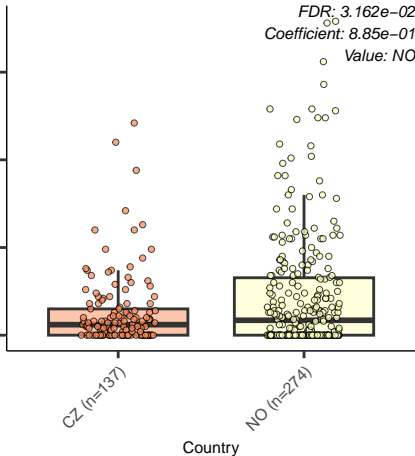
50

0

CZ (n=137)

NO (n=274)

Country



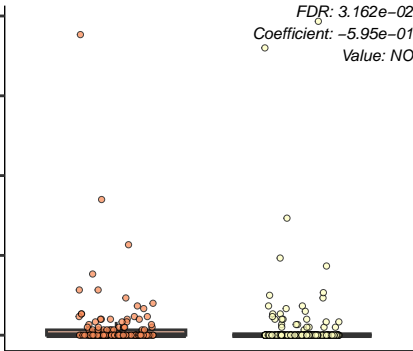
X.Eubacterium._brachy_group

FDR: 3.162e-02
Coefficient: -5.95e-01
Value: NO

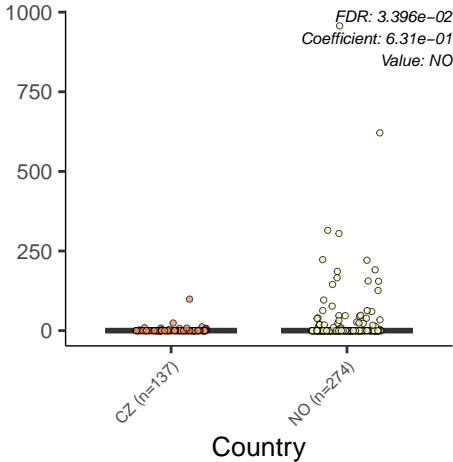
CZ (n=137)

NO (n=274)

Country



Tannerellaceae



f__Lachnospiraceae.g__unassigned

FDR: 3.406e-02
Coefficient: 8.79e-01
Value: NO

7500

5000

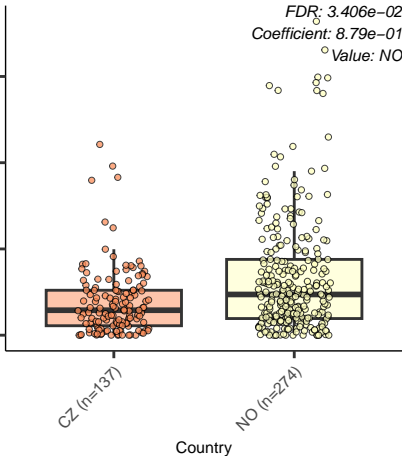
2500

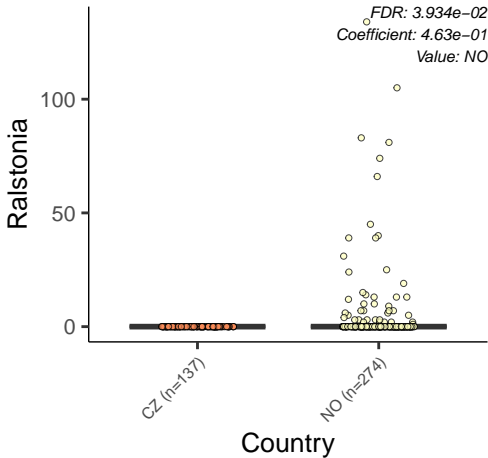
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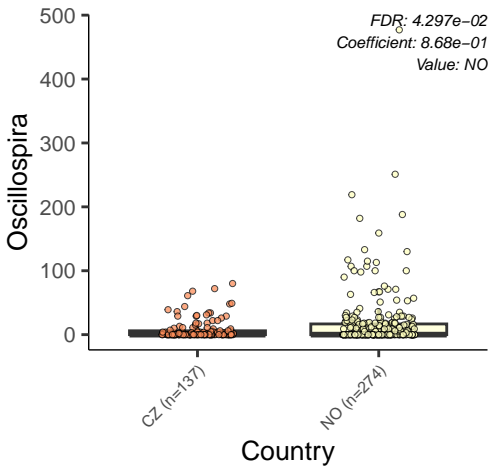
CZ (n=137)

NO (n=274)

Country







Acidaminococcus

5000
4000
3000
2000
1000
0

FDR: 4.636e-02
Coefficient: -7.00e-01
Value: NO

CZ (n=137)

NO (n=274)

Country

