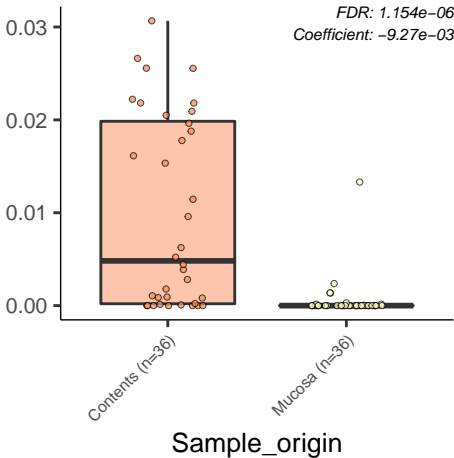


Lysinibacillus



f_Bacillaceae

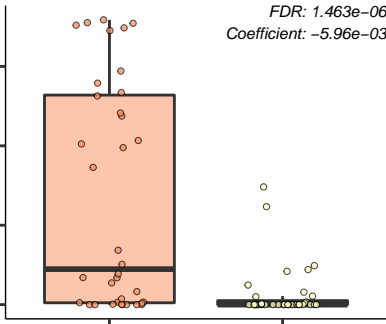
FDR: $1.463e-06$
Coefficient: $-5.96e-03$

0.015
0.010
0.005
0.000

Contents (n=36)

Mucosa (n=36)

Sample_origin



Brevinema andersonii

0.6
0.4
0.2
0.0

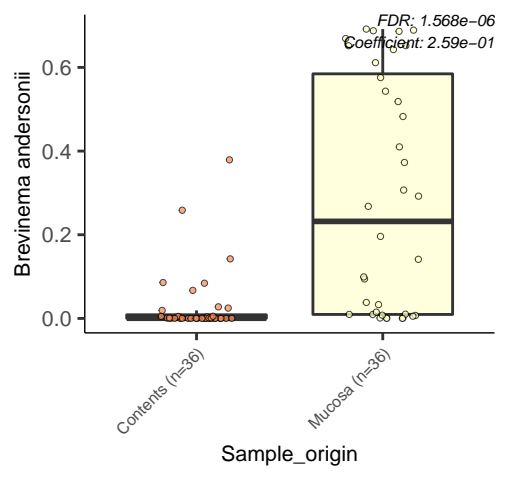
Contents (n=36)

Mucosa (n=36)

Sample_origin

FDR: $1.568e-06$

Coefficient: $2.59e-01$



f_Lachnospiraceae

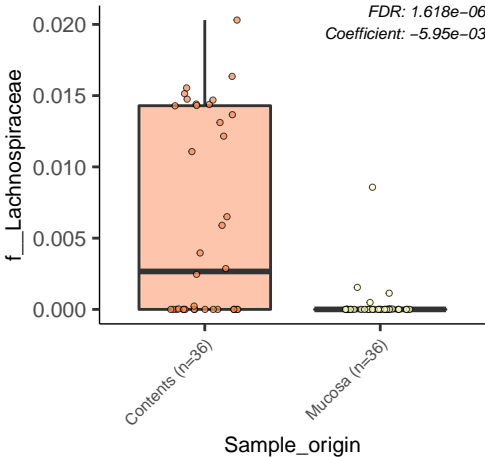
0.020
0.015
0.010
0.005
0.000

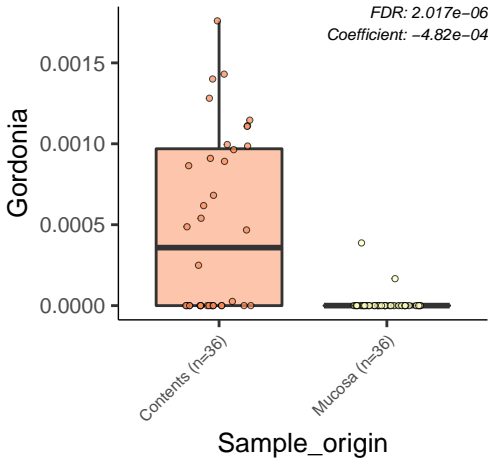
FDR: 1.618e-06
Coefficient: -5.95e-03

Contents (n=36)

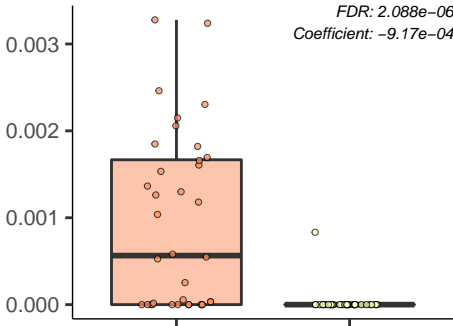
Mucosa (n=36)

Sample_origin





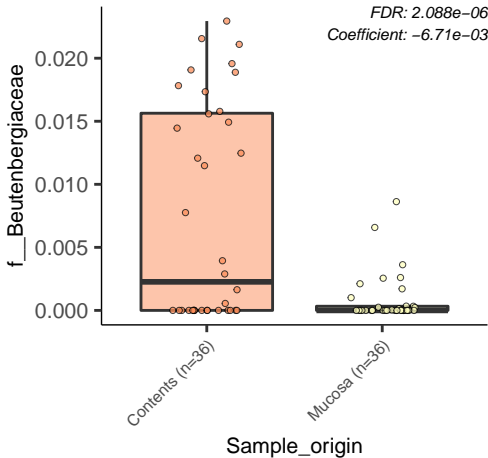
Coefficient: $-9.17e-04$

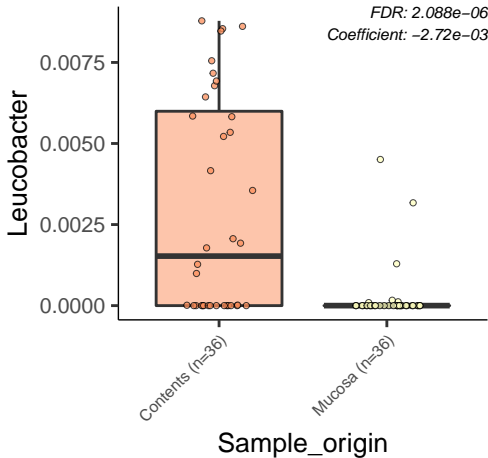


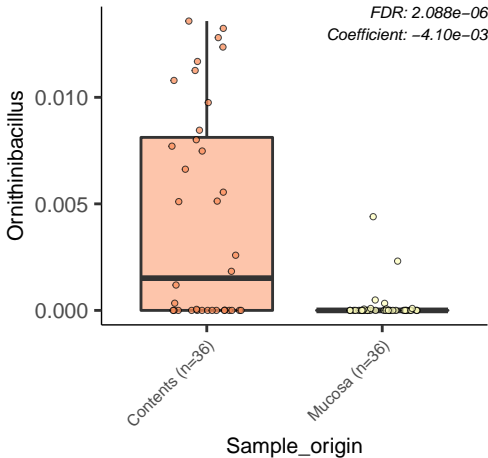
Contents (n=36)

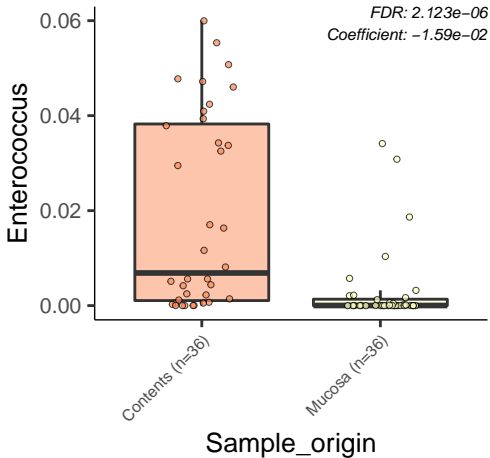
Mucosa (n=36)

Sample_origin









Oceanobacillus

FDR: $7.918e-06$

Coefficient: $-1.99e-02$

0.06

0.04

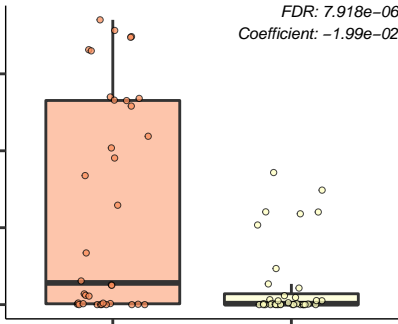
0.02

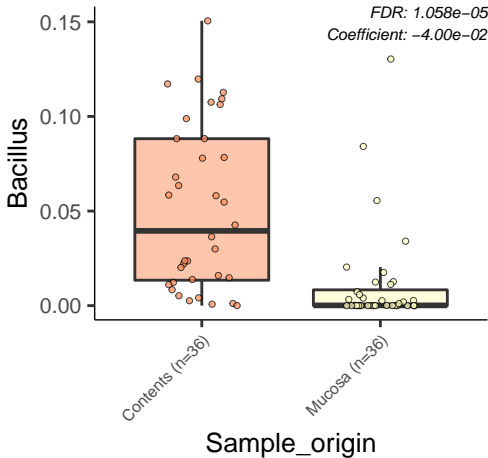
0.00

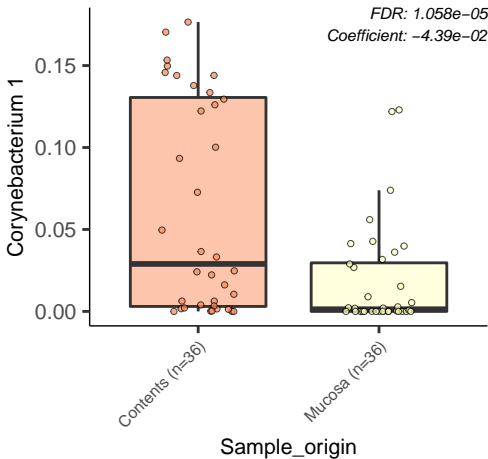
Contents (n=36)

Mucosa (n=36)

Sample_origin







Gracilibacillus

FDR: 1.058e-05
Coefficient: -7.81e-04

0.002

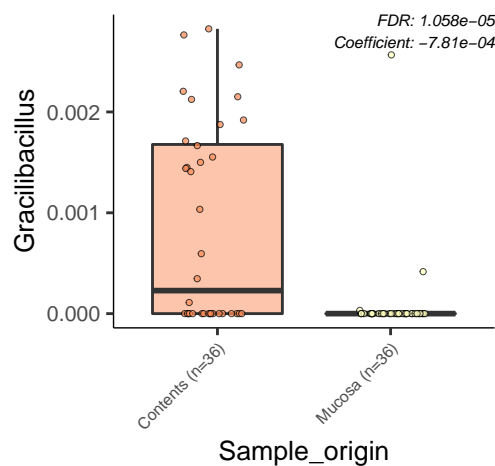
0.001

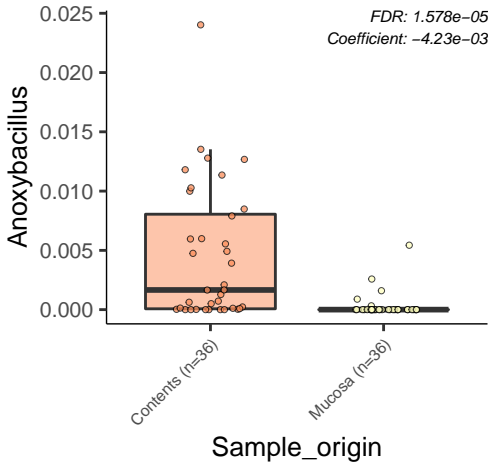
0.000

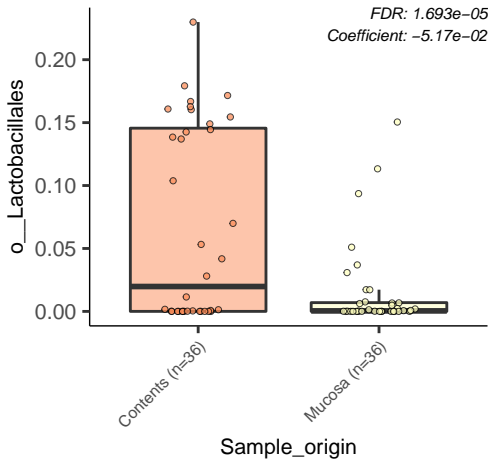
Contents (n=36)

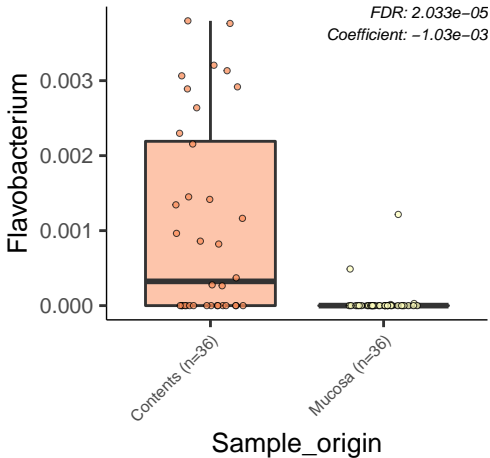
Mucosa (n=36)

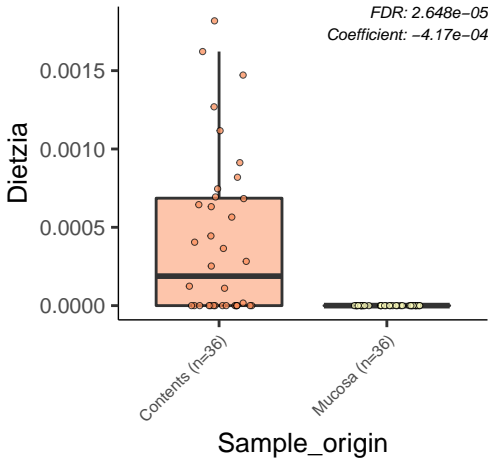
Sample_origin

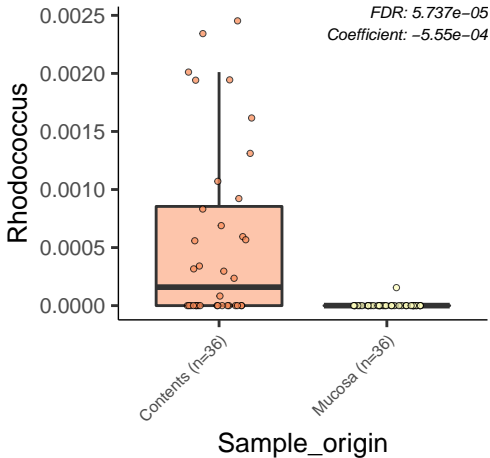












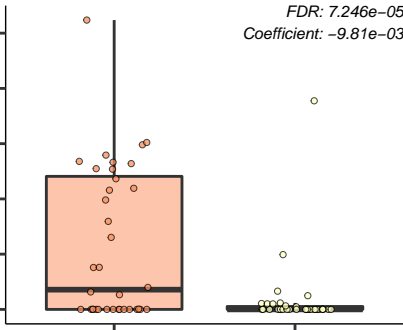
f_Enterococcaceae

FDR: $7.246e-05$
Coefficient: $-9.81e-03$

Contents (n=36)

Mucosa (n=36)

Sample_origin



Tepidimicrobium

FDR: 3.637e-04

Coefficient: -7.09e-03

0.03

0.02

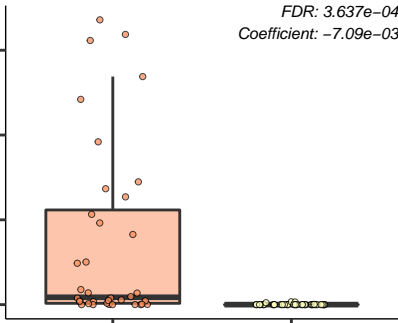
0.01

0.00

Contents (n=36)

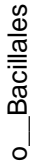
Mucosa (n=36)

Sample_origin



FDR: 4.081e-04

Coefficient: $-1.03e-03$



0.006

0.004

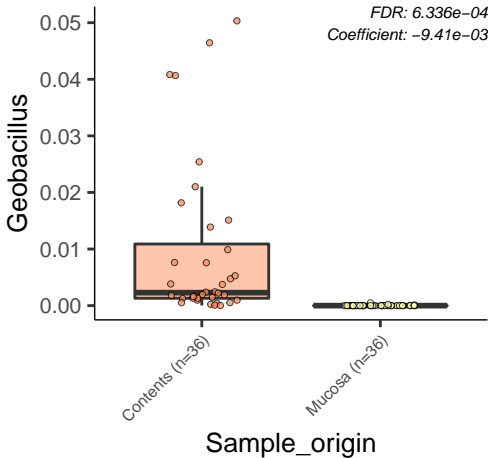
0.002

0.000

Contents (n=36)

Mucosa (n=36)

Sample_origin



f_Spirochaetaceae

FDR: $7.656e-04$
Coefficient: $1.50e-01$

0.6

0.4

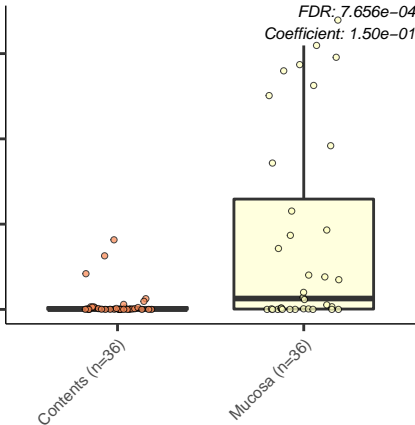
0.2

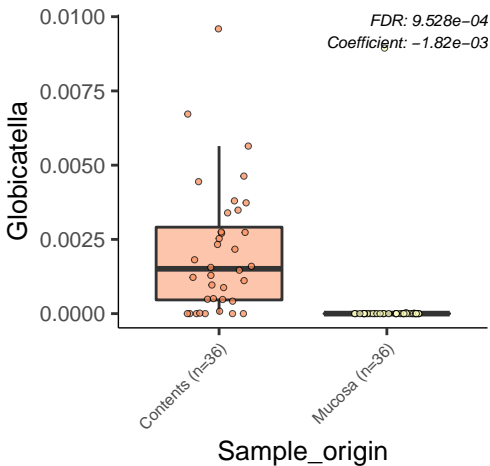
0.0

Contents (n=36)

Mucosa (n=36)

Sample_origin





Coefficient: $-2.79e-03$

Sample_origin

Peptostreptococcus

FDR: 2.202e-03

Coefficient: -4.36e-03

0.02

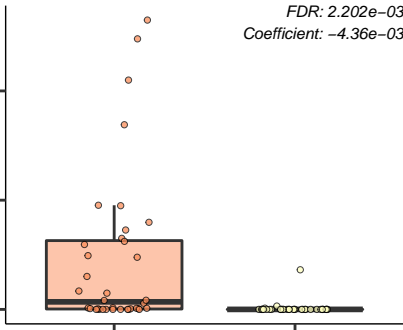
0.01

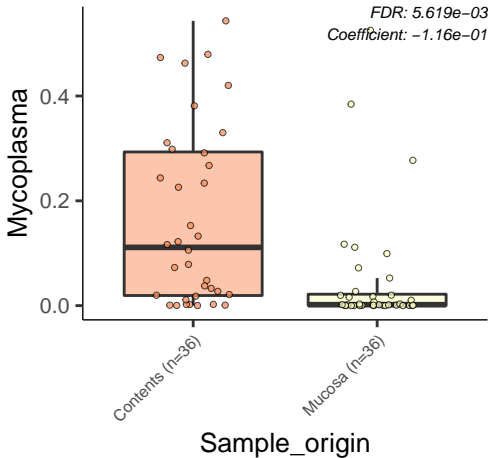
0.00

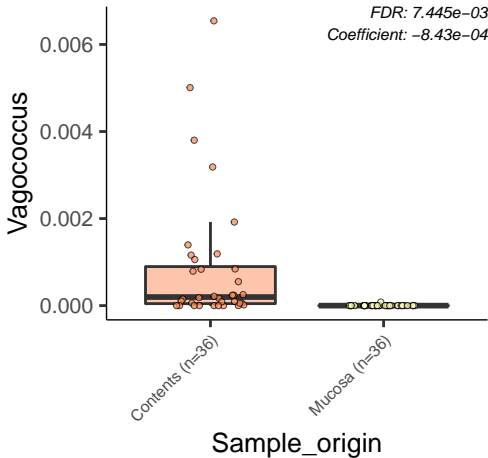
Contents (n=36)

Mucosa (n=36)

Sample_origin







Spingobacterium

FDR: 1.053e-02

Coefficient: -1.89e-03

0.015

0.010

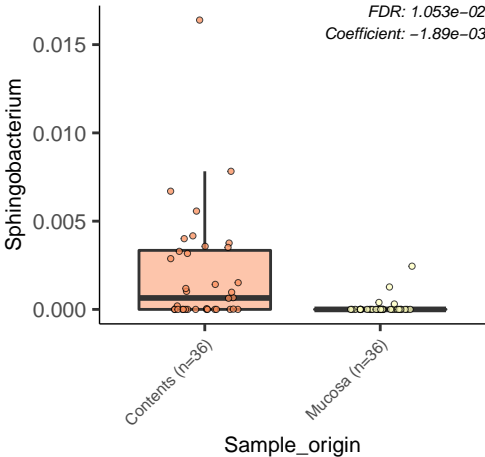
0.005

0.000

Contents (n=36)

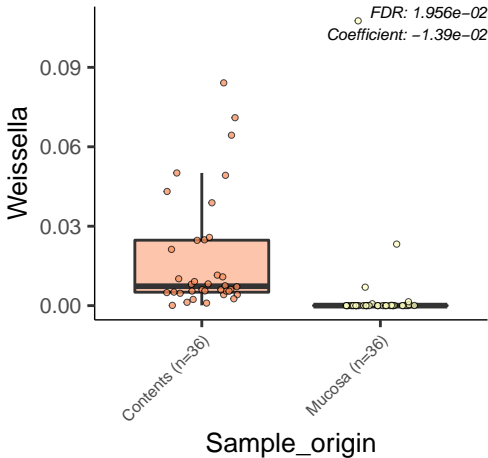
Mucosa (n=36)

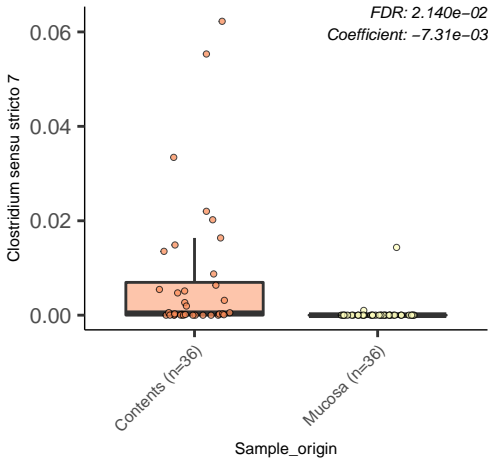
Sample_origin



Coefficient: $-1.59e-03$

Sample_origin





Staphylococcus

FDR: 2.207e-02
Coefficient: -3.76e-03

0.02

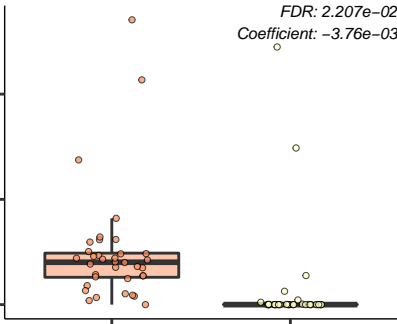
0.01

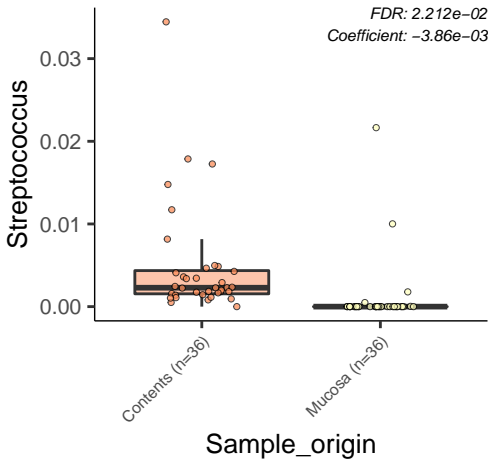
0.00

Contents (n=36)

Mucosa (n=36)

Sample_origin





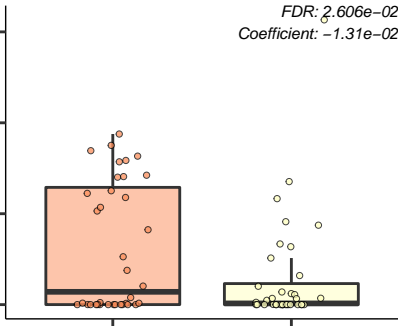
Actinomycetes

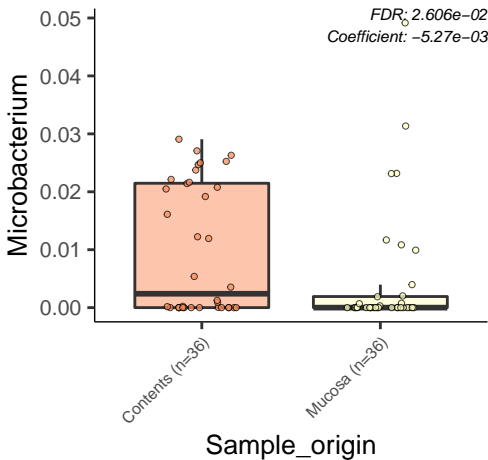
FDR: 2.606e-02
Coefficient: -1.31e-02

Contents (n=36)

Mucosa (n=36)

Sample_origin





Photobacterium

FDR: 2.606e-02

Coefficient: -3.48e-02

0.4

0.3

0.2

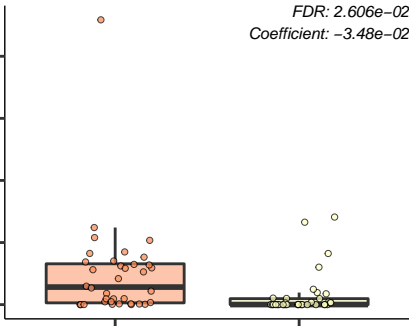
0.1

0.0

Contents (n=36)

Mucosa (n=36)

Sample_origin



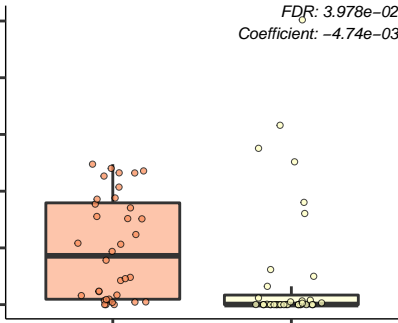
Brevibacterium

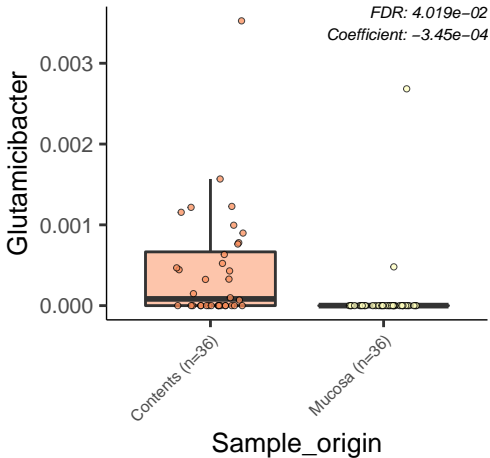
FDR: 3.978e-02
Coefficient: -4.74e-03

Contents (n=36)

Mucosa (n=36)

Sample_origin





f__Corynebacteriaceae

FDR: $5.544e-02$
Coefficient: $-1.25e-03$

0.010

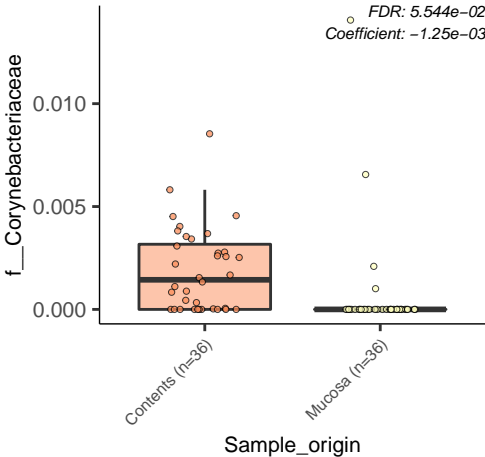
0.005

0.000

Contents (n=36)

Mucosa (n=36)

Sample_origin



p_RsaHF231

FDR: $6.288e-02$
Coefficient: $-9.38e-03$

Contents (n=36)

Mucosa (n=36)

Sample_origin

