

Gordonibacter

FDR: $7.284e-05$

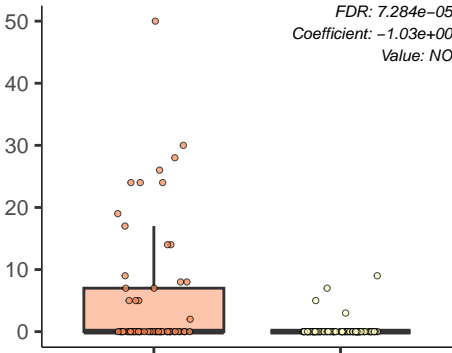
Coefficient: $-1.03e+00$

Value: NO

CZ (n=61)

NO (n=82)

Country



Peptostreptococcus

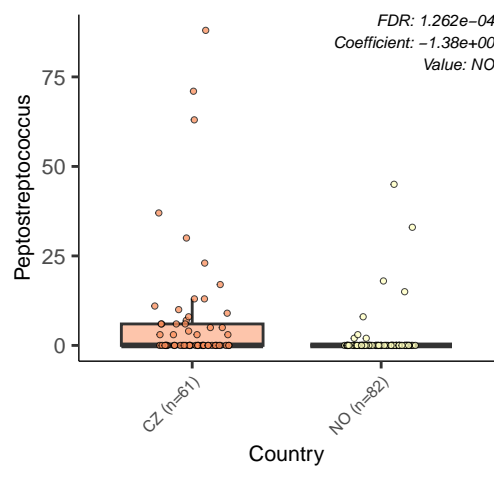
FDR: 1.262e-04
Coefficient: -1.38e+00
Value: NO

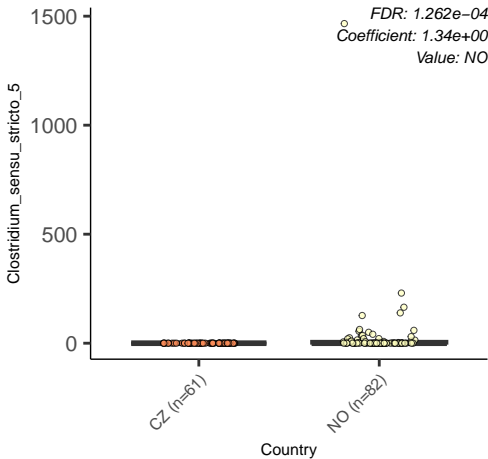
75
50
25
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CZ (n=61)

NO (n=82)

Country





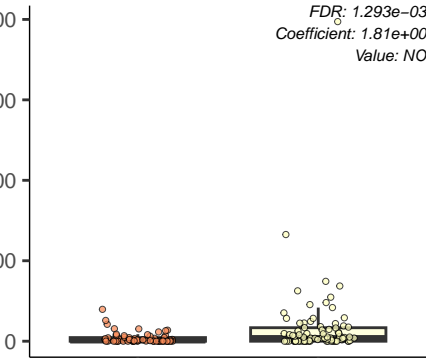
Christensenellaceae_R.7_group

FDR: 1.293e-03
Coefficient: 1.81e+00
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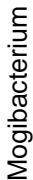
CZ (n=61)

NO (n=82)

Country



Value: NO



100

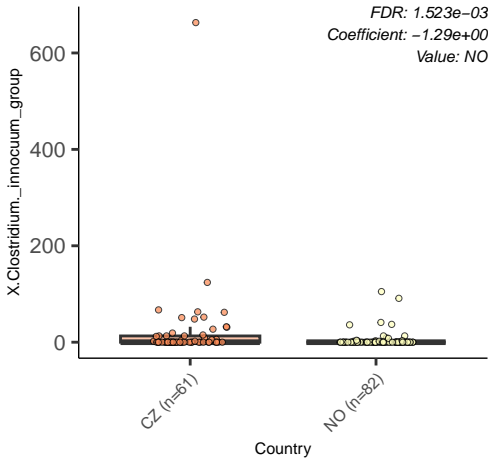
50

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

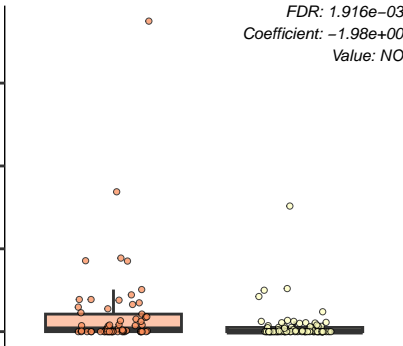
250

0

CZ (n=61)

NO (n=82)

Country



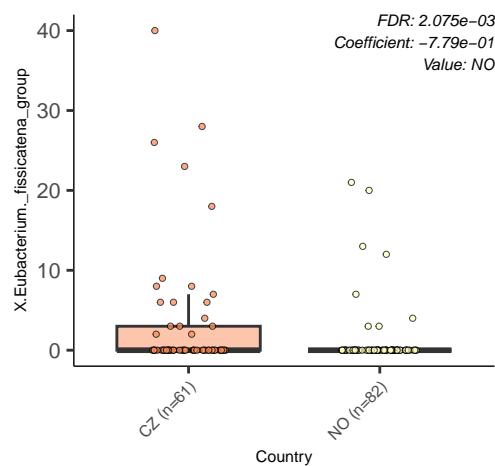
X.Eubacterium._fissicatena_group

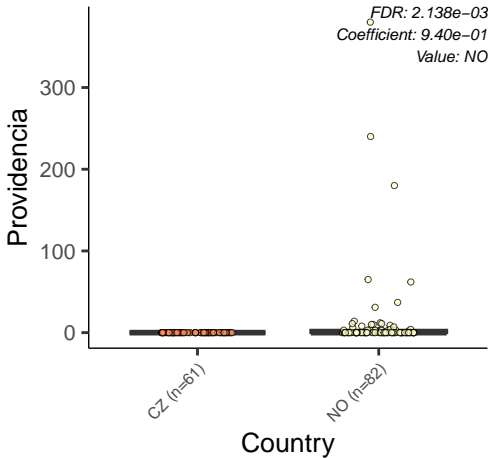
FDR: 2.075e-03
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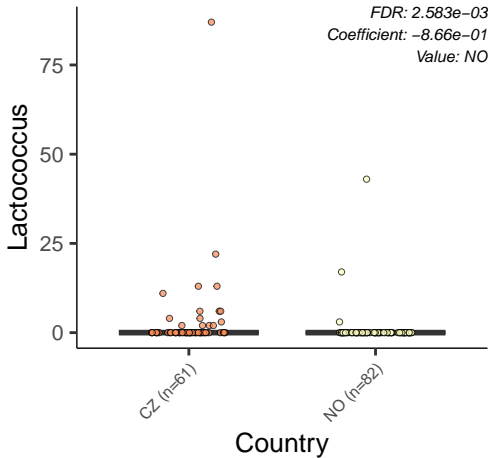
CZ (n=61)

NO (n=82)

Country







Streptococcus

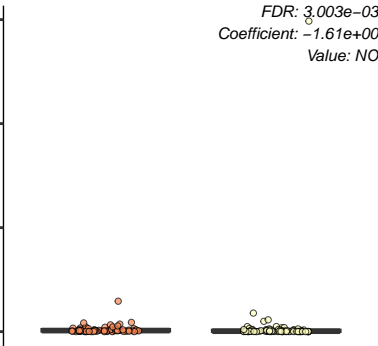
30000
20000
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CZ (n=61)

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

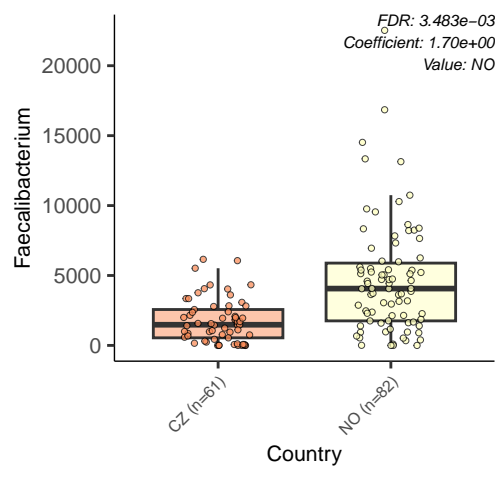
5000

0

CZ (n=61)

NO (n=82)

Country



Pseudomonas

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

1000

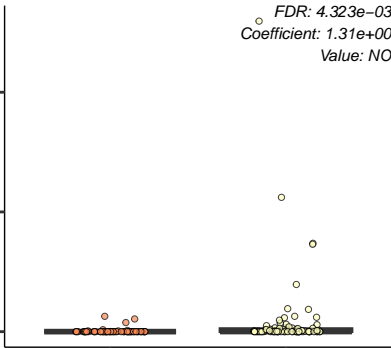
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0

CZ (n=61)

NO (n=82)

Country



Bifidobacterium

FDR: 4.414e-03

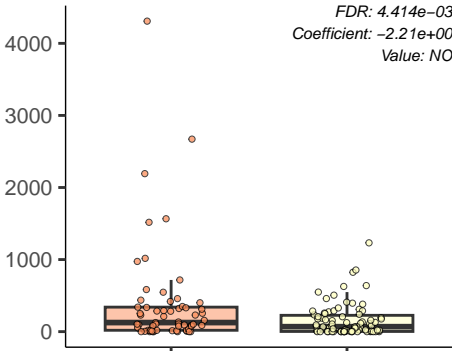
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CZ (n=61)

NO (n=82)

Country



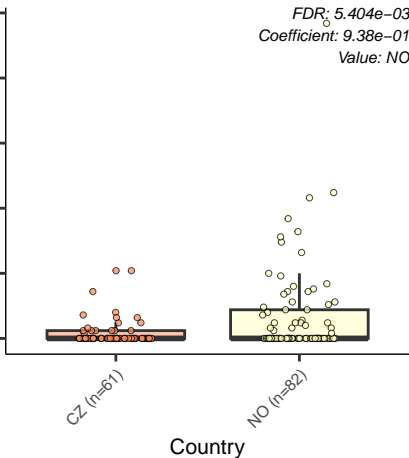
Family_XIII_UCG.001

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Coefficient: $9.38e-01$
Value: NO

CZ (n=61)

NO (n=82)

Country



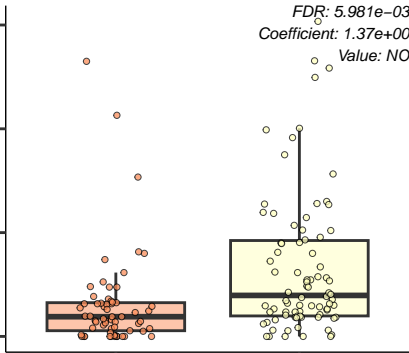
Lachnospiraceae_UCG.004

FDR: 5.981e-03
Coefficient: 1.37e+00
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CZ (n=61)

NO (n=82)

Country

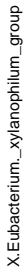


Value: NO



NO (n=82)

Country



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

CZ (n=61)

NO (n=82)

Country

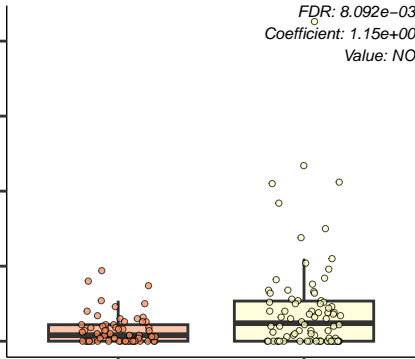
f__Ruminococcaceae.g__uncultured

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

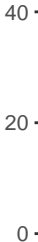
CZ (n=61)

NO (n=82)

Country



Value: NO

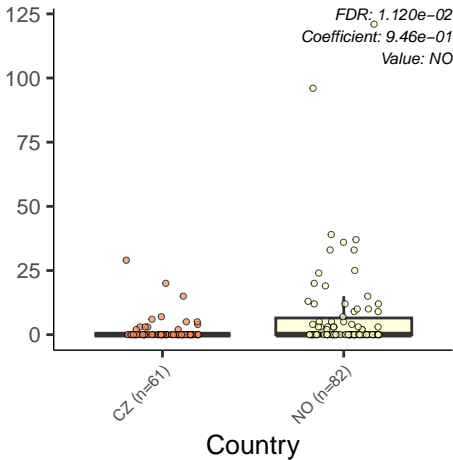


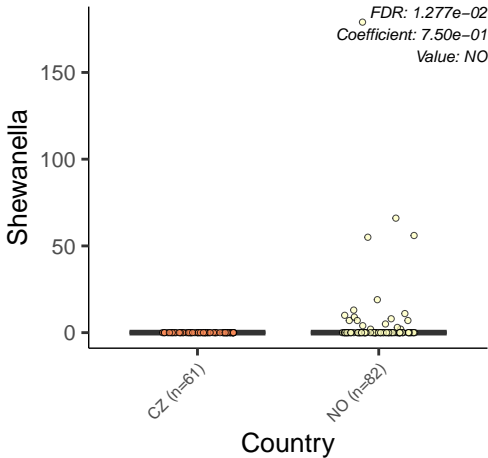
CZ (n=61)

NO (n=82)

Country

Anaerotruncus





Parabacteroides

1000

500

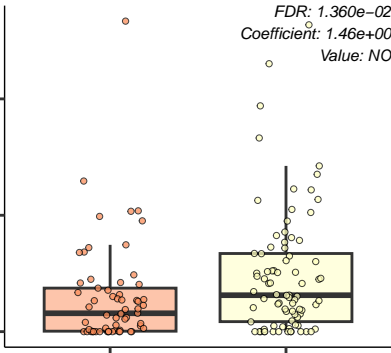
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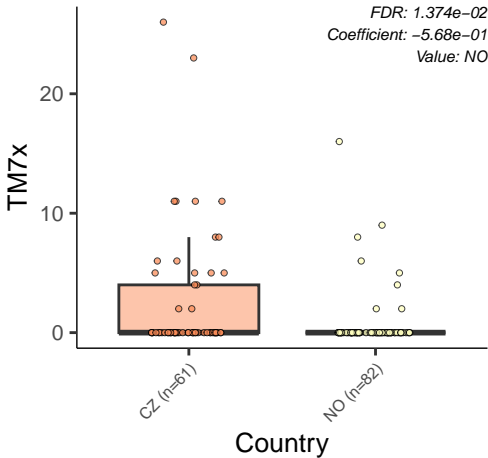
CZ (n=61)

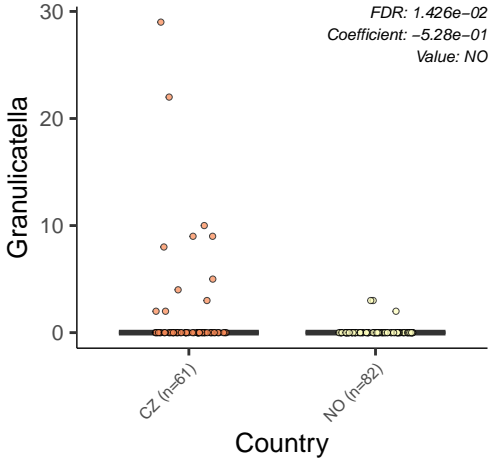
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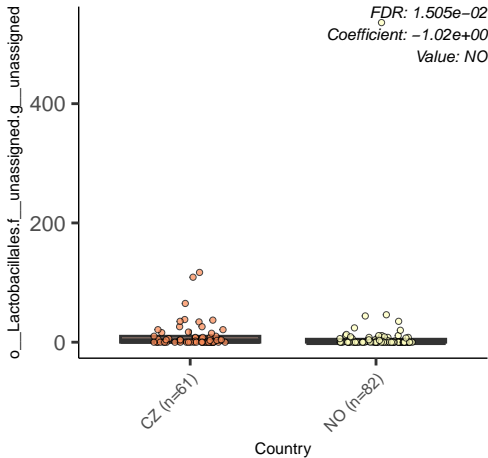
Country

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

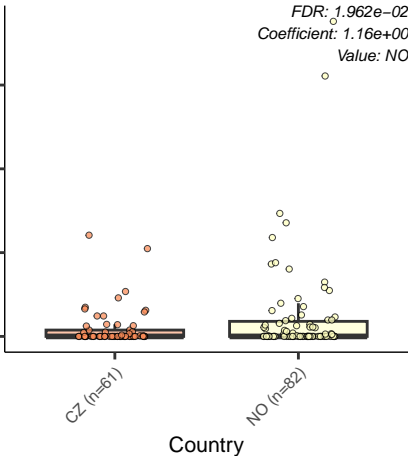
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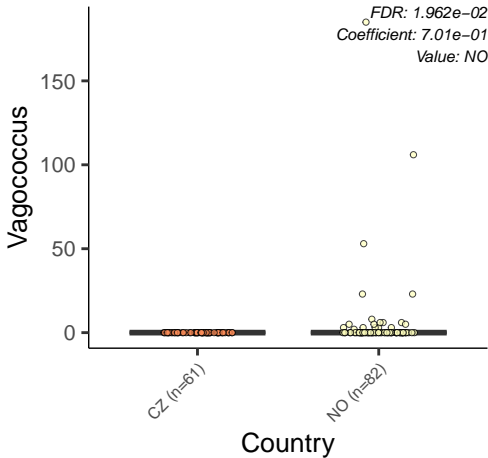
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CZ (n=61)

NO (n=82)

Country







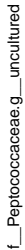
FDR: 1.974e-02

Coefficient: -8.11e-01

Value: NO



Country



FDR: 2.058e-02
Coefficient: 8.98e-01
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Country

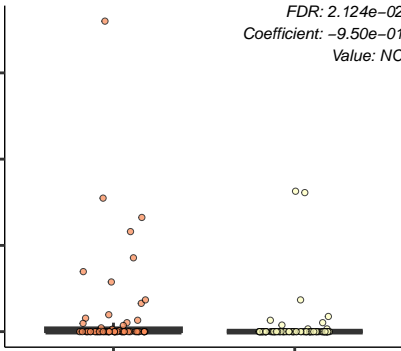
Acidaminococcus

FDR: $2.124e-02$
Coefficient: $-9.50e-01$
Value: NO

CZ (n=61)

NO (n=82)

Country



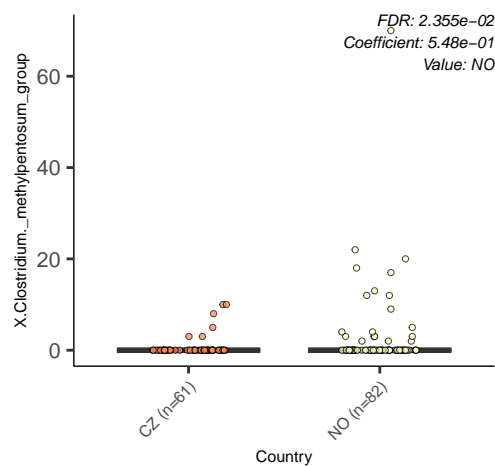
X.Clostridium._methylpentosum_group

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

CZ (n=61)

NO (n=82)

Country



Lachnospira

FDR: $2.455e-02$
Coefficient: $1.35e+00$
Value: NO

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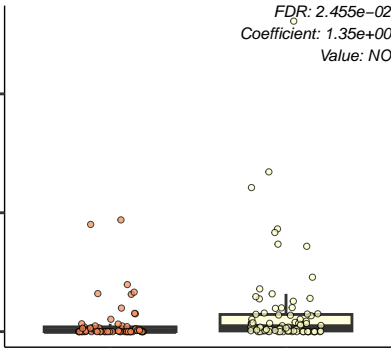
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CZ (n=61)

NO (n=82)

Country



Burkholderia.Caballeronia.Paraburkholderia

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

150

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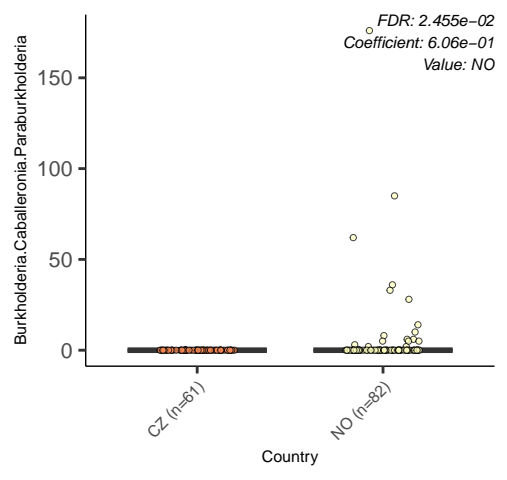
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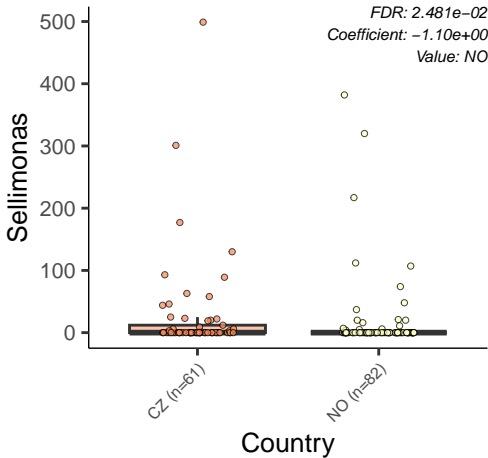
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CZ (n=61)

NO (n=82)

Country





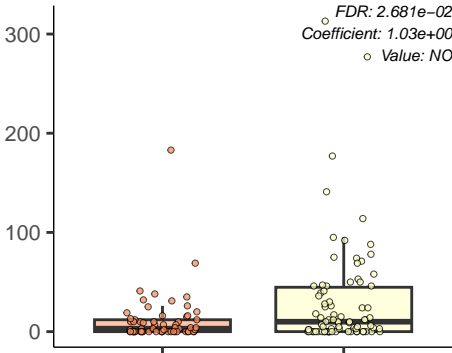
GCA.900066575

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

GZ (n=61)

NO (n=82)

Country



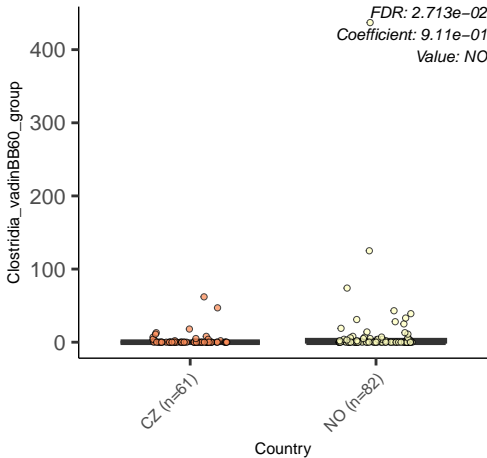
Clostridia_vadinBB60_group

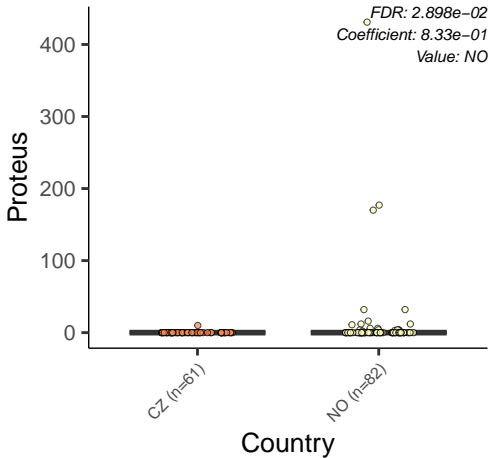
FDR: 2.713e-02
Coefficient: 9.11e-01
Value: NO

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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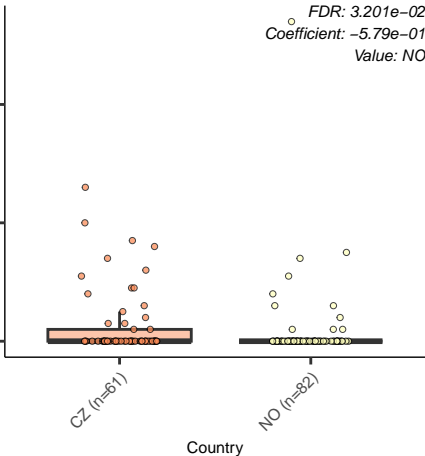
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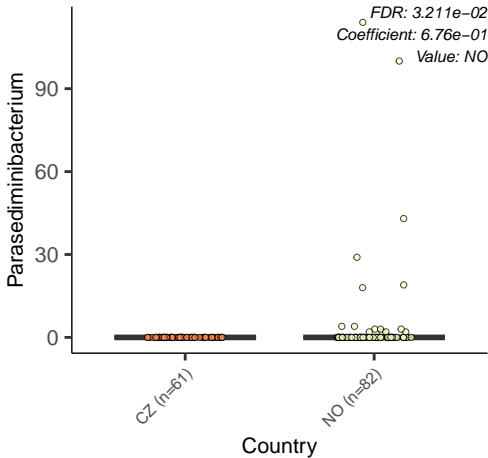
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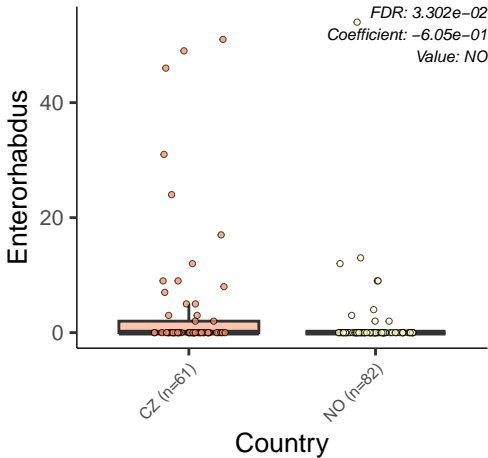
CZ (n=61)

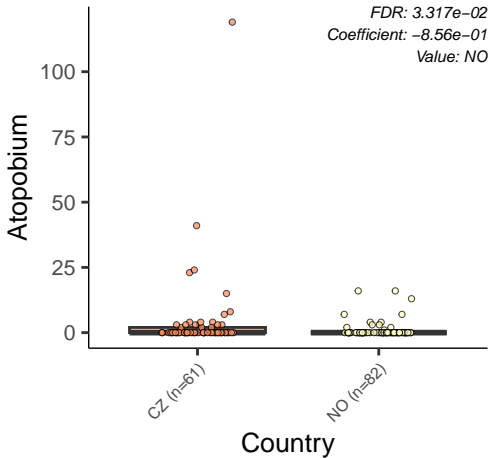
NO (n=82)

Country

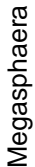








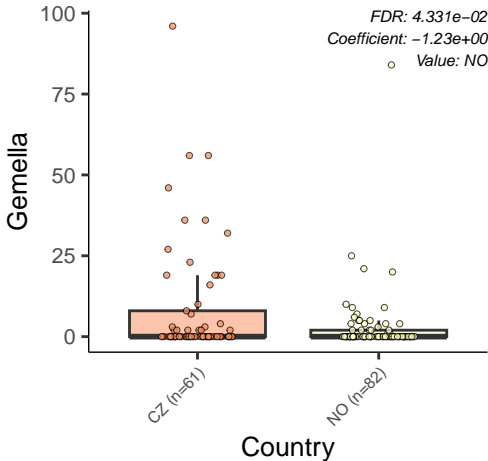
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CZ (n=61)

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

