

Plesiomonas

FDR: 1.365e-09

Coefficient: -2.69e+00

Value: A- T- P+

2000

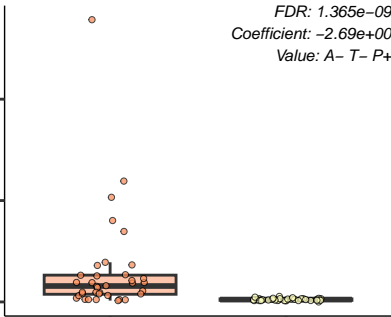
1000

0

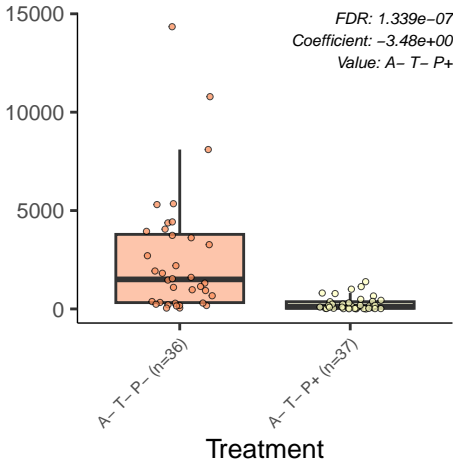
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Culicoidibacter



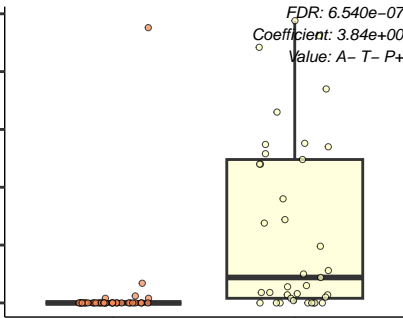
X67.14_Genus

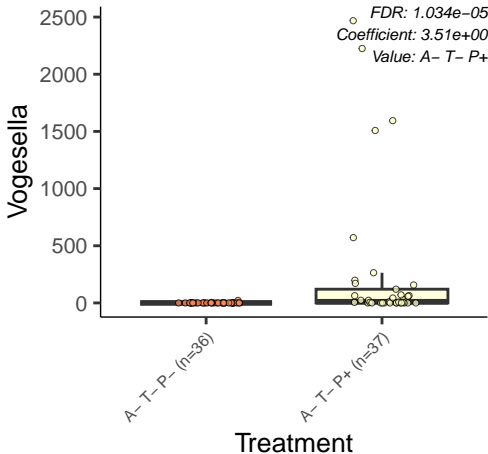
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

FDR: 6.540e-07
Coefficient: 3.84e+00
Value: A- T- P+





Polymorphobacter

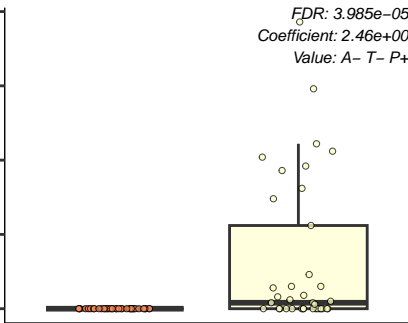
200
150
100
50
0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

FDR: 3.985e-05
Coefficient: 2.46e+00
Value: A- T- P+



Rhodocyclaceae_Genus

FDR: 1.010e-04
Coefficient: 2.38e+00
○ Value: A- T- P+

400

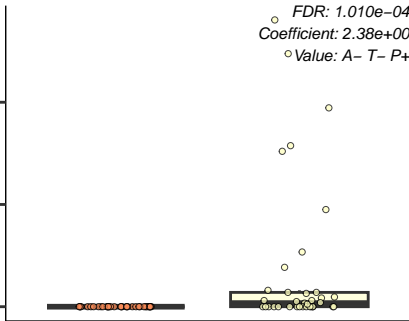
200

0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



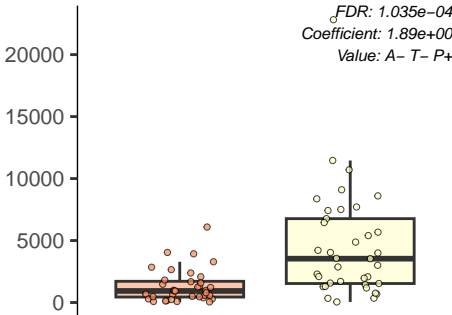
Pseudomonas

FDR: $1.035e-04$
Coefficient: $1.89e+00$
Value: A- T- P+

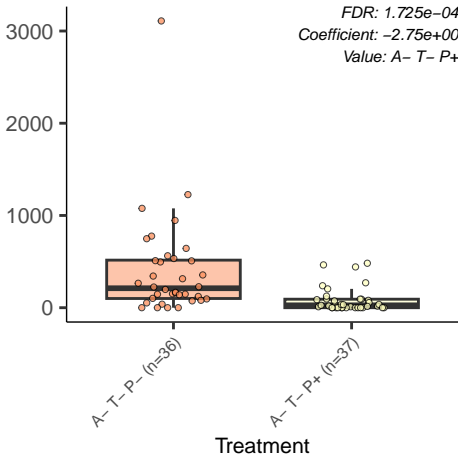
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Paenirhodobacter



Candidatus_Megaira

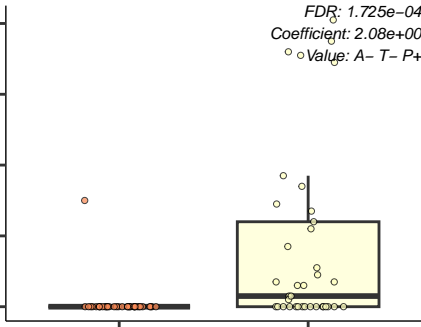
80
60
40
20
0

FDR: 1.725e-04
Coefficient: 2.08e+00
Value: A- T- P+

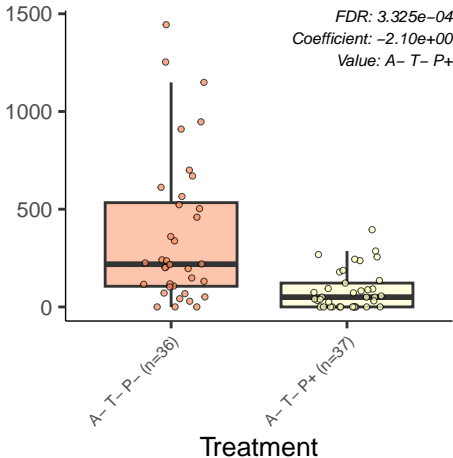
A- T- P- (n=36)

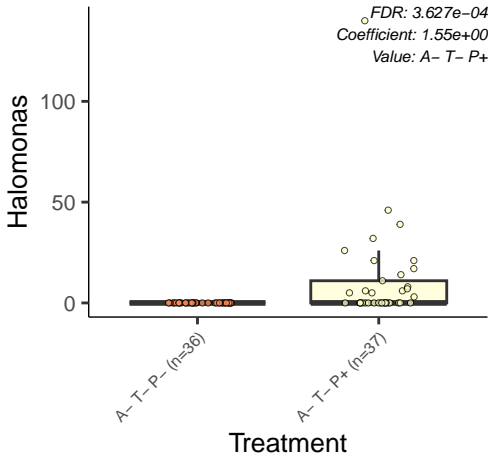
A- T- P+ (n=37)

Treatment



Gemmobacter





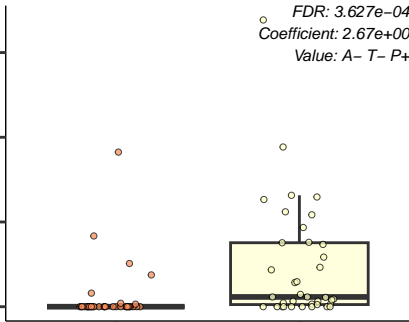
Vampirovibrionaceae_Genus

FDR: 3.627e-04
Coefficient: 2.67e+00
Value: A- T- P+

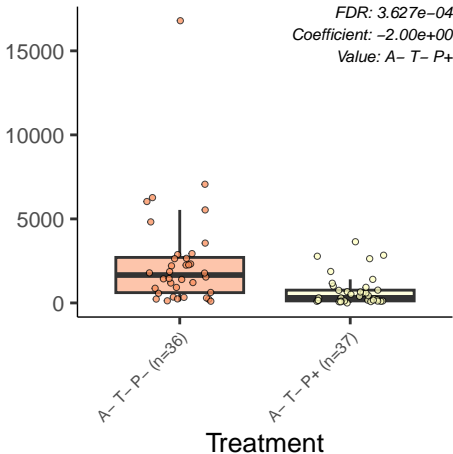
A- T- P- (n=36)

A- T- P+ (n=37)

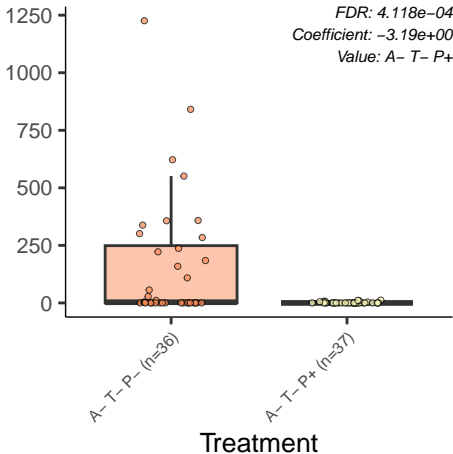
Treatment



Paucibacter



Agromyces



Acinetobacter

1000

500

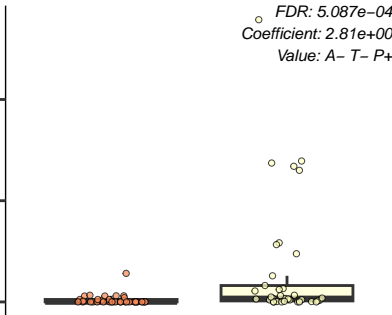
0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

FDR: 5.087e-04
Coefficient: 2.81e+00
Value: A- T- P+





FDR: 7.456e-04

Coefficient: 1.24e+00

Value: A- T- P+

A-T-P-(n=36)

A-T-P+ (n=37)

Treatment

Cloacibacterium

FDR: 5.678e-03

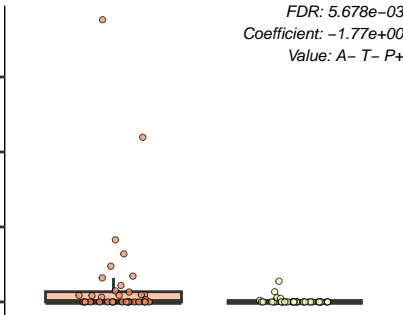
Coefficient: -1.77e+00

Value: A- T- P+

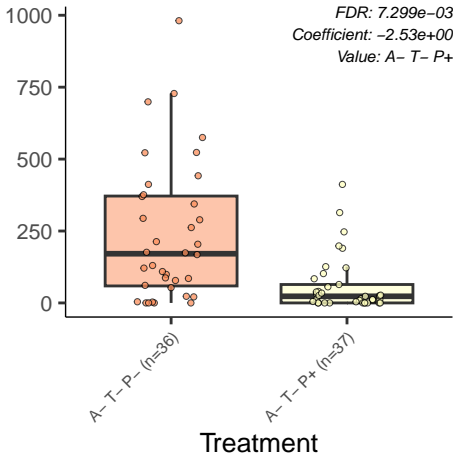
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Tundrisphaera



Chitinilyticum

FDR: 8.698e-03

Coefficient: -1.63e+00

Value: A- T- P+

4000

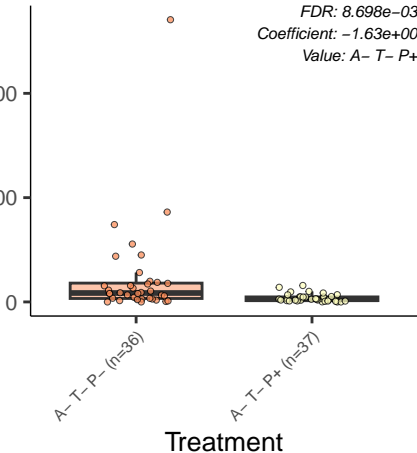
2000

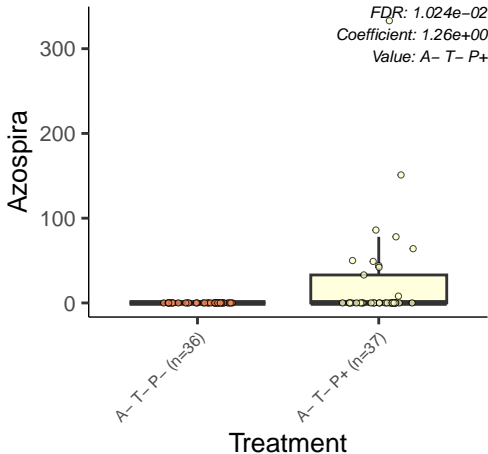
0

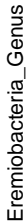
A-T-P- (n=36)

A-T-P+ (n=37)

Treatment







FDR: 1.024e-02

Coefficient: $-1.05e+00$

Value: A- T- P+



Treatment

Hyphomicrobiales_Incertae_Sedis_Genus

FDR: 1.051e-02
Coefficient: 2.92e+00
Value: A- T- P+

3000

2000

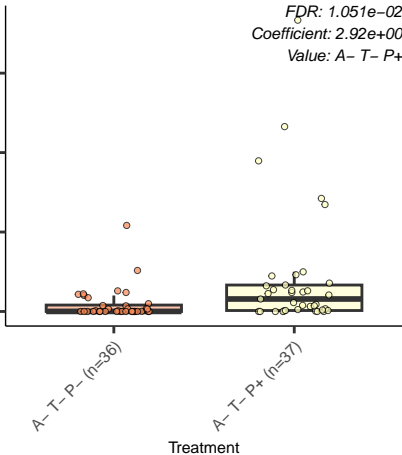
1000

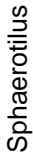
0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





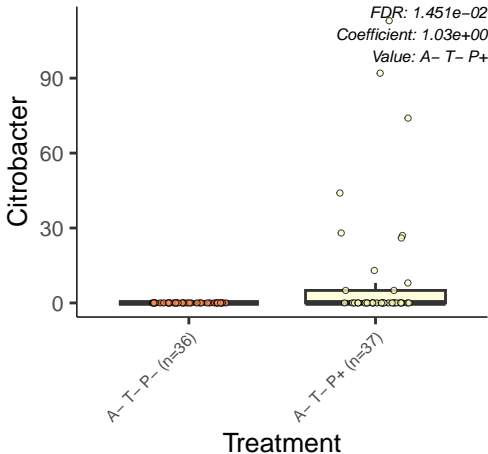
FDR: 1.275e-02

Coefficient: $-2.77e+00$

Value: A- T- P+



Treatment



Rubrivivax

FDR: 1.549e-02
Coefficient: 2.13e+00
Value: A- T- P+

1000

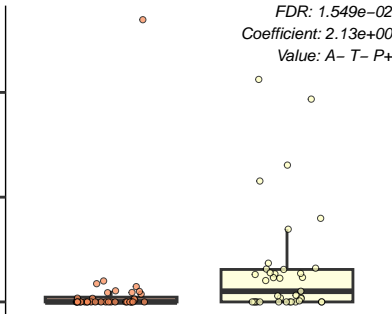
500

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Pirellulaceae_Genus

100
75
50
25
0

FDR: 1.594e-02
Coefficient: 8.73e-01
Value: A- T- P+

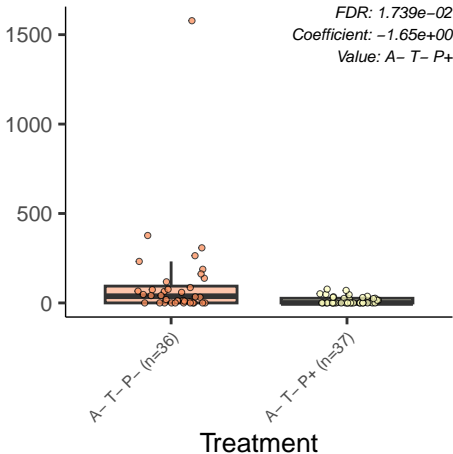
A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Rhizobium





FDR: 2.202e-02

Coefficient: 6.98e-01

Value: A- T- P+



Treatment

Dinghuibacter

FDR: 3.899e-02

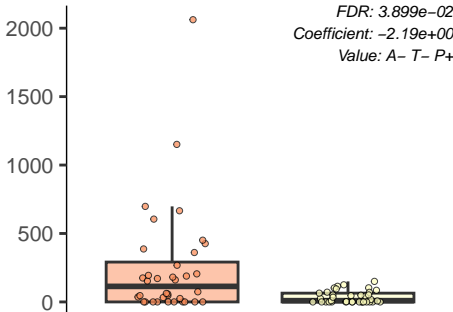
Coefficient: -2.19e+00

Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



Aurantisolimonas

1500

1000

500

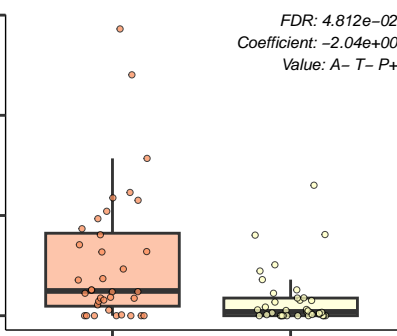
0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment

FDR: 4.812e-02
Coefficient: -2.04e+00
Value: A- T- P+



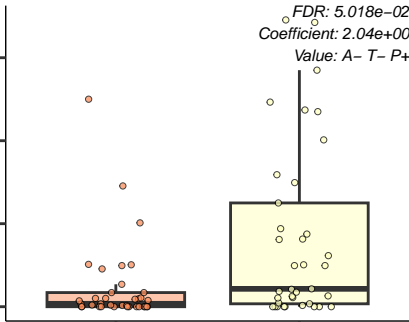
Gemmataceae_Genus

FDR: 5.018e-02
Coefficient: 2.04e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment



Sporomusaceae_Genus

FDR: 5.517e-02

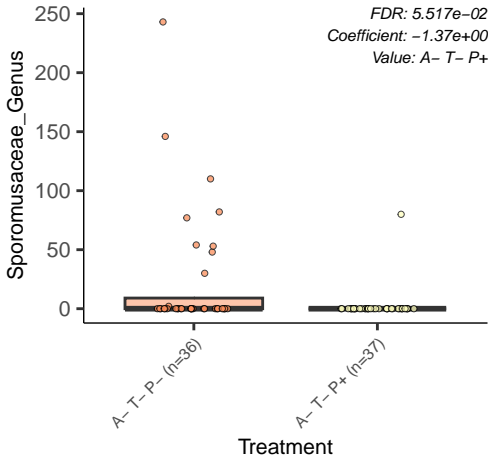
Coefficient: -1.37e+00

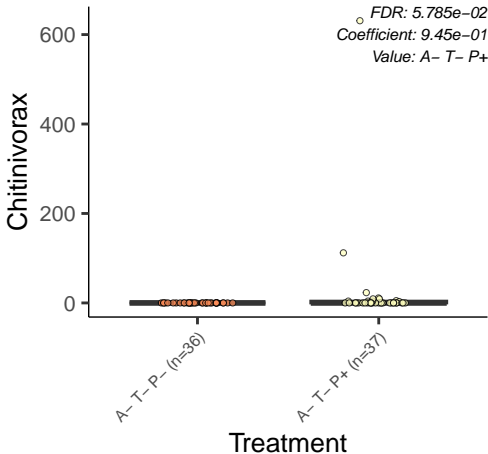
Value: A- T- P+

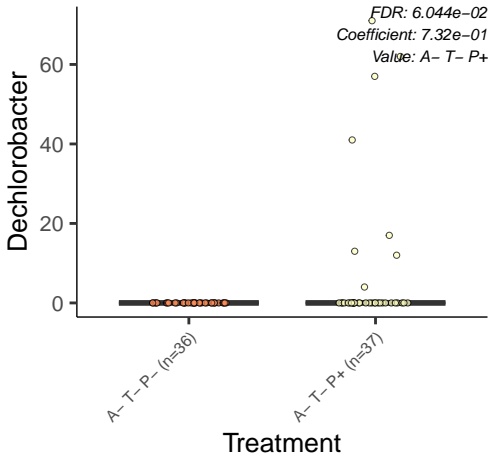
A- T- P- (n=36)

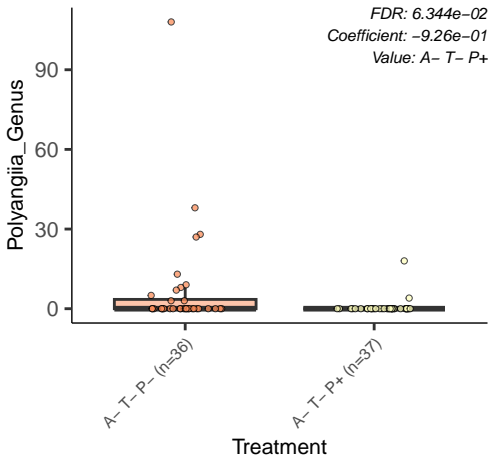
A- T- P+ (n=37)

Treatment











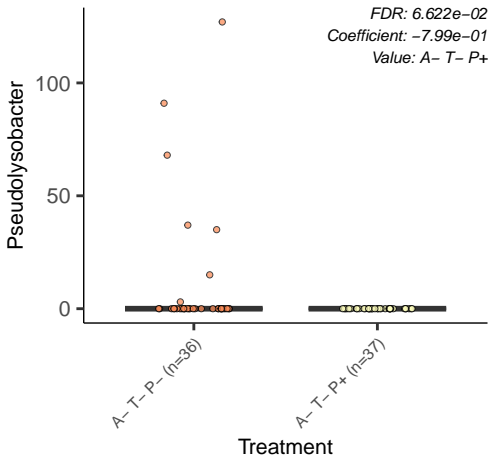
FDR: 6.622e-02

Coefficient: $-1.11\text{e}+00$

Value: A- T- P+



Treatment





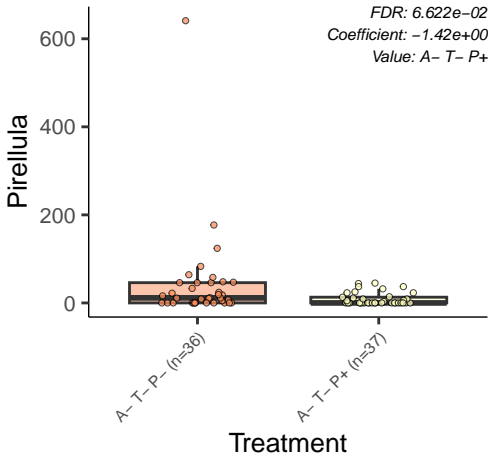
FDR: 6.622e-02

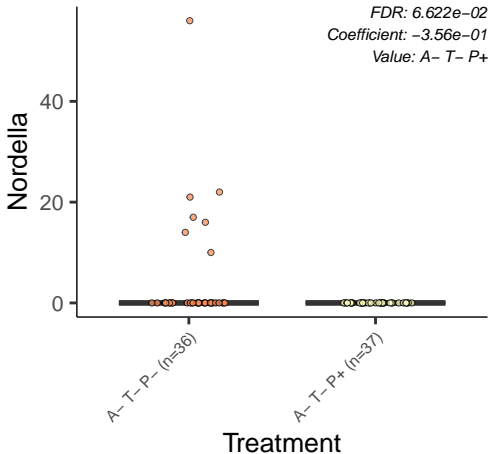
Coefficient: 1.18e+00

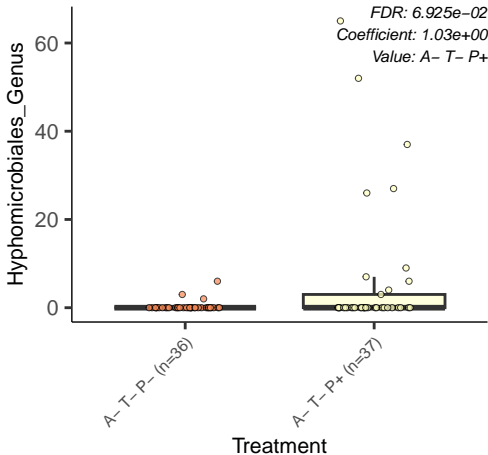
Value: A- T- P+

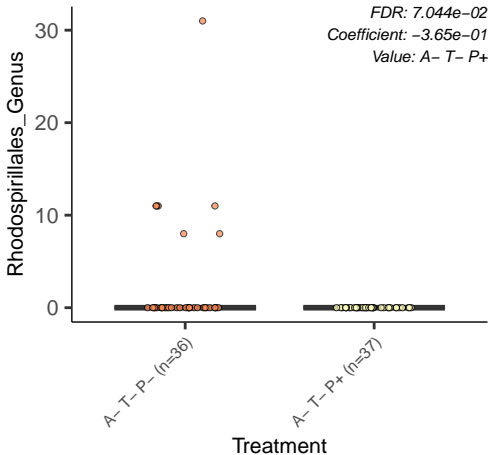


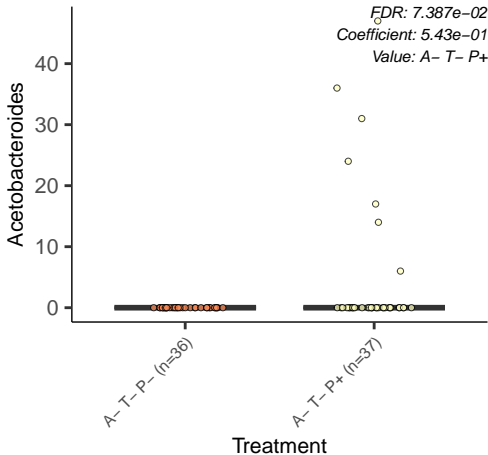
Treatment

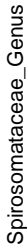












FDR: 7.814e-02
Coefficient: 7.65e-01
Value: A- T- P+



Treatment

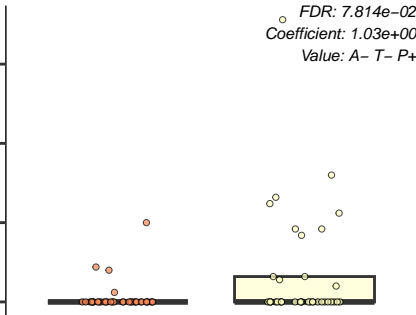
Mesorhizobium

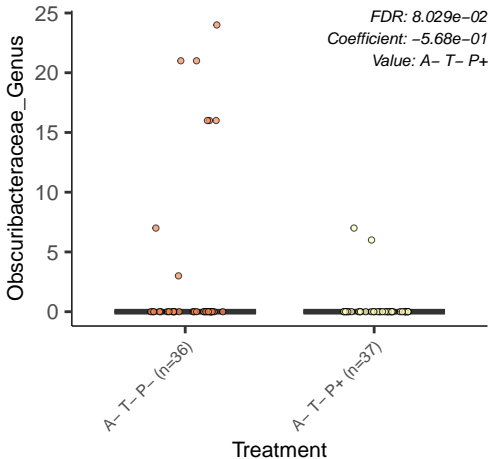
FDR: 7.814e-02
Coefficient: 1.03e+00
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment





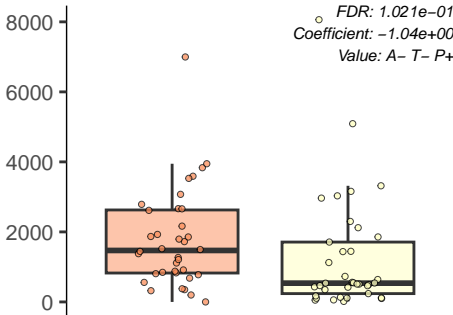
Crenobacter

FDR: 1.021e-01
Coefficient: -1.04e+00
Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment





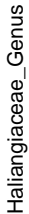
FDR: 1.021e-01

Coefficient: 1.38e+00

Value: A- T- P+



Treatment



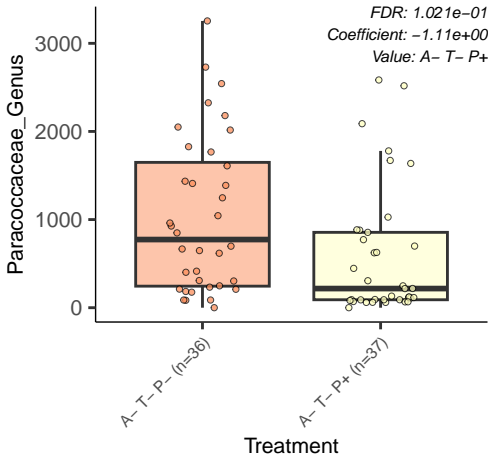
FDR: 1.021e-01

Coefficient: $-2.33e+00$

Value: A- T- P+



Treatment



Caulobacter

FDR: 1.126e-01

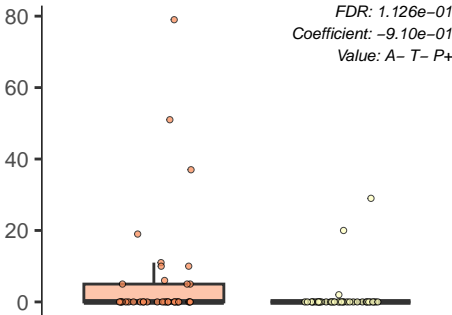
Coefficient: -9.10e-01

Value: A- T- P+

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



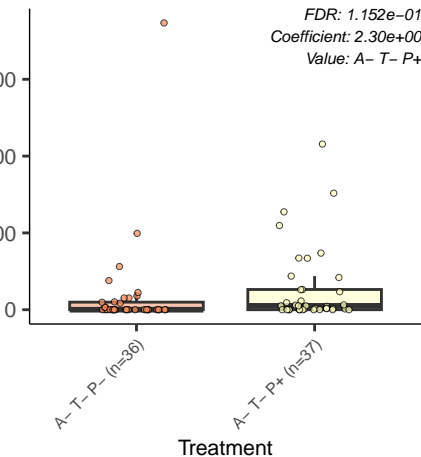
env.OPS_17_Genus

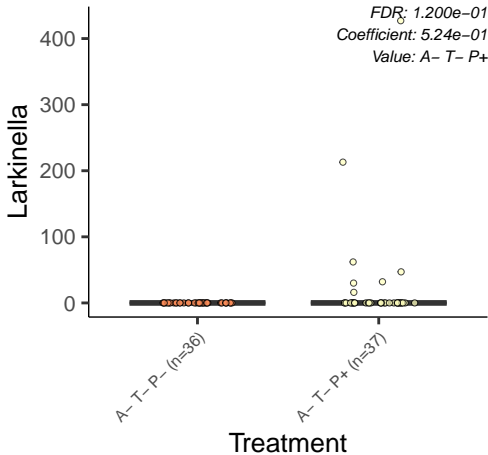
FDR: 1.152e-01
Coefficient: 2.30e+00
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment







FDR: 1.200e-01

Coefficient: $-5.46e-01$

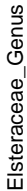
Value: A- T- P+



A-T-P-(n=36)

A-T-P+ (n=37)

Treatment



FDR: 1.200e-01

Coefficient: $-1.74e-01$

Value: A- T- P+



Treatment

Nocardia

FDR: 1.260e-01
Coefficient: 1.90e+00
Value: A- T- P+

2000

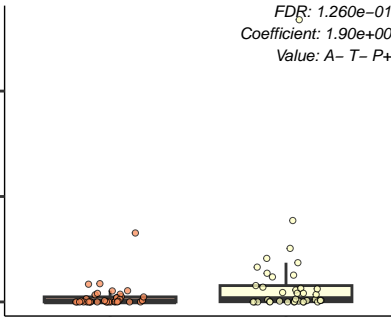
1000

0

A-T-P- (n=36)

A-T-P+ (n=37)

Treatment



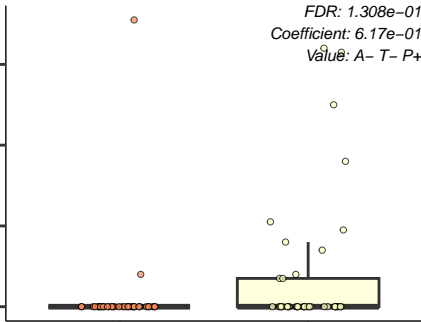
Pseudorhodoplanes

FDR: 1.308e-01
Coefficient: 6.17e-01
Value: A- T- P+

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





FDR: 1.389e-01

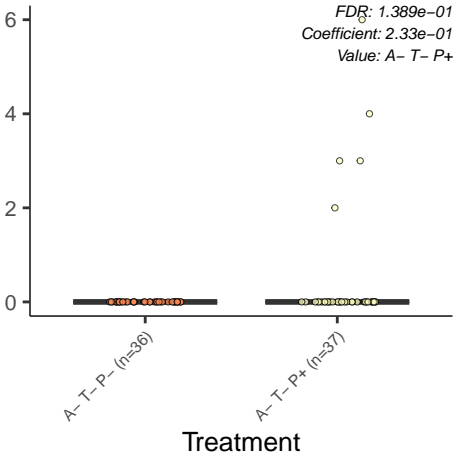
Coefficient: $-1.22e+00$

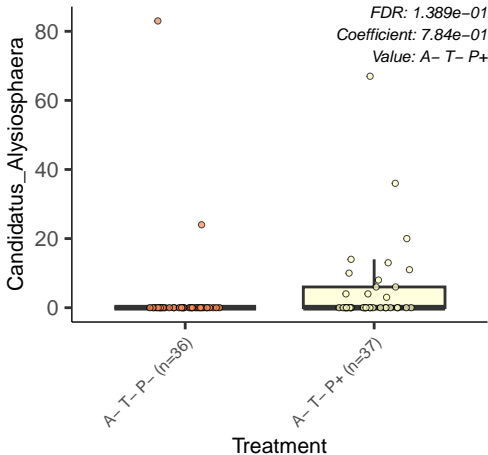
Value: A- T- P+

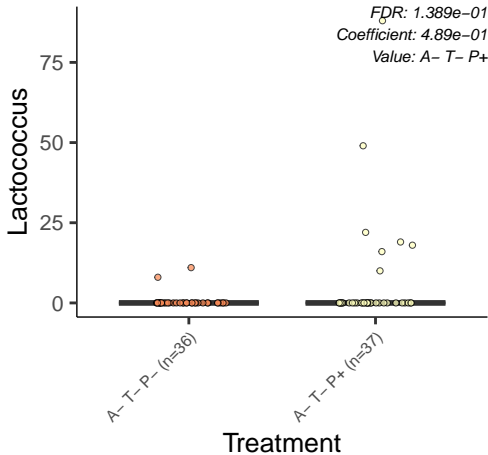


Treatment

Novispirillum







Candidatus_Protochlamydia

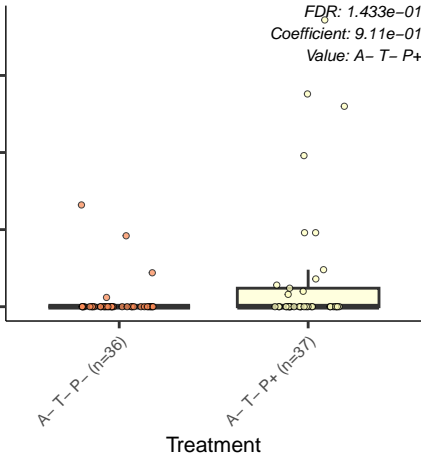
FDR: 1.433e-01
Coefficient: 9.11e-01
Value: A- T- P+

75
50
25
0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment





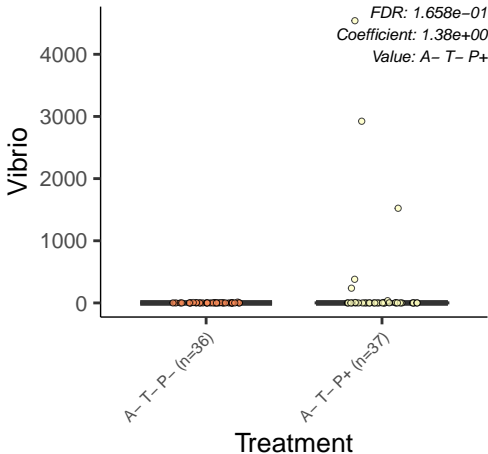
FDR: 1.658e-01

Coefficient: $-1.20e+00$

Value: A- T- P+



Treatment



Rhodococcus

1000

500

0

A-T-P- (n=36)

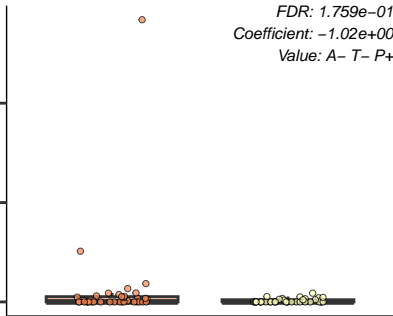
A-T-P+ (n=37)

Treatment

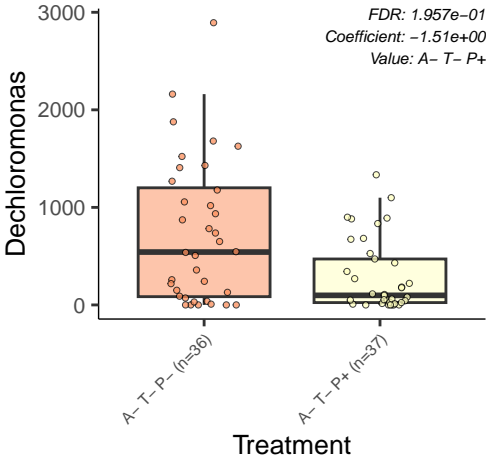
FDR: 1.759e-01

Coefficient: -1.02e+00

Value: A- T- P+



Dechloromonas





FDR: 2.113e-01

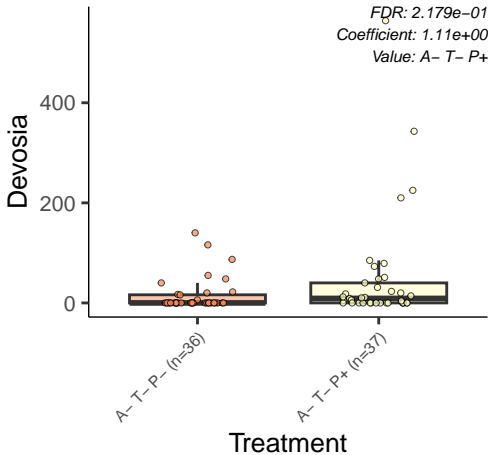
Coefficient: $-6.60e-01$

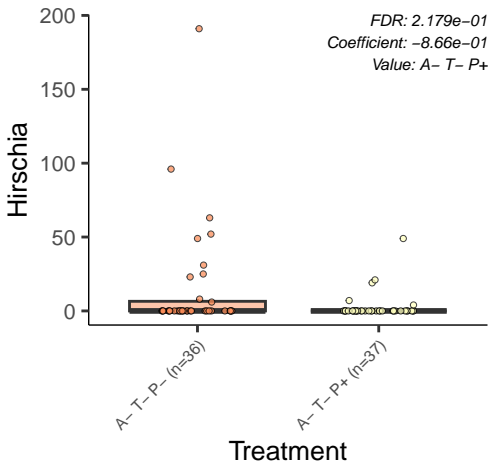
Value: A- T- P+

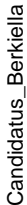
A-T-P-(n=36)

A-T-P+ (n=37)

Treatment







FDR: 2.219e-01

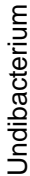
Coefficient: 6.96e-01

Value: A- T- P+

A-T-P-(n=36)

A-T-P+ (n=37)

Treatment



FDR: 2.219e-01

Coefficient: $-1.61e+00$

Value: A- T- P+



Treatment



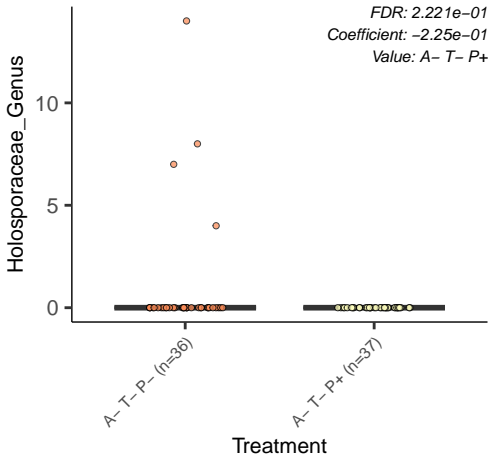
FDR: 2.221e-01

Coefficient: $-1.29e+00$

Value: A- T- P+



Treatment



Fluviicola

3000

2000

1000

0

A-T-P- (n=36)

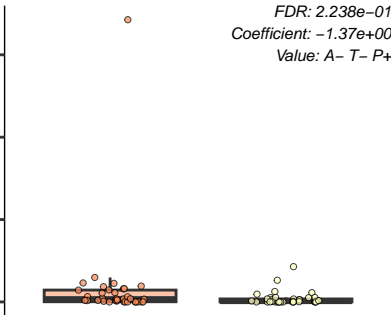
A-T-P+ (n=37)

Treatment

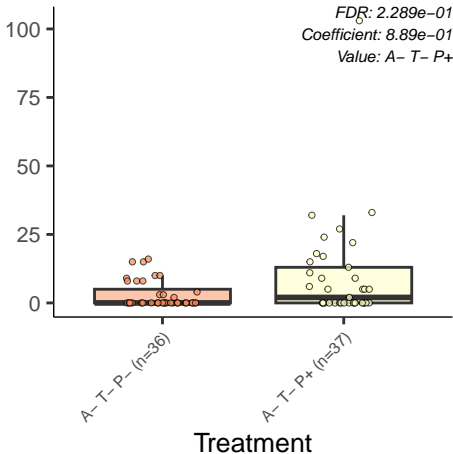
FDR: 2.238e-01

Coefficient: -1.37e+00

Value: A- T- P+



Roseicella



NS9_marine_group_Genus

FDR: 2.480e-01
Coefficient: 1.23e+00
Value: A- T- P+

200

100

0

A- T- P- (n=36)

A- T- P+ (n=37)

Treatment

