

Gordonibacter

FDR: $7.284e-05$

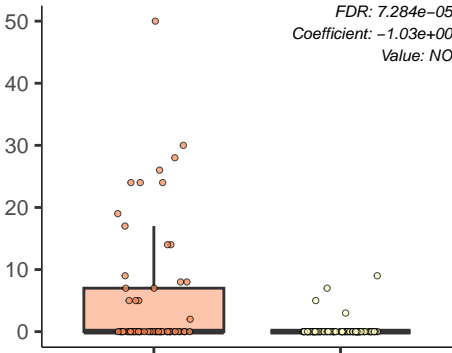
Coefficient: $-1.03e+00$

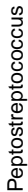
Value: NO

CZ (n=61)

NO (n=82)

Country





FDR: 1.262e-04

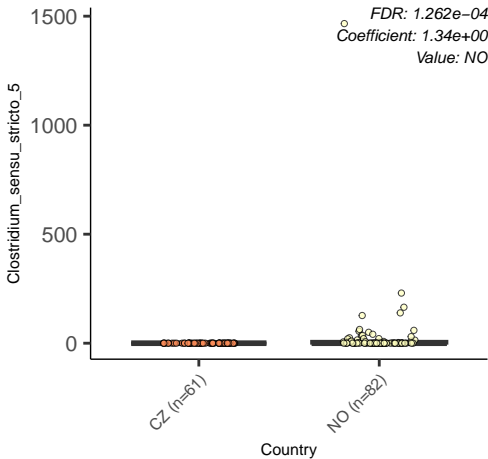
Coefficient: $-1.38e+00$

Value: NO

CZ (n=61)

NO (n=82)

Country



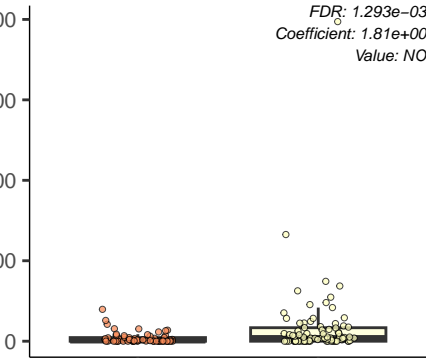
Christensenellaceae_R.7_group

FDR: 1.293e-03
Coefficient: 1.81e+00
Value: NO

CZ (n=61)

NO (n=82)

Country



Mogibacterium

FDR: $1.332e-03$
Coefficient: $-1.04e+00$
Value: NO

100

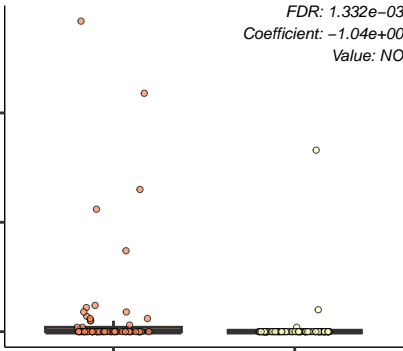
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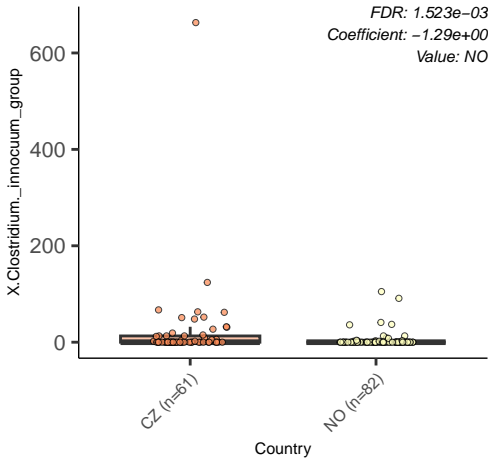
0

CZ (n=61)

NO (n=82)

Country





Value: NO



150

50

0

CZ (n=61)

NO (n=82)

Country

Eggerthella

FDR: 1.916e-03

Coefficient: -1.98e+00

Value: NO

750

500

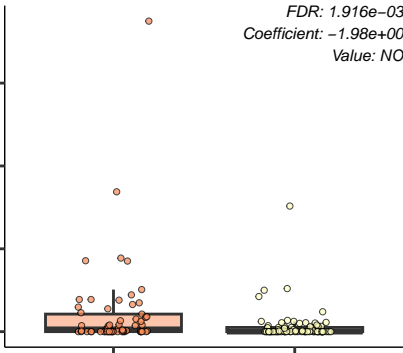
250

0

CZ (n=61)

NO (n=82)

Country



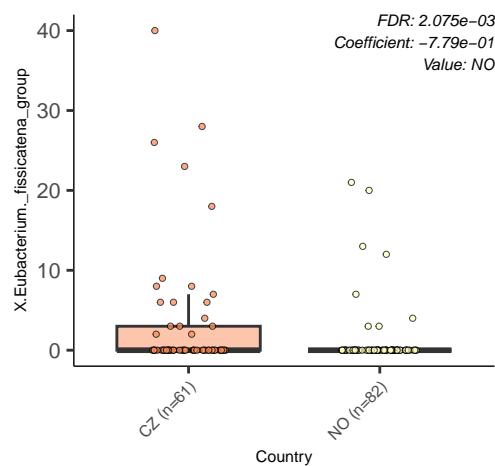
X.Eubacterium._fissicatena_group

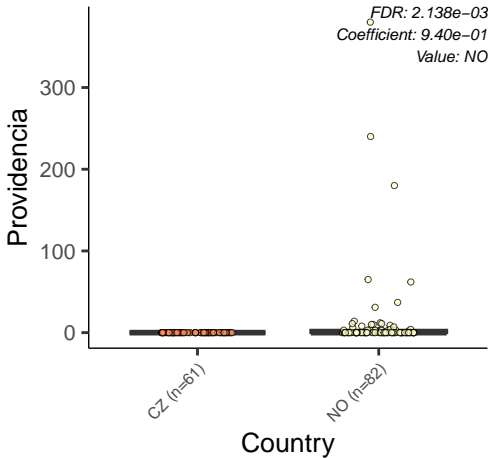
FDR: 2.075e-03
Coefficient: -7.79e-01
Value: NO

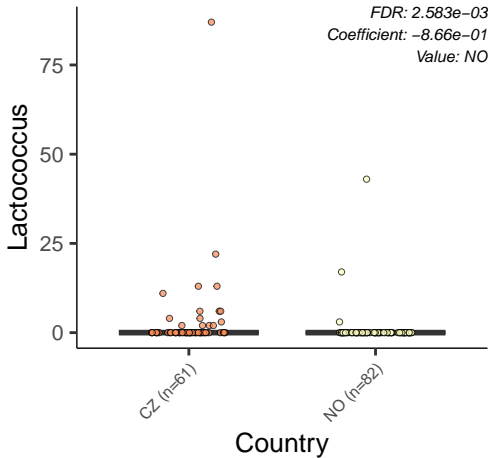
CZ (n=61)

NO (n=82)

Country







Streptococcus

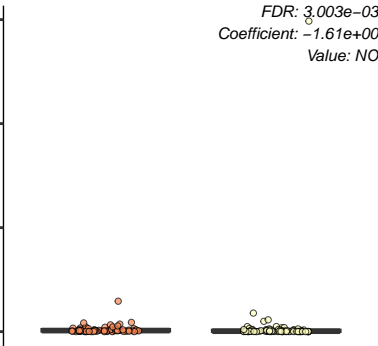
30000
20000
10000
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CZ (n=61)

NO (n=82)

Country

FDR: 3.003e-03
Coefficient: -1.61e+00
Value: NO



Faecalibacterium

FDR: $3.483e-03$
Coefficient: $1.70e+00$
Value: NO

20000

15000

10000

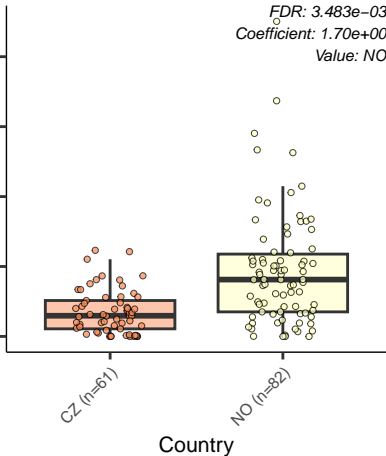
5000

0

CZ (n=61)

NO (n=82)

Country



Pseudomonas

FDR: 4.323e-03
Coefficient: 1.31e+00
Value: NO

1000

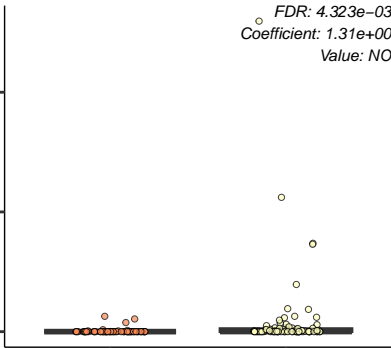
500

0

CZ (n=61)

NO (n=82)

Country



Bifidobacterium

FDR: 4.414e-03

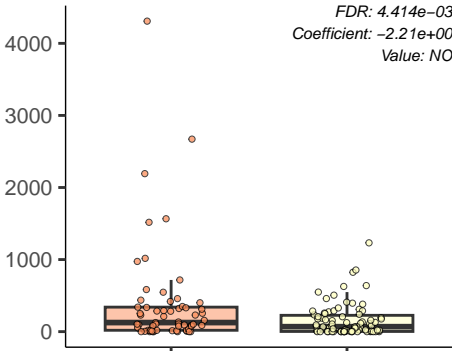
Coefficient: -2.21e+00

Value: NO

CZ (n=61)

NO (n=82)

Country



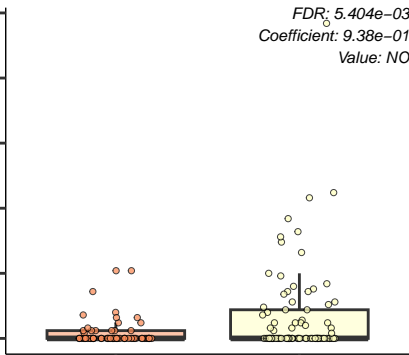
Family_XIII_UCG.001

FDR: $5.404e-03$
Coefficient: $9.38e-01$
Value: NO

CZ (n=61)

NO (n=82)

Country



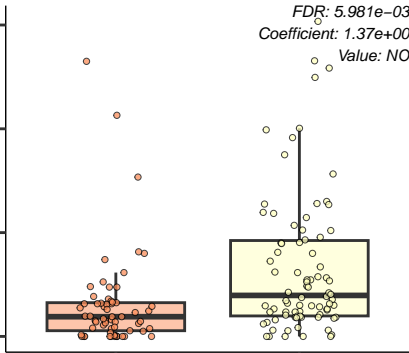
Lachnospiraceae_UCG.004

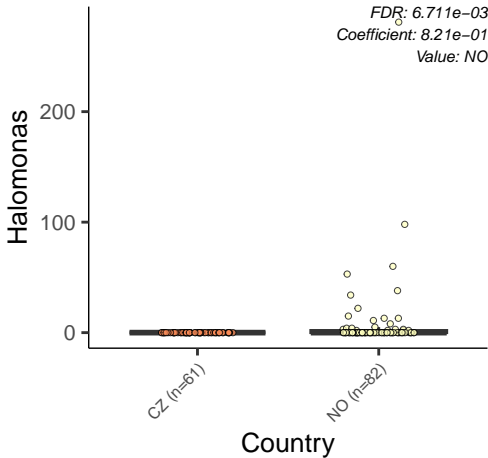
FDR: 5.981e-03
Coefficient: 1.37e+00
Value: NO

CZ (n=61)

NO (n=82)

Country





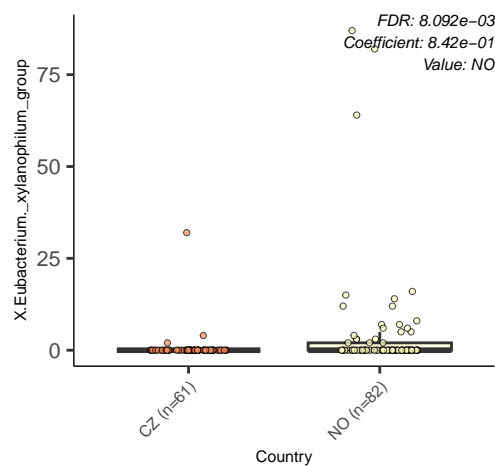
X.Eubacterium._xylanophilum_group

FDR: 8.092e-03
Coefficient: 8.42e-01
Value: NO

CZ (n=61)

NO (n=82)

Country



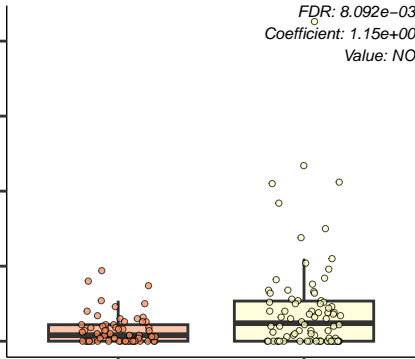
f__Ruminococcaceae.g__uncultured

FDR: $8.092e-03$
Coefficient: $1.15e+00$
Value: NO

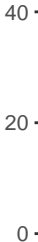
CZ (n=61)

NO (n=82)

Country



Value: NO

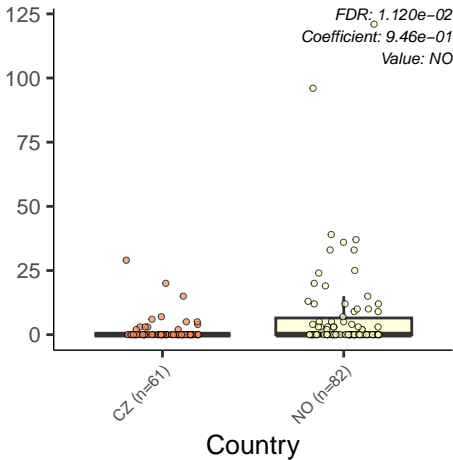


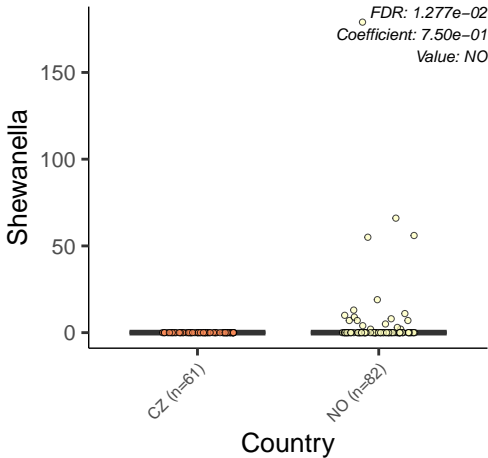
CZ (n=61)

NO (n=82)

Country

Anaerotruncus





Parabacteroides

1000

500

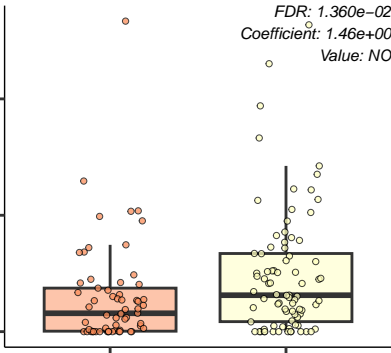
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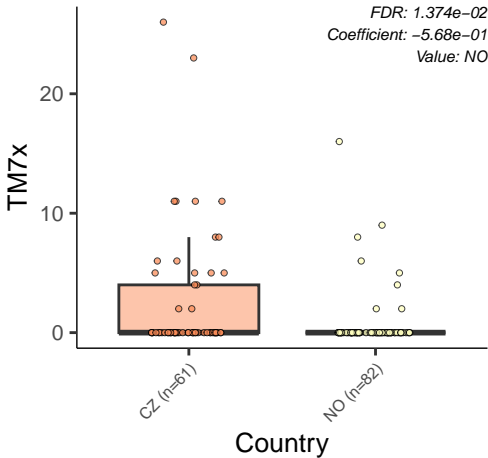
CZ (n=61)

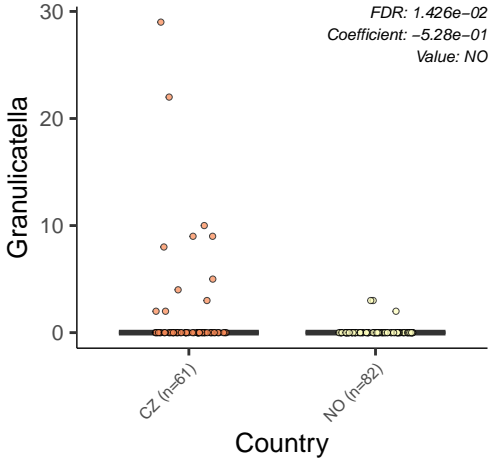
NO (n=82)

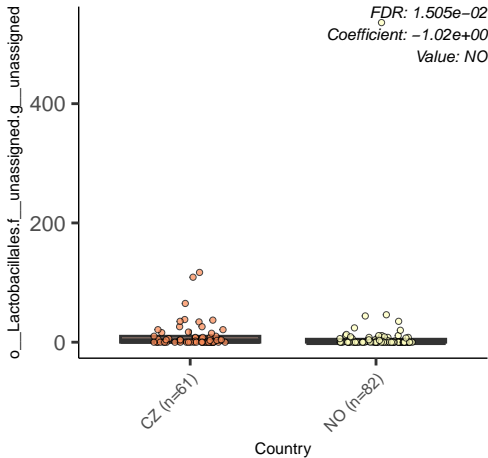
Country

FDR: 1.360e-02
Coefficient: 1.46e+00
Value: NO









Tannerellaceae

FDR: 1.656e-02
Coefficient: 7.29e-01
Value: NO

CZ (n=61)

NO (n=82)

Country

300

200

100

0

Clostridia_UCG.014

FDR: 1.962e-02
Coefficient: 1.16e+00
Value: NO

750

500

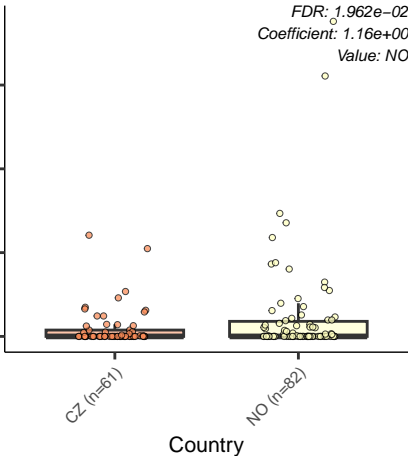
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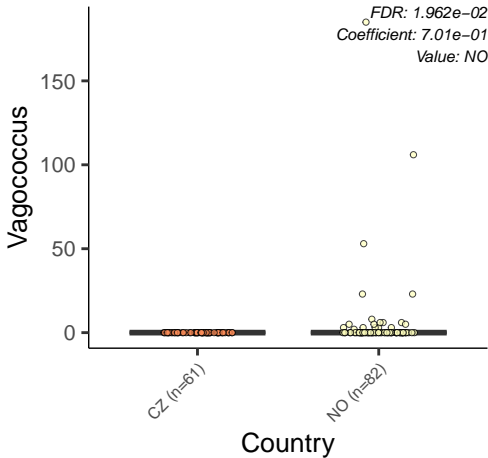
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CZ (n=61)

NO (n=82)

Country







FDR: 1.974e-02

Coefficient: -8.11e-01

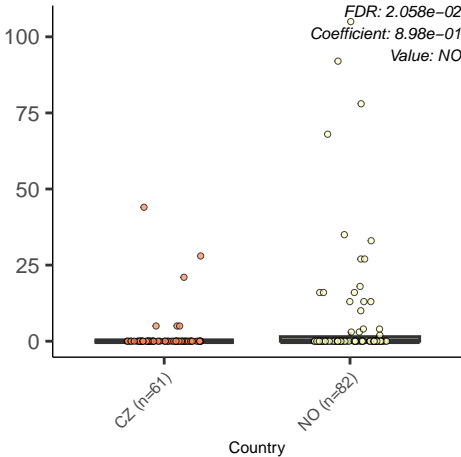
Value: NO



Country

f__Peptococcaceae.g__uncultured

FDR: 2.058e-02
Coefficient: 8.98e-01
Value: NO





FDR: 2.124e-02

Coefficient: -9.50e-01

Value: NO

CZ (n=61)

NO (n=82)

Country

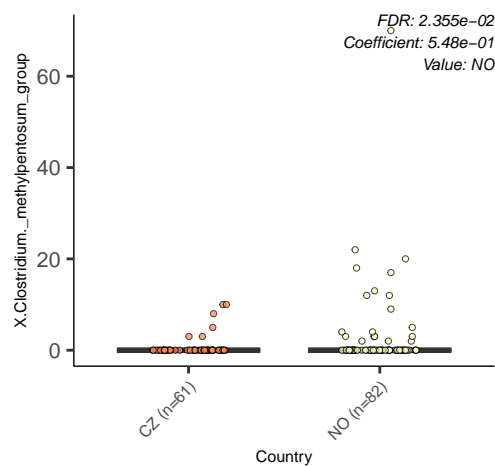
X.Clostridium._methylpentosum_group

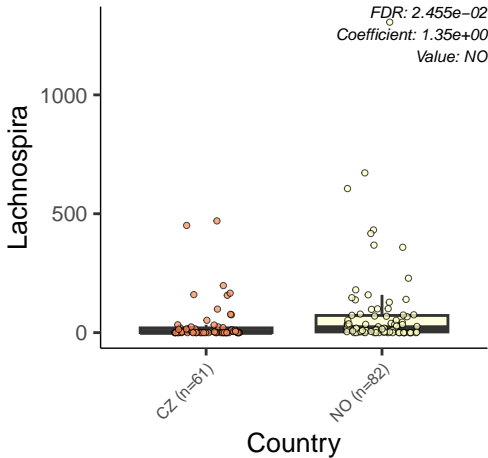
FDR: 2.355e-02
Coefficient: 5.48e-01
Value: NO

CZ (n=61)

NO (n=82)

Country





Burkholderia.Caballeronia.Paraburkholderia

FDR: 2.455e-02
Coefficient: 6.06e-01
Value: NO

150

100

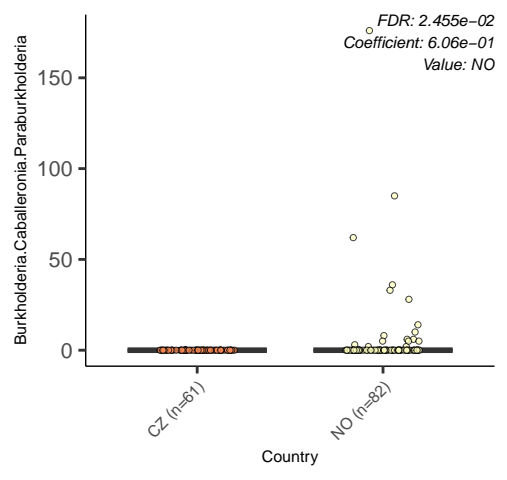
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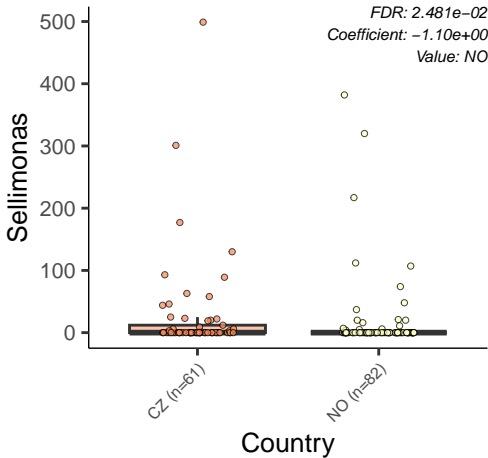
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CZ (n=61)

NO (n=82)

Country





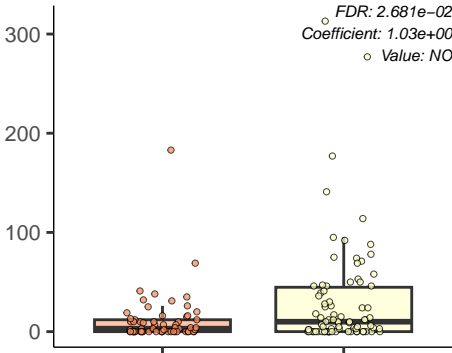
GCA.900066575

FDR: 2.681e-02
Coefficient: 1.03e+00
○ Value: NO

GZ (n=61)

NO (n=82)

Country



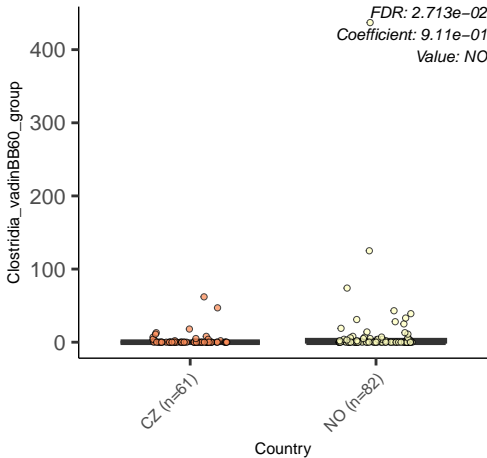
Clostridia_vadinBB60_group

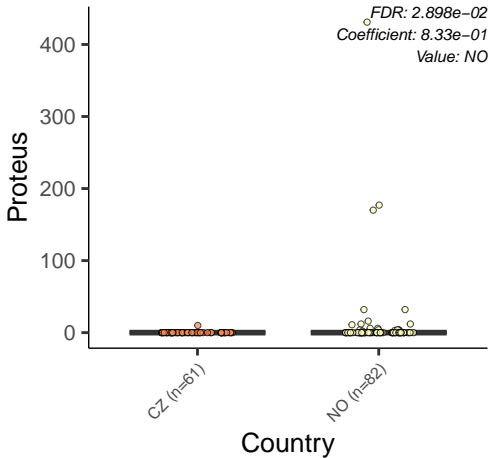
FDR: 2.713e-02
Coefficient: 9.11e-01
Value: NO

CZ (n=61)

NO (n=82)

Country





X.Eubacterium._brachy_group

FDR: 3.201e-02
Coefficient: -5.79e-01
Value: NO

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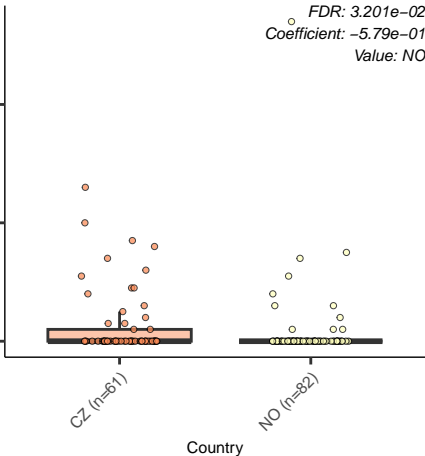
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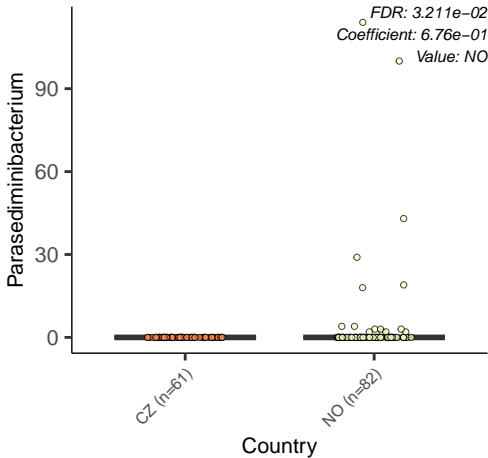
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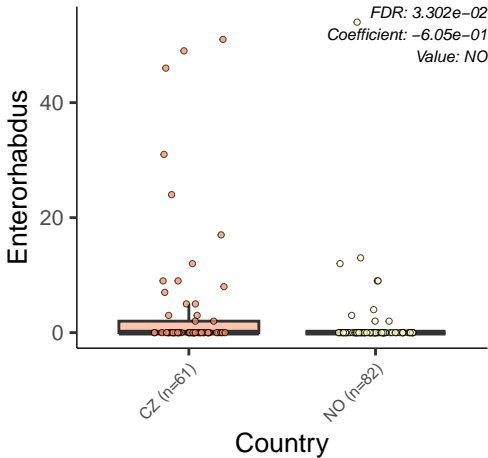
CZ (n=61)

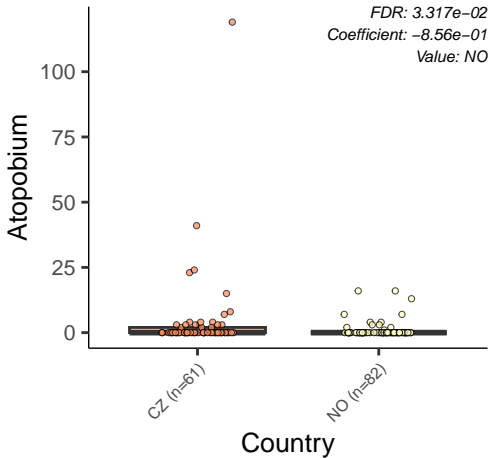
NO (n=82)

Country

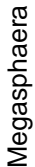








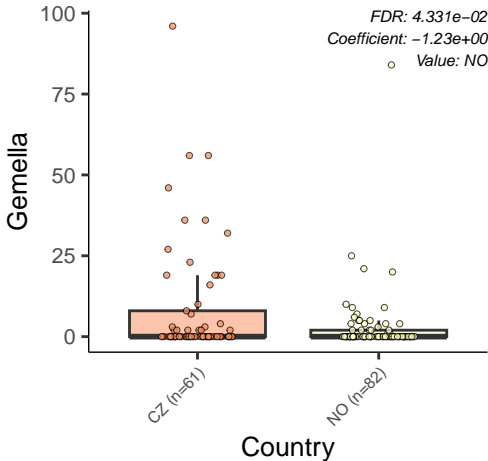
Value: NO



CZ (n=61)

NO (n=82)

Country



Flavonifractor

FDR: 4.565e-02

Coefficient: -1.08e+00

Value: NO

600

400

200

0

CZ (n=61)

NO (n=82)

Country

