

Collinsella

FDR: 1.238e-05

Coefficient: -2.96e+00

Value: NO

4000

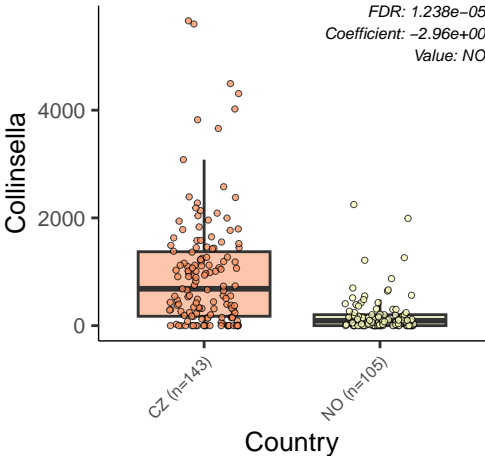
2000

0

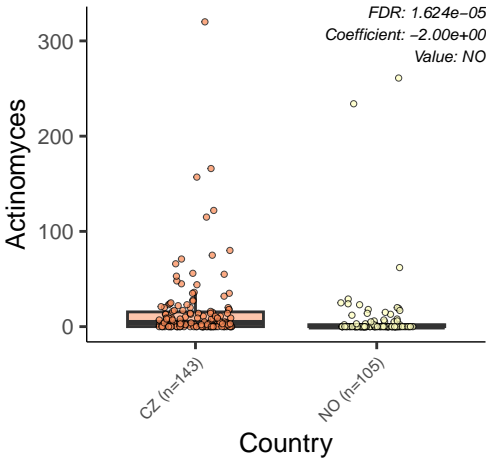
CZ (n=143)

NO (n=105)

Country



Actinomycetes



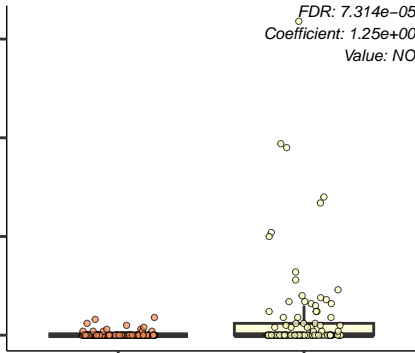
X.Eubacterium._xylanophilum_group

FDR: $7.314e-05$
Coefficient: $1.25e+00$
Value: NO

CZ (n=143)

NO (n=105)

Country



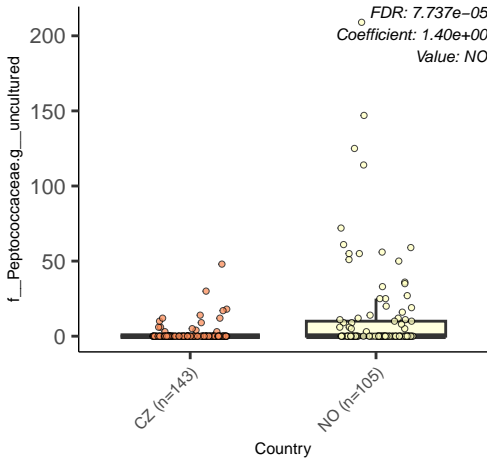
f__Peptococcaceae.g__uncultured

FDR: 7.737e-05
Coefficient: 1.40e+00
Value: NO

CZ (n=143)

NO (n=105)

Country



Gordonibacter

FDR: 1.532×10^{-4}

Coefficient: -1.54×10^0

Value: NO

200

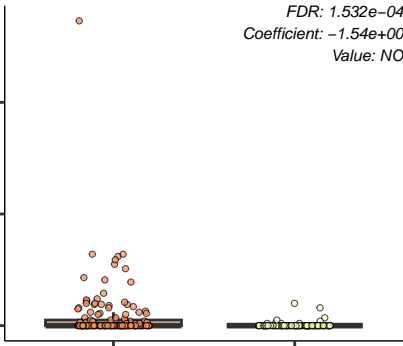
100

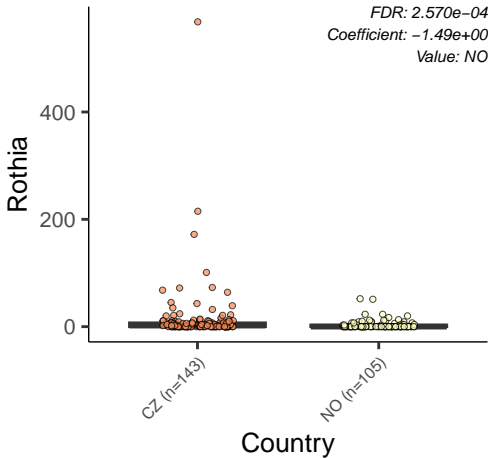
0

CZ (n=143)

NO (n=105)

Country





Bifidobacterium

FDR: 2.843e-04
Coefficient: -2.78e+00
Value: NO

4000

3000

2000

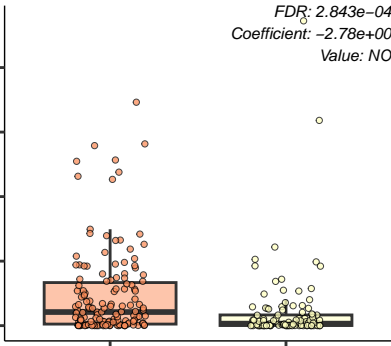
1000

0

CZ (n=143)

NO (n=105)

Country



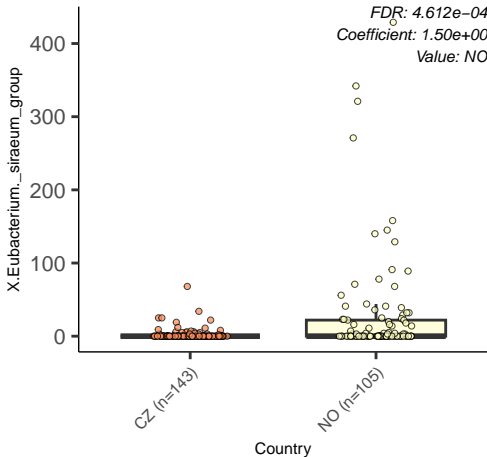
X.Eubacterium._siraenum_group

FDR: 4.612e-04
Coefficient: 1.50e+00
Value: NO

CZ (n=143)

NO (n=105)

Country



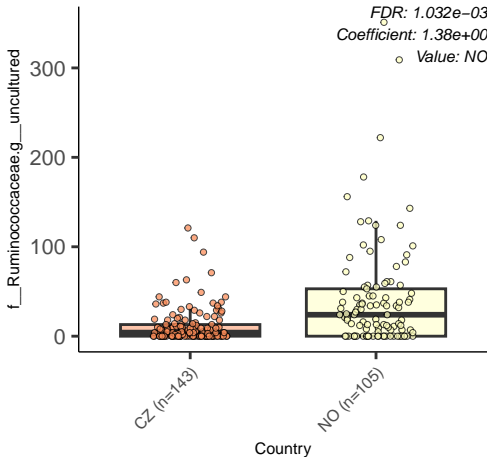
f_Ruminococcaceae.g__uncultured

FDR: 1.032e-03
Coefficient: 1.38e+00
Value: NO

CZ (n=143)

NO (n=105)

Country



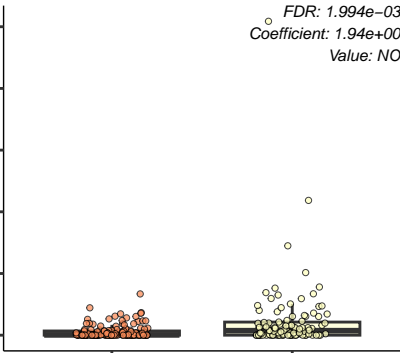
Christensenellaceae_R.7_group

FDR: 1.994e-03
Coefficient: 1.94e+00
Value: NO

CZ (n=143)

NO (n=105)

Country



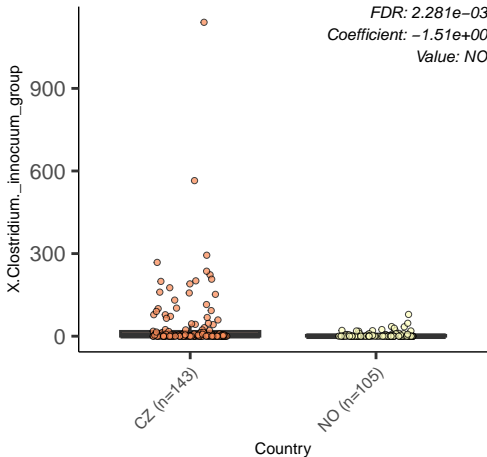
X.Clostridium._innocuum_group

FDR: 2.281e-03
Coefficient: -1.51e+00
Value: NO

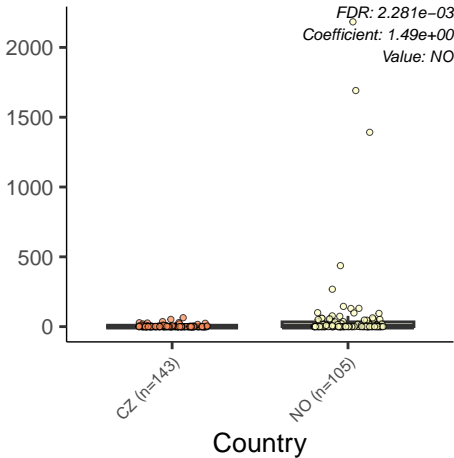
CZ (n=143)

NO (n=105)

Country



Eisenbergiella



Senegalimassilia

FDR: 3.713e-03
Coefficient: -1.44e+00
Value: NO

400

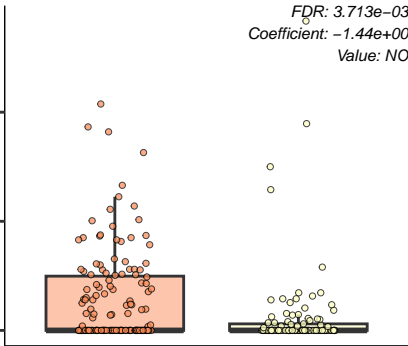
200

0

CZ (n=143)

NO (n=105)

Country



Streptococcus

10000

7500

5000

2500

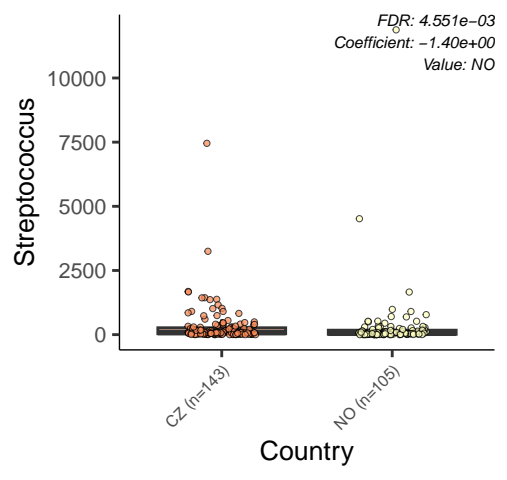
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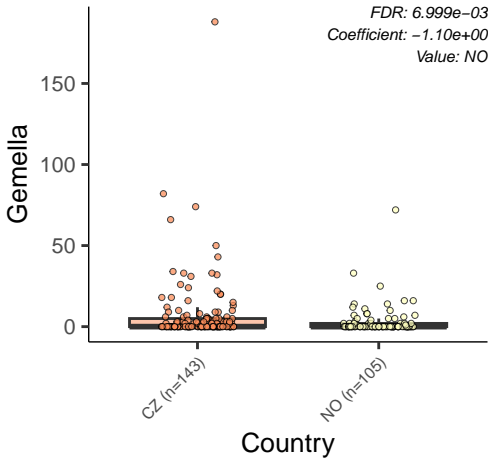
CZ (n=143)

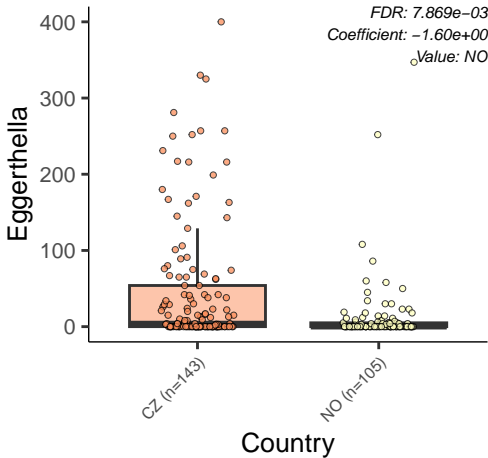
NO (n=105)

Country

FDR: 4.551e-03
Coefficient: -1.40e+00
Value: NO

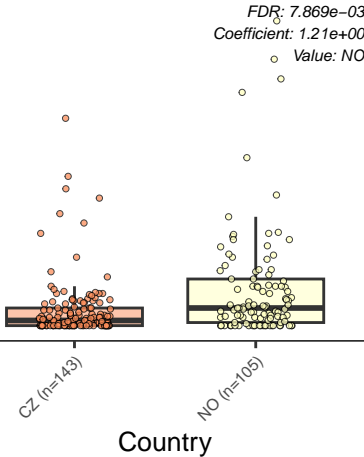






Marvinbryantia

FDR: $7.869e-03$
Coefficient: $1.21e+00$
Value: NO



Mogibacterium

FDR: 1.005e-02

Coefficient: -9.20e-01

Value: NO

200

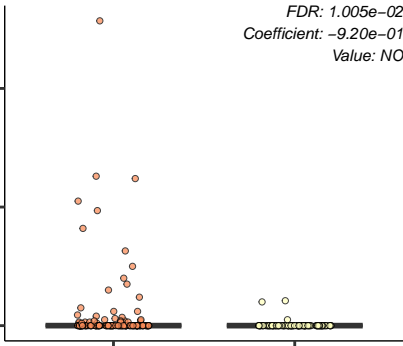
100

0

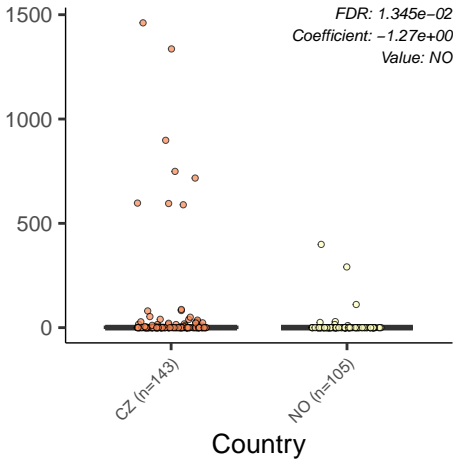
CZ (n=143)

NO (n=105)

Country



Megasphaera



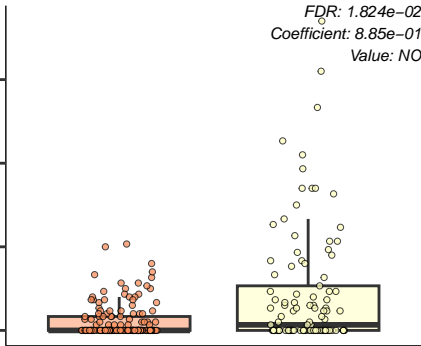
Family_XIII_UCG.001

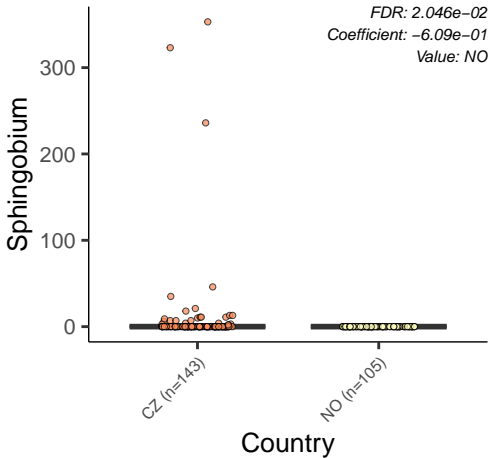
FDR: 1.824e-02
Coefficient: 8.85e-01
Value: NO

CZ (n=143)

NO (n=105)

Country





Faecalitalea

FDR: 2.249e-02

Coefficient: -1.01e+00

Value: NO

400

300

200

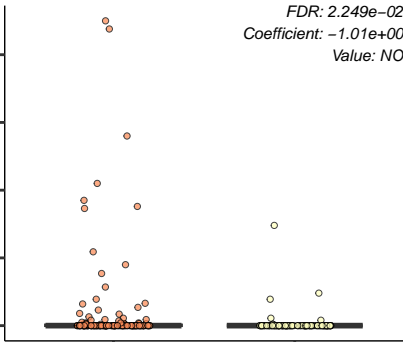
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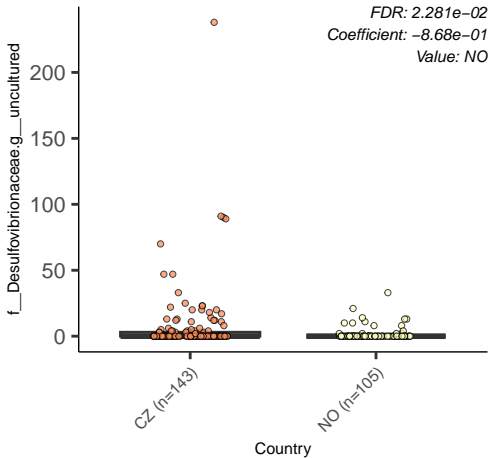
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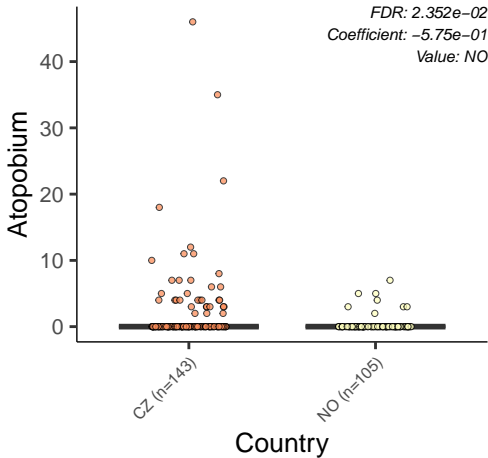
CZ (n=143)

NO (n=105)

Country







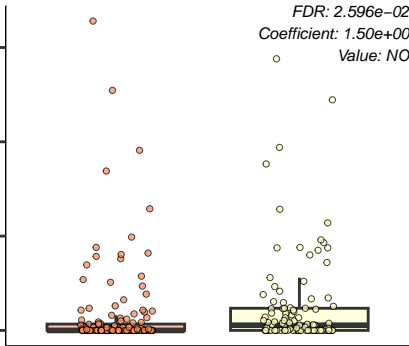
Erysipelatoclostridium

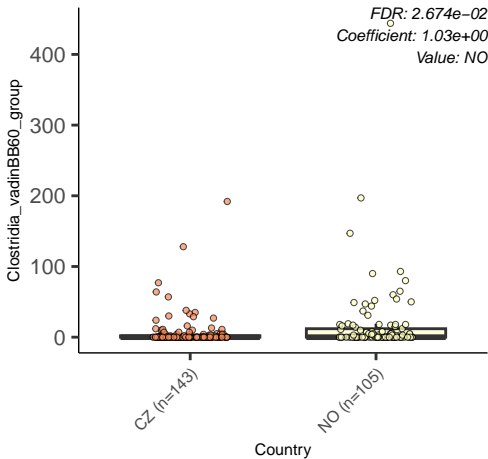
FDR: 2.596e-02
Coefficient: 1.50e+00
Value: NO

CZ (n=143)

NO (n=105)

Country





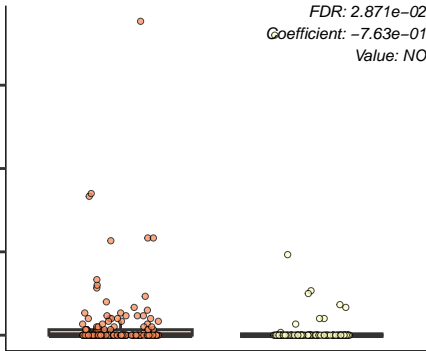
X.Eubacterium._brachy_group

FDR: 2.871e-02
Coefficient: -7.63e-01
Value: NO

CZ (n=143)

NO (n=105)

Country



X.Eubacterium._hallii_group

1500

1000

500

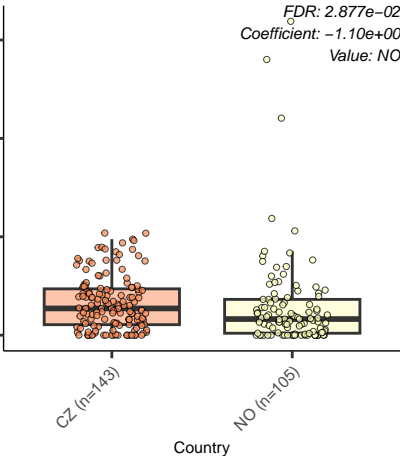
0

CZ (n=143)

NO (n=105)

Country

FDR: 2.877e-02
Coefficient: -1.10e+00
Value: NO



Peptostreptococcus

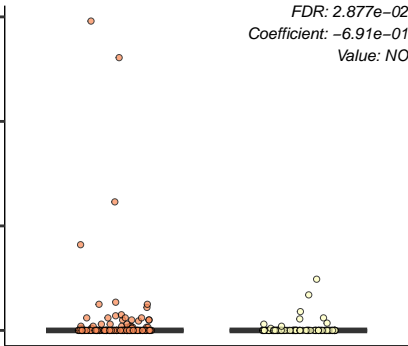
300
200
100
0

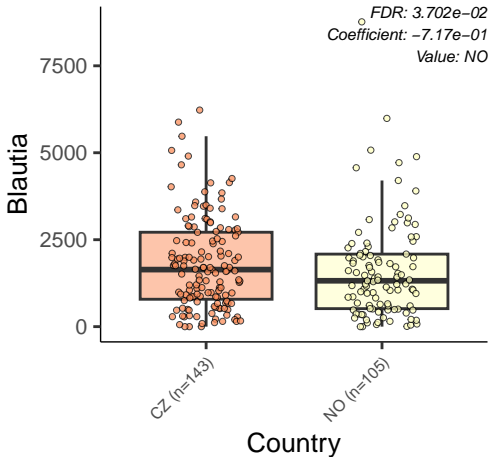
FDR: 2.877e-02
Coefficient: -6.91e-01
Value: NO

CZ (n=143)

NO (n=105)

Country





Methylobacterium.Methylorubrum

FDR: 3.769e-02
Coefficient: 3.35e-01
Value: NO

75

50

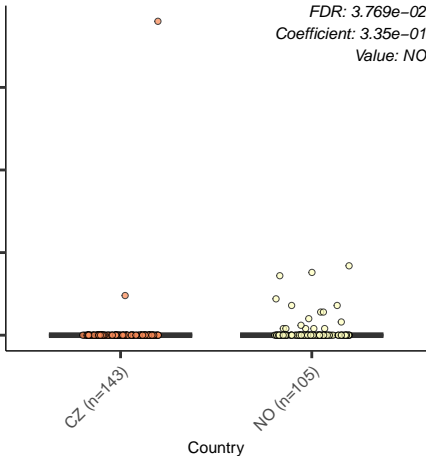
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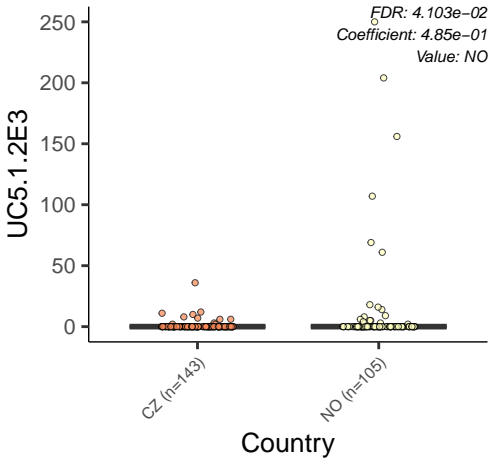
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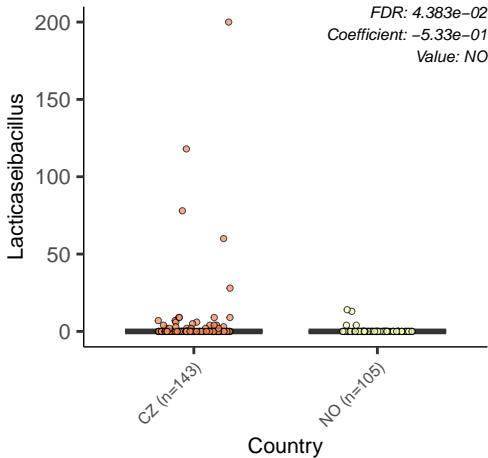
CZ (n=143)

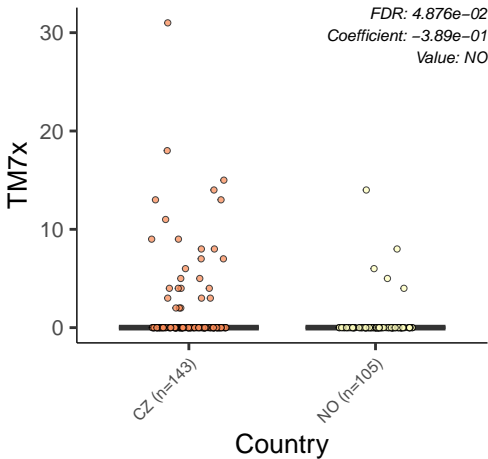
NO (n=105)

Country









Anaerotruncus

