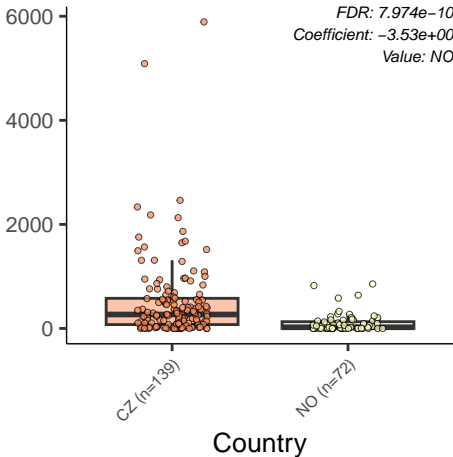
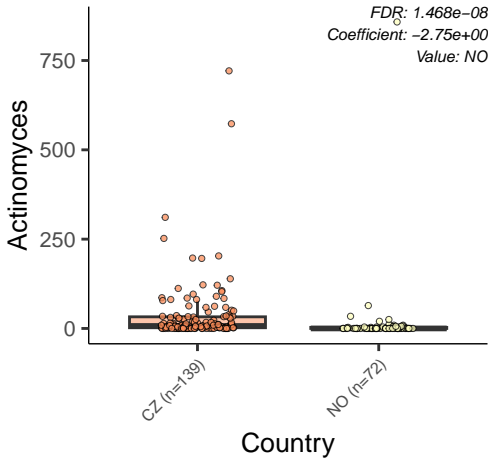


Bifidobacterium





Collinsella

FDR: 3.602e-08

Coefficient: -3.29e+00

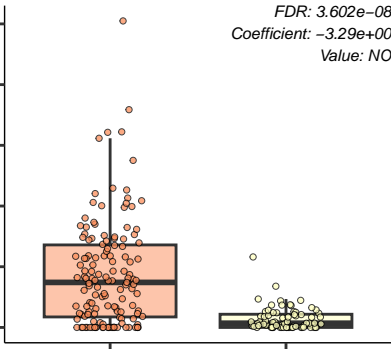
Value: NO

5000
4000
3000
2000
1000
0

CZ (n=139)

NO (n=72)

Country



Streptococcus

12000

9000

6000

3000

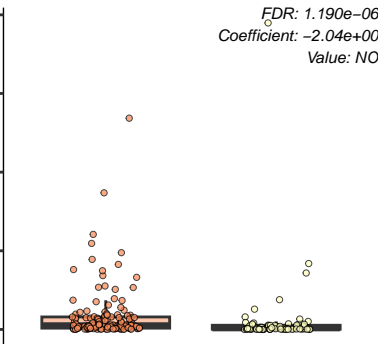
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CZ (n=139)

NO (n=72)

Country

FDR: $1.190e-06$
Coefficient: $-2.04e+00$
Value: NO



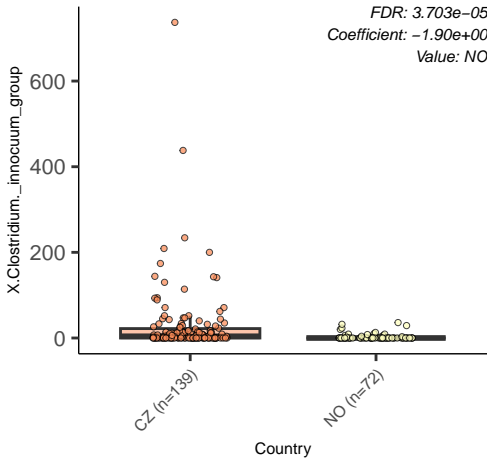
X.Clostridium._innocuum_group

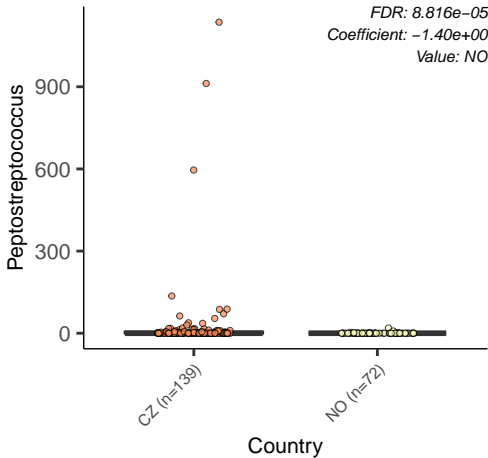
FDR: 3.703e-05
Coefficient: -1.90e+00
Value: NO

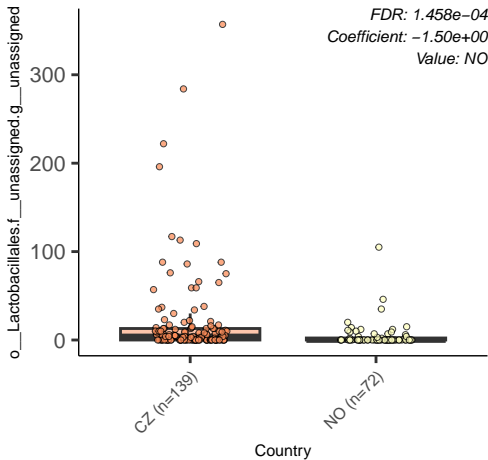
CZ (n=139)

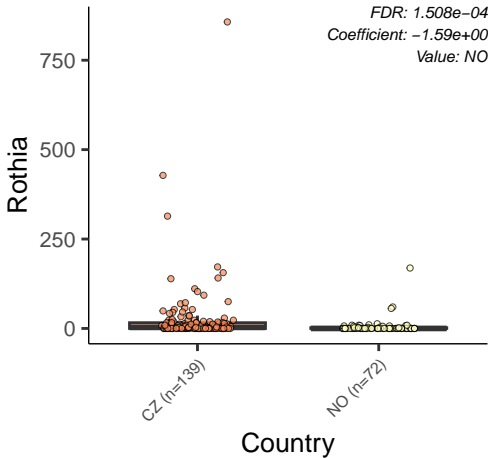
NO (n=72)

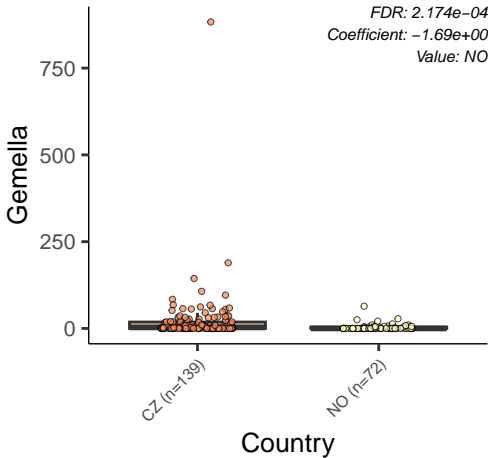
Country



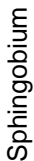








Value: NO



200

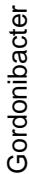
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0

CZ (n=139)

NO (n=72)

Country



FDR: 3.585e-04

Coefficient: -9.08e-01

Value: NO

CZ (n=139)

NO (n=72)

Country

Acidaminococcus

FDR: 6.187e-04
Coefficient: -1.89e+00
Value: NO

CZ (n=139)

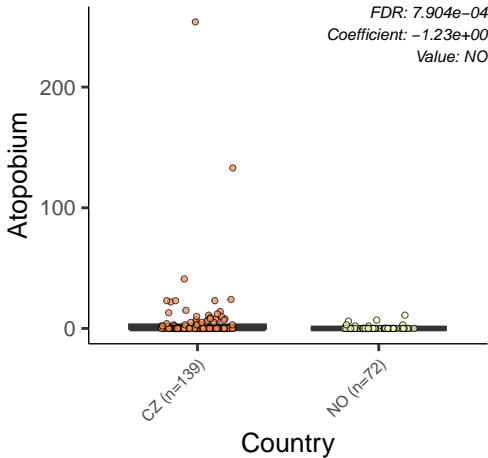
NO (n=72)

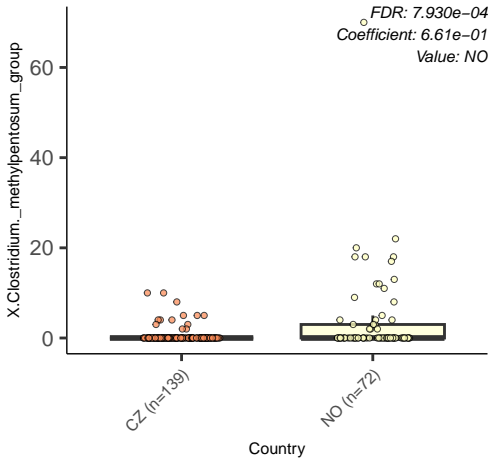
Country

1000

500

0





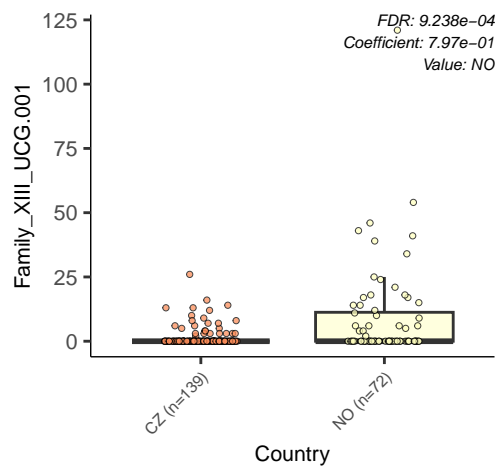
Family_XIII_UCG.001

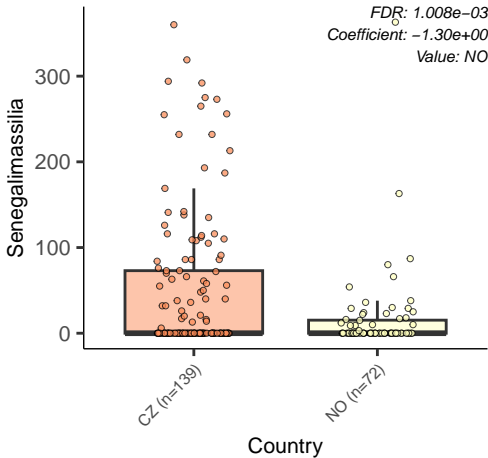
FDR: 9.238e-04
Coefficient: 7.97e-01
Value: NO

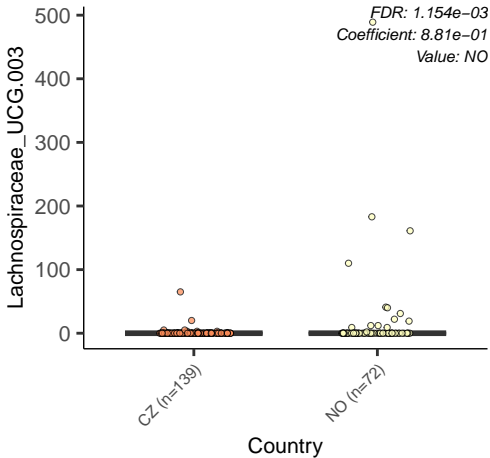
CZ (n=139)

NO (n=72)

Country







Erysipelatoclostridium

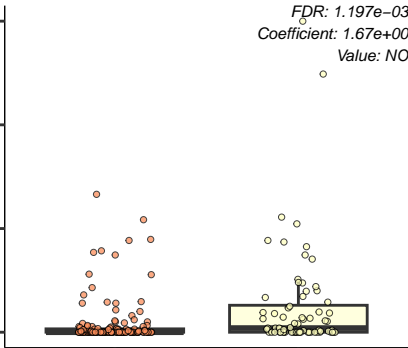
600
400
200
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FDR: 1.197e-03
Coefficient: 1.67e+00
Value: NO

CZ (n=139)

NO (n=72)

Country



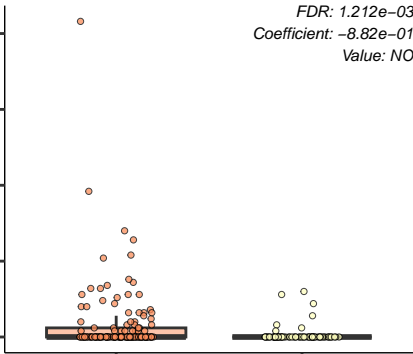
X.Eubacterium._brachy_group

FDR: 1.212e-03
Coefficient: -8.82e-01
Value: NO

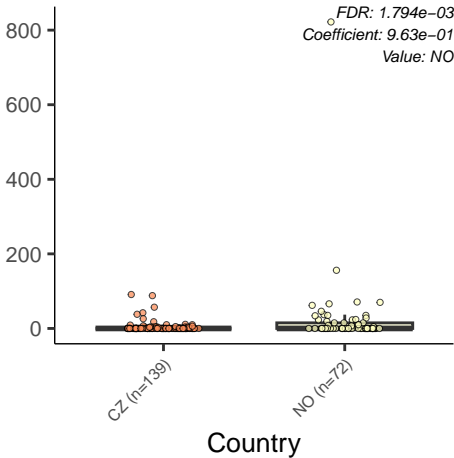
CZ (n=139)

NO (n=72)

Country



Eisenbergiella



f__Peptococcaceae.g__uncultured

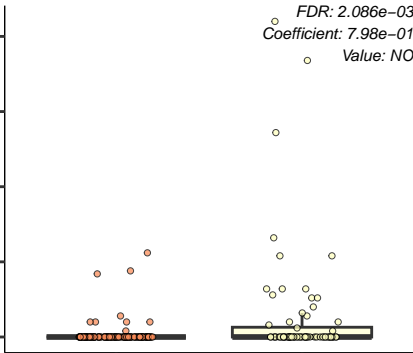
FDR: 2.086e-03
Coefficient: 7.98e-01
Value: NO

100
75
50
25
0

CZ (n=139)

NO (n=72)

Country



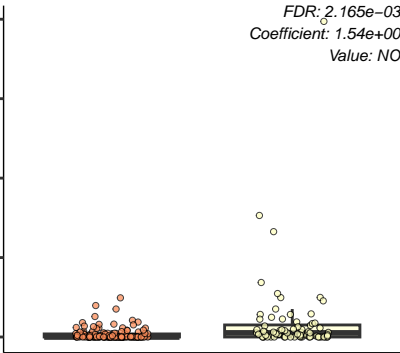
Christensenellaceae_R.7_group

FDR: 2.165e-03
Coefficient: 1.54e+00
Value: NO

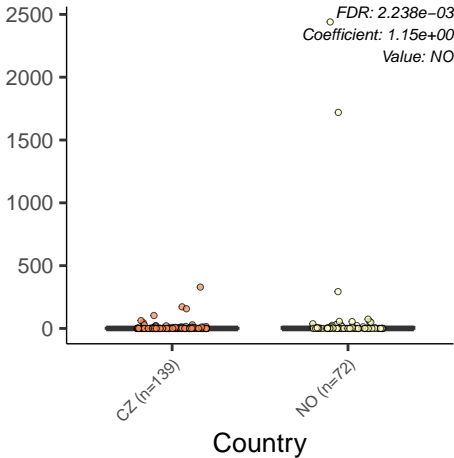
CZ (n=139)

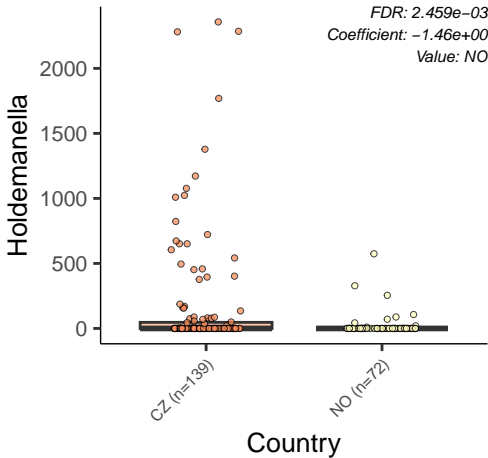
NO (n=72)

Country

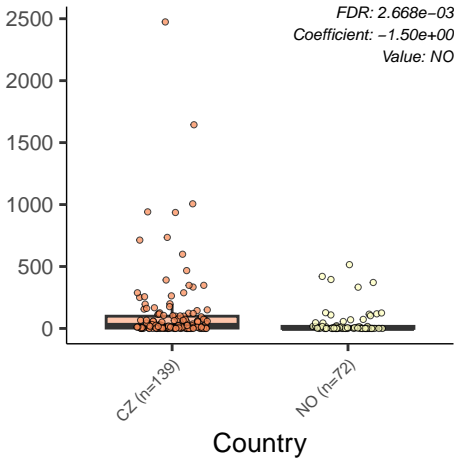


Hungatella





Haemophilus



Escherichia.Shigella

FDR: $3.429e-03$
Coefficient: $1.57e+00$
Value: NO

30000

20000

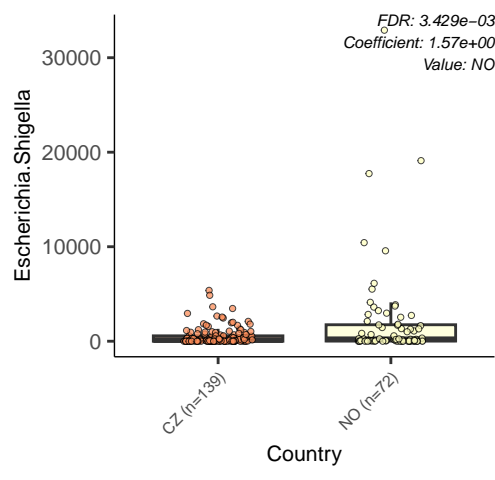
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0

CZ (n=139)

NO (n=72)

Country



CAG.56

FDR: 4.291e-03
Coefficient: 1.49e+00
Value: NO

750

500

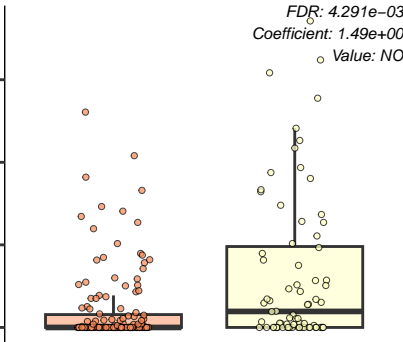
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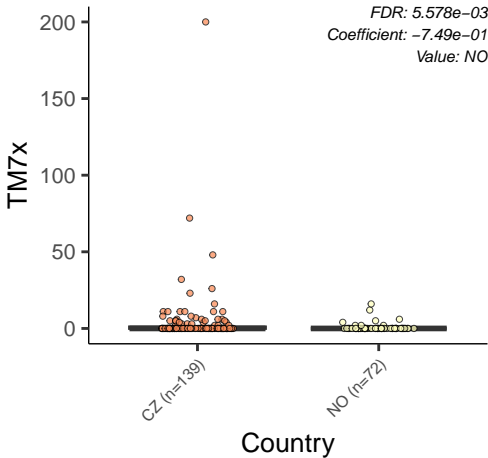
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CZ (n=139)

NO (n=72)

Country





Mogibacterium

FDR: 5.830e-03

Coefficient: -9.18e-01

Value: NO

200

150

100

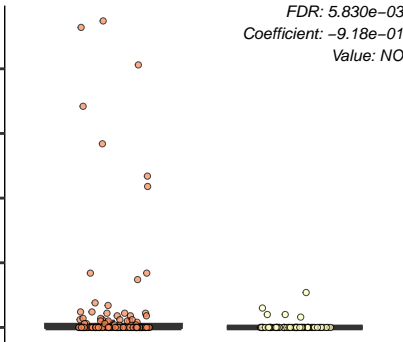
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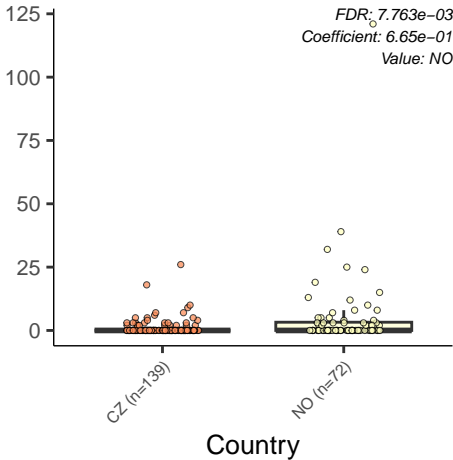
CZ (n=139)

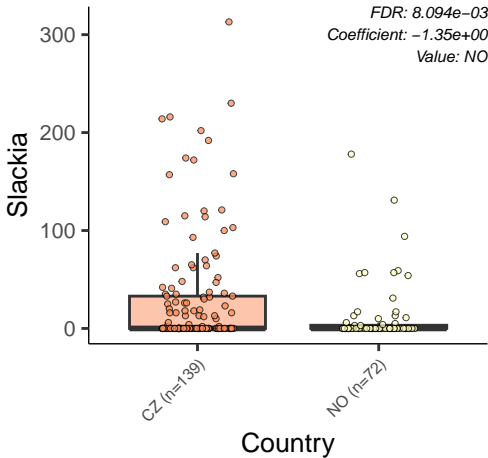
NO (n=72)

Country



Anaerotruncus





Romboutsia

1500

1000

500

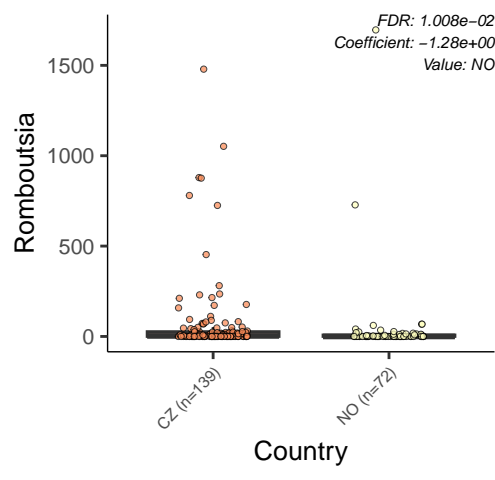
0

FDR: 1.008e-02
Coefficient: -1.28e+00
Value: NO

CZ (n=139)

NO (n=72)

Country



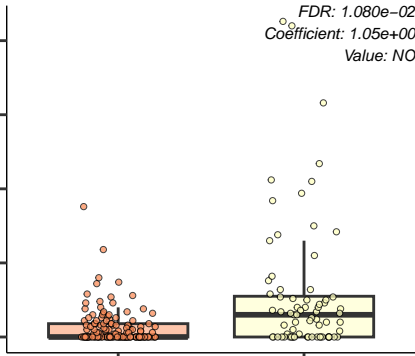
f__Ruminococcaceae.g__uncultured

FDR: 1.080e-02
Coefficient: 1.05e+00
Value: NO

CZ (n=139)

NO (n=72)

Country



Marvinbryantia

FDR: 1.200e-02
Coefficient: 1.00e+00
Value: NO

200

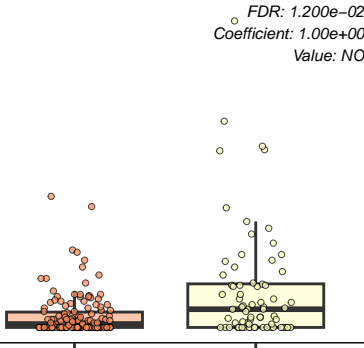
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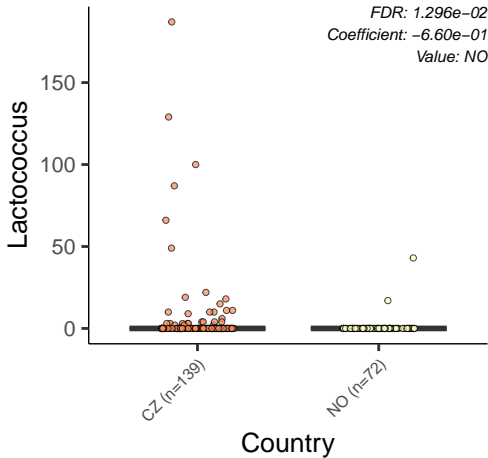
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CZ (n=139)

NO (n=72)

Country





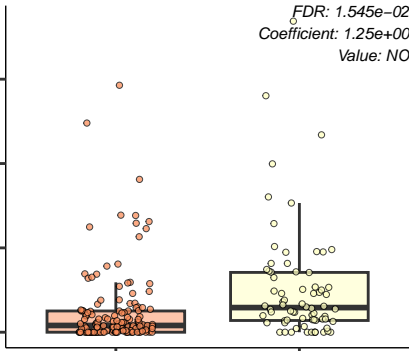
Lachnospiraceae_UCG.004

FDR: 1.545e-02
Coefficient: 1.25e+00
Value: NO

CZ (n=139)

NO (n=72)

Country



Corynebacterium

100

50

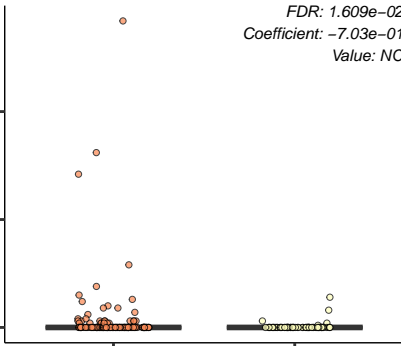
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CZ (n=139)

NO (n=72)

Country

FDR: $1.609e-02$
Coefficient: $-7.03e-01$
Value: NO



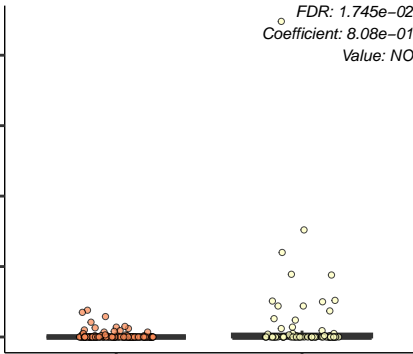
X.Eubacterium._siraeum_group

FDR: 1.745e-02
Coefficient: 8.08e-01
Value: NO

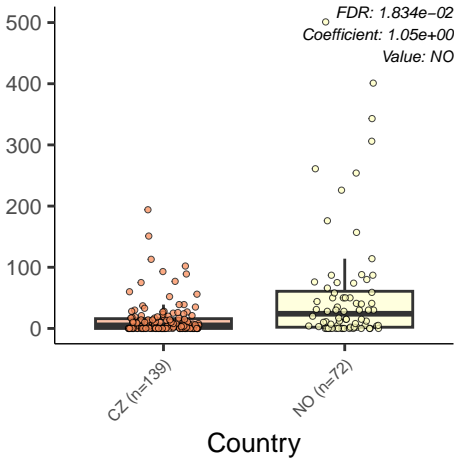
CZ (n=139)

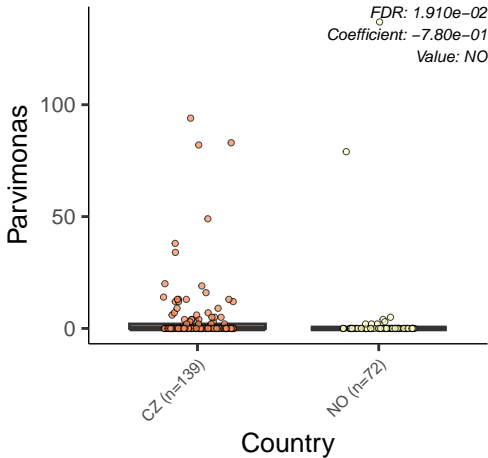
NO (n=72)

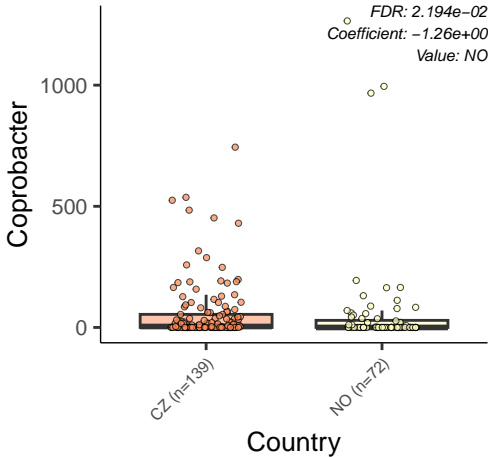
Country



Intestinimonas







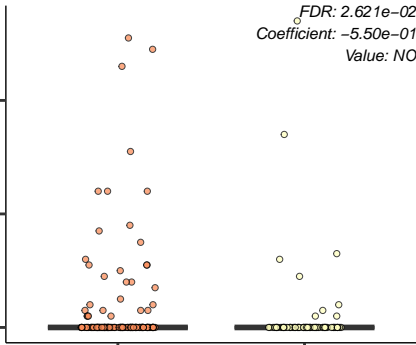
Enterorhabdus

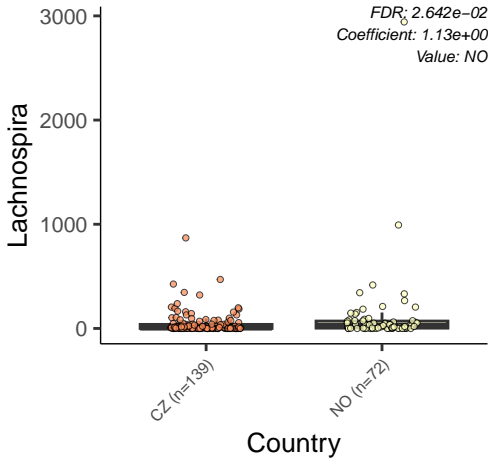
FDR: 2.621e-02
Coefficient: -5.50e-01
Value: NO

CZ (n=139)

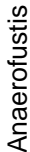
NO (n=72)

Country





Value: NO



CZ (n=139)

NO (n=72)

Country

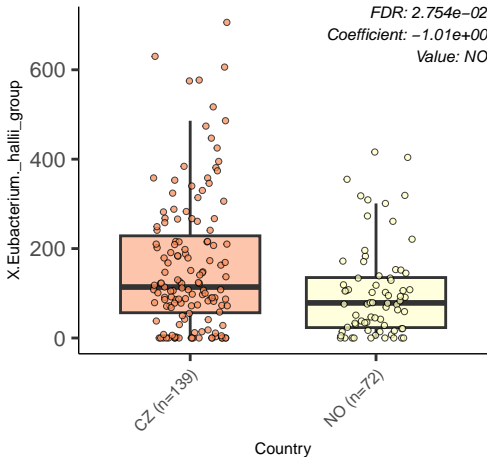
X.Eubacterium._hallii_group

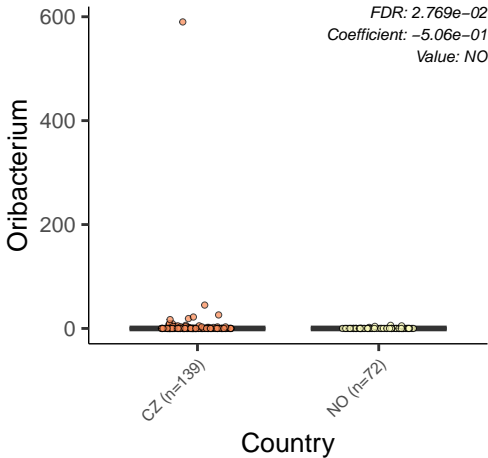
FDR: 2.754e-02
Coefficient: -1.01e+00
Value: NO

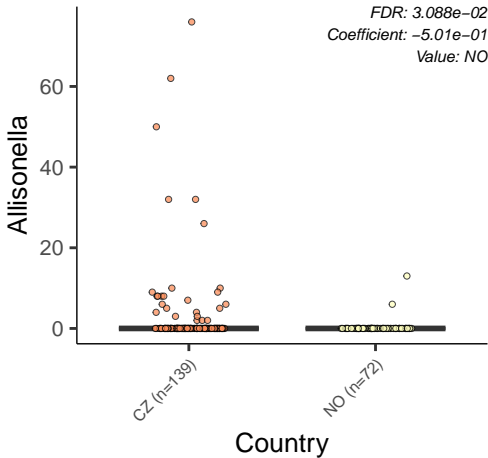
CZ (n=139)

NO (n=72)

Country







Megasphaera

FDR: 3.259e-02

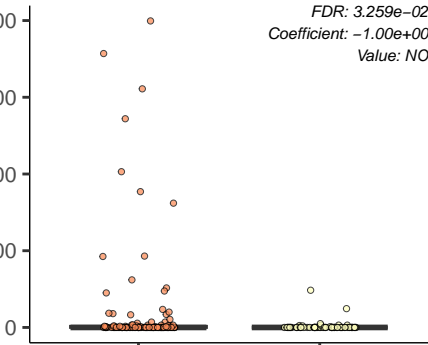
Coefficient: -1.00e+00

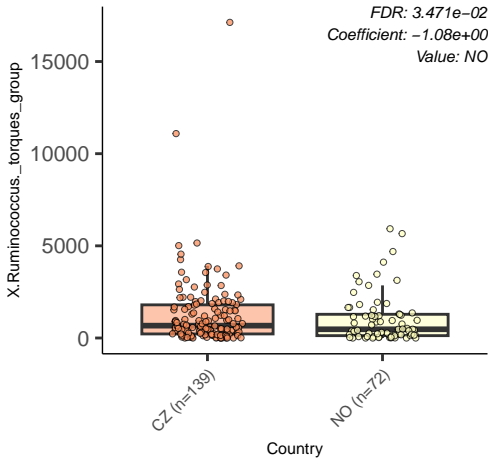
Value: NO

CZ (n=139)

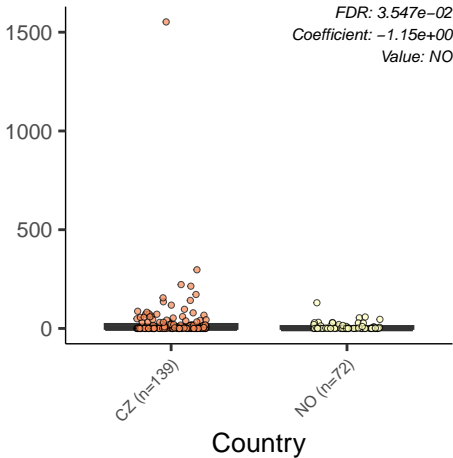
NO (n=72)

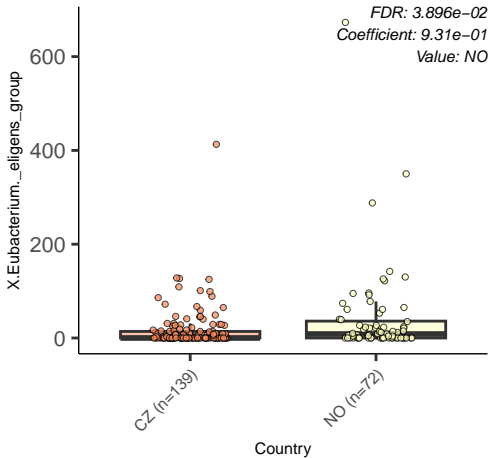
Country





Eggerthella





Flavonifactor

FDR: 4.008e-02

Coefficient: -8.89e-01

Value: NO

CZ (n=139)

NO (n=72)

Country

