

Bifidobacterium

4000

2000

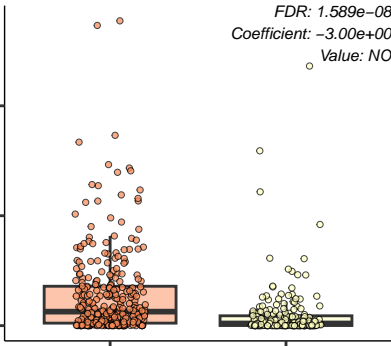
0

CZ (n=304)

NO (n=207)

Country

FDR: 1.589e-08
Coefficient: -3.00e+00
Value: NO



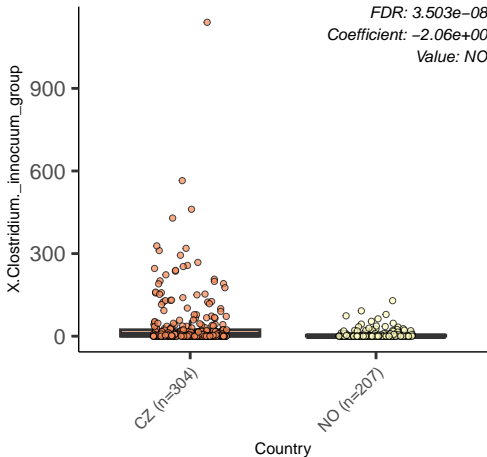
X.Clostridium._innocuum_group

FDR: 3.503e-08
Coefficient: -2.06e+00
Value: NO

CZ (n=304)

NO (n=207)

Country



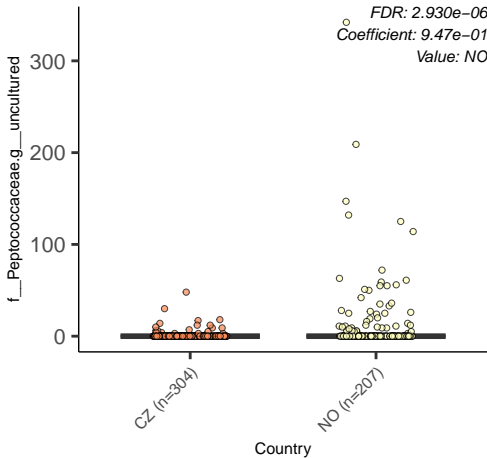
f__Peptococcaceae.g__uncultured

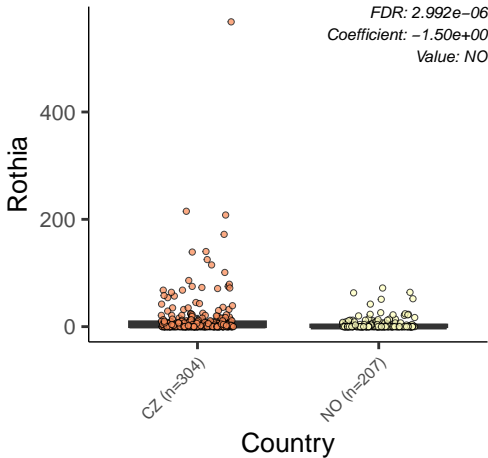
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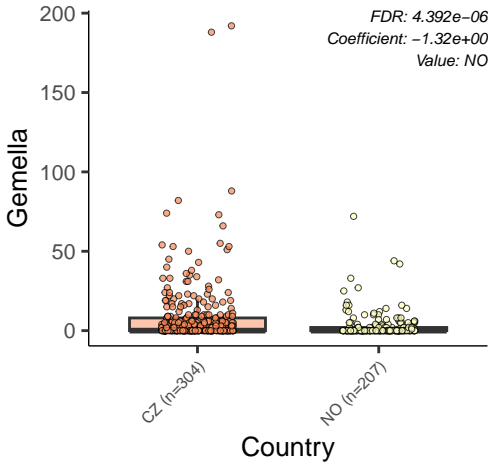
CZ (n=304)

NO (n=207)

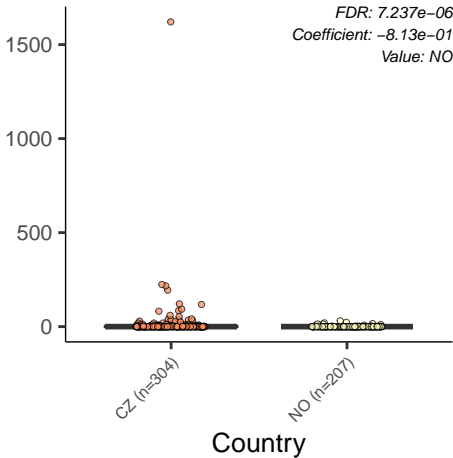
Country







Corynebacterium



Streptococcus

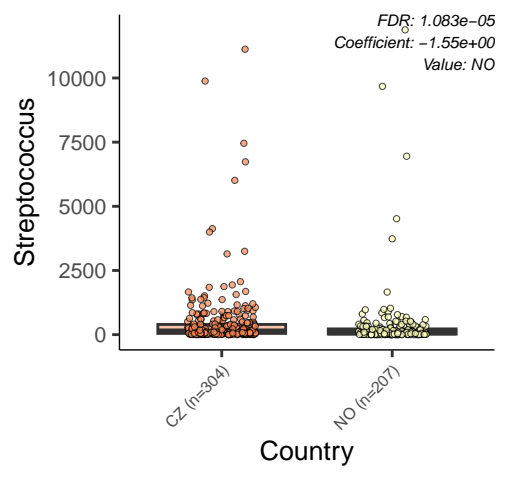
10000
7500
5000
2500
0

CZ (n=304)

NO (n=207)

Country

FDR: 1.083e-05
Coefficient: -1.55e+00
Value: NO



Gordonibacter

FDR: 1.131e-05

Coefficient: -1.12e+00

Value: NO

200

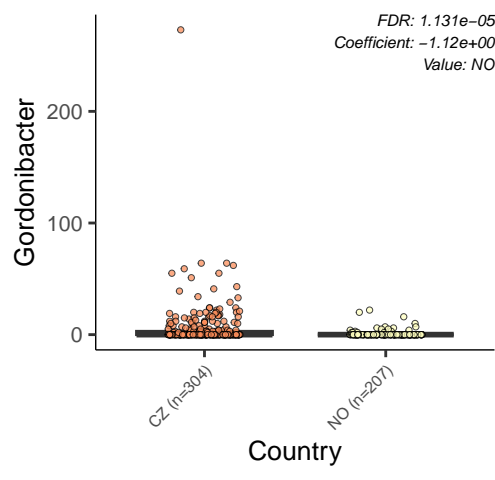
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CZ (n=304)

NO (n=207)

Country



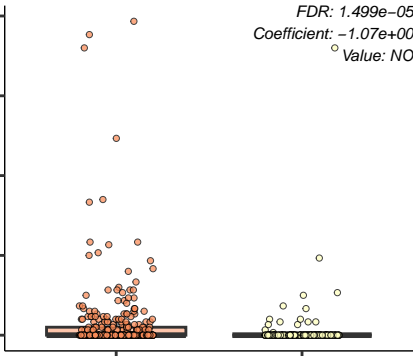
X.Eubacterium._brachy_group

FDR: 1.499e-05
Coefficient: -1.07e+00
Value: NO

CZ (n=304)

NO (n=207)

Country



Senegalimassilia

FDR: 3.654e-05
Coefficient: -2.02e+00
Value: NO

400

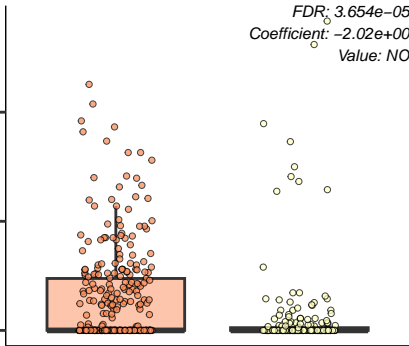
200

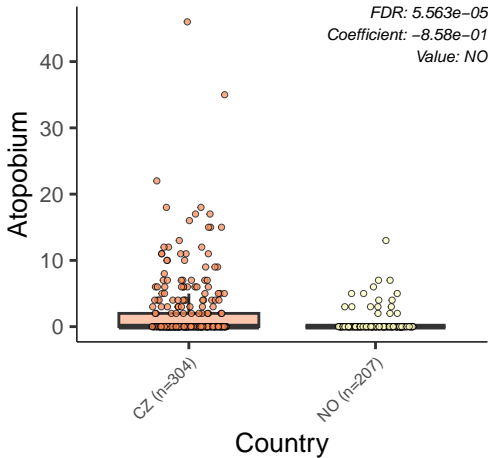
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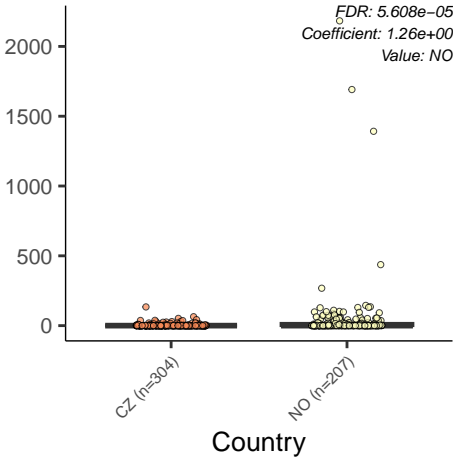
NO (n=207)

Country





Eisenbergiella



Mogibacterium

FDR: 1.200e-04

Coefficient: -1.00e+00

Value: NO

300

200

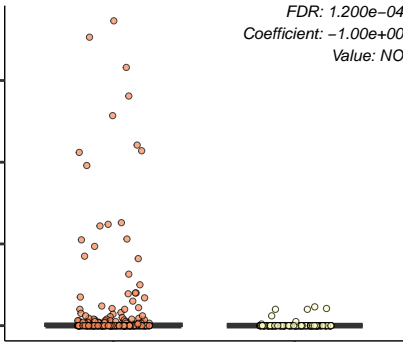
100

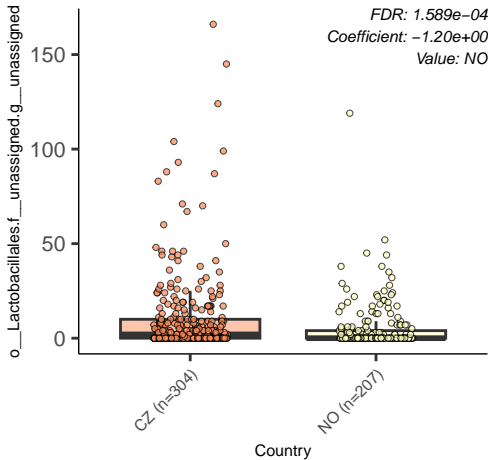
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CZ (n=304)

NO (n=207)

Country





Family_XIII_UCG.001

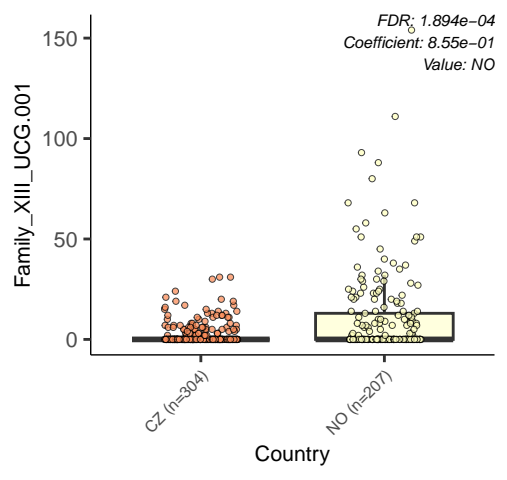
FDR: 1.894e-04
Coefficient: 8.55e-01
Value: NO

CZ (n=304)

NO (n=207)

Country

150
100
50
0



Erysipelatoclostridium

600

400

200

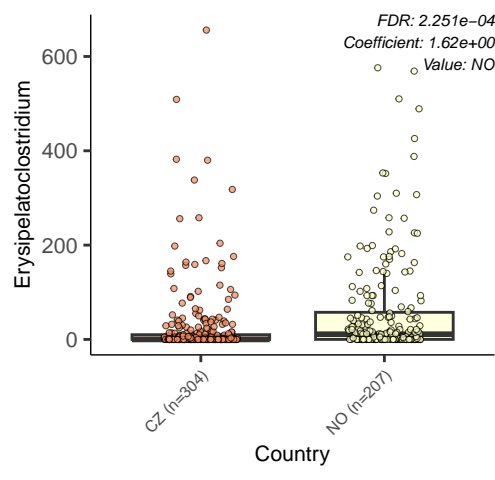
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CZ (n=304)

NO (n=207)

Country

FDR: 2.251e-04
Coefficient: 1.62e+00
Value: NO



Marvinbryantia

FDR: $2.623e-04$
Coefficient: $1.20e+00$
Value: NO

400

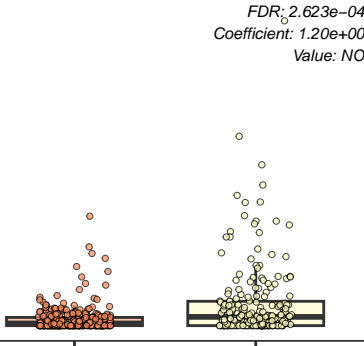
200

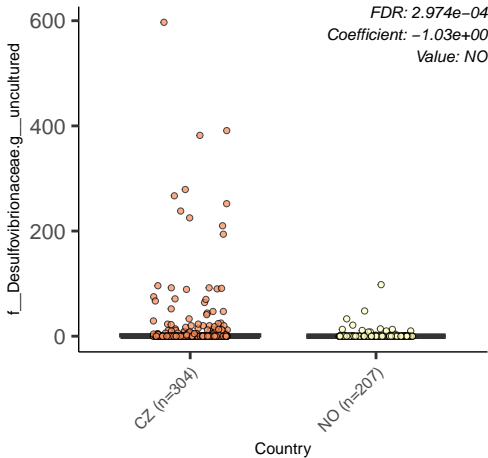
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CZ (n=304)

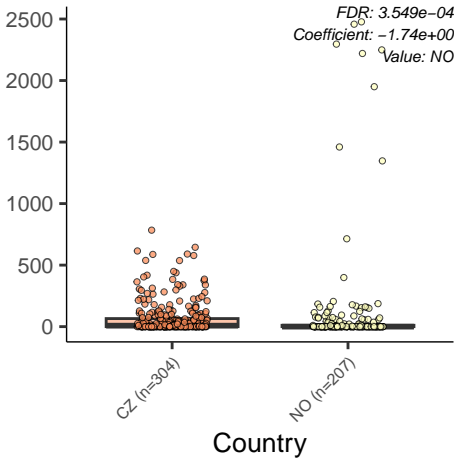
NO (n=207)

Country

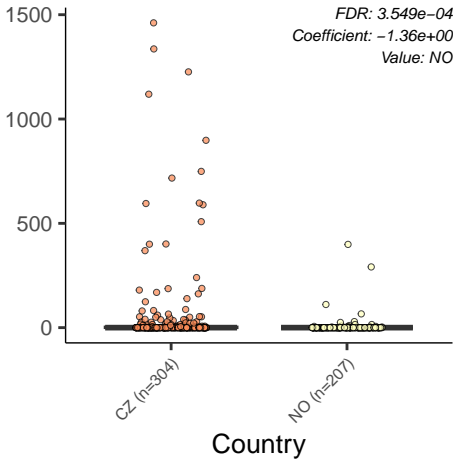




Coprobacter



Megasphaera



Acidaminococcus

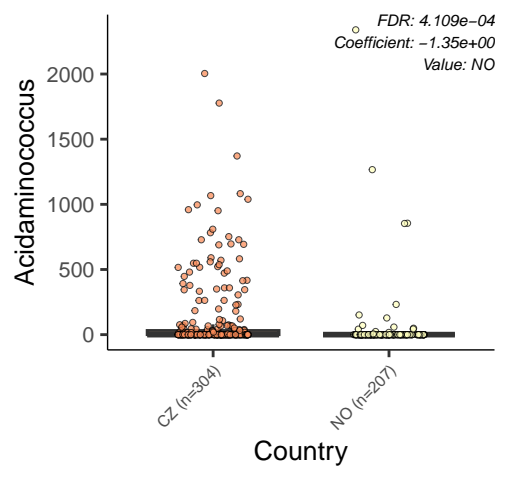
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Coefficient: -1.35e+00
Value: NO

CZ (n=304)

NO (n=207)

Country

2000
1500
1000
500
0



Holdemanella

FDR: 4.235e-04

Coefficient: -1.59e+00

Value: NO

2000

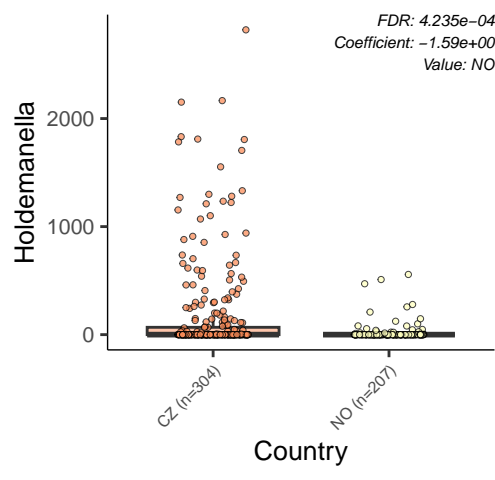
1000

0

CZ (n=304)

NO (n=207)

Country



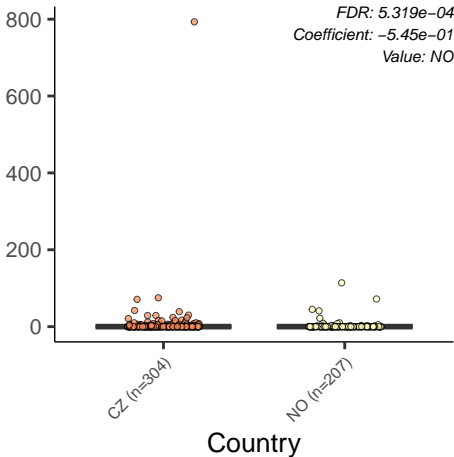
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NO (n=207)

Country

Staphylococcus



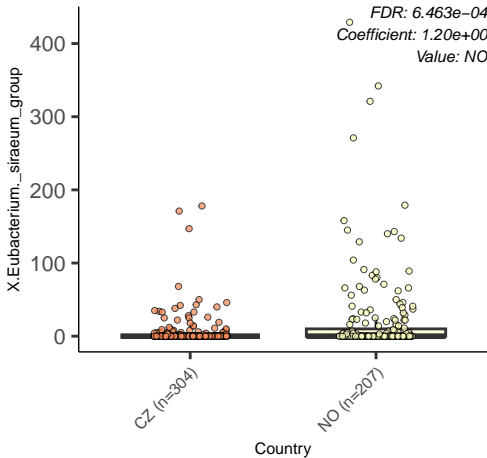
X.Eubacterium._siraеum_group

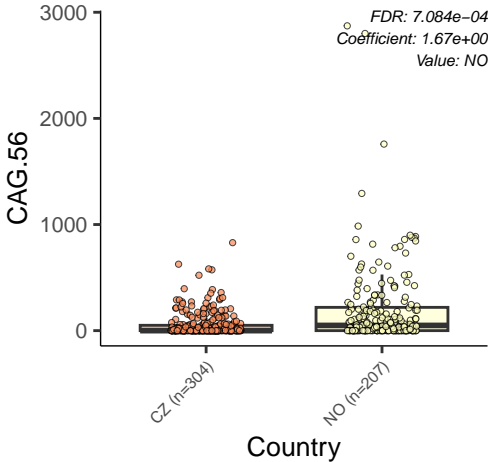
FDR: 6.463e-04
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Value: NO

CZ (n=304)

NO (n=207)

Country





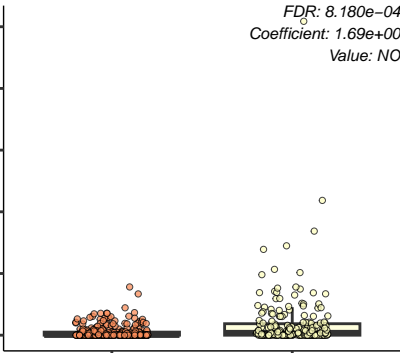
Christensenellaceae_R.7_group

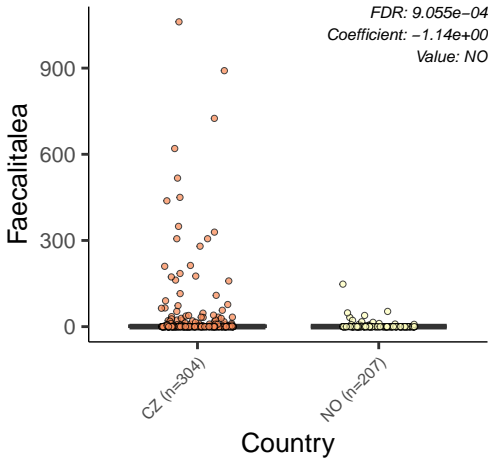
FDR: 8.180e-04
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CZ (n=304)

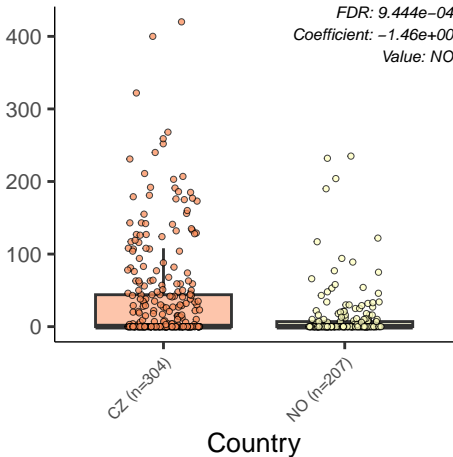
NO (n=207)

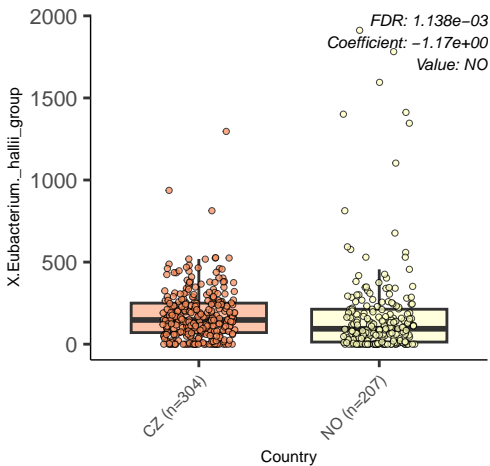
Country

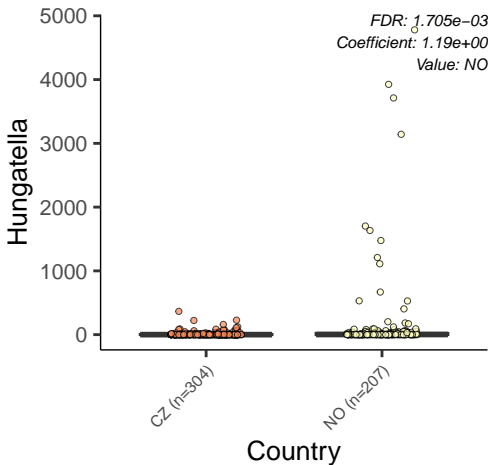


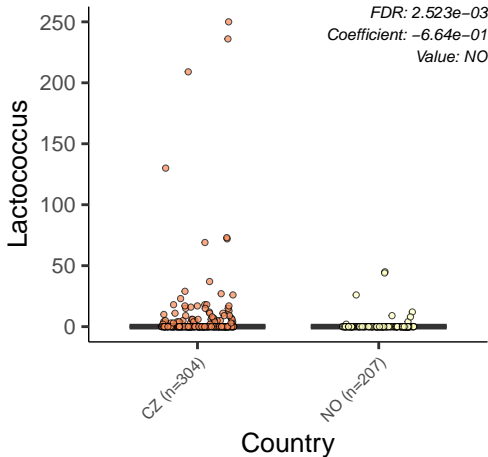


Slackia









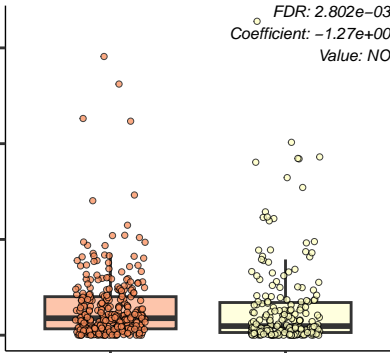
X.Ruminococcus._torques_group

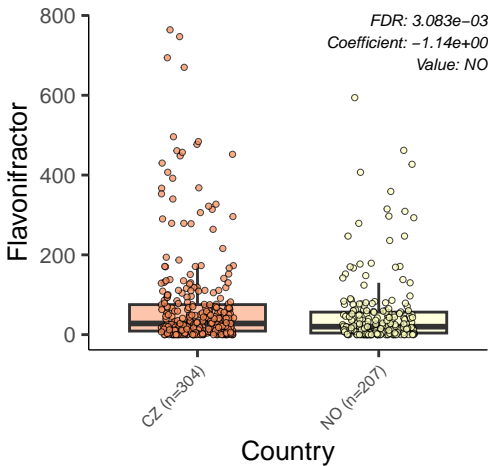
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Coefficient: -1.27e+00
Value: NO

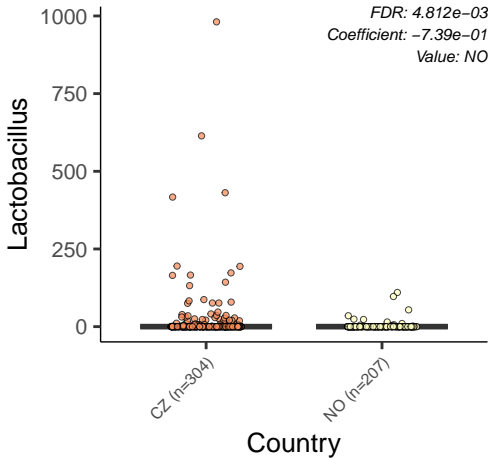
CZ (n=304)

NO (n=207)

Country







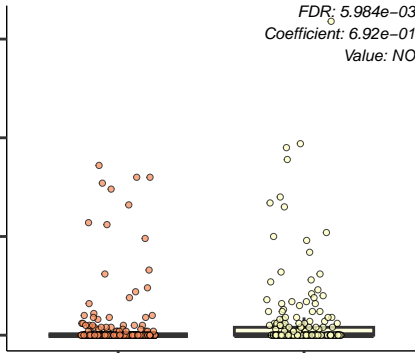
X.Eubacterium._xylanophilum_group

FDR: 5.984e-03
Coefficient: 6.92e-01
Value: NO

CZ (n=304)

NO (n=207)

Country



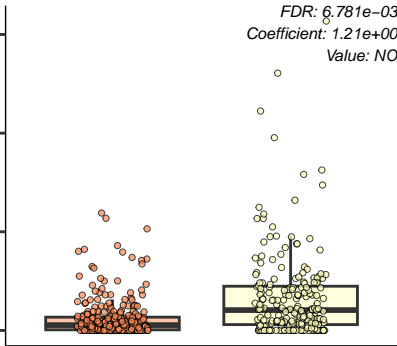
Lachnospiraceae_UCG.004

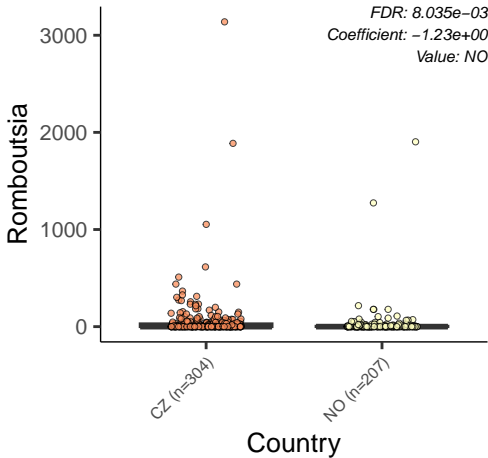
FDR: $6.781e-03$
Coefficient: $1.21e+00$
Value: NO

CZ (n=304)

NO (n=207)

Country





Eggerthella

600

400

200

0

CZ (n=304)

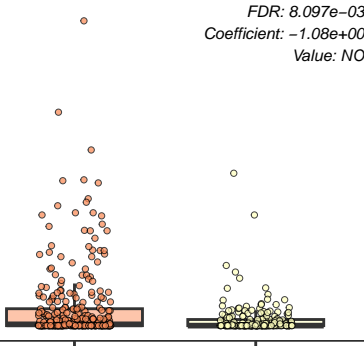
NO (n=207)

Country

FDR: 8.097e-03

Coefficient: -1.08e+00

Value: NO



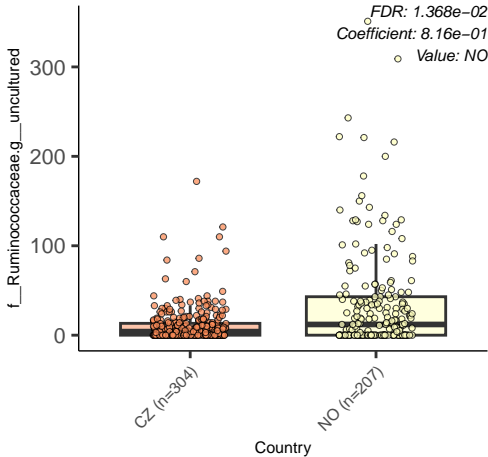
f_Ruminococcaceae.g__uncultured

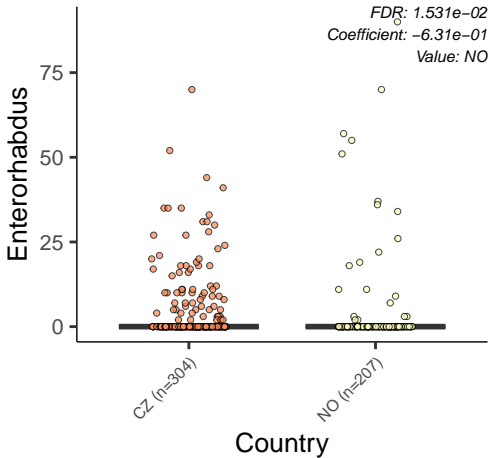
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Value: NO

CZ (n=304)

NO (n=207)

Country





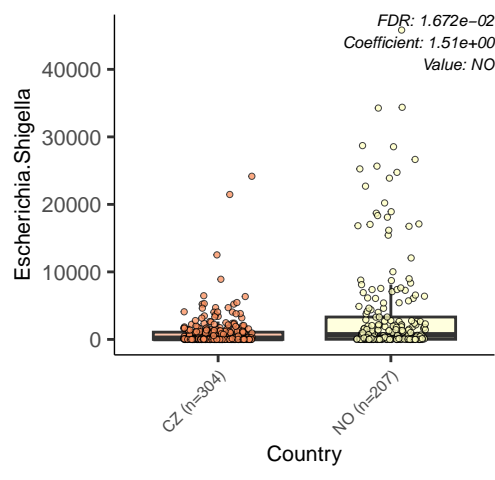
Escherichia.Shigella

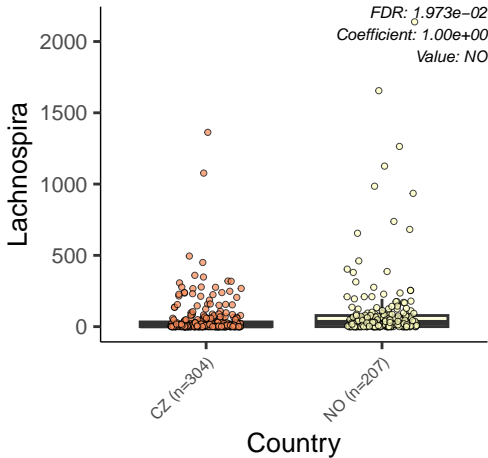
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CZ (n=304)

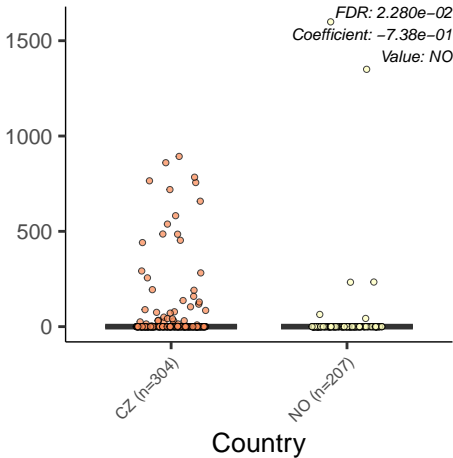
NO (n=207)

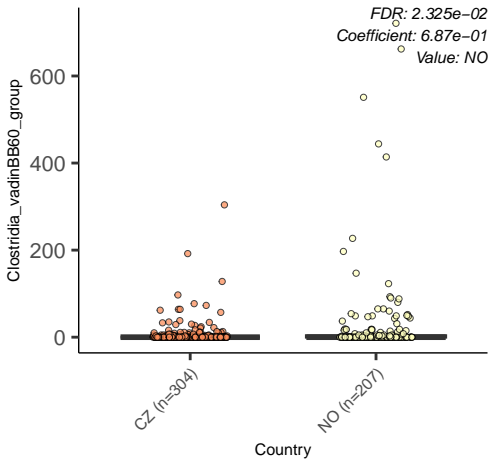
Country



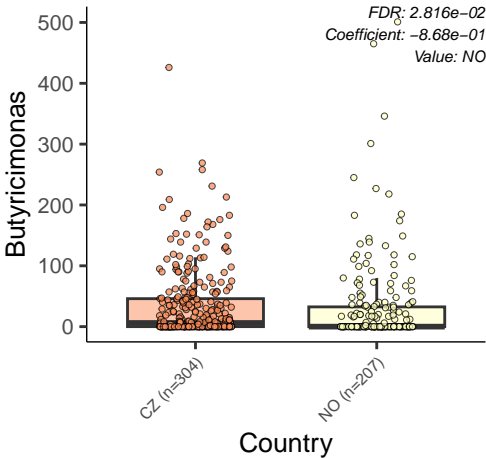


Catenibacterium





Butyricimonas



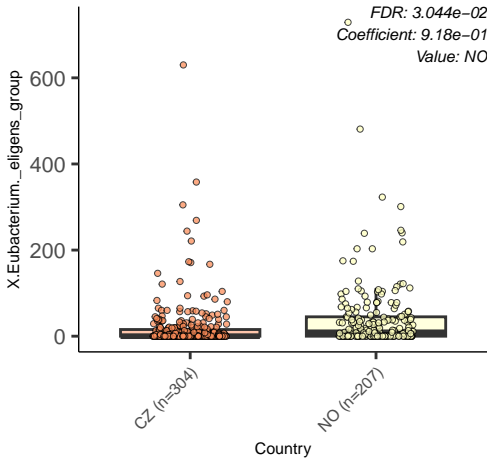
X.Eubacterium._elgens_group

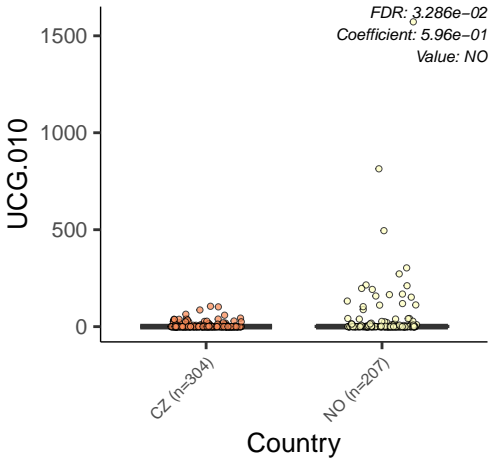
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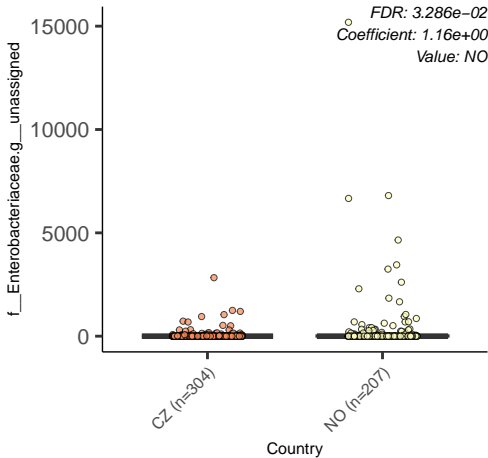
CZ (n=304)

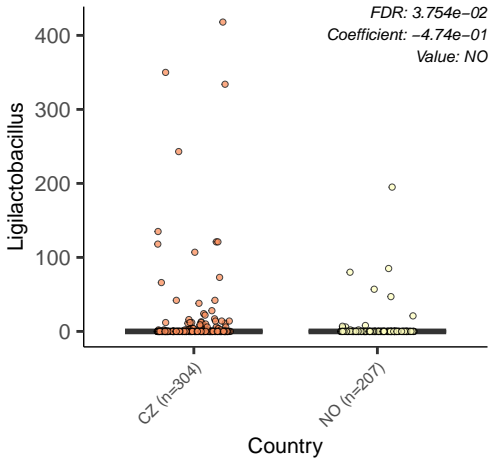
NO (n=207)

Country









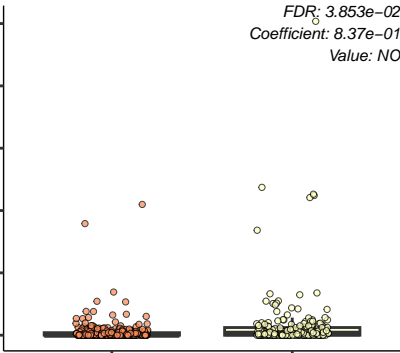
Lachnospiraceae_NK4A136_group

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Coefficient: 8.37e-01
Value: NO

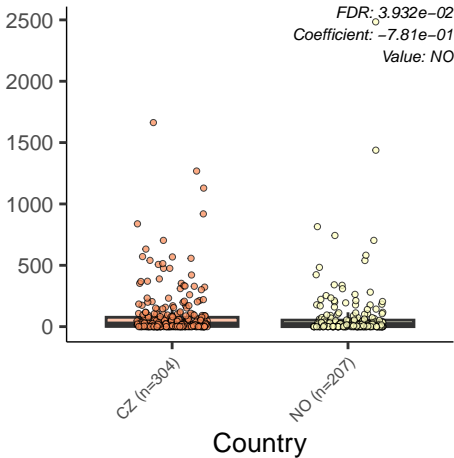
CZ (n=304)

NO (n=207)

Country



Haemophilus



Faecalibacterium

FDR: 3.982e-02
Coefficient: 9.36e-01
Value: NO

20000

15000

10000

5000

0

CZ (n=304)

NO (n=207)

Country

