

Cloacibacterium

FDR: 6.881e-07

Coefficient: -2.51e+00

Value: A+ T+ P+

200

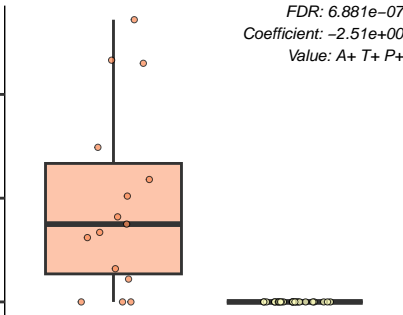
100

0

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment



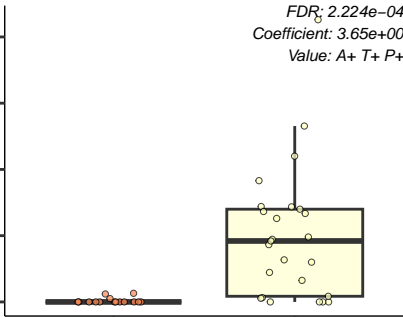
Flavobacterium

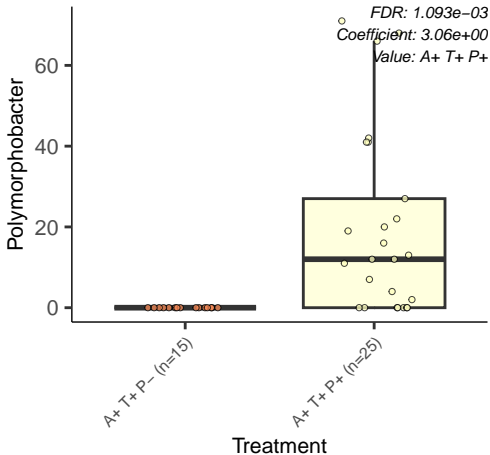
FDR: 2.224e-04
Coefficient: 3.65e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





Rhodococcus

750

500

250

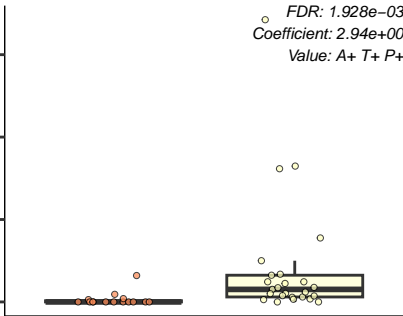
0

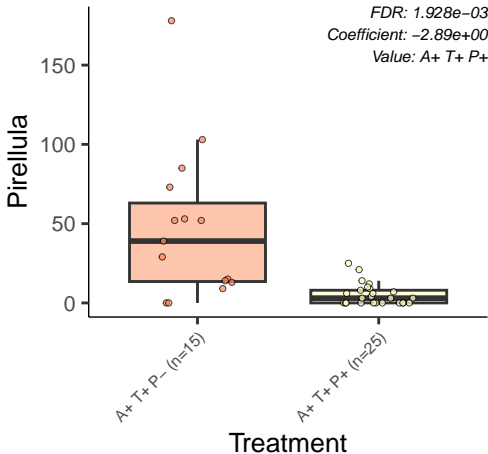
A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment

FDR: 1.928e-03
Coefficient: 2.94e+00
Value: A+ T+ P+





Pseudobdellovibrionaceae_Genus

FDR: 5.945e-03

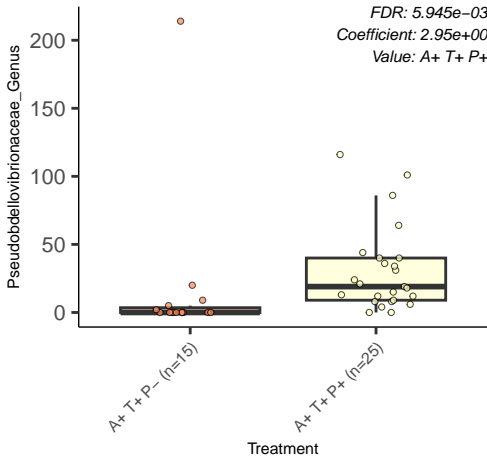
Coefficient: 2.95e+00

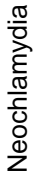
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





FDR: 5.945e-03

Coefficient: $-2.97e+00$

Value: $A+ T+ P+$



Treatment

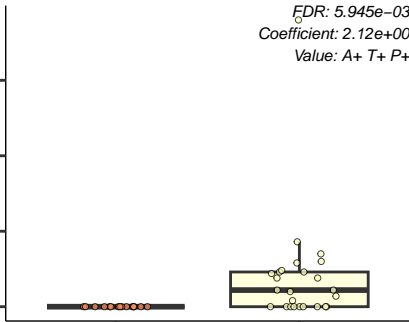
Candidatus_Megaira

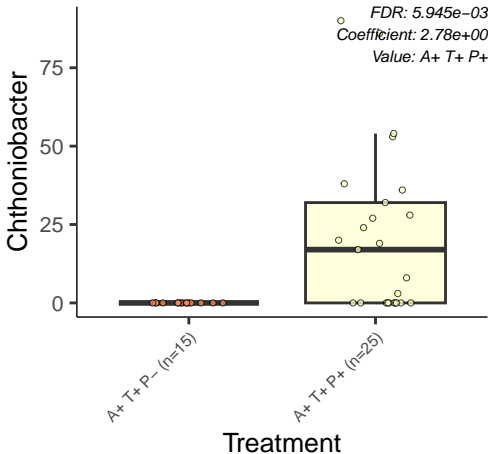
FDR: 5.945e-03
Coefficient: 2.12e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





Niveibacterium

FDR: 6.605e-03

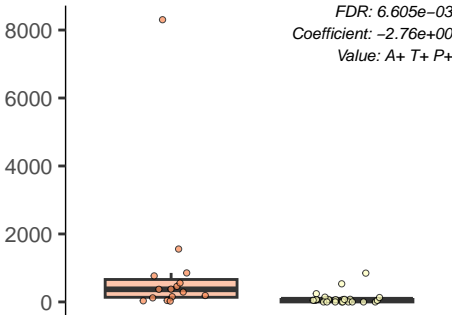
Coefficient: -2.76e+00

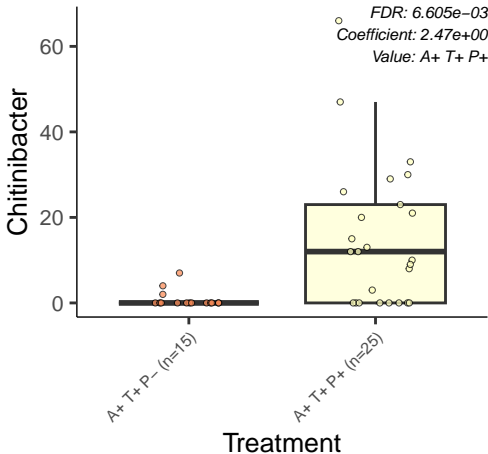
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment

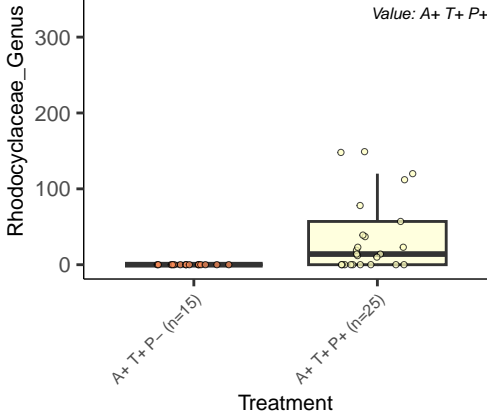


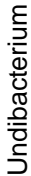


FDR: 1.402e-02

Coefficient: 1.94e+00

Value: A+ T+ P+





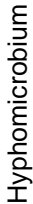
FDR: 2.408e-02

Coefficient: $-3.49e+00$

Value: A+ T+ P+



Treatment



FDR: 2.528e-02

Coefficient: $-1.64e+00$

Value: A+ T+ P+



Treatment

Pseudomonas

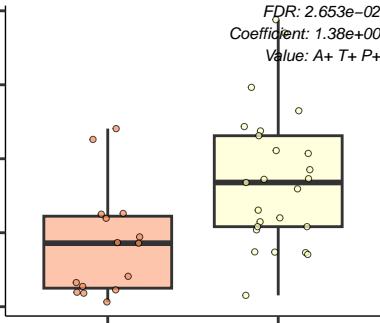
10000
7500
5000
2500
0

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment

FDR: 2.653e-02
Coefficient: 1.38e+00
Value: A+ T+ P+



Dinghuibacter

FDR: 2.653e-02

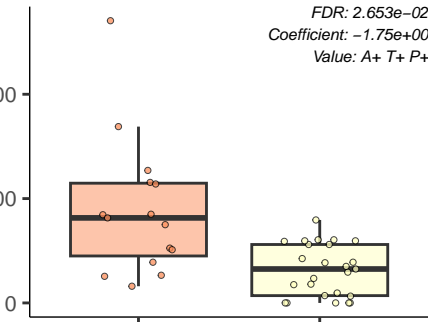
Coefficient: -1.75e+00

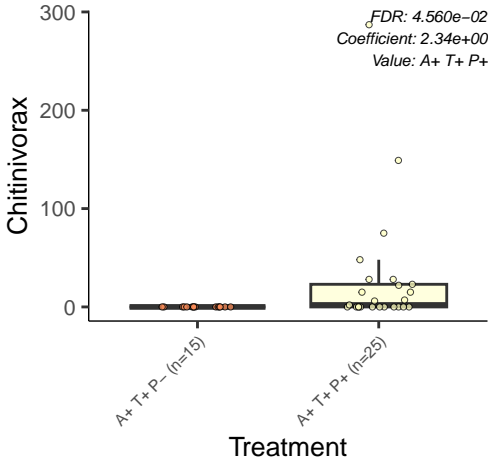
Value: A+ T+ P+

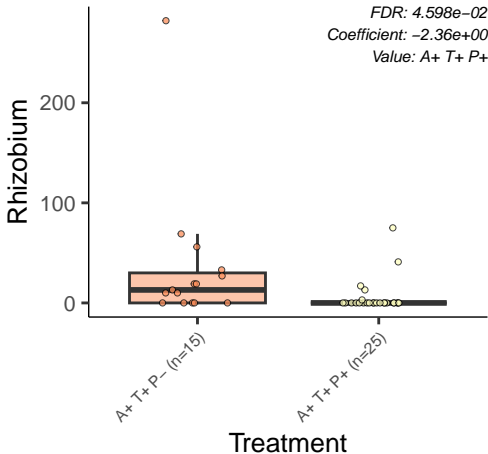
A+ T+ P- (n=15)

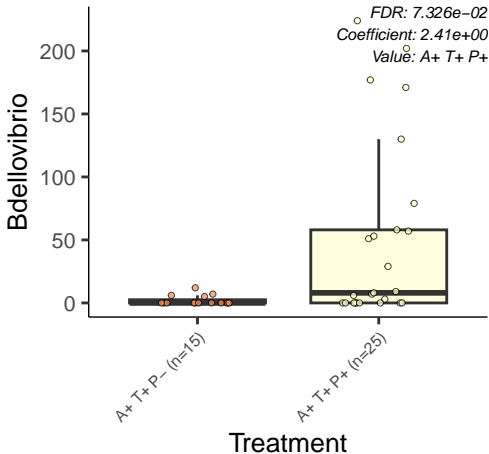
A+ T+ P+ (n=25)

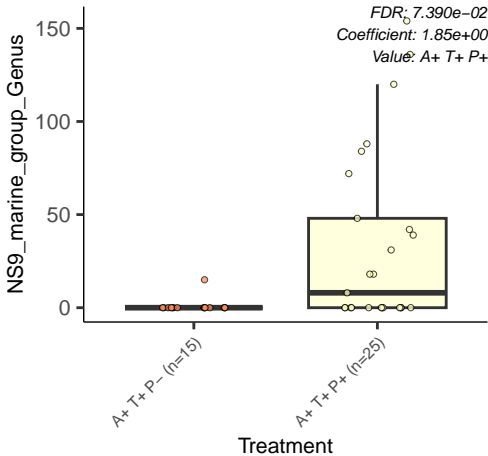
Treatment

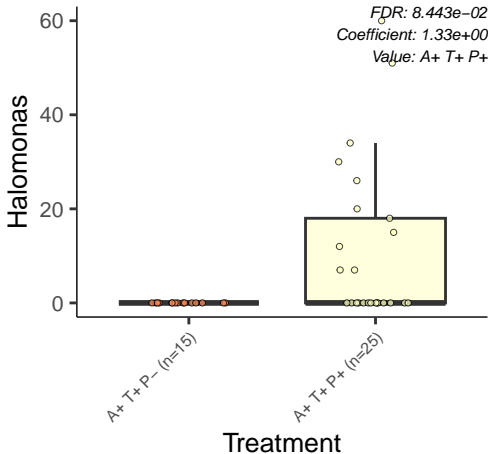




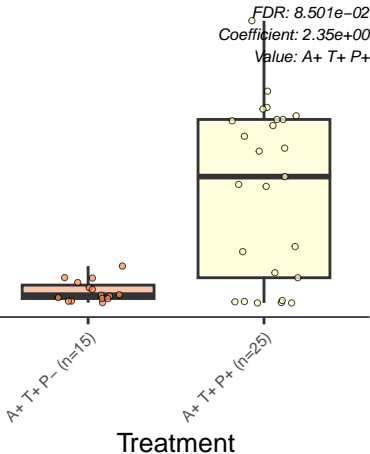








Shewanella





○ *FDR: 8.501e-02*
Coefficient: 1.94e+00
Value: A+ T+ P+



Treatment

Pirellulaceae_Genus

FDR: 8.501e-02
Coefficient: 1.33e+00
Value: A+ T+ P+

20

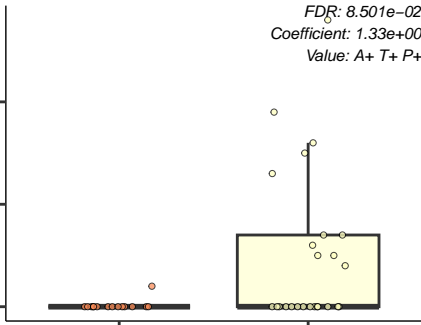
10

0

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment



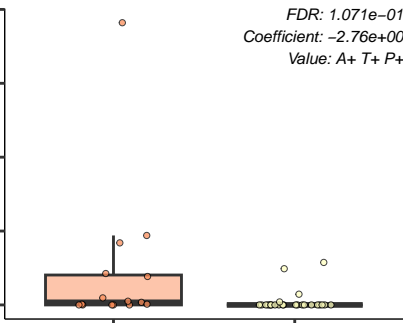
X11.24_Genus

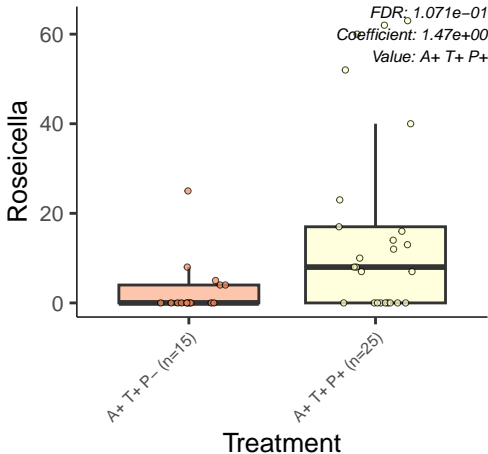
FDR: 1.071e-01
Coefficient: -2.76e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





Rhizobiaceae_Genus

FDR: 1.101e-01

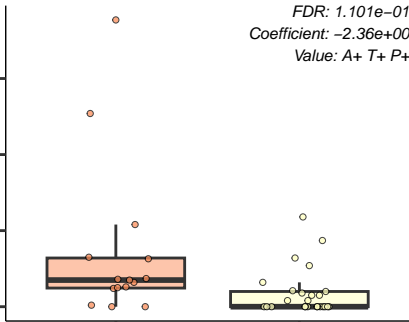
Coefficient: -2.36e+00

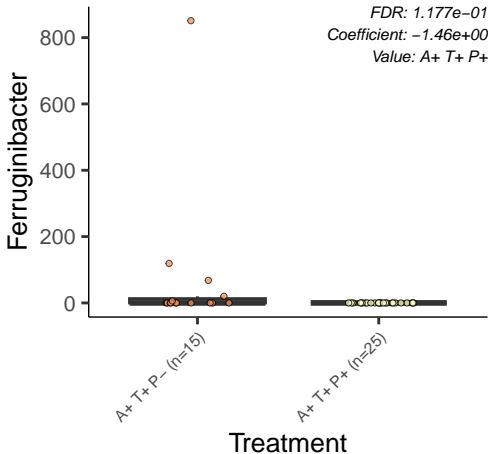
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





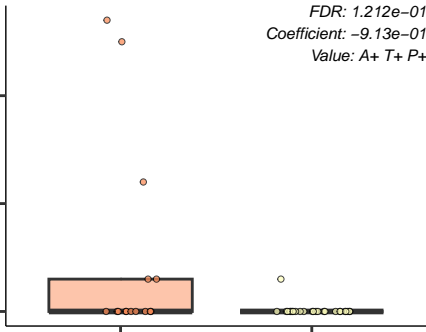
Thermodesulfobacteriota_Genus

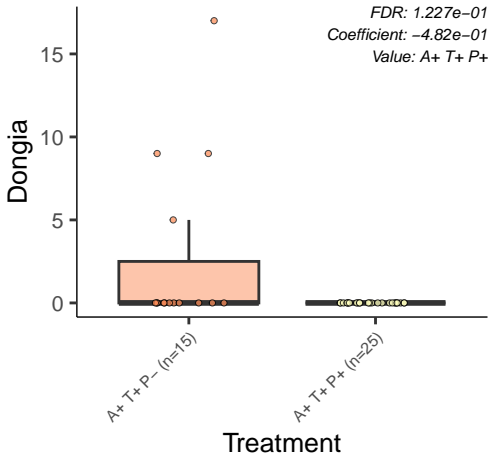
FDR: 1.212e-01
Coefficient: -9.13e-01
Value: A+ T+ P+

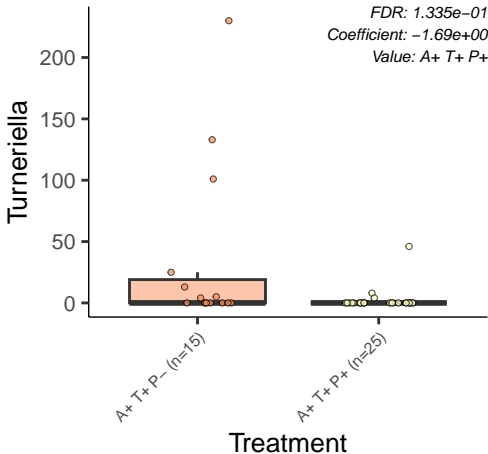
A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment









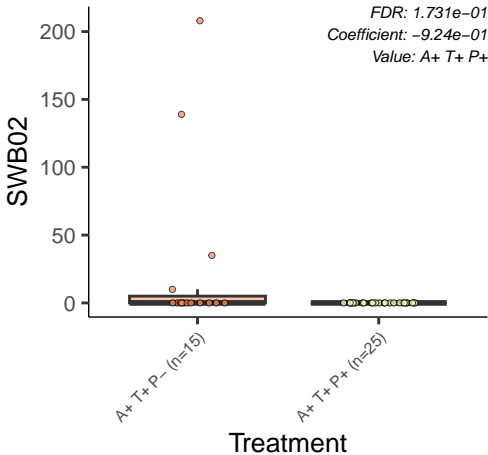
FDR: 1.650e-01

Coefficient: $-1.21e+00$

Value: A+ T+ P+



Treatment



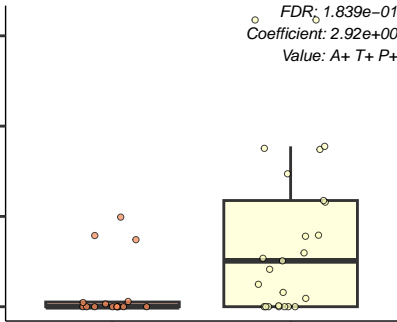
env.OPS_17_Genus

FDR: 1.839e-01
Coefficient: 2.92e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment



Saprospiraceae_Genus

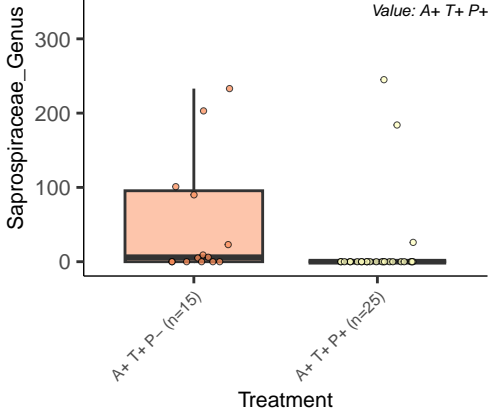
400
300
200
100
0

FDR: 1.847e-01
Coefficient: -1.86e+00
Value: A+ T+ P+

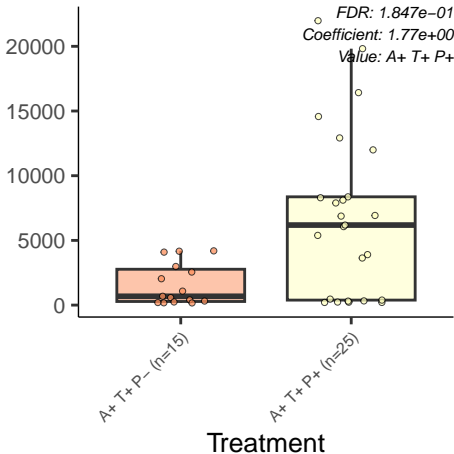
A+ T+ P- (n=15)

A+ T+ P+ (n=25)

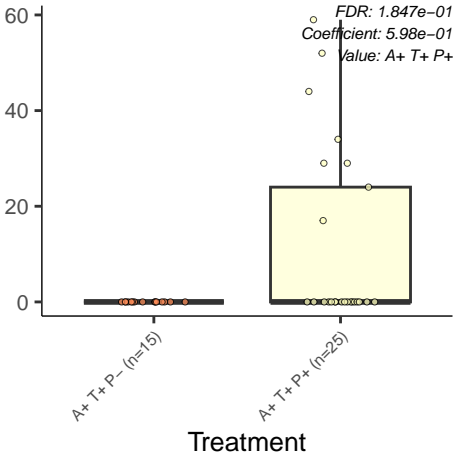
Treatment

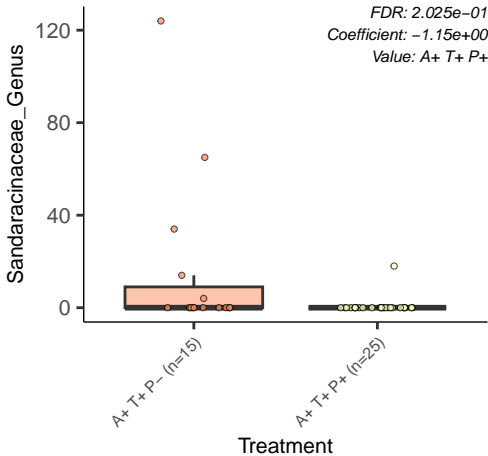


Cetobacterium



Pararhodobacter





Pedomicrobium

FDR: 2.143e-01

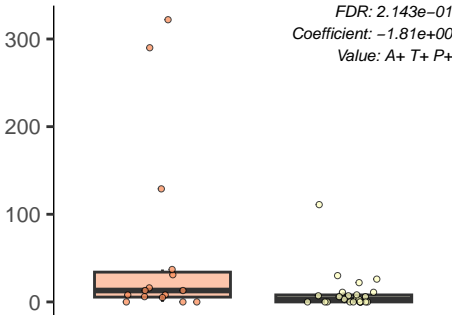
Coefficient: -1.81e+00

Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment



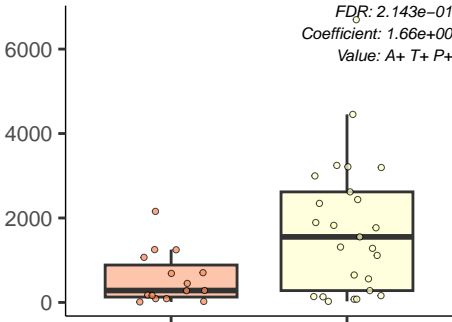
Crenobacter

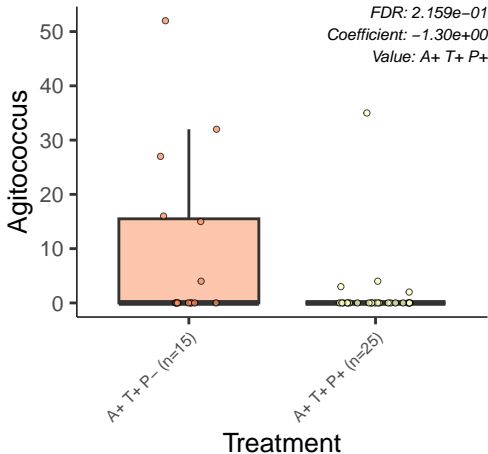
FDR: 2.143e-01
Coefficient: 1.66e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment





Paucibacter

FDR: 2.188e-01

Coefficient: -1.45e+00

Value: A+ T+ P+

6000

4000

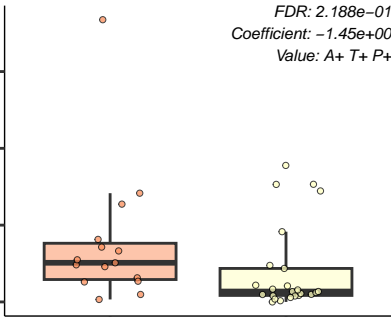
2000

0

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment



Aquihabitans

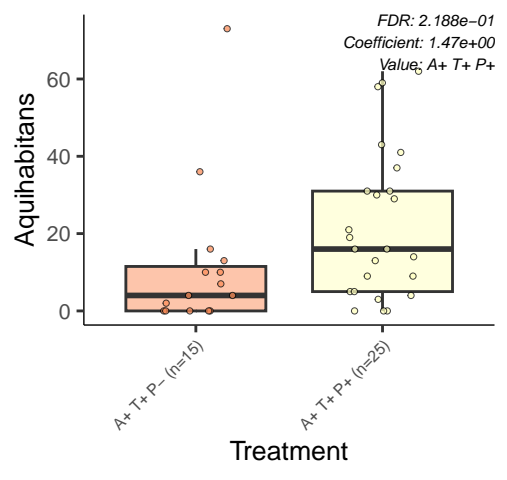
A+ T+ P- (n=15)

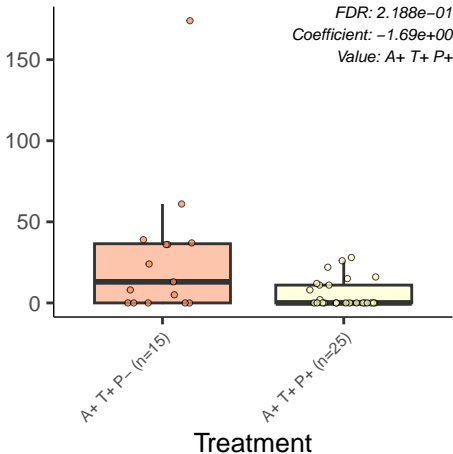
A+ T+ P+ (n=25)

Treatment

FDR: 2.188e-01
Coefficient: 1.47e+00
Value: A+ T+ P+

60
40
20
0





Staphylococcus

FDR: 2.188e-01

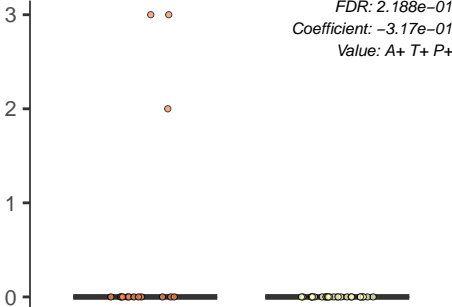
Coefficient: -3.17e-01

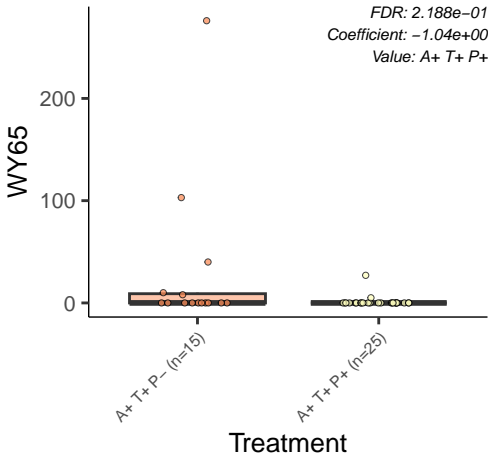
Value: A+ T+ P+

A+ T+ P- (n=15)

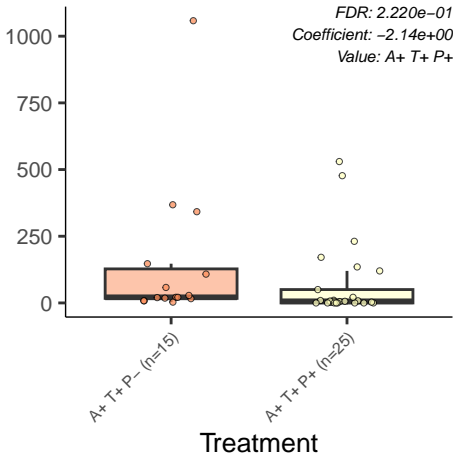
A+ T+ P+ (n=25)

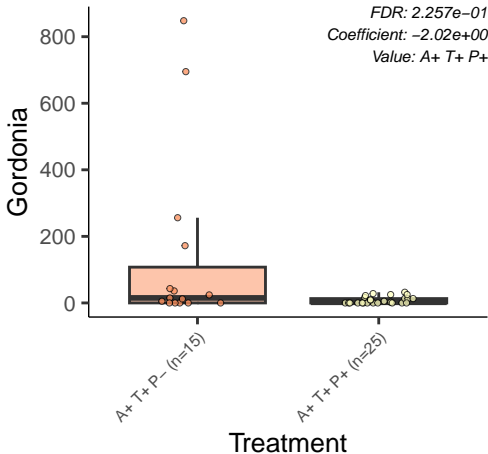
Treatment





Clostridium





Telmatocola

FDR: 2.269e-01

Coefficient: -1.88e+00

Value: A+ T+ P+

300

200

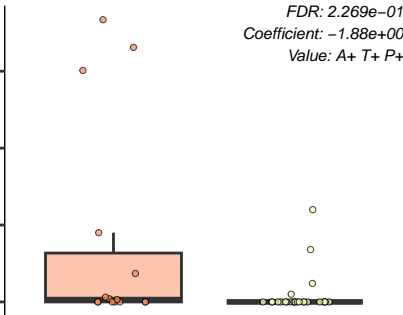
100

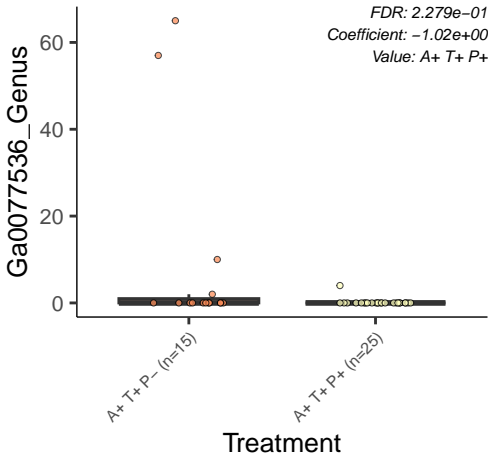
0

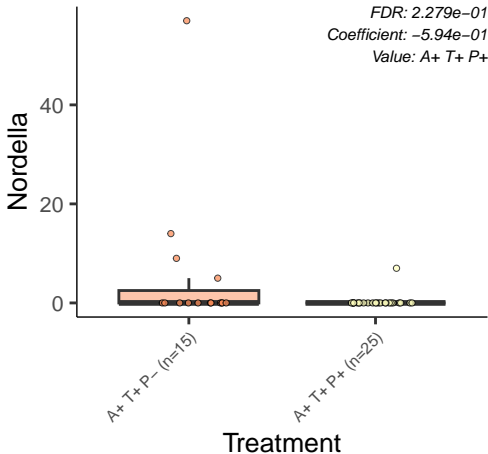
A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment







Chryseotalea

FDR: 2.382e-01
Coefficient: 2.42e+00
Value: A+ T+ P+

A+ T+ P- (n=15)

A+ T+ P+ (n=25)

Treatment

3000

2000

1000

0

