

Lysinibacillus

FDR: $1.154e-06$

Coefficient: $-9.27e-03$

0.03

0.02

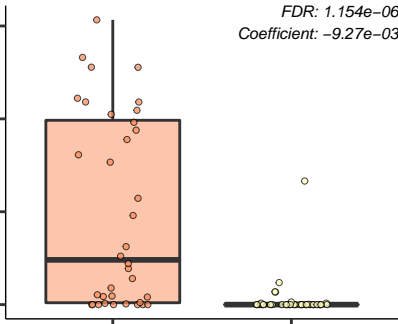
0.01

0.00

Contents (n=36)

Mucosa (n=36)

Sample_origin



f_Bacillaceae

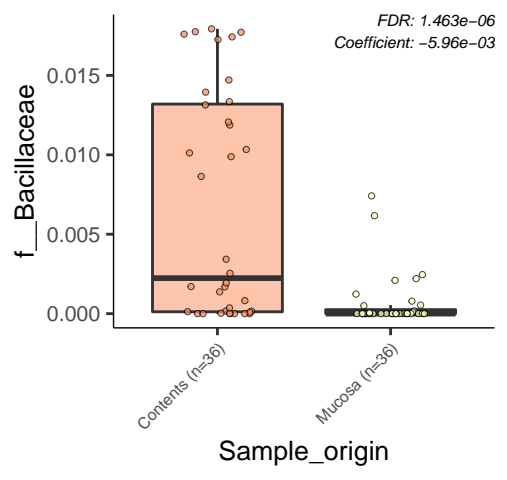
FDR: $1.463e-06$
Coefficient: $-5.96e-03$

Contents (n=36)

Mucosa (n=36)

Sample_origin

0.000
0.005
0.010
0.015



Brevinema andersonii

0.6
0.4
0.2
0.0

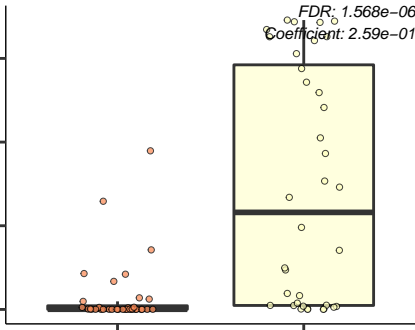
Contents (n=36)

Mucosa (n=36)

Sample_origin

FDR: 1.568e-06

Coefficient: 2.59e-01



f_Lachnospiraceae

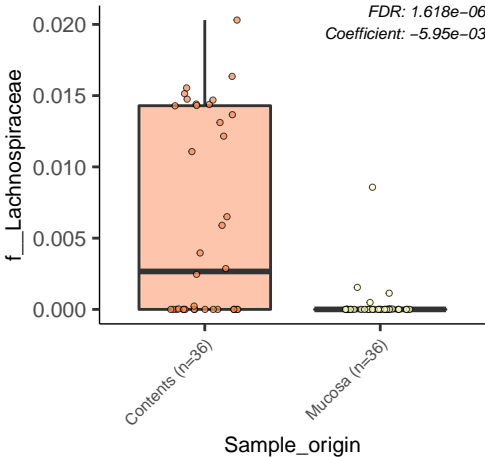
0.020
0.015
0.010
0.005
0.000

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

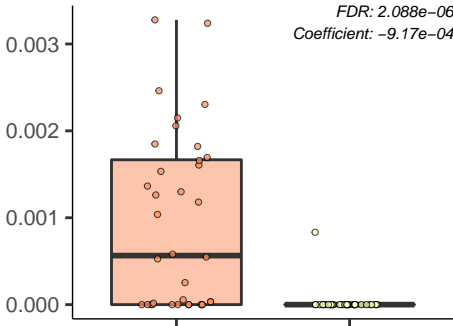
Contents (n=36)

Mucosa (n=36)

Sample_origin



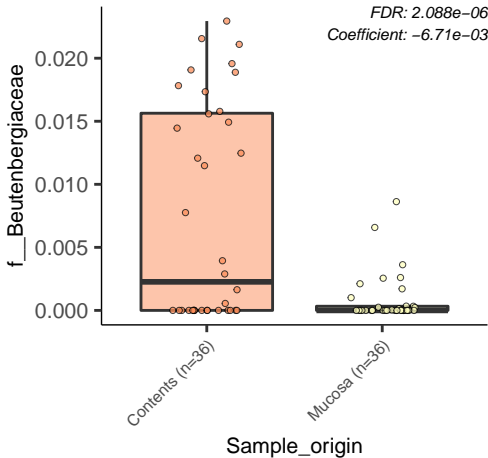
Coefficient: $-9.17e-04$

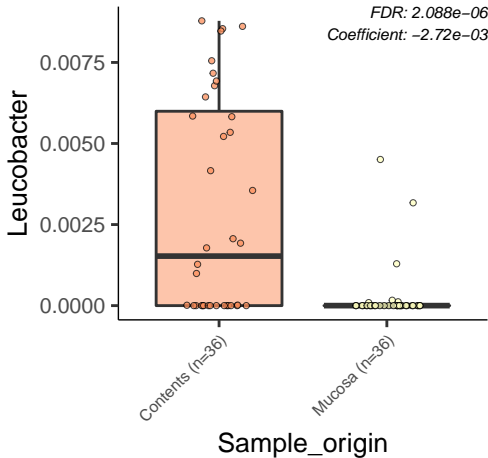


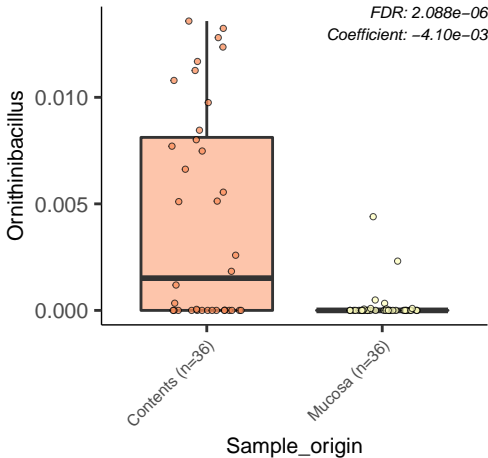
Contents (n=36)

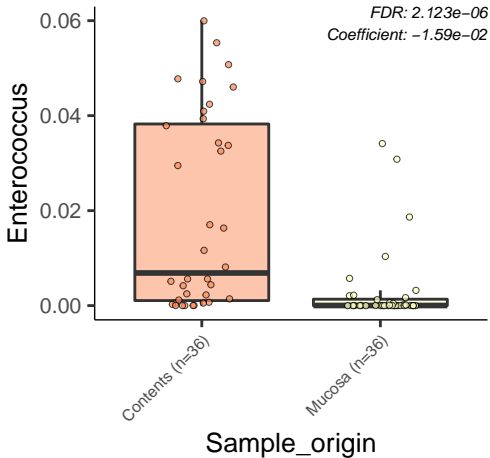
Mucosa (n=36)

Sample_origin









Oceanobacillus

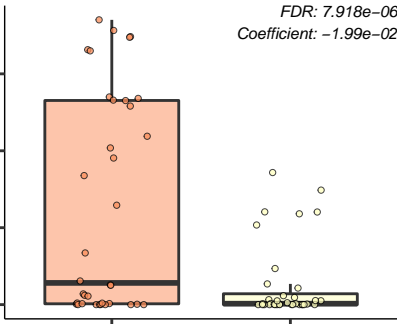
FDR: 7.918e-06
Coefficient: -1.99e-02

0.06
0.04
0.02
0.00

Contents (n=36)

Mucosa (n=36)

Sample_origin



Lactobacillus

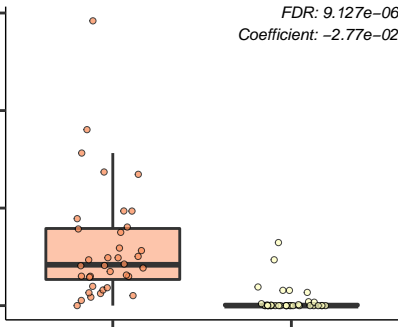
FDR: 9.127e-06

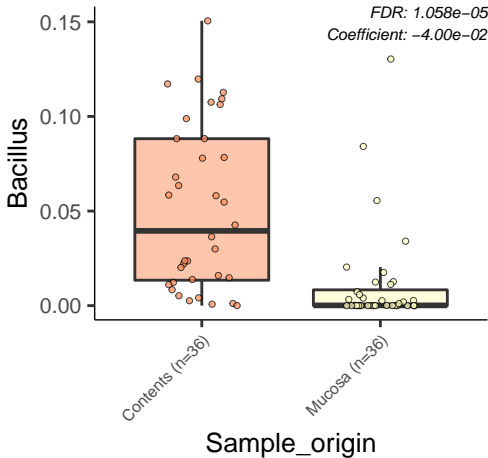
Coefficient: -2.77e-02

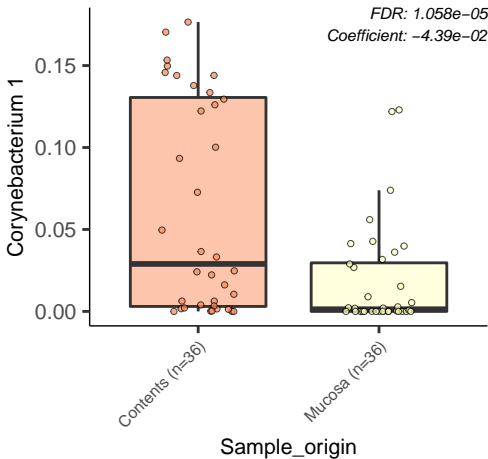
Contents (n=36)

Mucosa (n=36)

Sample_origin







Gracilibacillus

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

0.002

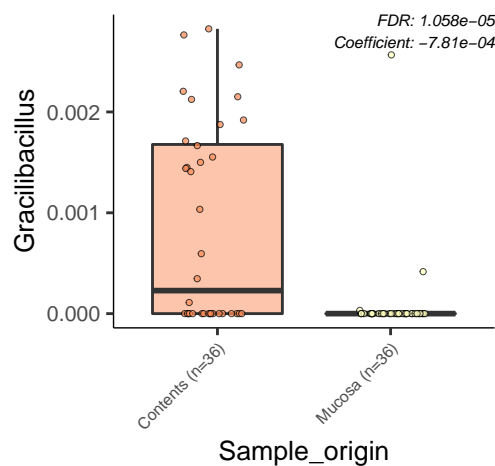
0.001

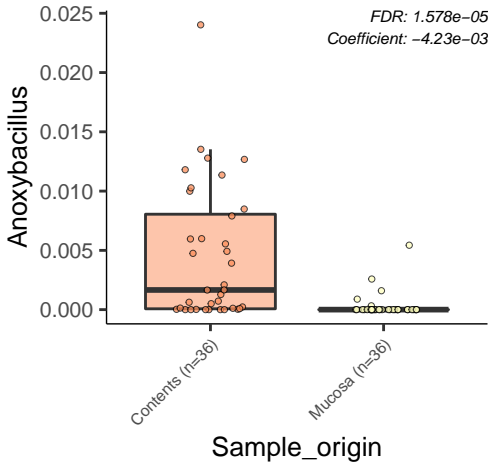
0.000

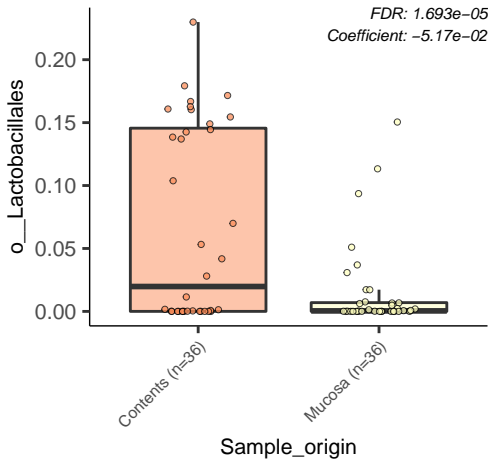
Contents (n=36)

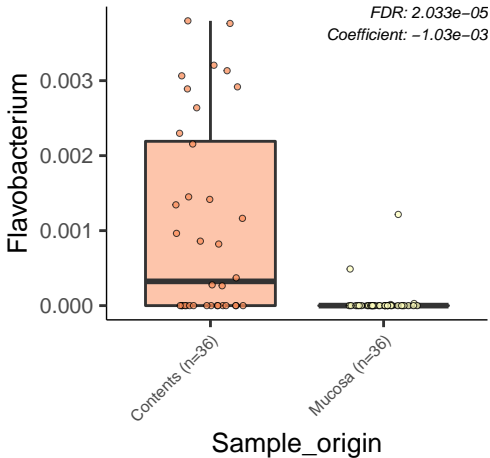
Mucosa (n=36)

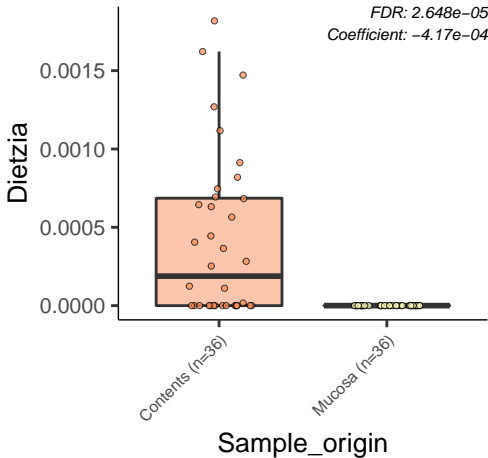
Sample_origin

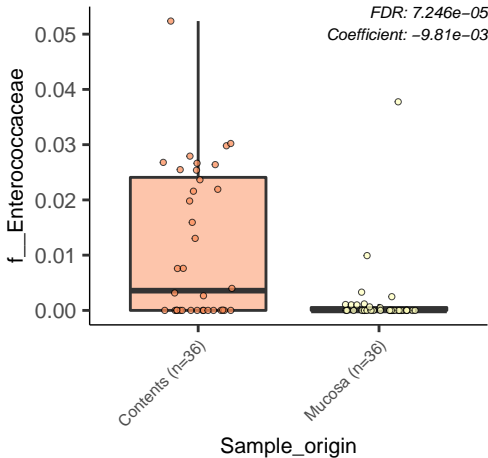












Tepidimicrobium

FDR: 3.637e-04

Coefficient: -7.09e-03

0.03

0.02

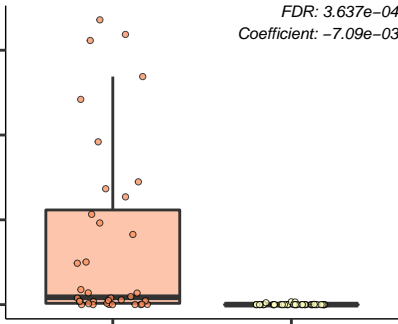
0.01

0.00

Contents (n=36)

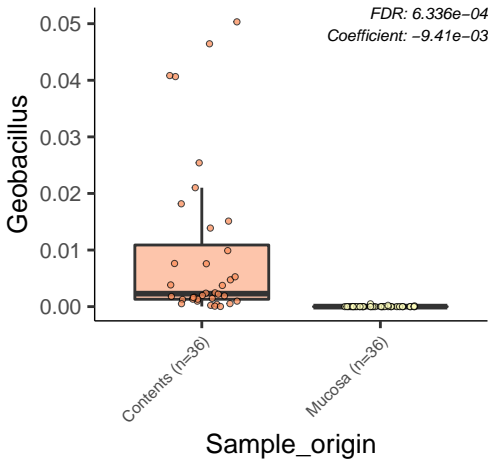
Mucosa (n=36)

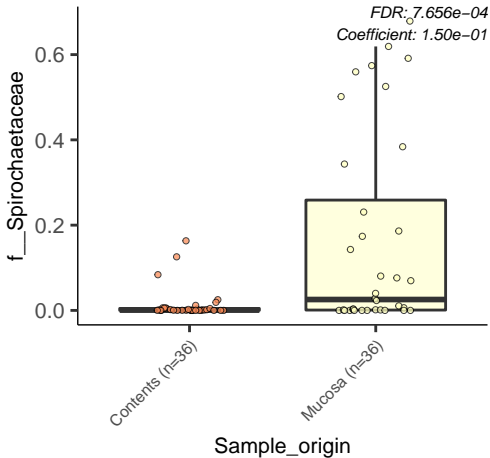
Sample_origin

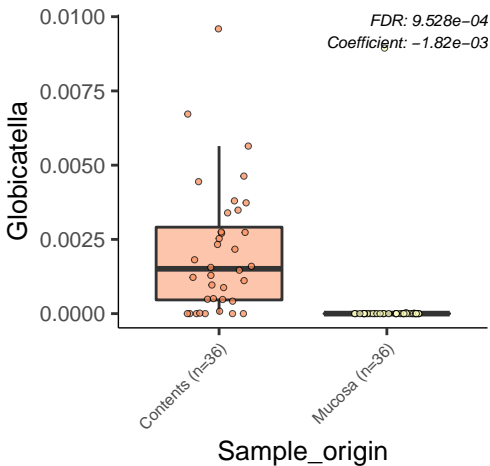


Coefficient: $-1.03e-03$

Sample_origin







Coefficient: $-2.79e-03$

Sample_origin

Mycoplasma

FDR: 5.619e-03

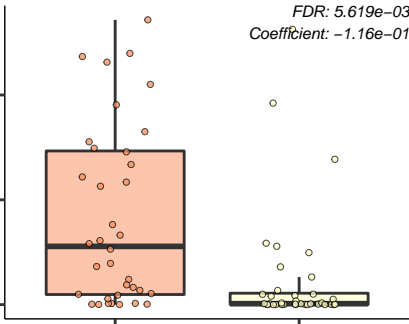
Coefficient: -1.16e-01

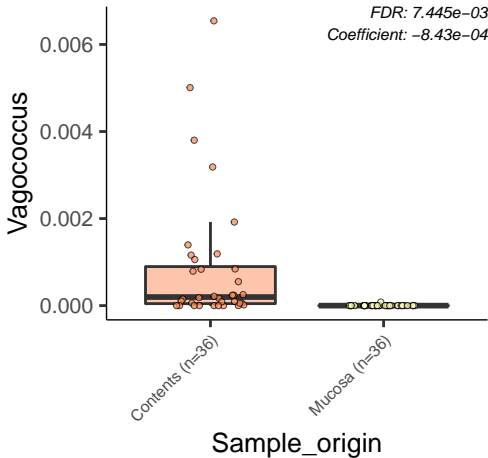
0.4
0.2
0.0

Contents (n=36)

Mucosa (n=36)

Sample_origin





Spingobacterium

FDR: 1.053e-02

Coefficient: -1.89e-03

0.015

0.010

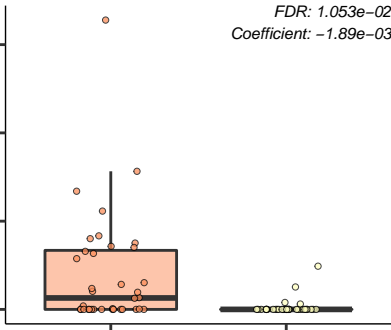
0.005

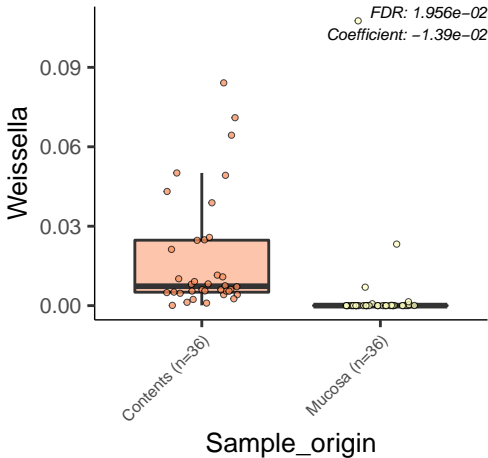
0.000

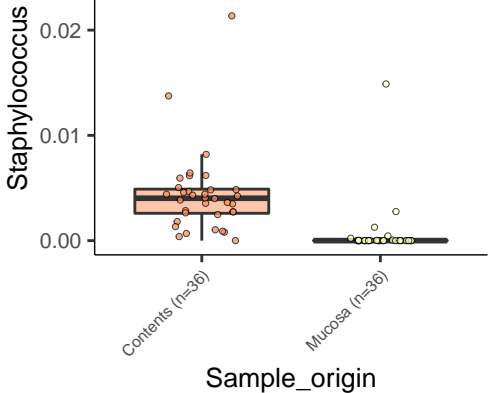
Contents (n=36)

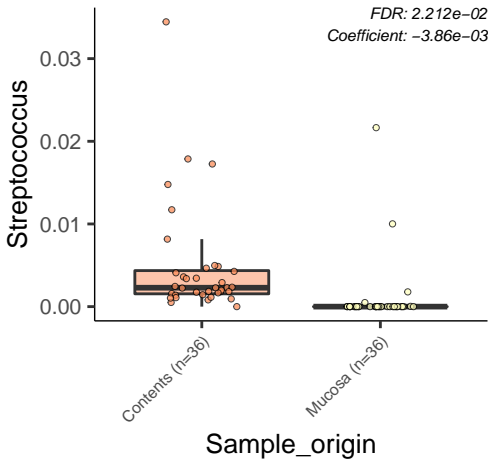
Mucosa (n=36)

Sample_origin









Actinomycetes

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

0.12

0.08

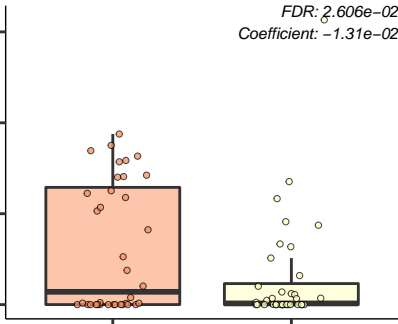
0.04

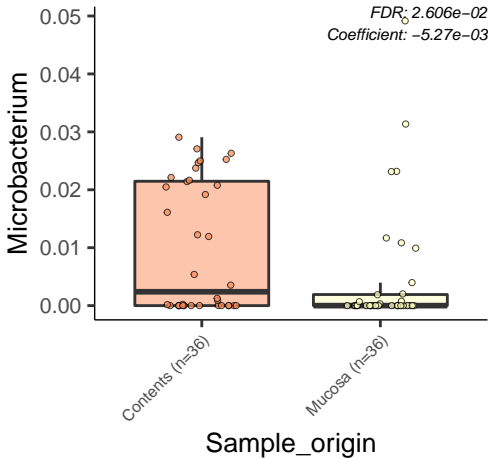
0.00

Contents (n=36)

Mucosa (n=36)

Sample_origin





Photobacterium

FDR: 2.606e-02

Coefficient: -3.48e-02

0.4

0.3

0.2

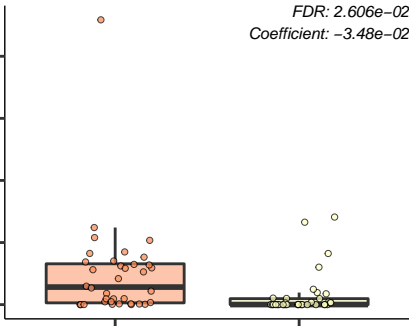
0.1

0.0

Contents (n=36)

Mucosa (n=36)

Sample_origin



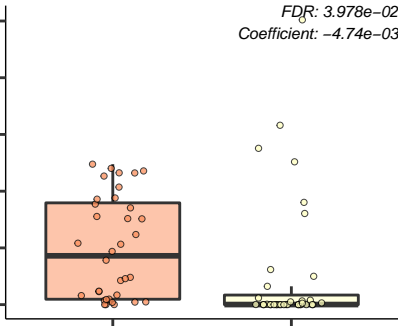
Brevibacterium

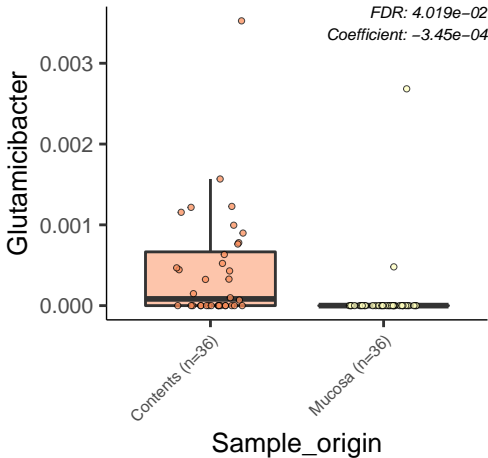
FDR: 3.978e-02
Coefficient: -4.74e-03

Contents (n=36)

Mucosa (n=36)

Sample_origin





f__Corynebacteriaceae

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

0.010

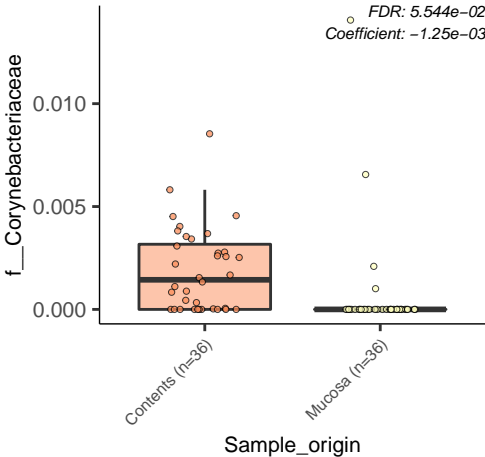
0.005

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

