

Plesiomonas

FDR: 1.691e-09

Coefficient: -2.69e+00

Value: A- T- P+

2000

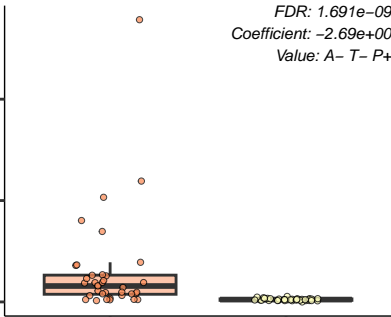
1000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



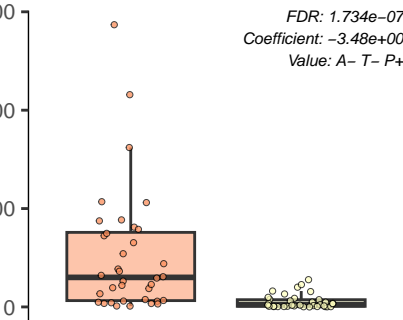
Culicoidibacter

A-T-P- (n=36)

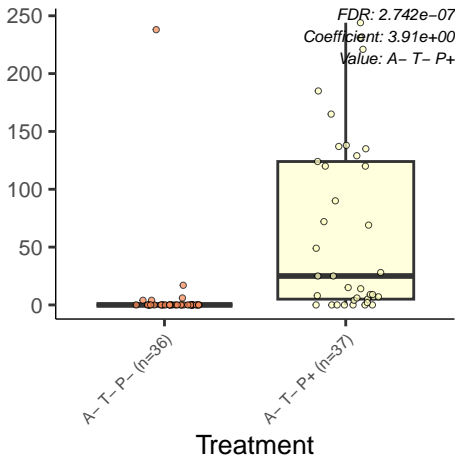
A-T-P+ (n=37)

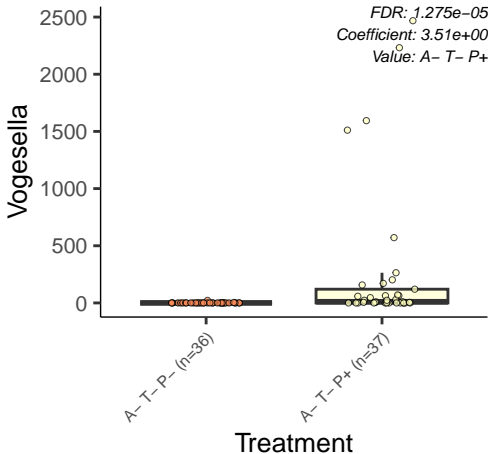
Treatment

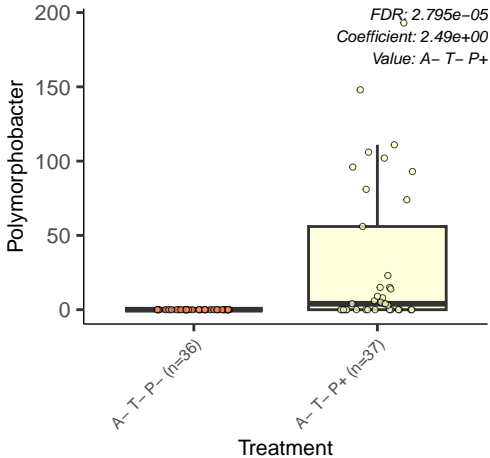
FDR: $1.734e-07$
Coefficient: $-3.48e+00$
Value: A- T- P+

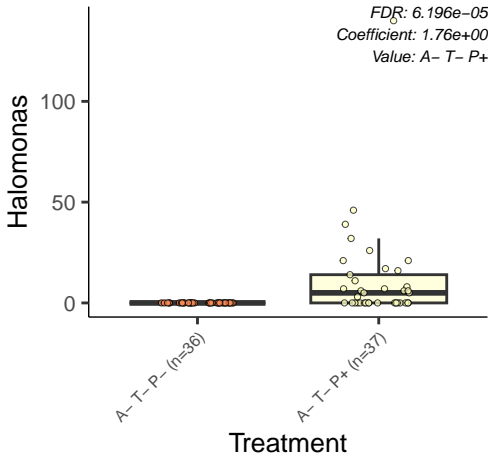


X67.14_Genus









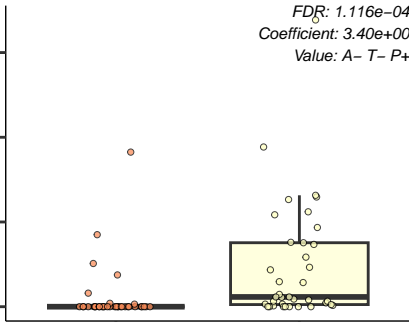
Vampirovibrionaceae_Genus

FDR: 1.116e-04
Coefficient: 3.40e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



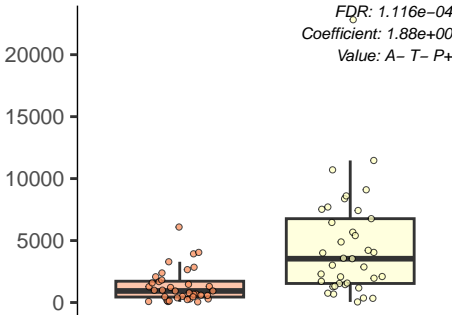
Pseudomonas

FDR: $1.116e-04$
Coefficient: $1.88e+00$
Value: A- T- P+

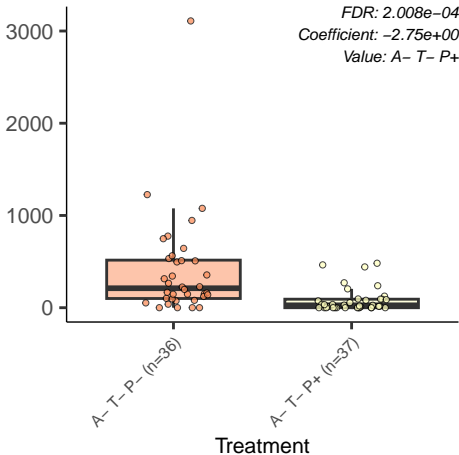
A-T-P- (n=36)

A-T-P+ (n=37)

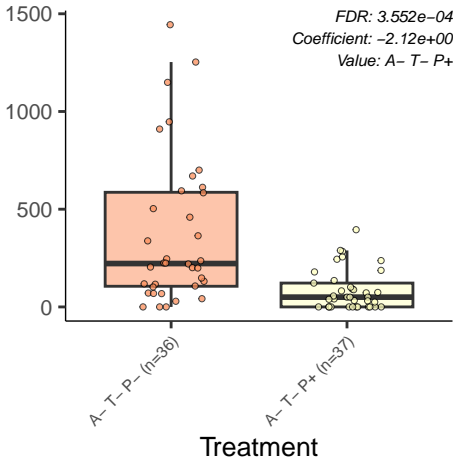
Treatment



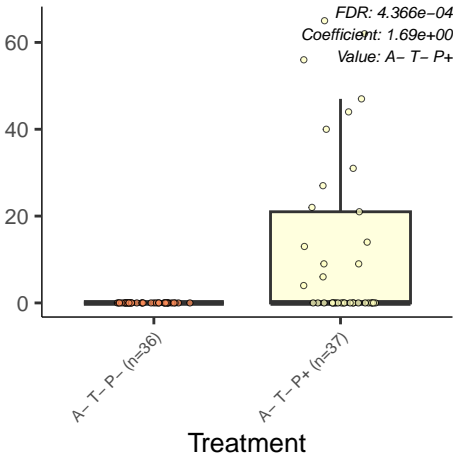
Paenirhodobacter



Gemmobacter



Rhodovarius



Paucibacter

FDR: 4.917e-04

Coefficient: -2.00e+00

Value: A- T- P+

15000

10000

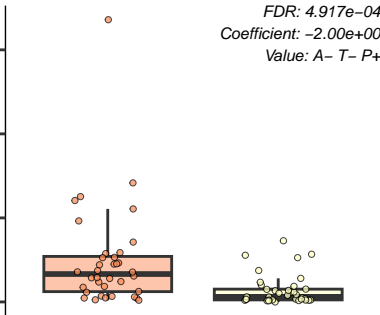
5000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



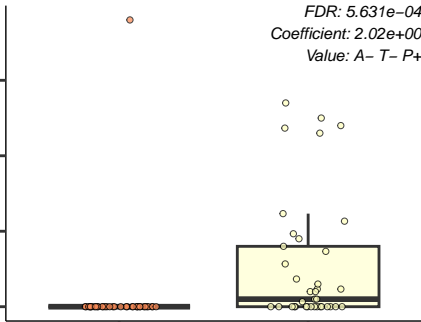
Candidatus_Megaira

FDR: 5.631e-04
Coefficient: 2.02e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Acinetobacter

1000

500

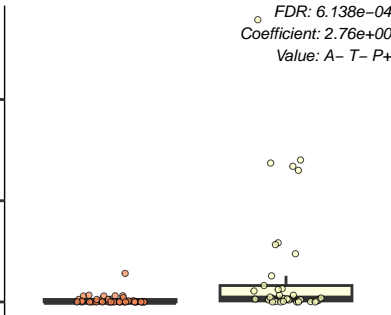
0

A-T-P- (n=36)

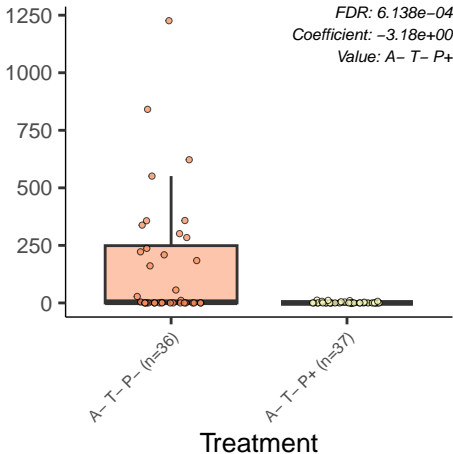
A-T-P+ (n=37)

Treatment

FDR: 6.138e-04
Coefficient: 2.76e+00
Value: A- T- P+



Agromyces



Rhodocyclaceae_Genus

FDR: 1.177e-03
Coefficient: 2.12e+00
Value: A- T- P+

400

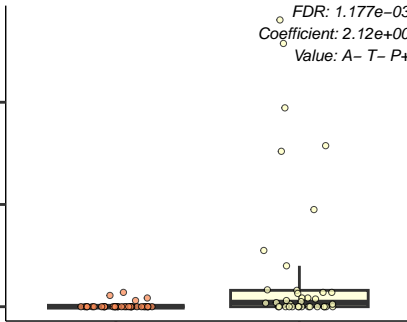
200

0

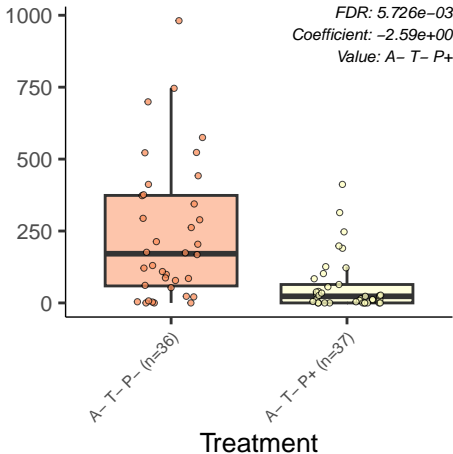
A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Tundrisphaera



Cloacibacterium

FDR: 6.709e-03

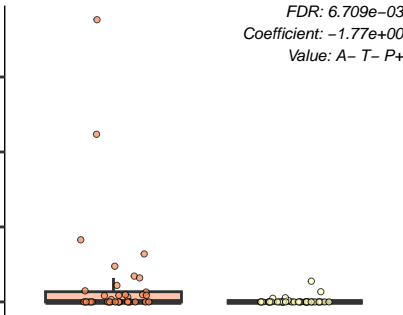
Coefficient: -1.77e+00

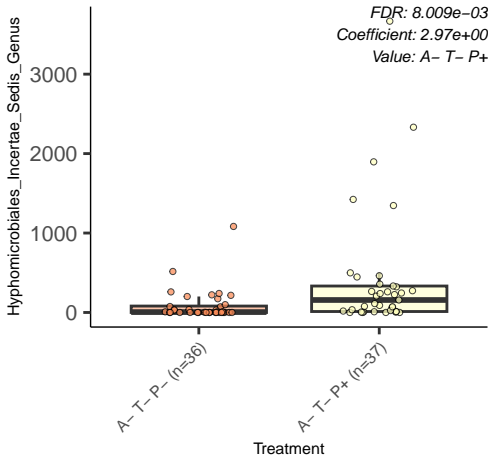
Value: A- T- P+

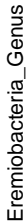
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment







FDR: 8.660e-03

Coefficient: $-1.09e+00$

Value: A- T- P+



Treatment

Chitinilyticum

FDR: 9.521e-03

Coefficient: -1.63e+00

Value: A- T- P+

4000

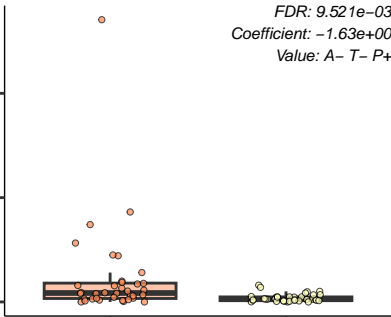
2000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Citrobacter

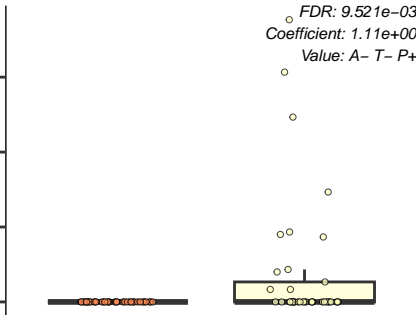
FDR: 9.521e-03
Coefficient: 1.11e+00
Value: A- T- P+

90
60
30
0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



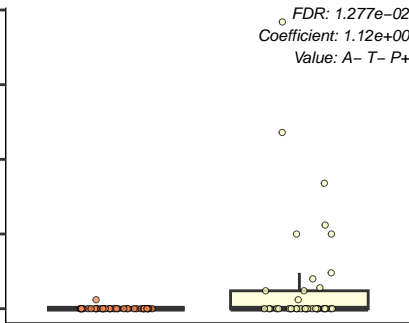
Pirellulaceae_Genus

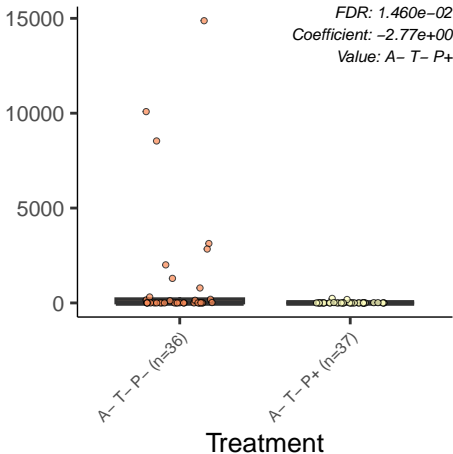
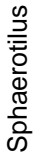
FDR: 1.277e-02
Coefficient: 1.12e+00
Value: A- T- P+

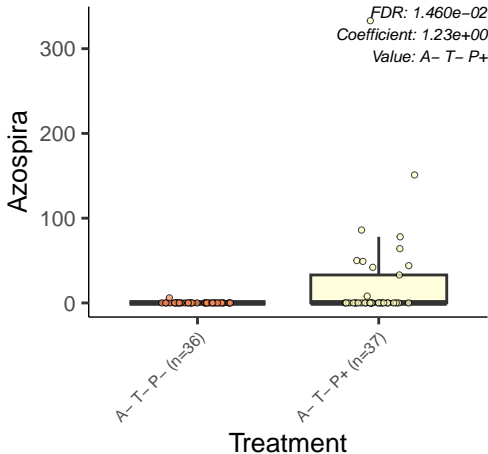
A- T- P- (n=36)

A- T- P+ (n=37)

Treatment







Rubrivivax

FDR: 1.850e-02
Coefficient: 2.13e+00
Value: A- T- P+

1000

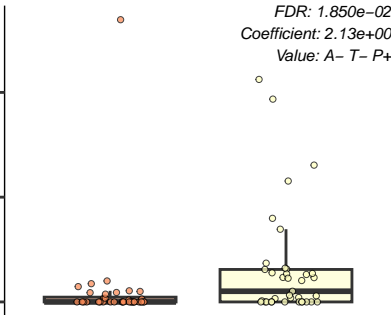
500

0

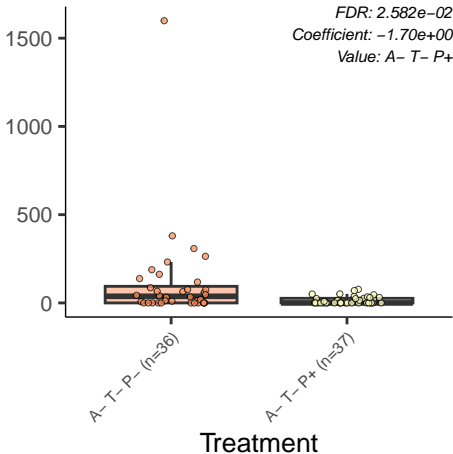
A-T-P- (n=36)

A-T-P+ (n=37)

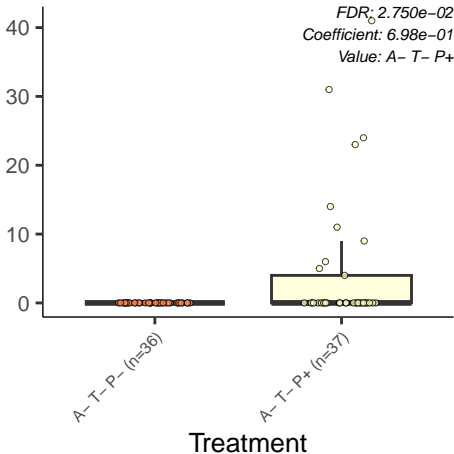
Treatment

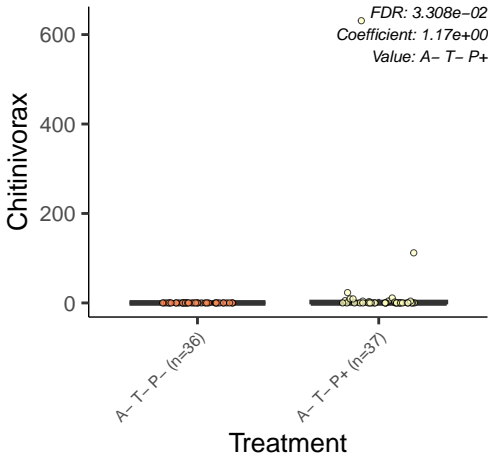


Rhizobium



Chthoniobacter





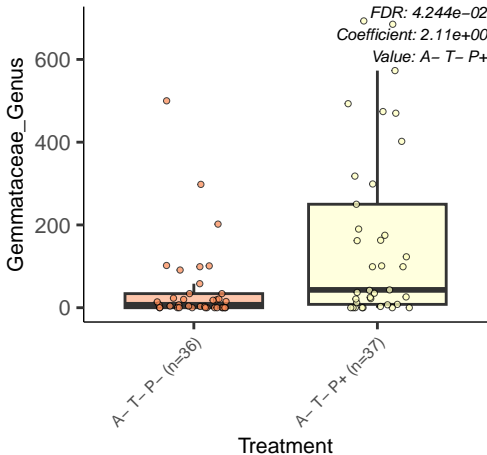
Gemmataceae_Genus

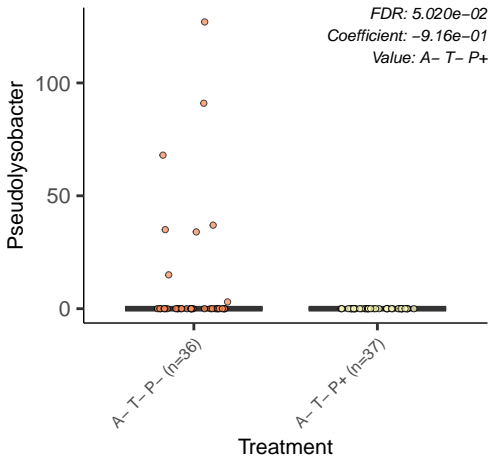
FDR: 4.244e-02
Coefficient: 2.11e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Dinghuibacter

FDR: 5.108e-02

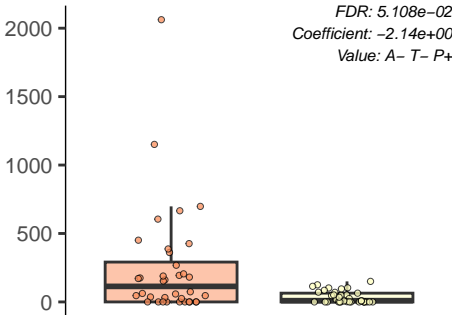
Coefficient: -2.14e+00

Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Aurantisolimonas

FDR: 5.441e-02

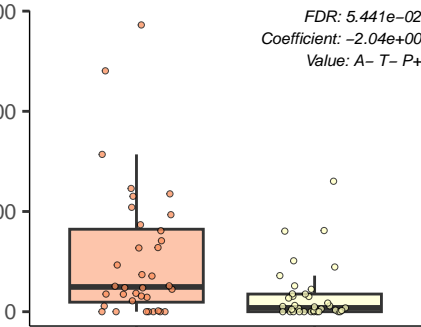
Coefficient: -2.04e+00

Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Sporomusaceae_Genus

FDR: 6.456e-02

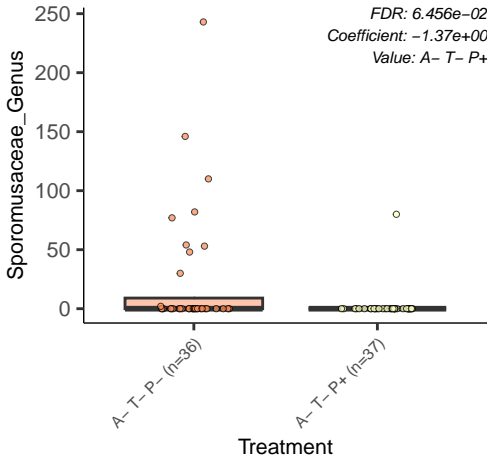
Coefficient: -1.37e+00

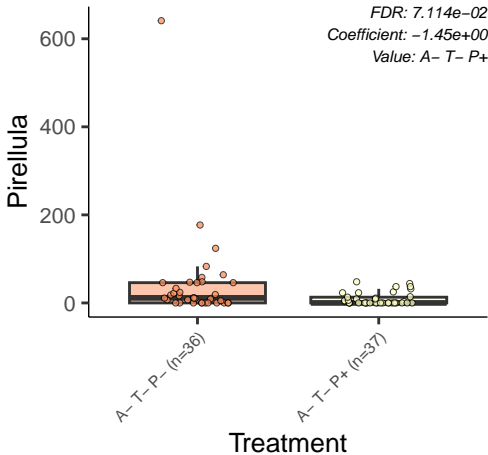
Value: A- T- P+

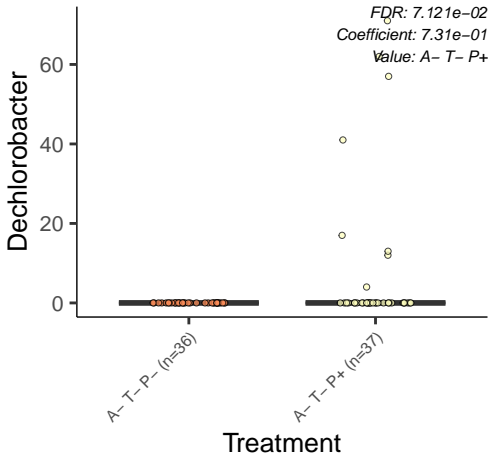
A- T- P- (n=36)

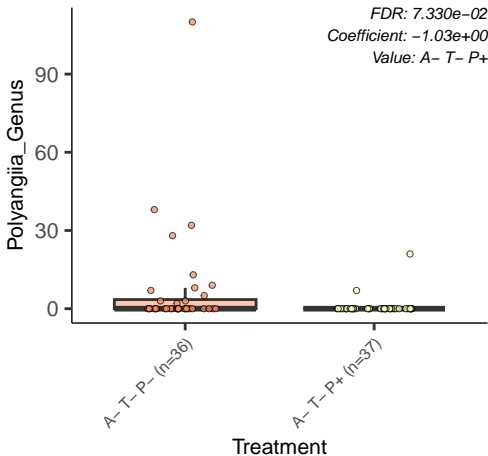
A- T- P+ (n=37)

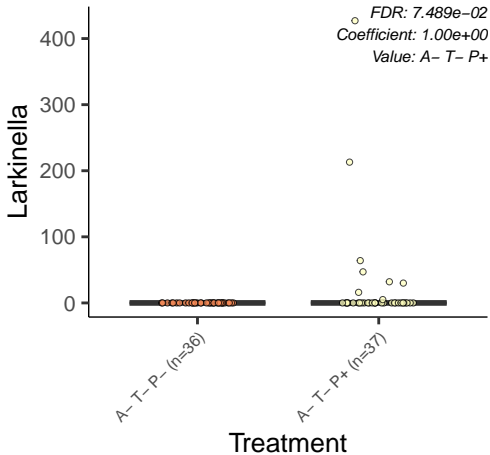
Treatment

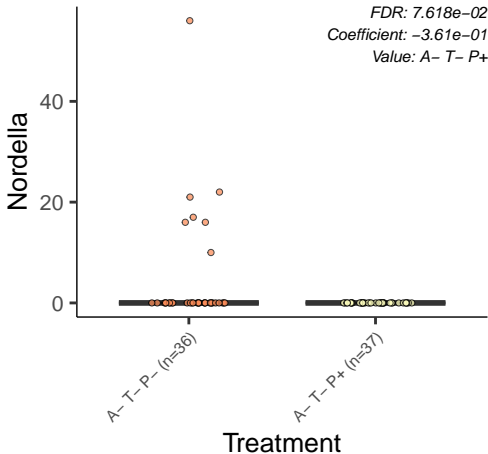














FDR: 8.266e-02

Coefficient: 1.18e+00

Value: A- T- P+



Treatment

Verrucomicrobiia_Genus

FDR: 8.703e-02

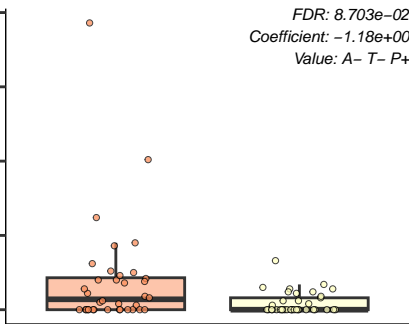
Coefficient: -1.18e+00

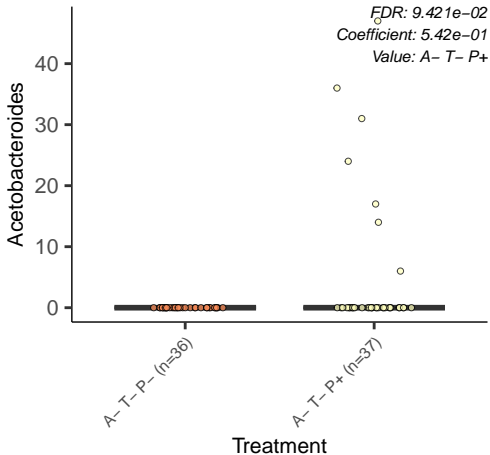
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Caulobacter

FDR: 9.536e-02

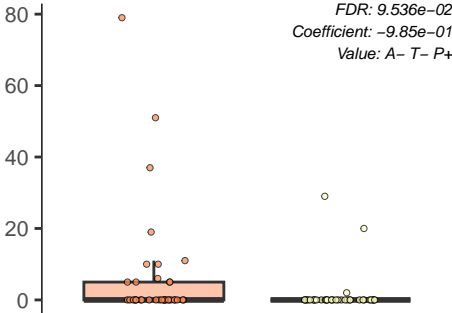
Coefficient: -9.85e-01

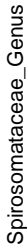
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



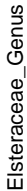


FDR: 9.536e-02
Coefficient: 7.64e-01
Value: A- T- P+

A-T-P-(n=36)

A-T-P+ (n=37)

Treatment



FDR: 9.536e-02

Coefficient: $-2.01e-01$

Value: A- T- P+



Treatment

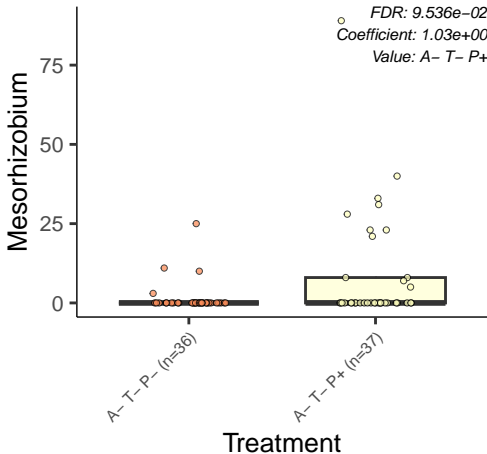
Mesorhizobium

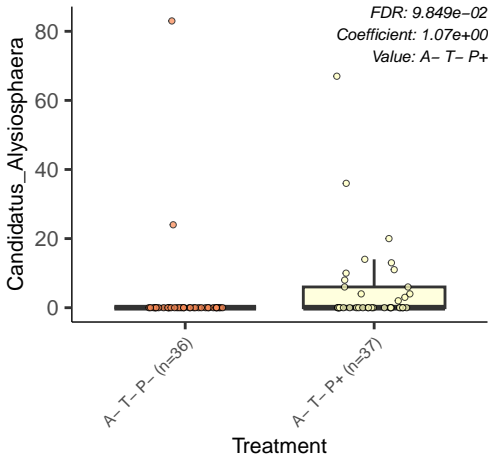
FDR: 9.536e-02
Coefficient: 1.03e+00
Value: A- T- P+

A-T-P- (n=36)

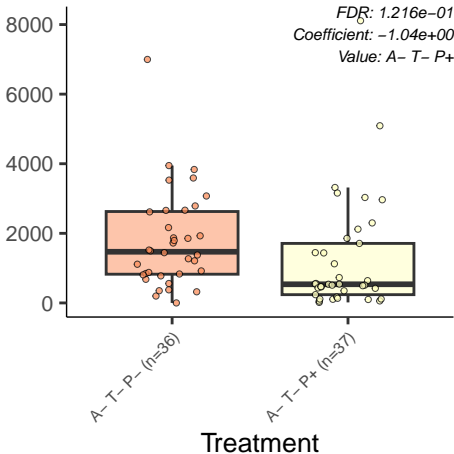
A-T-P+ (n=37)

Treatment





Crenobacter



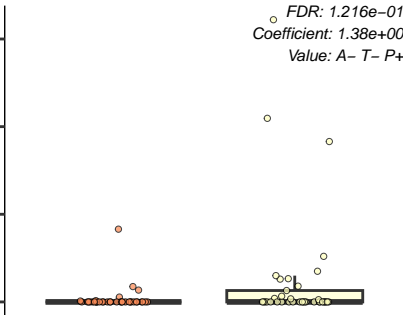
Luteolibacter

FDR: 1.216e-01
Coefficient: 1.38e+00
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Paracoccaceae_Genus

3000

2000

1000

0

A-T-P- (n=36)

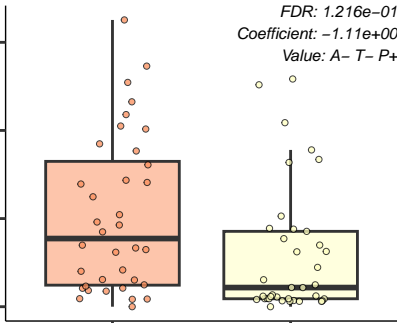
A-T-P+ (n=37)

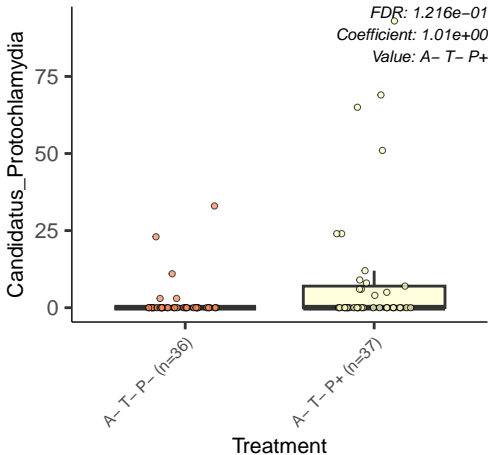
Treatment

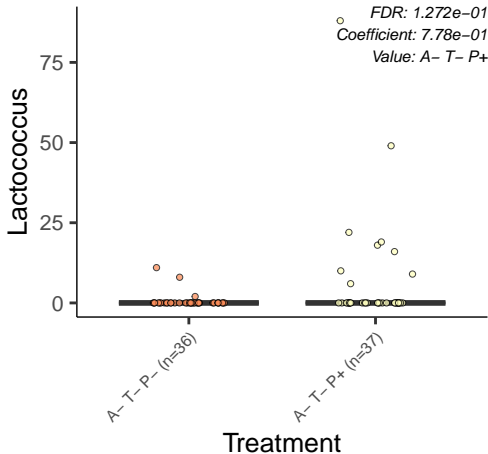
FDR: 1.216e-01

Coefficient: -1.11e+00

Value: A- T- P+







Haliangiaceae_Genus

FDR: 1.332e-01

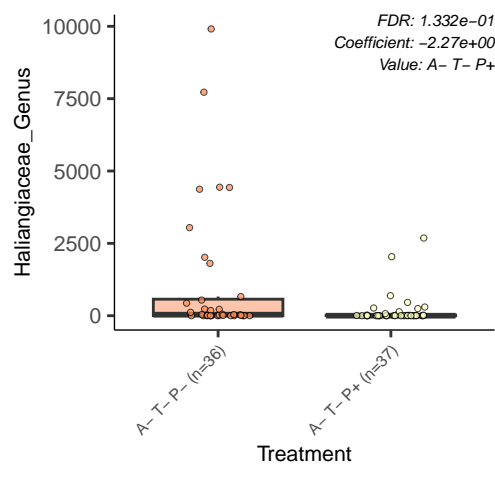
Coefficient: -2.27e+00

Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



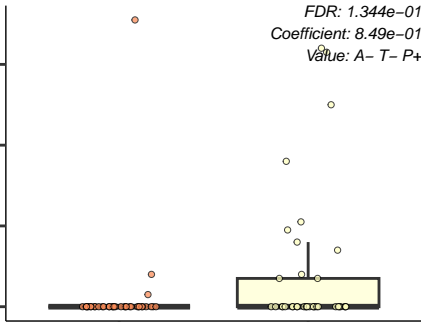
Pseudorhodoplanes

FDR: 1.344e-01
Coefficient: 8.49e-01
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



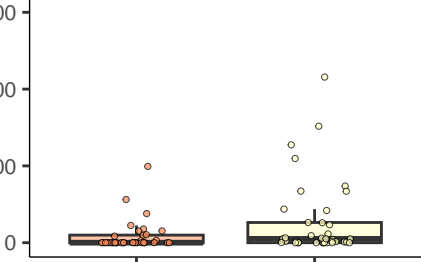
env.OPS_17_Genus

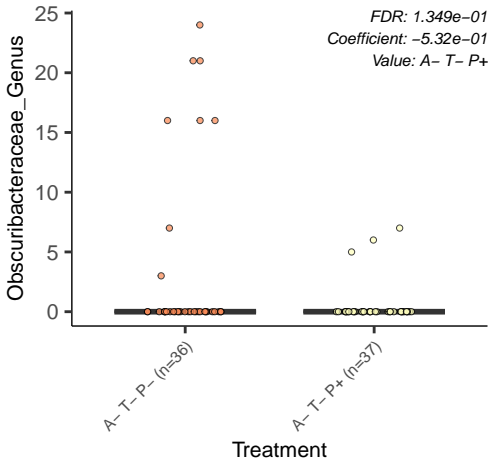
FDR: 1.349e-01
Coefficient: 2.30e+00
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment





Candidatus_Berkiella

FDR: 1.410e-01
Coefficient: 9.34e-01
Value: A- T- P+

100

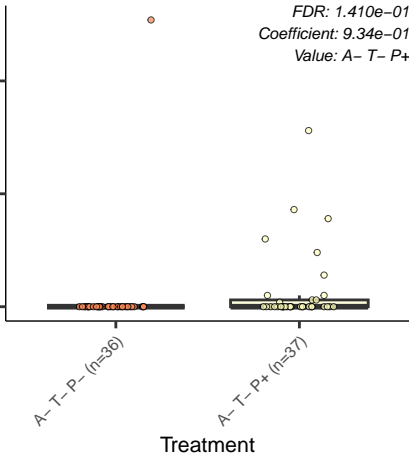
50

0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Phreatobacter

FDR: 1.562e-01

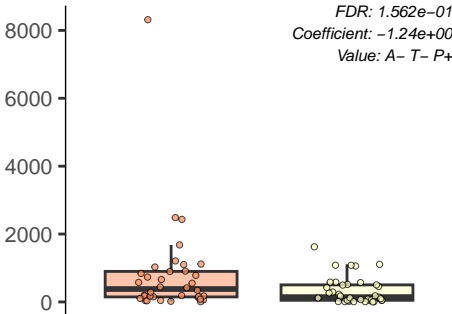
Coefficient: -1.24e+00

Value: A- T- P+

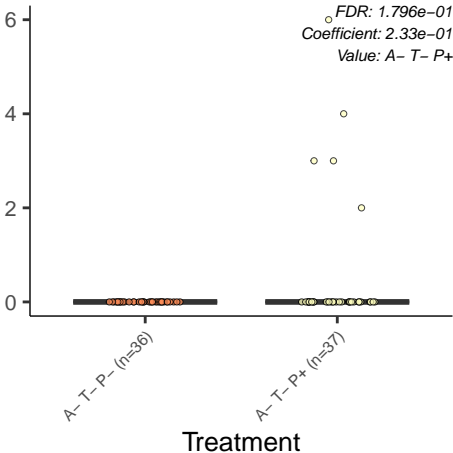
A-T-P- (n=36)

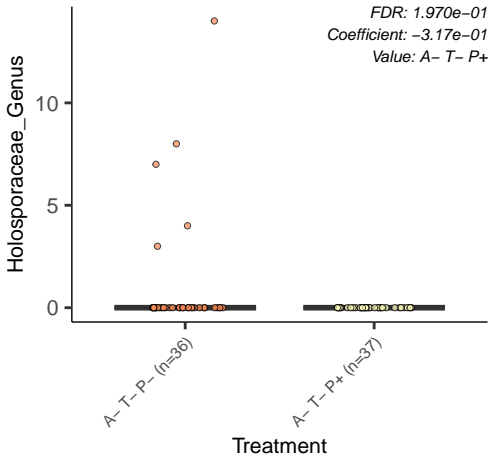
A-T-P+ (n=37)

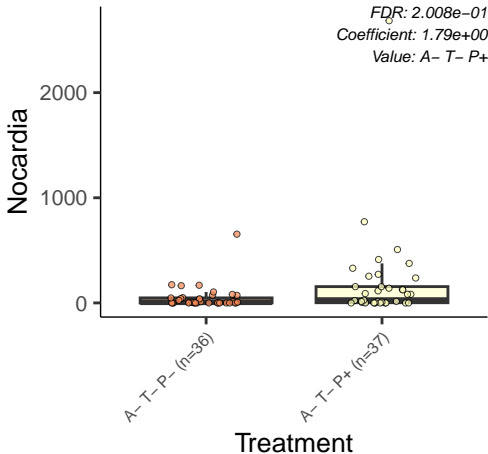
Treatment

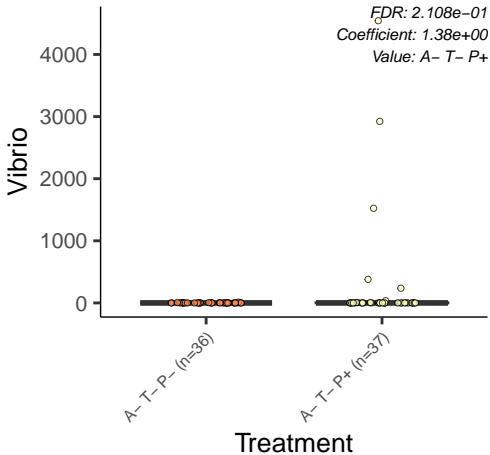


Novispirillum









Rhodococcus

1000

500

0

A-T-P- (n=36)

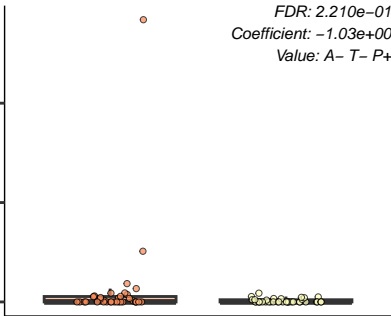
A-T-P+ (n=37)

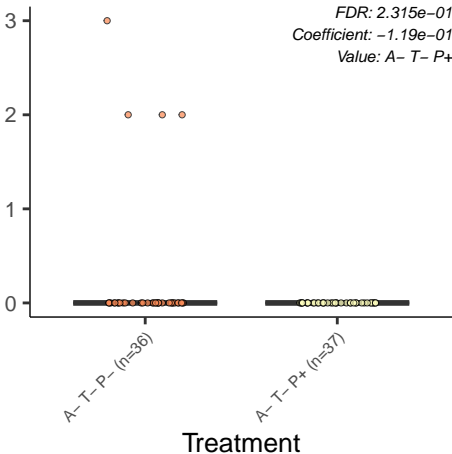
Treatment

FDR: 2.210e-01

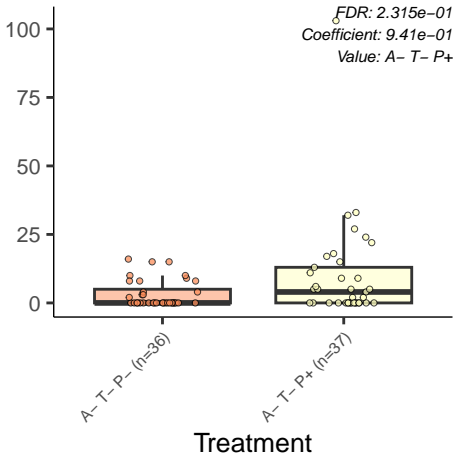
Coefficient: -1.03e+00

Value: A- T- P+

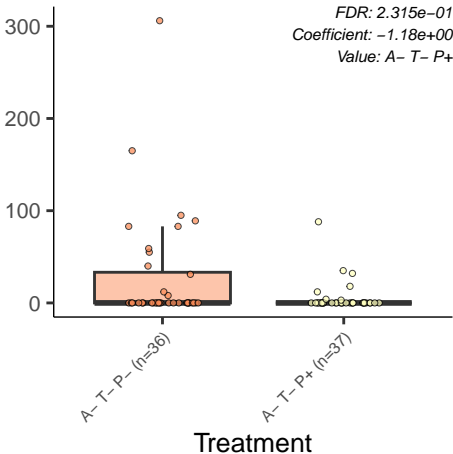




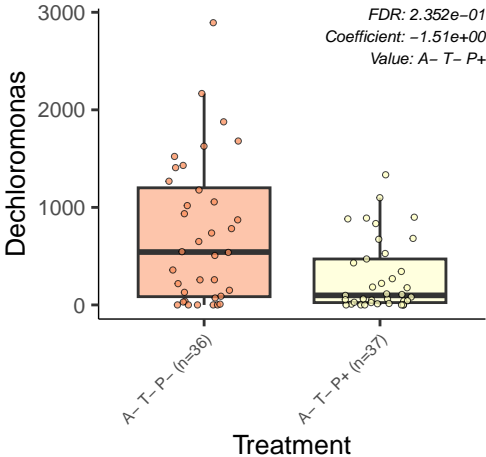
Roseicella

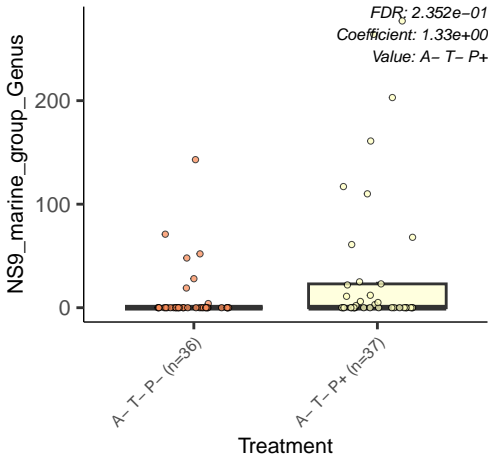


Deinococcus



Dechloromonas





Value: A- T- P+

