

Plesiomonas

FDR: 1.691e-09

Coefficient: -2.69e+00

Value: A- T- P+

2000

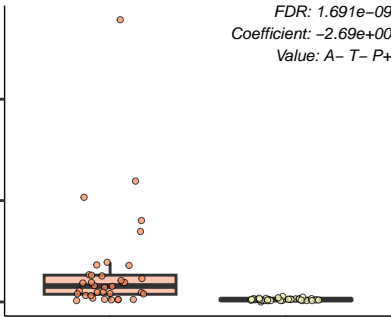
1000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



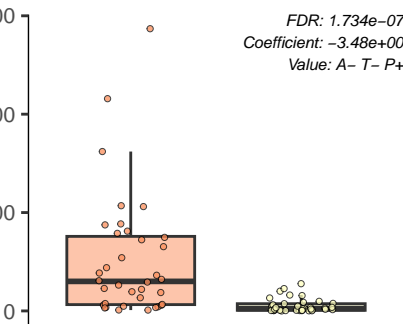
Culicoidibacter

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

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



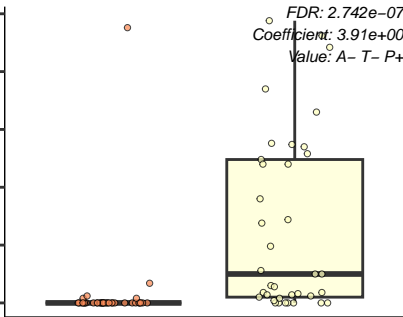
X67.14_Genus

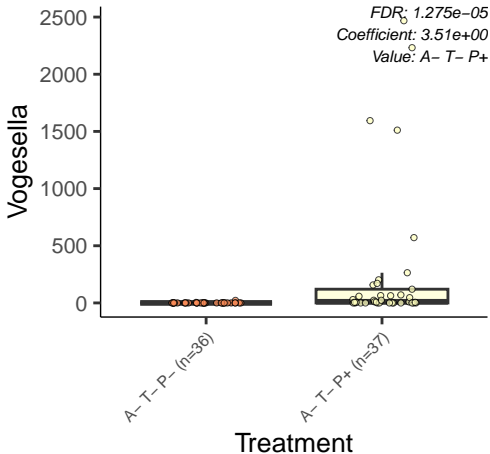
FDR: 2.742e-07
Coefficient: 3.91e+00
Value: A- T- P+

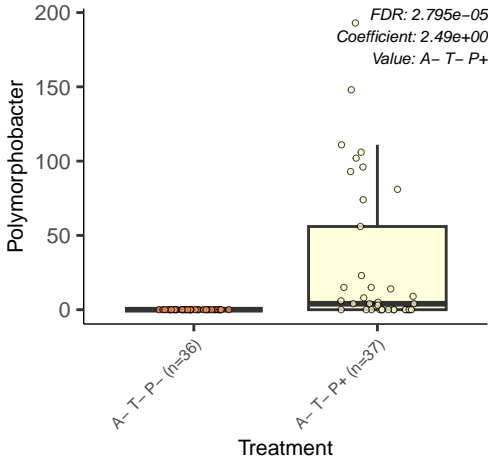
A-T-P- (n=36)

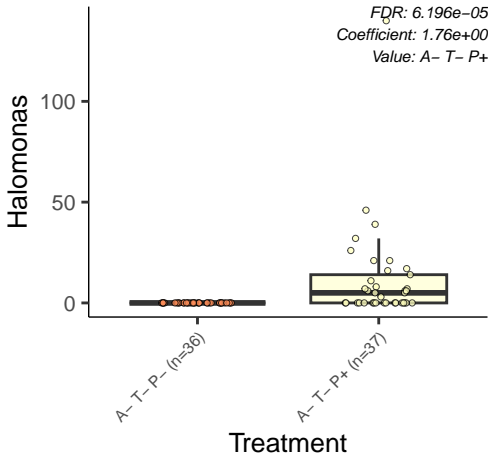
A-T-P+ (n=37)

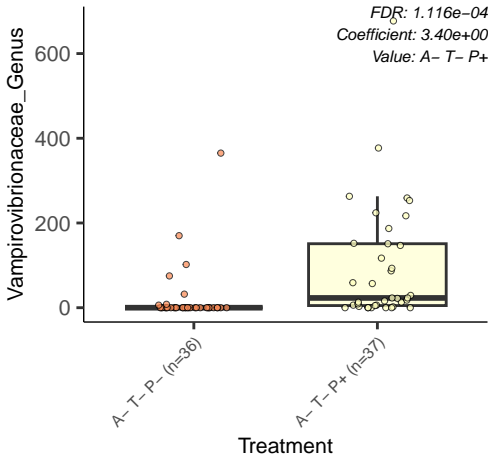
Treatment











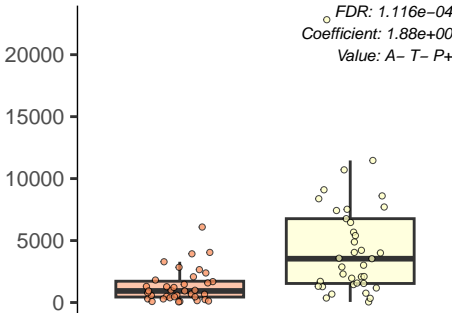
Pseudomonas

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

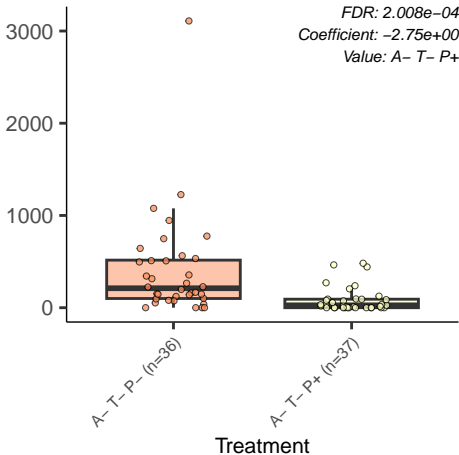
A-T-P- (n=36)

A-T-P+ (n=37)

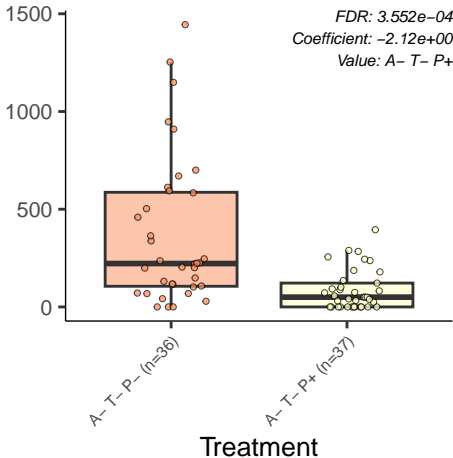
Treatment



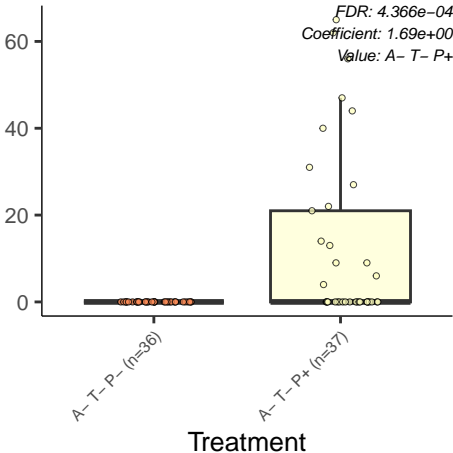
Paenirhodobacter



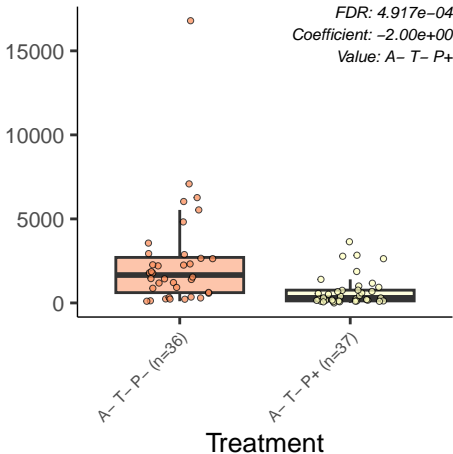
Gemmobacter



Rhodovarius



Paucibacter



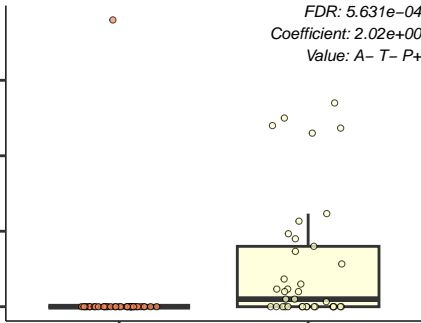
Candidatus_Megaira

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

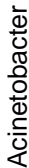
A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



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



1000

500

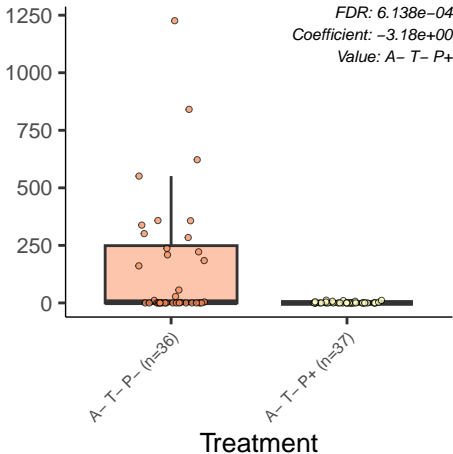
0

A-T-P-(n=36)

A-T-P+ (n=37)

Treatment

Agromyces



Rhodocyclaceae_Genus

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

400

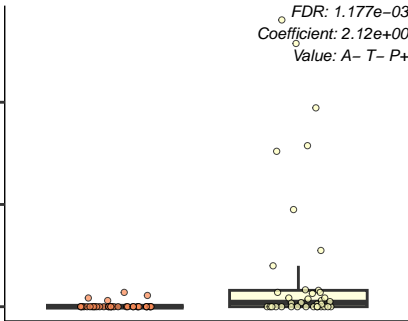
200

0

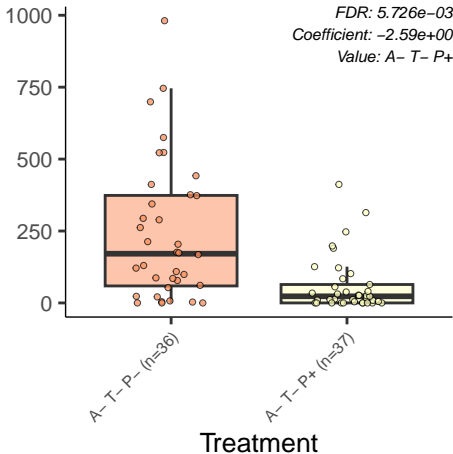
A- T- P- (n=36)

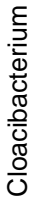
A- T- P+ (n=37)

Treatment



Tundrisphaera





FDR: 6.709e-03

Coefficient: $-1.77e+00$

Value: A- T- P+



Treatment

Hyphomicrobiales_Incertae_Sedis_Genus

FDR: 8.009e-03
Coefficient: 2.97e+00
Value: A- T- P+

3000

2000

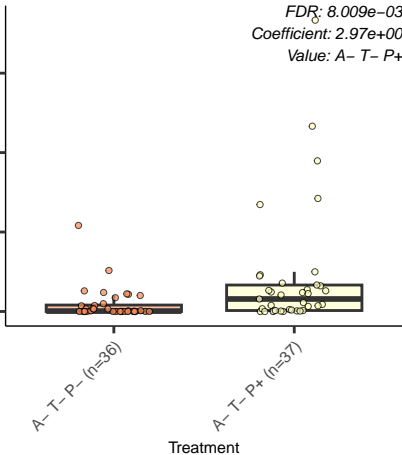
1000

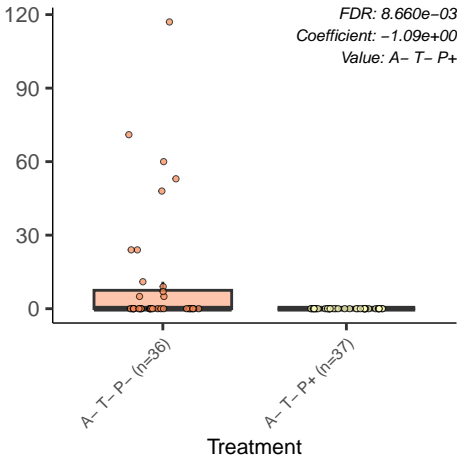
0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Chitinilyticum

FDR: 9.521e-03

Coefficient: -1.63e+00

Value: A- T- P+

4000

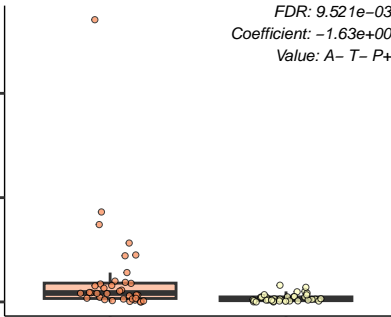
2000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Citrobacter

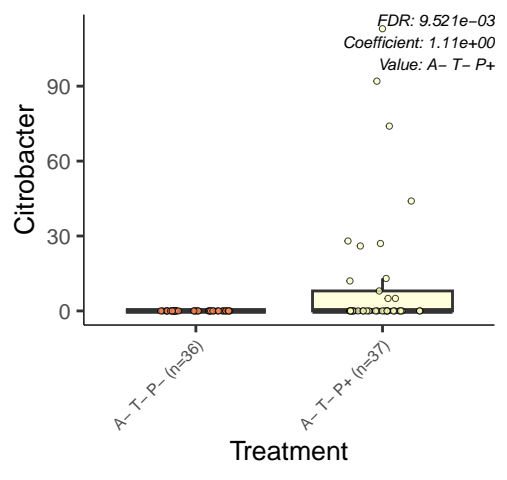
90
60
30
0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

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



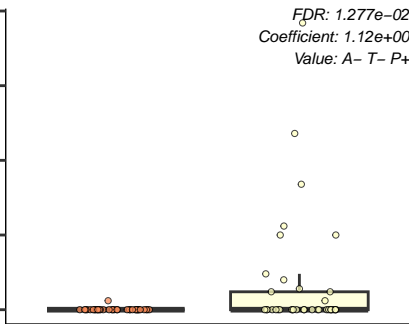
Pirellulaceae_Genus

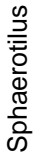
FDR: 1.277e-02
Coefficient: 1.12e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





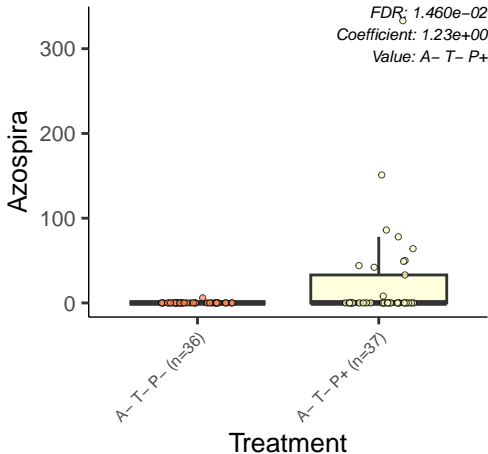
FDR: 1.460e-02

Coefficient: $-2.77e+00$

Value: A- T- P+



Treatment



Rubrivivax

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

1000

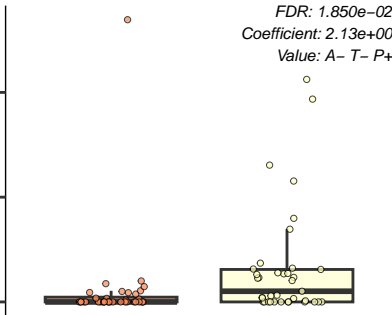
500

0

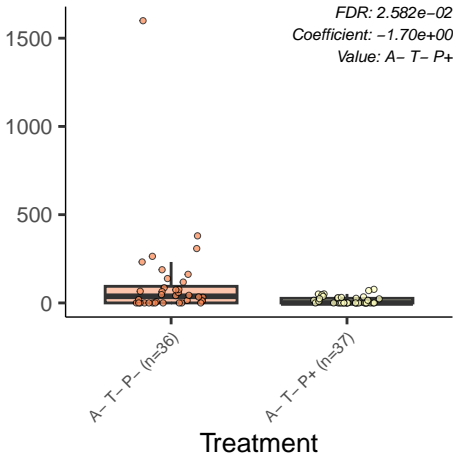
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Rhizobium





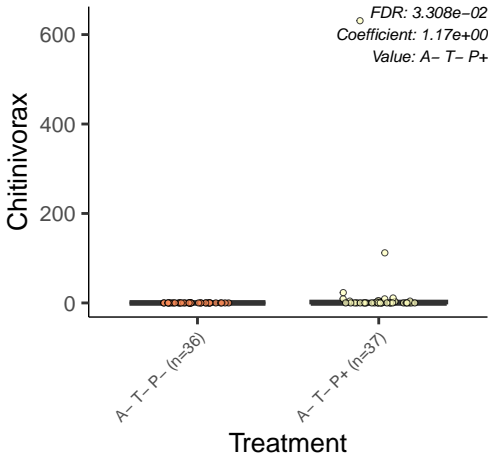
FDR: 2.750e-02

Coefficient: 6.98e-01

Value: A- T- P+



Treatment



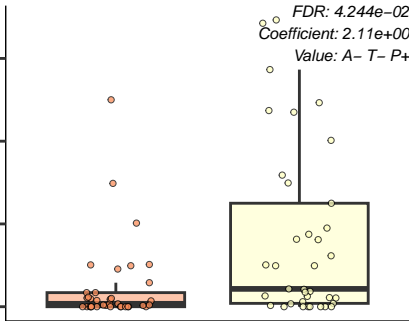
Gemmataceae_Genus

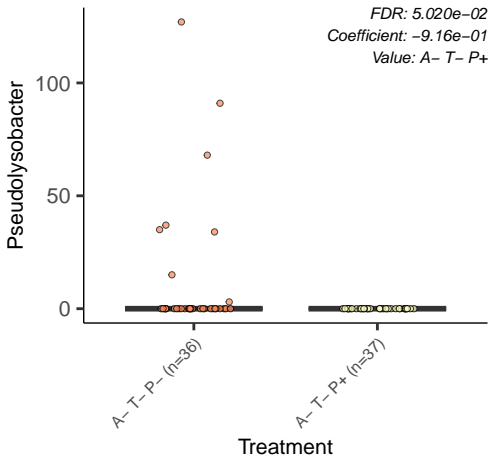
FDR: 4.244e-02
Coefficient: 2.11e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Dinghuibacter

FDR: 5.108e-02

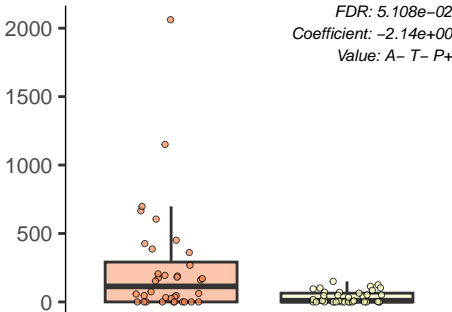
Coefficient: -2.14e+00

Value: A- T- P+

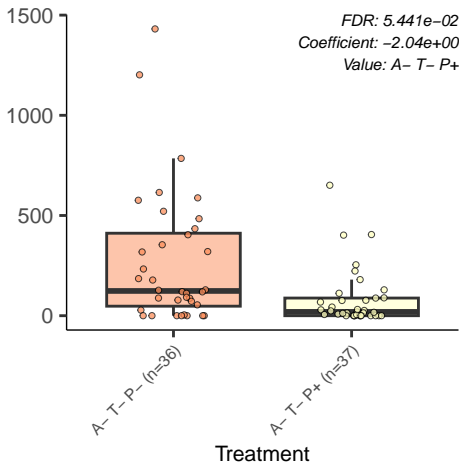
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Aurantisolimonas



Sporomusaceae_Genus

FDR: 6.456e-02

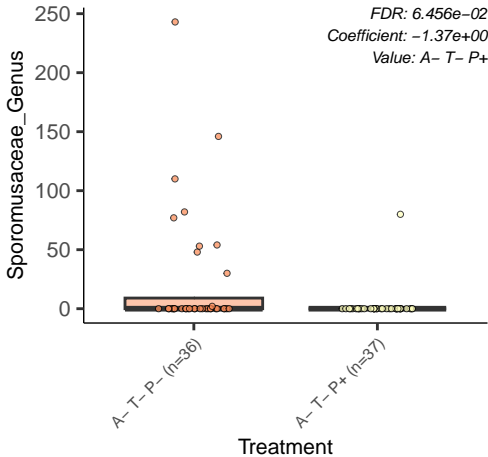
Coefficient: -1.37e+00

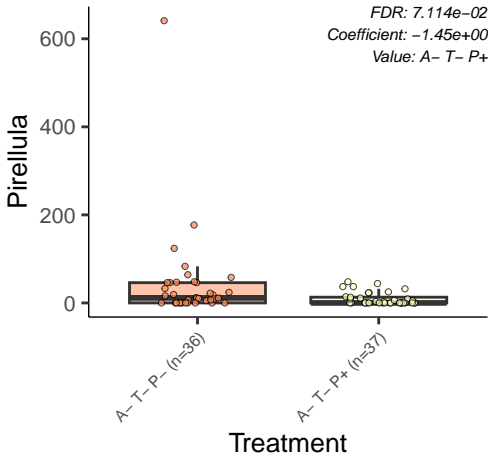
Value: A- T- P+

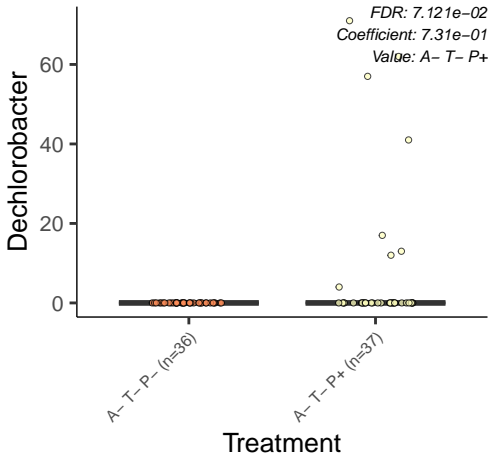
A- T- P- (n=36)

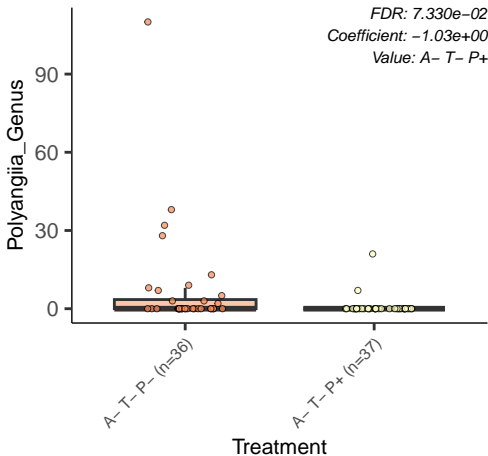
A- T- P+ (n=37)

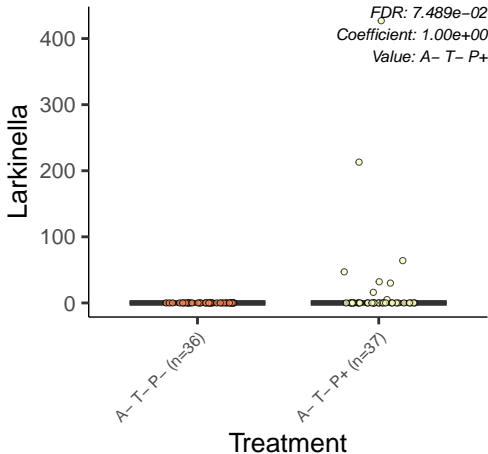
Treatment

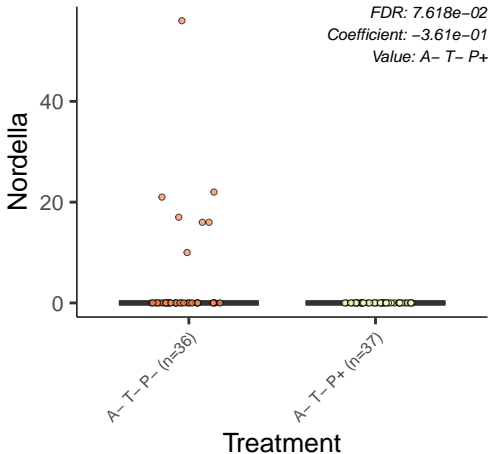














FDR: 8.266e-02

Coefficient: 1.18e+00

Value: A- T- P+



Treatment

Verrucomicrobiia_Genus

FDR: 8.703e-02

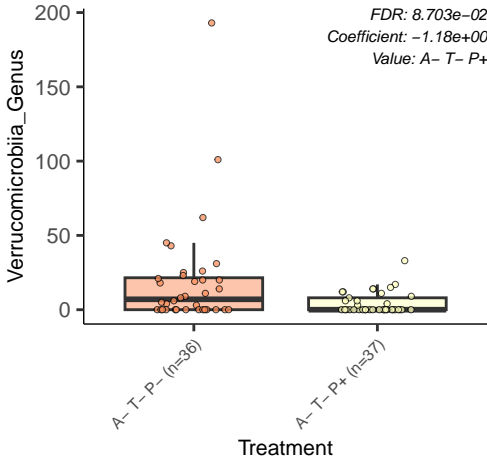
Coefficient: -1.18e+00

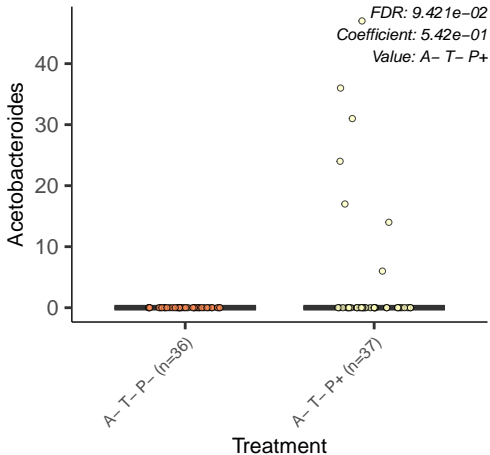
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Caulobacter

FDR: 9.536e-02

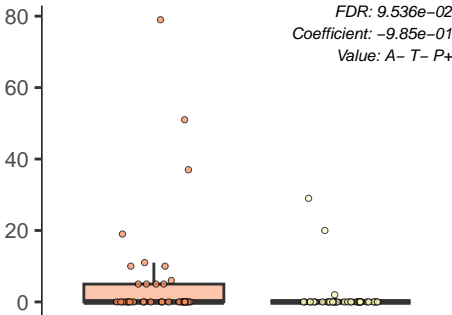
Coefficient: -9.85e-01

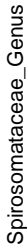
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment





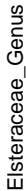
FDR: 9.536e-02

Coefficient: 7.64e-01

Value: A- T- P+



Treatment



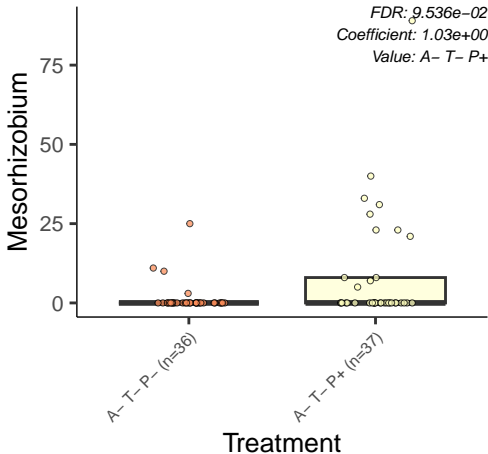
FDR: 9.536e-02

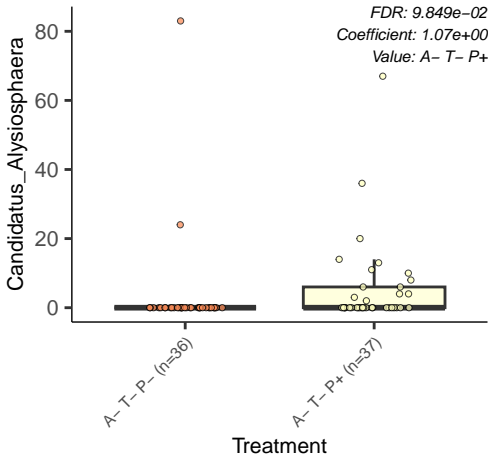
Coefficient: $-2.01e-01$

Value: A- T- P+

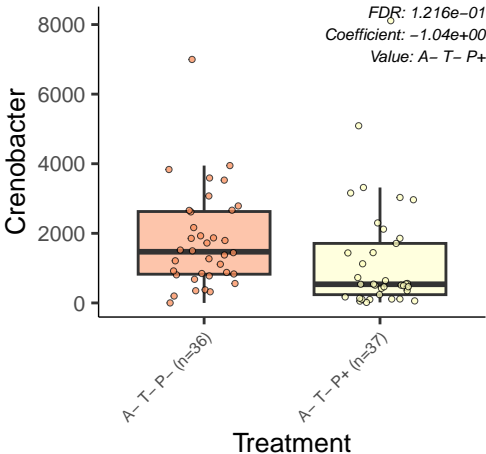


Treatment

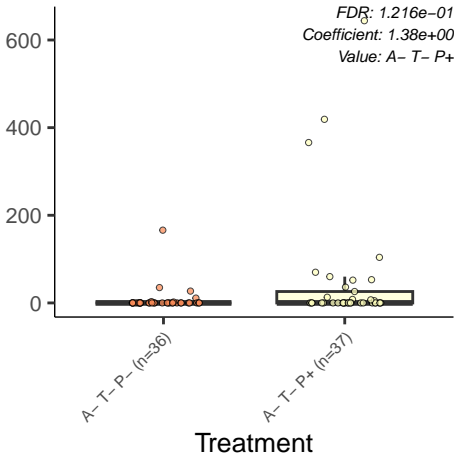




Crenobacter



Luteolibacter



Paracoccaceae_Genus

3000

2000

1000

0

A-T-P- (n=36)

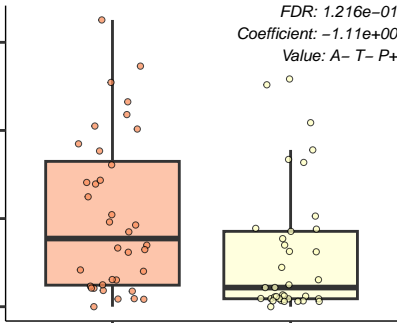
A-T-P+ (n=37)

Treatment

FDR: 1.216e-01

Coefficient: -1.11e+00

Value: A- T- P+



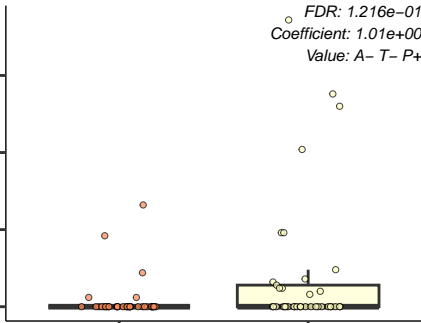
Candidatus_Protochlamydia

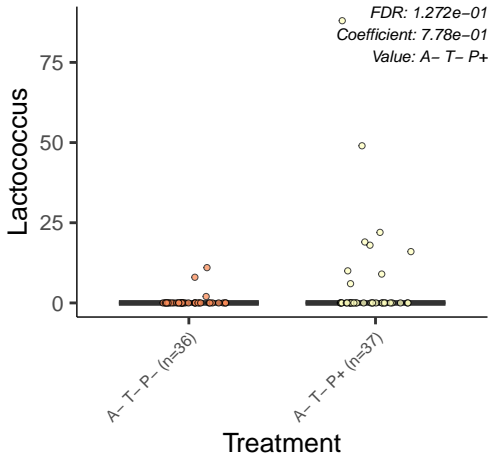
FDR: 1.216e-01
Coefficient: 1.01e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





Haliangiaceae_Genus

FDR: 1.332e-01

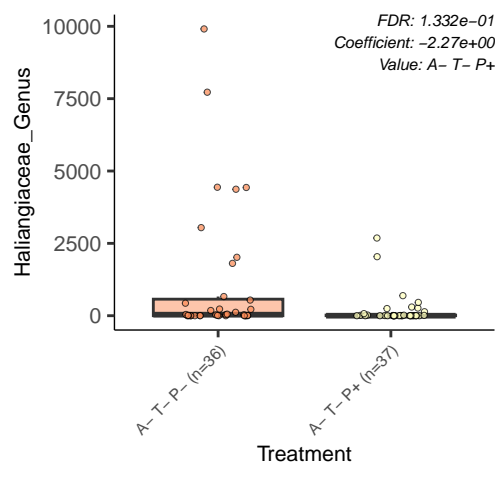
Coefficient: -2.27e+00

Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Pseudorhodoplanes

FDR: 1.344e-01

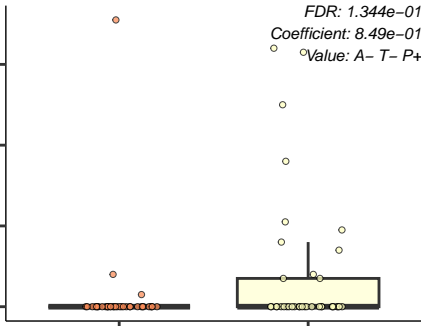
Coefficient: 8.49e-01

Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



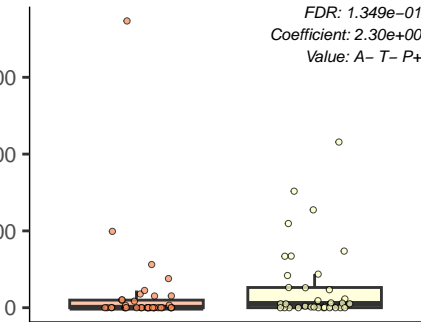
env.OPS_17_Genus

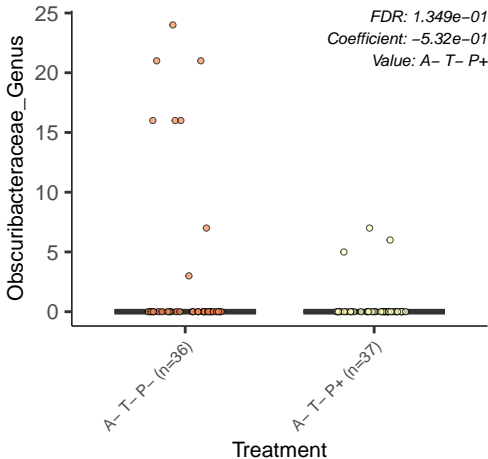
FDR: 1.349e-01
Coefficient: 2.30e+00
Value: A- T- P+

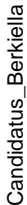
A- T- P- (n=36)

A- T- P+ (n=37)

Treatment







FDR: 1.410e-01

Coefficient: 9.34e-01

Value: A- T- P+

A-T-P-(n=36)

A-T-P+ (n=37)

Treatment

Phreatobacter

FDR: 1.562e-01

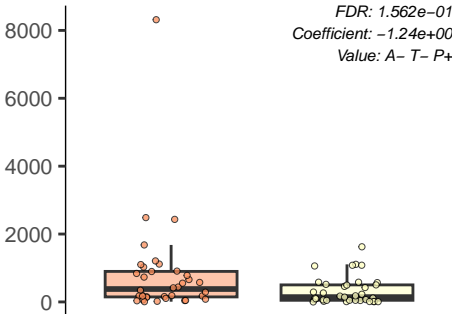
Coefficient: -1.24e+00

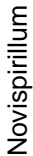
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



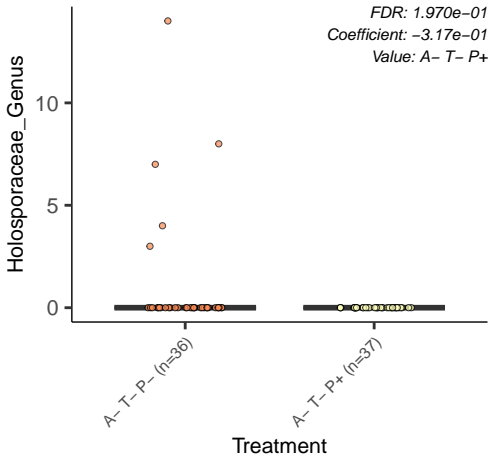


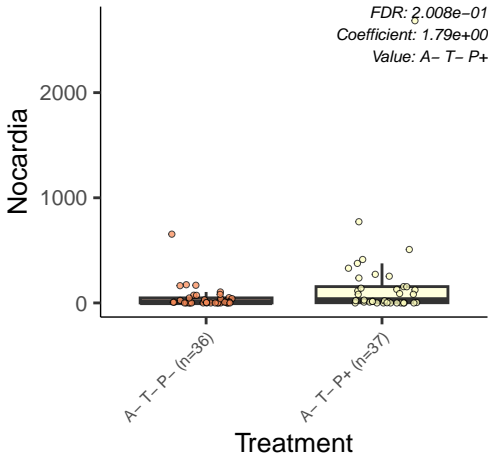
FDR: 1.796e-01
Coefficient: 2.33e-01
Value: A- T- P+

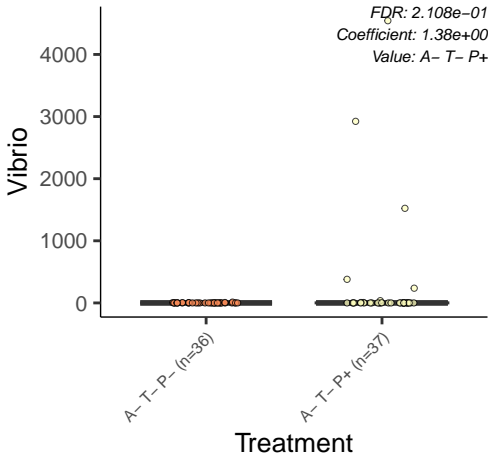
A-T-P-(n=36)

A-T-P+ (n=37)

Treatment







Rhodococcus

1000

500

0

A-T-P- (n=36)

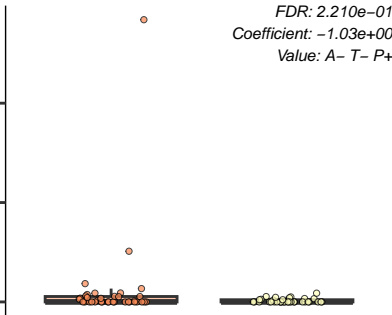
A-T-P+ (n=37)

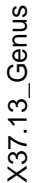
Treatment

FDR: 2.210e-01

Coefficient: -1.03e+00

Value: A- T- P+





FDR: 2.315e-01

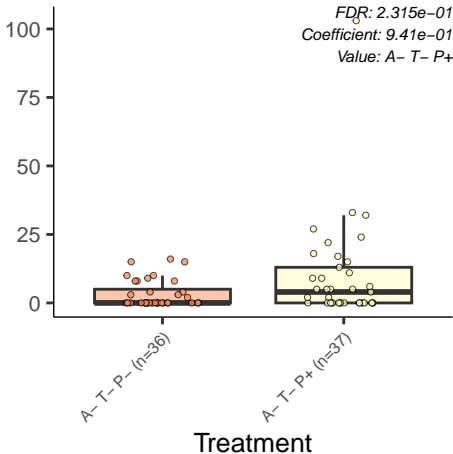
Coefficient: $-1.19e-01$

Value: A- T- P+



Treatment

Roseicella





FDR: 2.315e-01

Coefficient: $-1.18e+00$

Value: A- T- P+



Treatment

Dechloromonas

