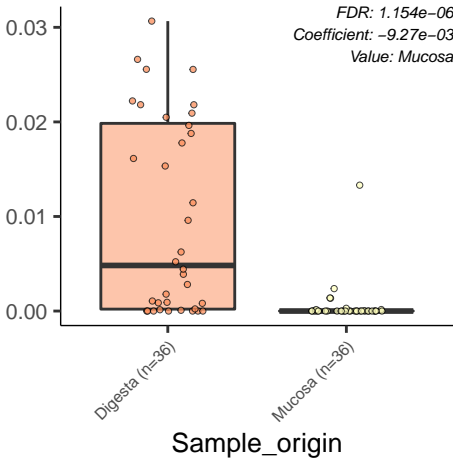


Lysinibacillus



f_Bacillaceae

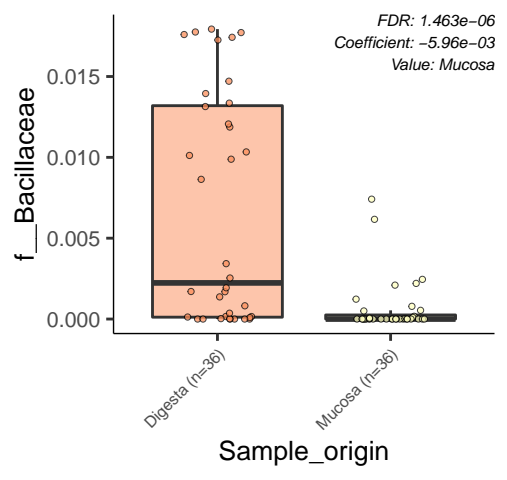
FDR: 1.463e-06
Coefficient: -5.96e-03
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin

0.015
0.010
0.005
0.000



Brevinema andersonii

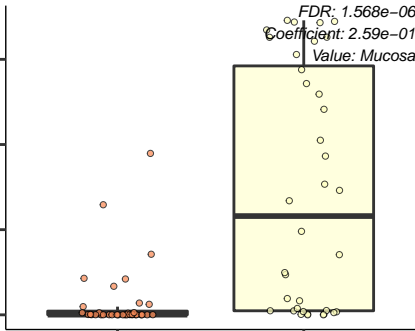
0.6
0.4
0.2
0.0

Digesta (n=36)

Mucosa (n=36)

Sample_origin

FDR: 1.568e-06
Coefficient: 2.59e-01
Value: Mucosa



f_Lachnospiraceae

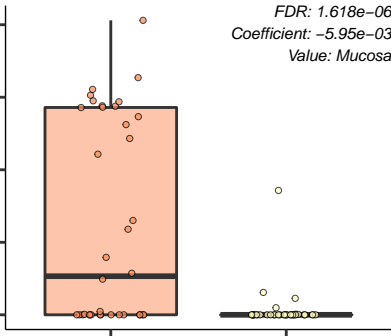
0.020
0.015
0.010
0.005
0.000

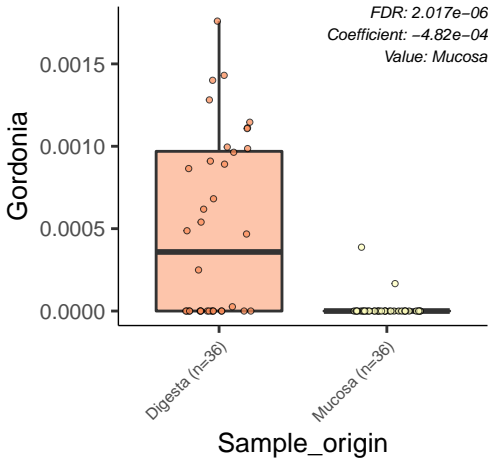
FDR: 1.618e-06
Coefficient: -5.95e-03
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin



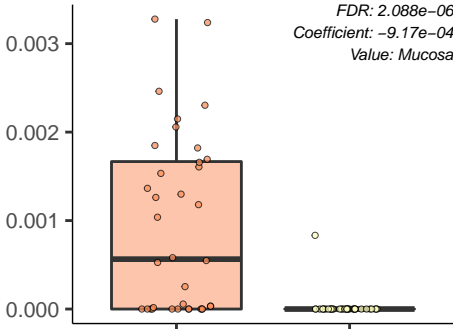




FDR: 2.088e-06

Coefficient: $-9.17e-04$

Value: Mucosa



Digesta (n=36)

Mucosa (n=36)

Sample_origin

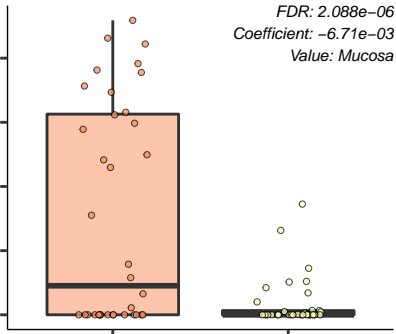
f_ Beutenbergiaceae

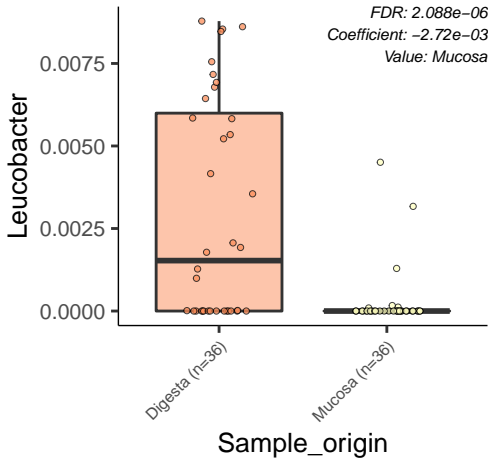
FDR: 2.088e-06
Coefficient: -6.71e-03
Value: Mucosa

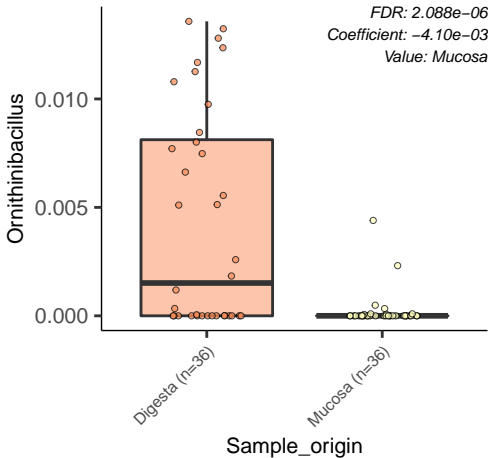
Digesta (n=36)

Mucosa (n=36)

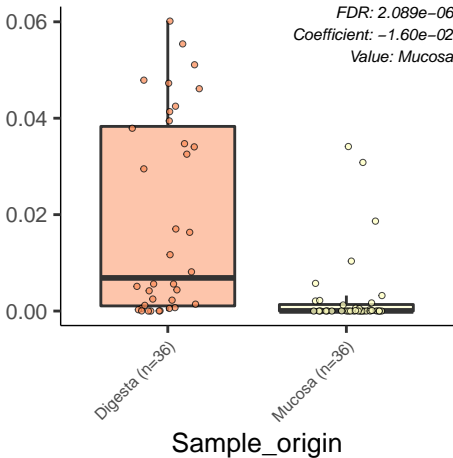
Sample_origin







Enterococcus



Oceanobacillus

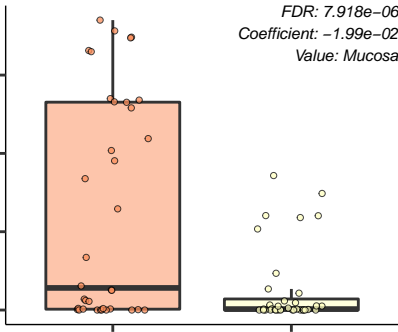
FDR: 7.918e-06
Coefficient: -1.99e-02
Value: Mucosa

0.06
0.04
0.02
0.00

Digesta (n=36)

Mucosa (n=36)

Sample_origin



Lactobacillus

FDR: 9.127e-06

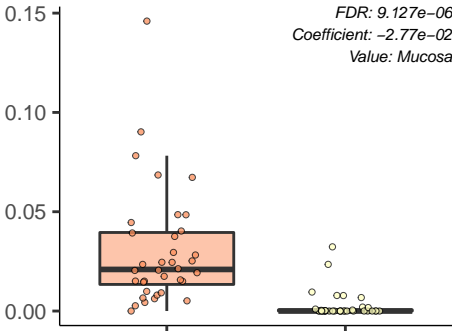
Coefficient: -2.77e-02

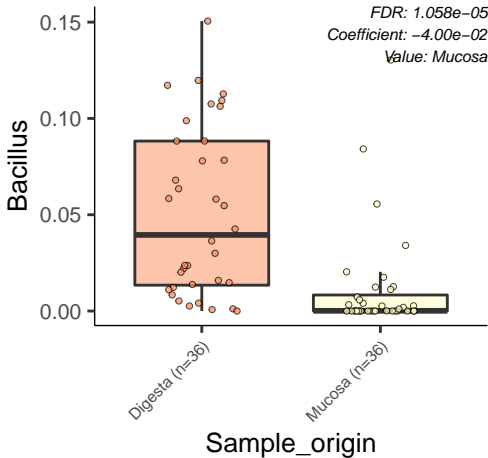
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin





Corynebacterium 1

FDR: $1.058e-05$
Coefficient: $-4.39e-02$
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin

0.15

0.10

0.05

0.00

Gracilibacillus

FDR: 1.058e-05

Coefficient: -7.81e-04

Value: Mucosa

0.002

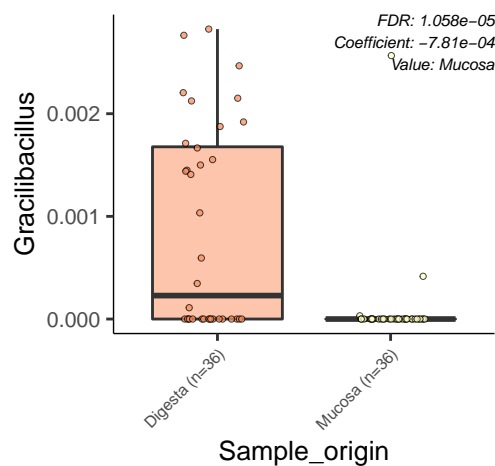
0.001

0.000

Digesta (n=36)

Mucosa (n=36)

Sample_origin



Anoxybacillus

FDR: 1.578e-05

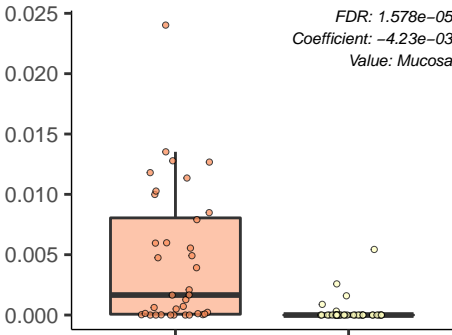
Coefficient: -4.23e-03

Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin



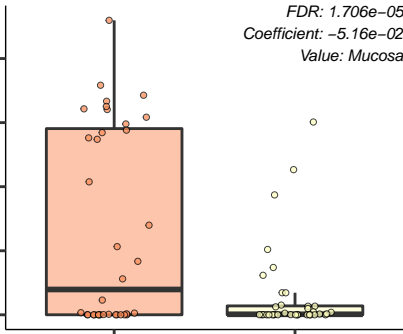
o_Lactobacillales

FDR: 1.706e-05
Coefficient: -5.16e-02
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin



Flavobacterium

FDR: 2.033e-05

Coefficient: -1.03e-03

Value: Mucosa

0.003

0.002

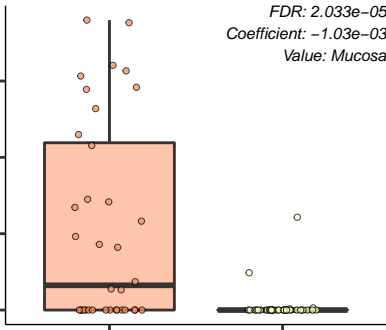
0.001

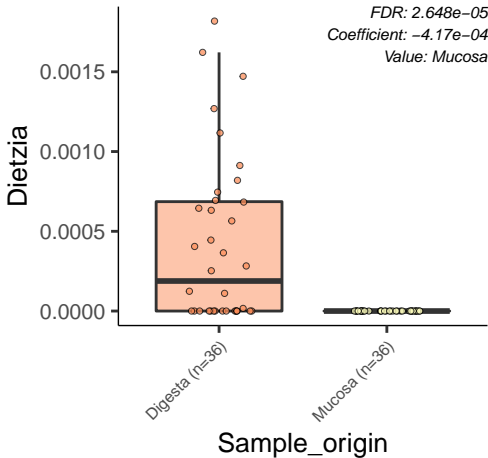
0.000

Digesta (n=36)

Mucosa (n=36)

Sample_origin





Rhodococcus

FDR: 5.737e-05

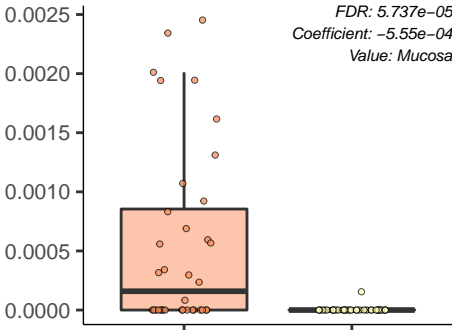
Coefficient: -5.55e-04

Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin



f_Enterococcaceae

FDR: 7.246×10^{-5}

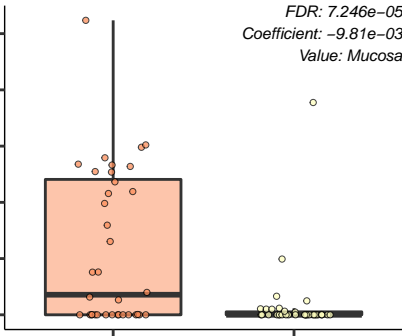
Coefficient: -9.81×10^{-3}

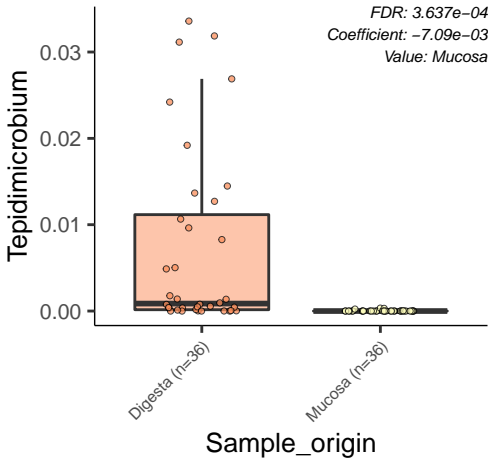
Value: Mucosa

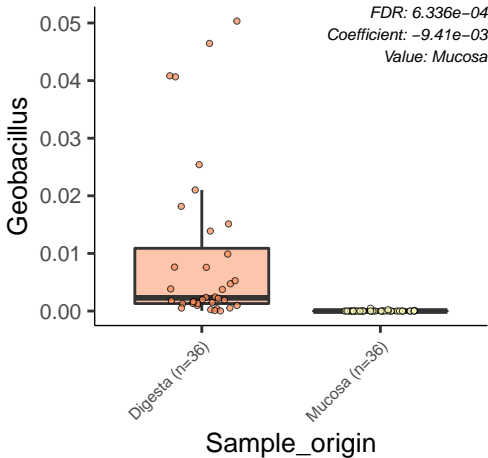
Digesta (n=36)

Mucosa (n=36)

Sample_origin







f_Spirochaetaceae

0.6

0.4

0.2

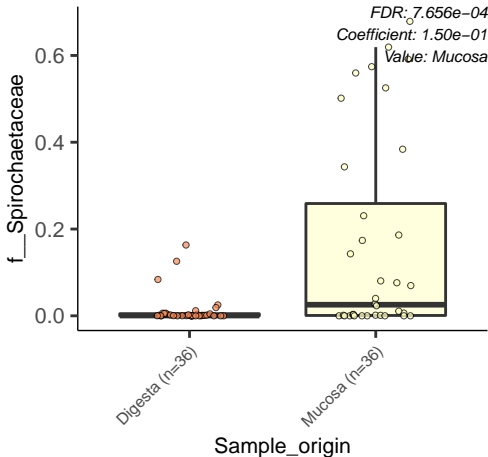
0.0

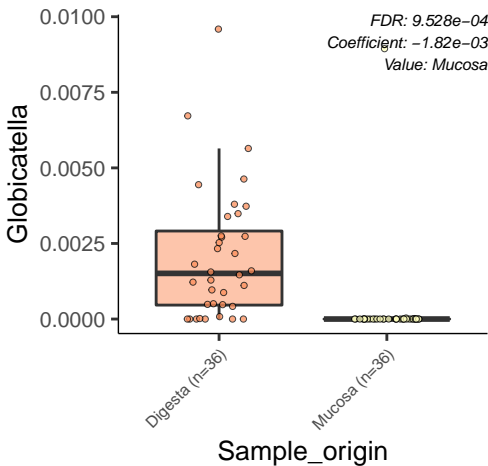
Digesta (n=36)

Mucosa (n=36)

Sample_origin

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





Peptostreptococcus

FDR: 2.202e-03

Coefficient: -4.36e-03

Value: Mucosa

0.02

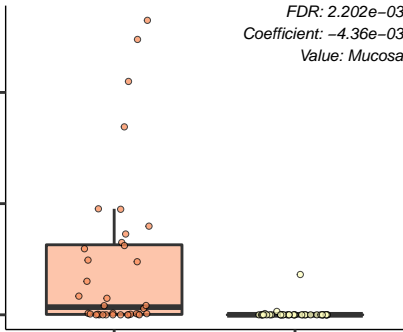
0.01

0.00

Digesta (n=36)

Mucosa (n=36)

Sample_origin



Mycoplasma

FDR: 5.619e-03

Coefficient: -1.16e-01

Value: Mucosa

0.4

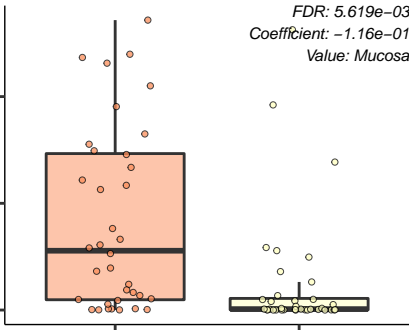
0.2

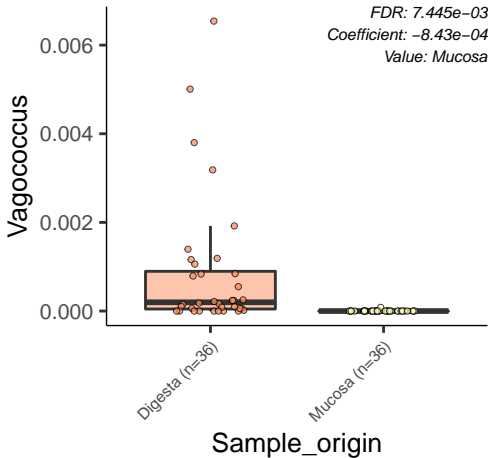
0.0

Digesta (n=36)

Mucosa (n=36)

Sample_origin





Value: Mucosa



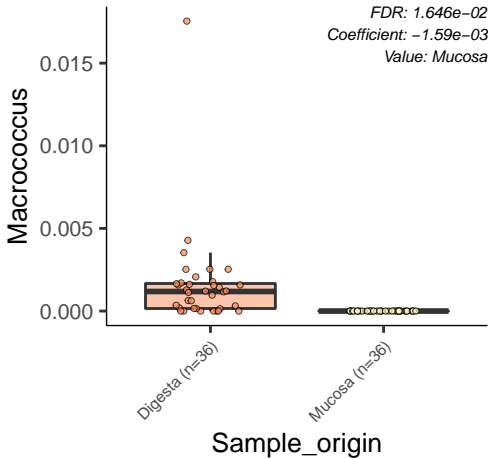
Sample_origin

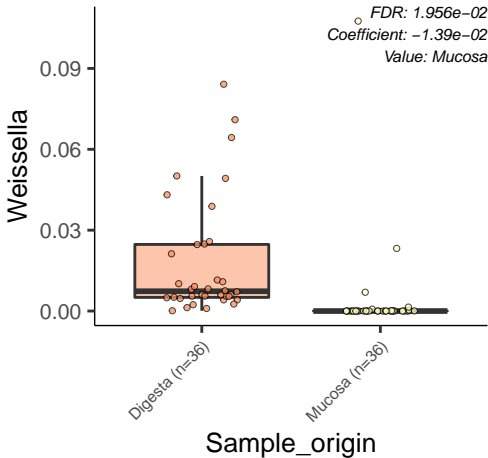
Value: Mucosa

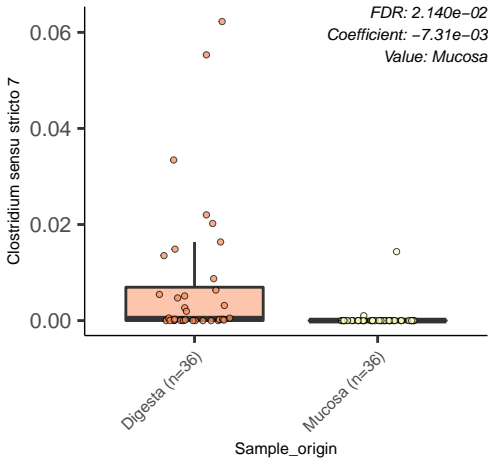


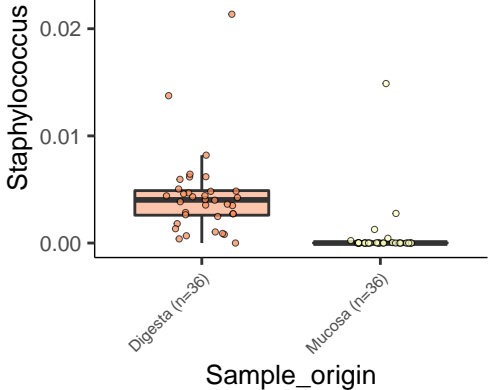
Mucosa (n=36)

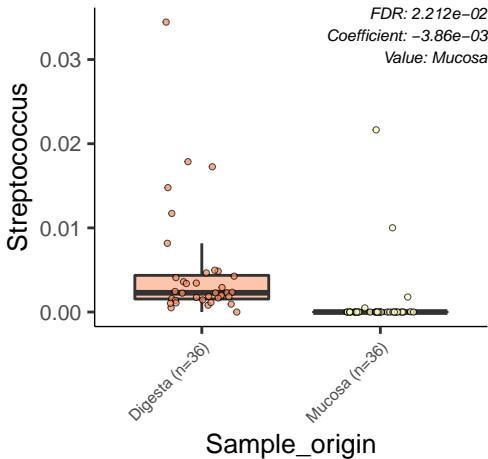
Sample_origin

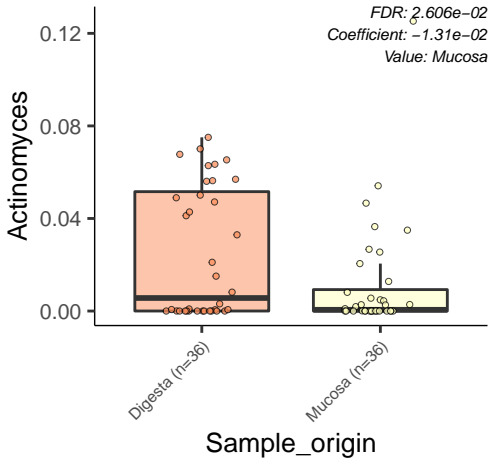


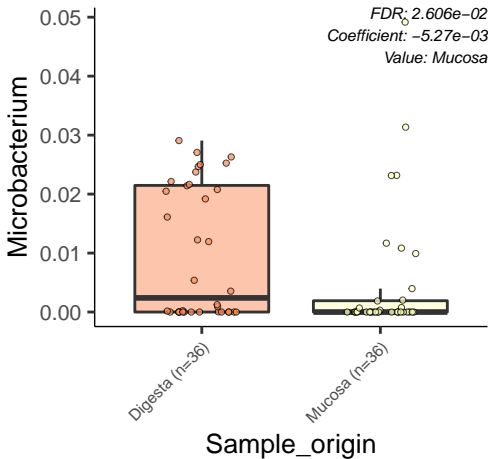












Photobacterium

FDR: 2.606e-02

Coefficient: -3.48e-02

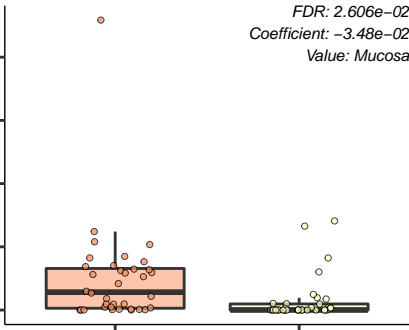
Value: Mucosa

Digesta (n=36)

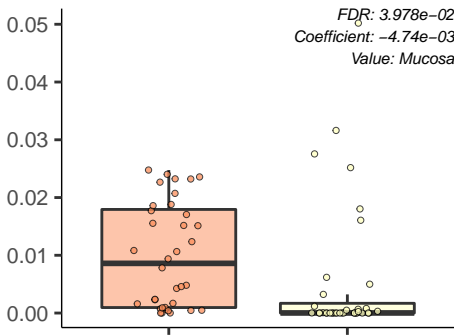
Mucosa (n=36)

Sample_origin

0.4
0.3
0.2
0.1
0.0



Brevibacterium

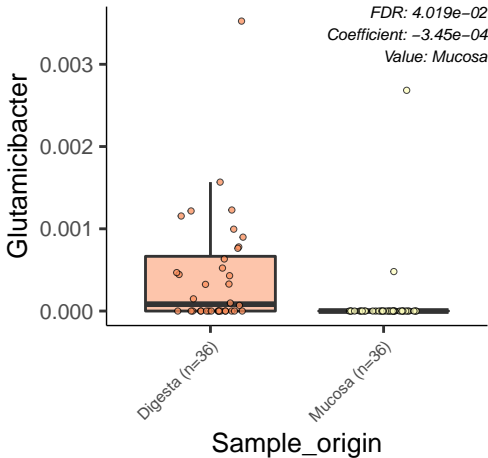


Digesta (n=36)

Mucosa (n=36)

Sample_origin

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



f_Corynebacteriaceae

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

0.010

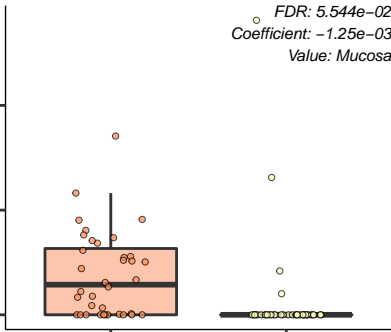
0.005

0.000

Digesta (n=36)

Mucosa (n=36)

Sample_origin



p_RsaHF231

FDR: 6.288e-02
Coefficient: -9.38e-03
Value: Mucosa

Digesta (n=36)

Mucosa (n=36)

Sample_origin

