

Cloacibacterium

FDR: 1.572e-26

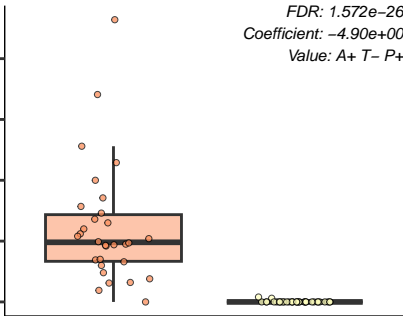
Coefficient: -4.90e+00

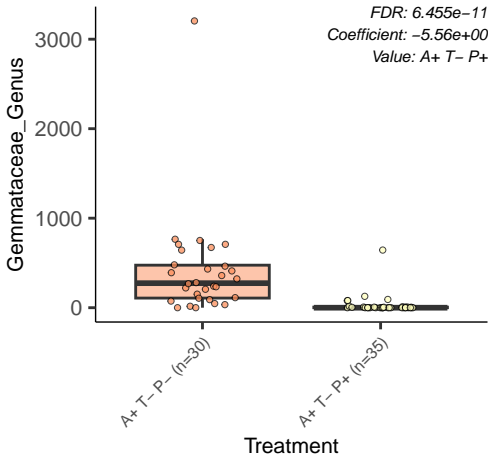
Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





Acinetobacter

FDR: $1.497e-10$
Coefficient: $5.00e+00$
Value: A+ T- P+

2000

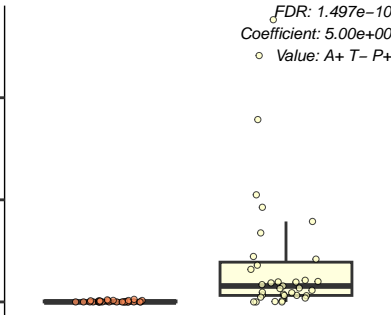
1000

0

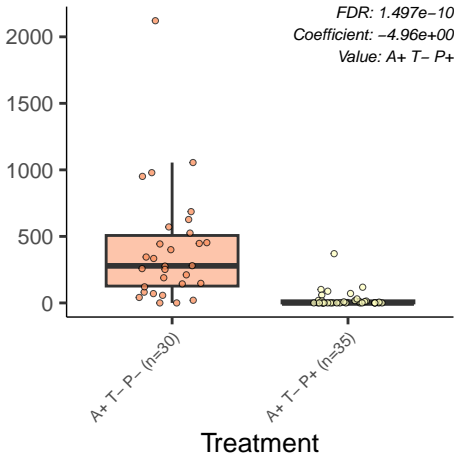
A+ T- P- (n=30)

A+ T- P+ (n=35)

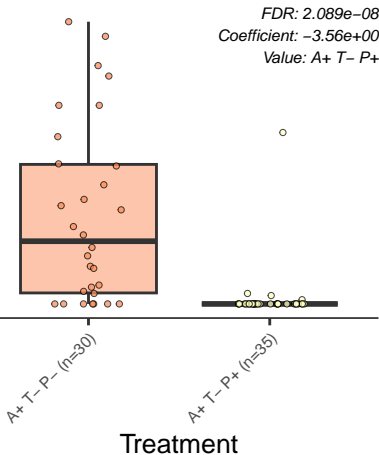
Treatment



Dinghuibacter



Caulobacter



Rhizobium

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: 1.170e-06
Coefficient: -2.36e+00
Value: A+ T- P+

200

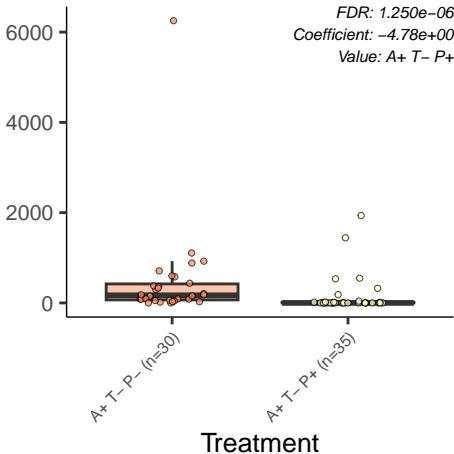
150

100

50

0

Niveibacterium



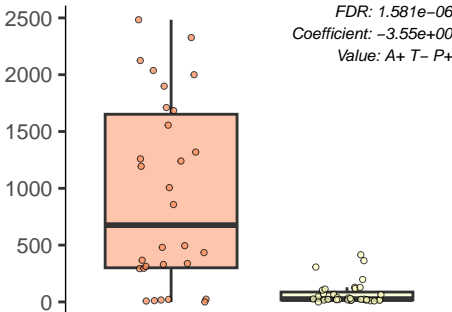
Plesiomonas

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: $1.581e-06$
Coefficient: $-3.55e+00$
Value: A+ T- P+



Roseomonas

FDR: 2.570e-06

Coefficient: -3.15e+00

Value: A+ T- P+

200

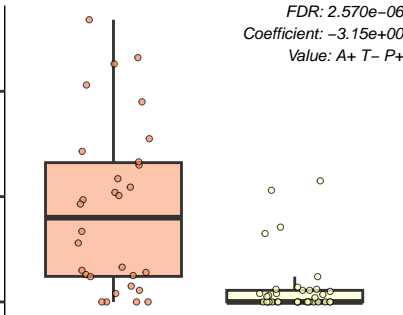
100

0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment



Paenirhodobacter

FDR: 6.596e-06

Coefficient: -2.61e+00

Value: A+ T- P+

900

600

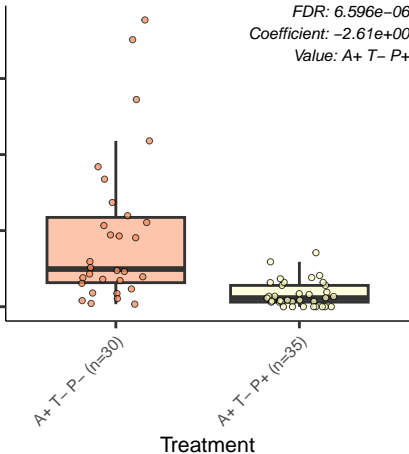
300

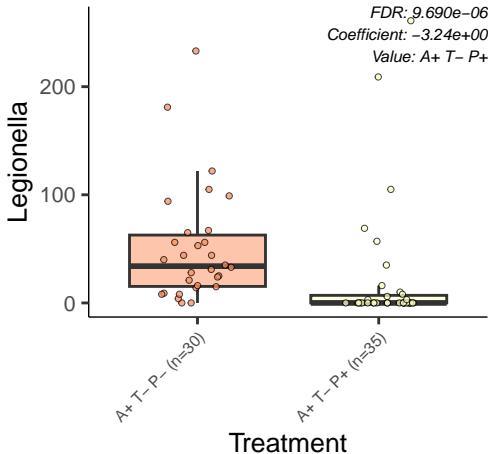
0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





Paracoccaceae_Genus

6000

4000

2000

0

FDR: 3.815e-05

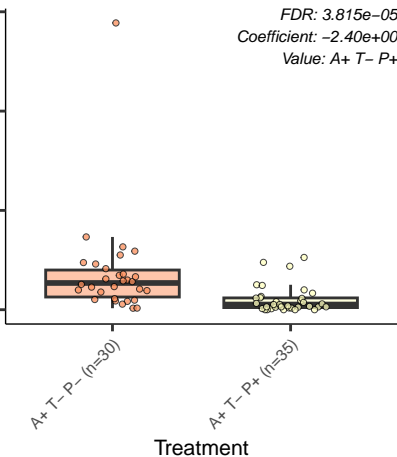
Coefficient: -2.40e+00

Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment



Tundrisphaera

FDR: 3.815e-05

Coefficient: -4.29e+00

Value: A+ T- P+

4000

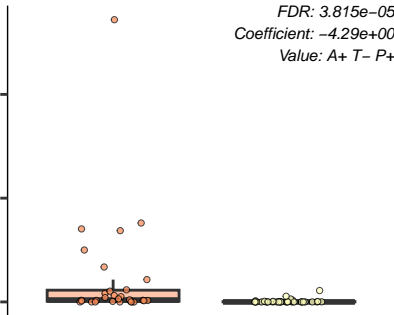
2000

0

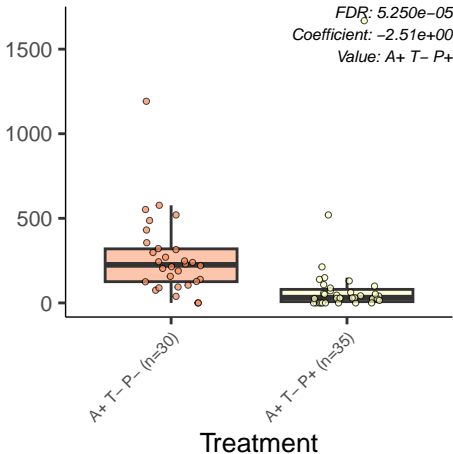
A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment



Gemmobacter



Aurantisolimonas

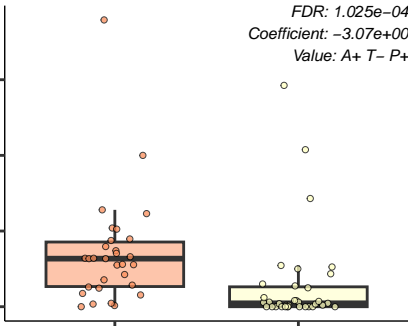
600
400
200
0

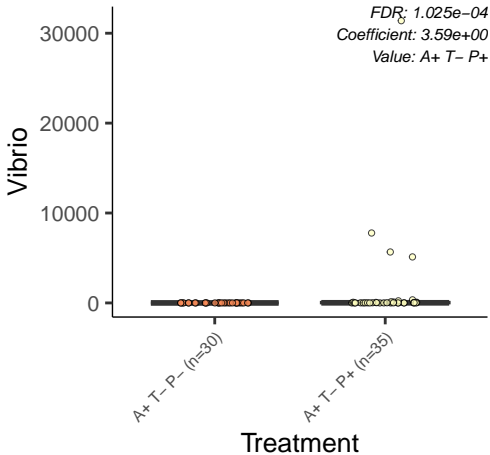
A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: 1.025×10^{-4}
Coefficient: -3.07×10^0
Value: A+ T- P+





Dechloromonas

2000

1000

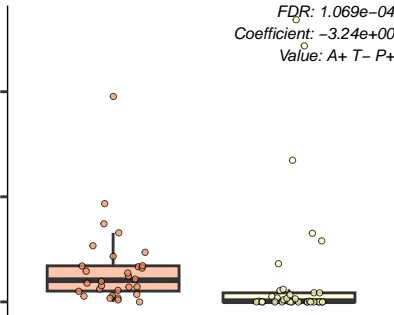
0

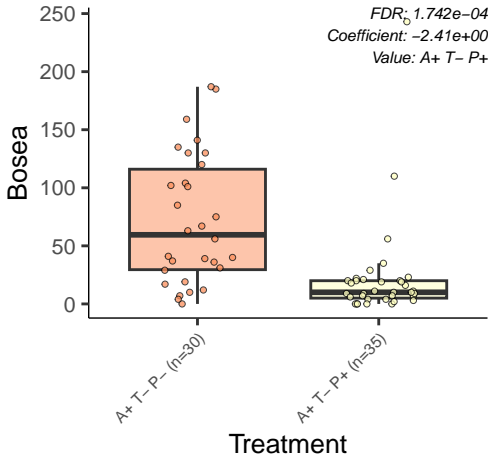
A+ T- P- (n=30)

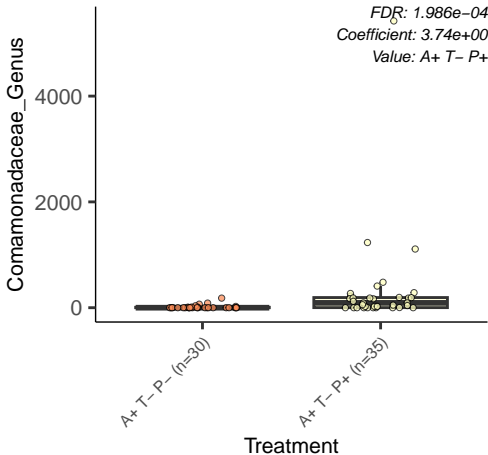
A+ T- P+ (n=35)

Treatment

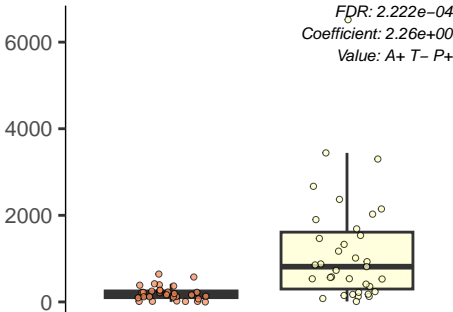
FDR: 1.069e-04
Coefficient: -3.24e+00
Value: A+ T- P+







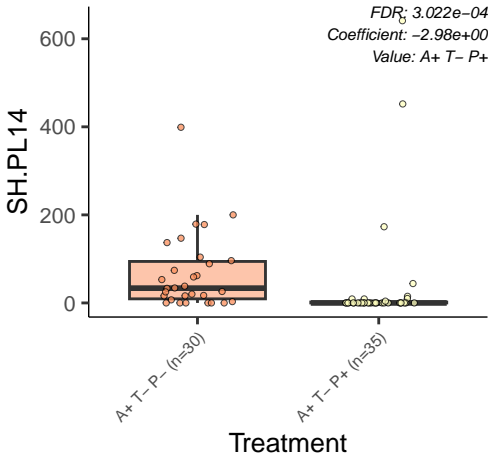
Shewanella



A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment



Gammaproteobacteria_Genus

FDR: 3.186e-04

Coefficient: -3.18e+00

Value: A+ T- P+

1500

1000

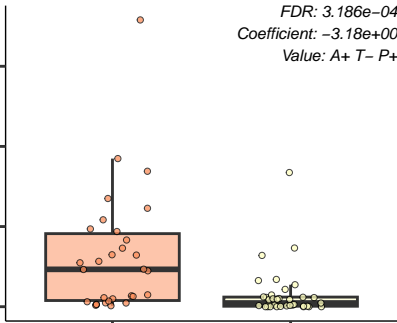
500

0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





FDR: 3.186e-04
 Coefficient: -2.65e+00
 Value: A+ T- P+



Treatment

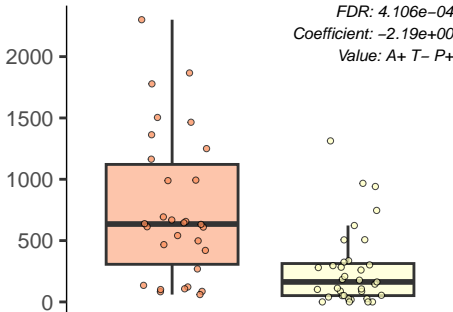
Phreatobacter

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: $4.106e-04$
Coefficient: $-2.19e+00$
Value: A+ T- P+



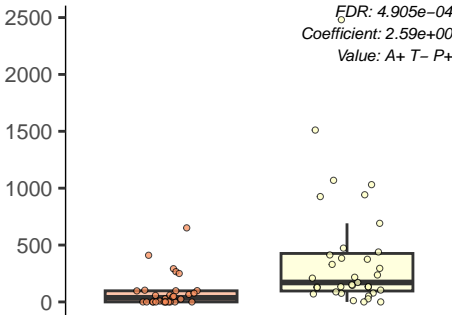
Flavobacterium

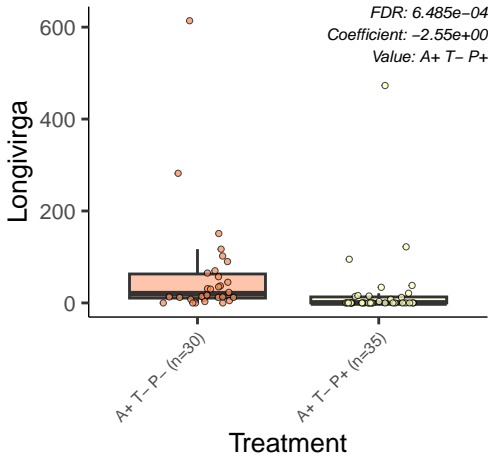
FDR: $4.905e-04$
Coefficient: $2.59e+00$
Value: A+ T- P+

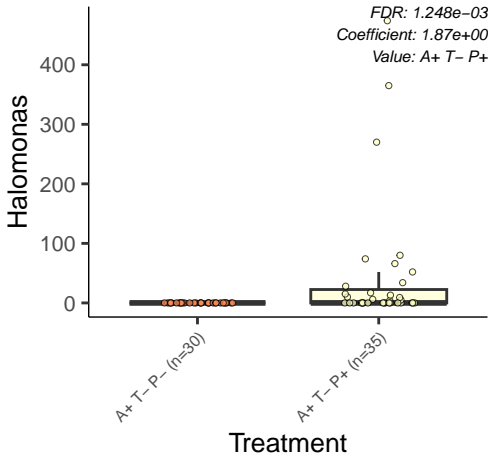
A+ T- P- (n=30)

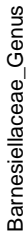
A+ T- P+ (n=35)

Treatment









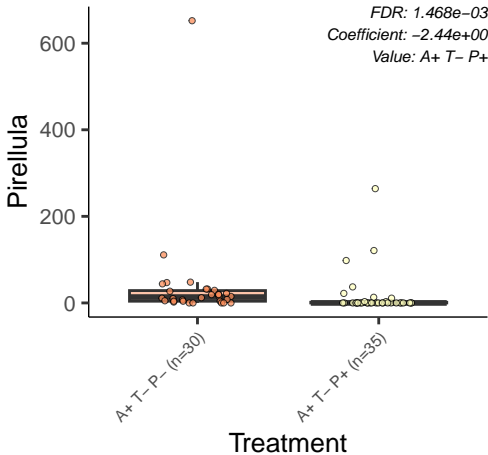
FDR: 1.463e-03

Coefficient: $-2.54e+00$

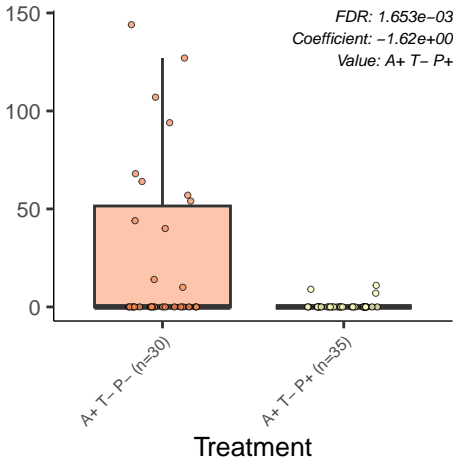
Value: $A+ T- P+$



Treatment



Luteolibacter



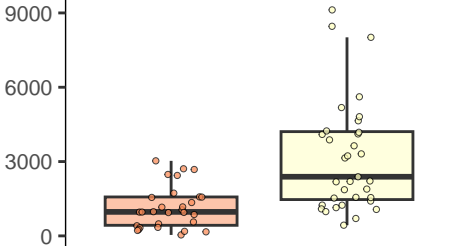
Pseudomonas

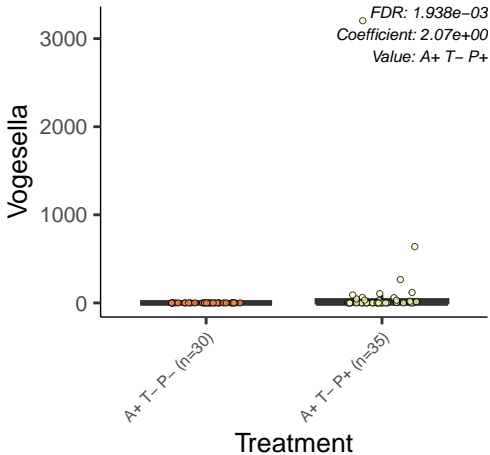
A+ T- P- (n=30)

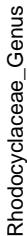
A+ T- P+ (n=35)

Treatment

FDR: 1.938e-03
Coefficient: 1.37e+00
Value: A+ T- P+







FDR: 4.110e-03

Coefficient: 1.84e+00

Value: $A+ T- P+$



Treatment

Pseudorhodoplanes

FDR: 4.409e-03

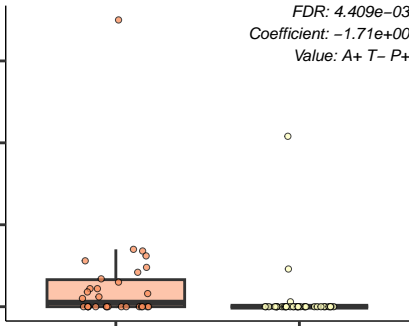
Coefficient: -1.71e+00

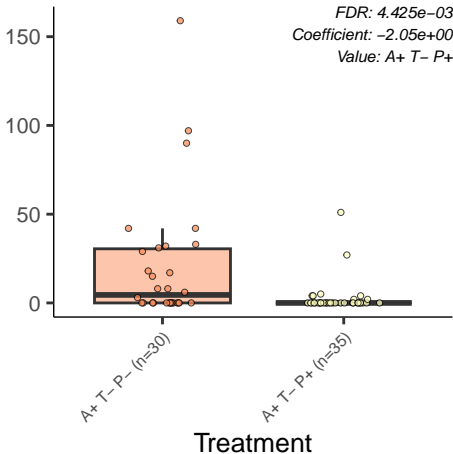
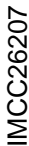
Value: A+ T- P+

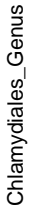
A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment







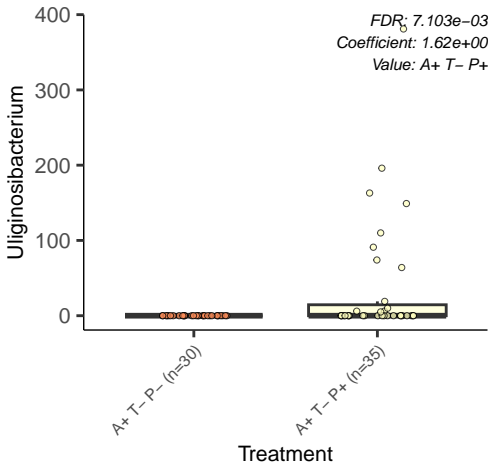
FDR: 6.260e-03

Coefficient: $-1.59e+00$

Value: $A+ T- P+$



Treatment



Chitinilyticum

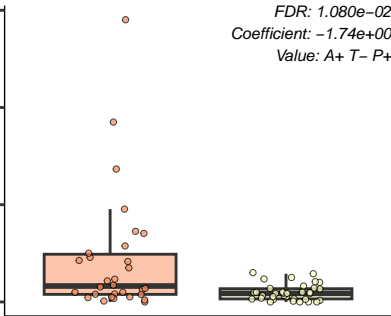
3000
2000
1000
0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: 1.080e-02
Coefficient: -1.74e+00
Value: A+ T- P+



Rhodovastum

FDR: 1.697e-02

Coefficient: -1.60e+00

Value: A+ T- P+

400

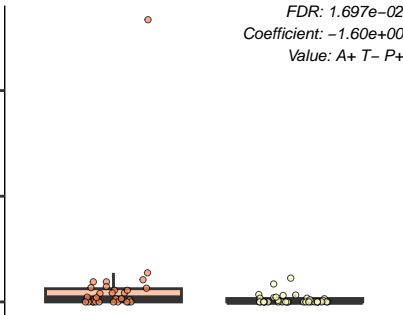
200

0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment



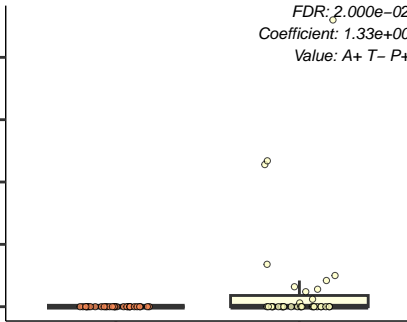
Candidatus_Megaira

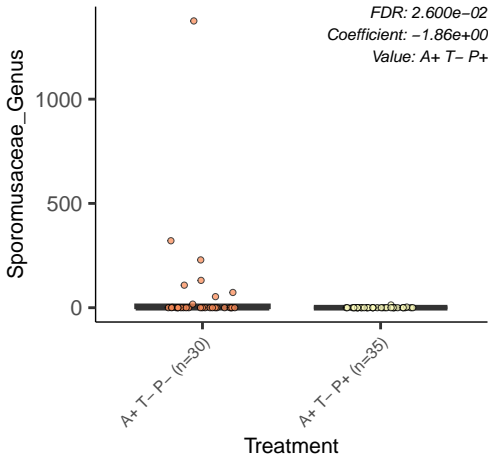
FDR: 2.000e-02
Coefficient: 1.33e+00
Value: A+ T- P+

A+ T- P- (n=30)

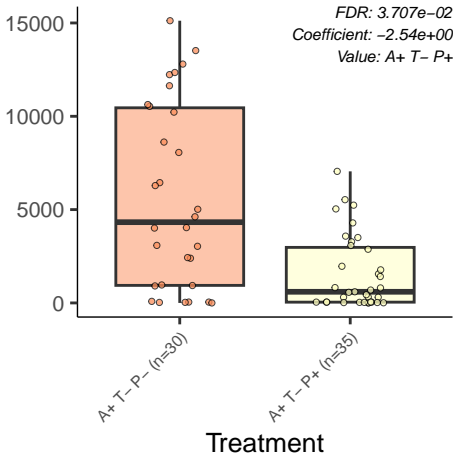
A+ T- P+ (n=35)

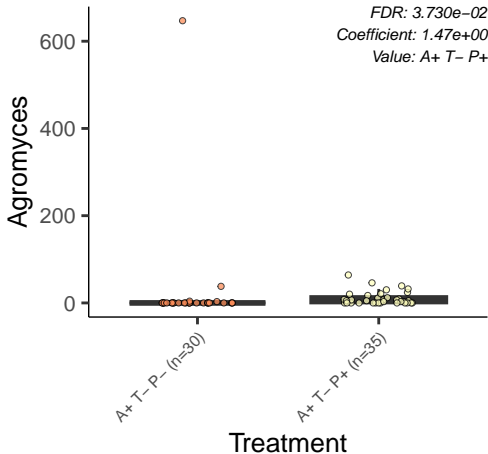
Treatment





Culicoidibacter





Rhizobiaceae_Genus

2000

1500

1000

500

0

A+ T- P- (n=30)

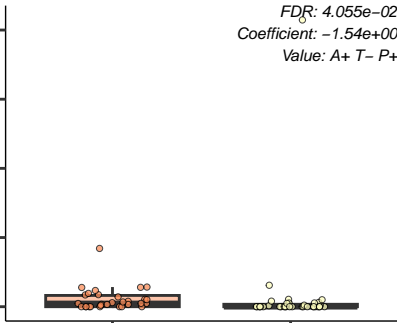
A+ T- P+ (n=35)

Treatment

FDR: $4.055e-02$

Coefficient: $-1.54e+00$

Value: A+ T- P+





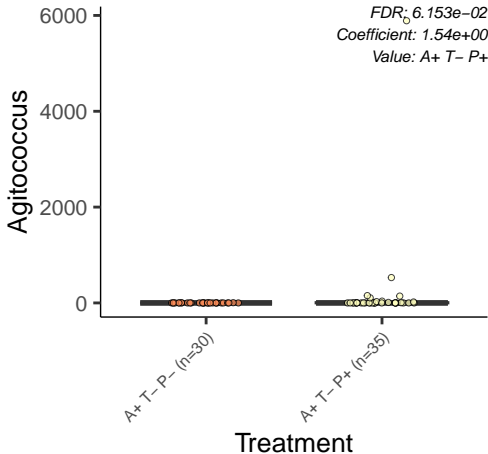
FDR: 4.055e-02

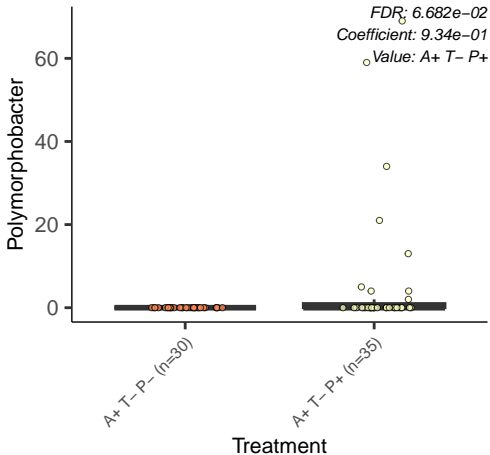
Coefficient: $-1.04e+00$

Value: $A+ T- P+$

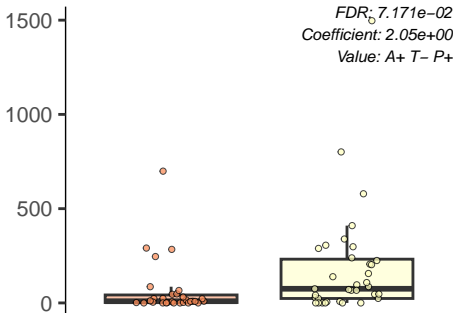


Treatment





Cellvibrio



A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

Pir4_lineage

FDR: $7.227e-02$

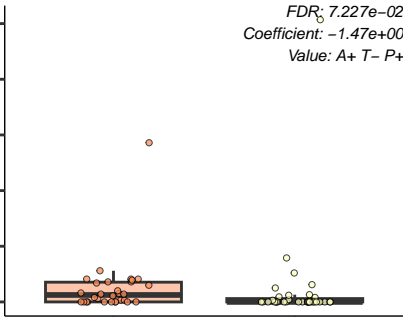
Coefficient: $-1.47e+00$

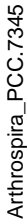
Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





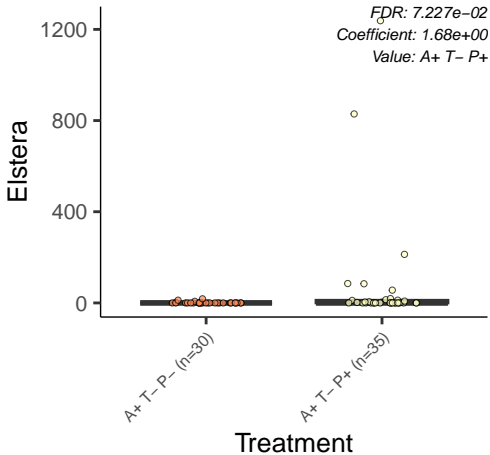
FDR: 7.227e-02

Coefficient: 8.98e-01

Value: A+ T- P+



Treatment



Microtrichales_Genus

FDR: 7.312e-02

Coefficient: -1.40e+00

Value: A+ T- P+

200

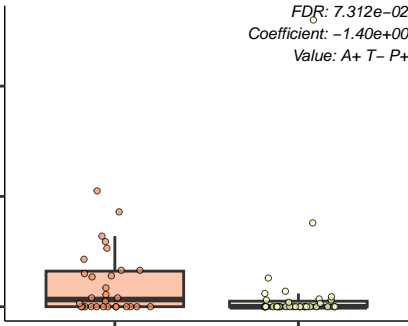
100

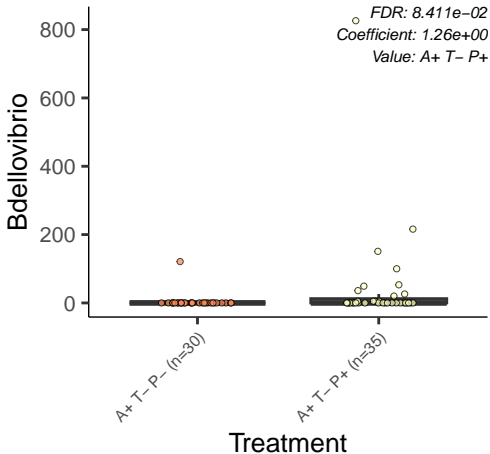
0

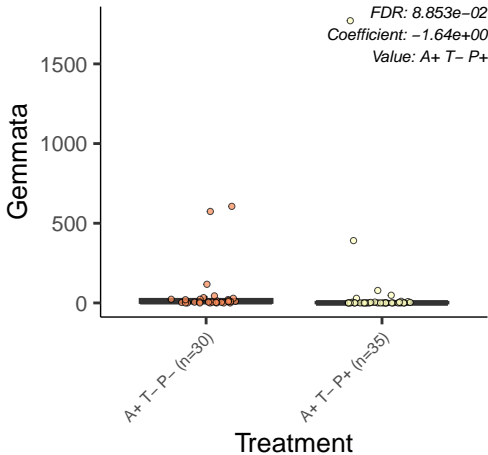
A+ T- P- (n=30)

A+ T- P+ (n=35)

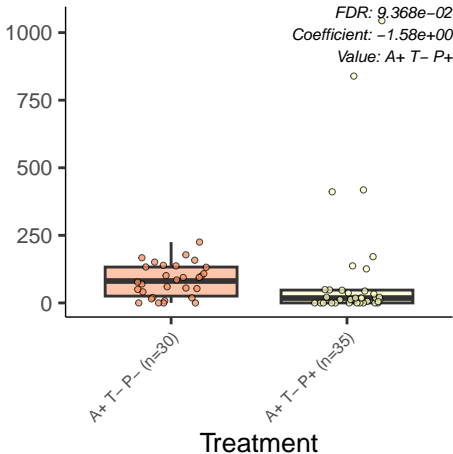
Treatment

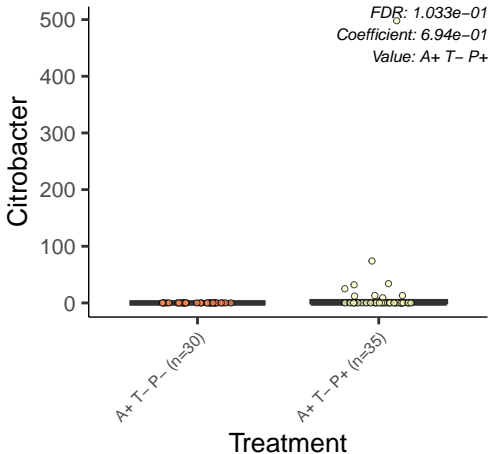






Ensifer





Crenobacter

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

FDR: 1.103e-01
Coefficient: -1.20e+00
Value: A+ T- P+

6000

4000

2000

0

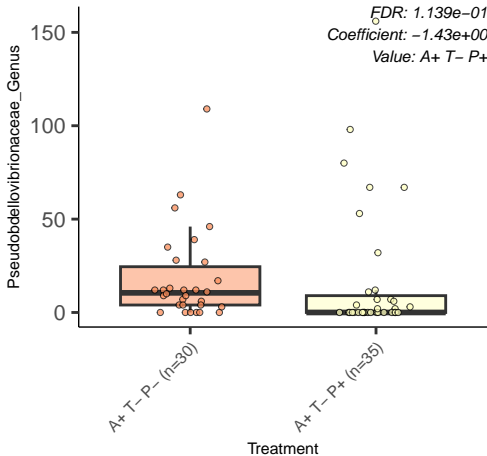
Pseudobdellovibrionaceae_Genus

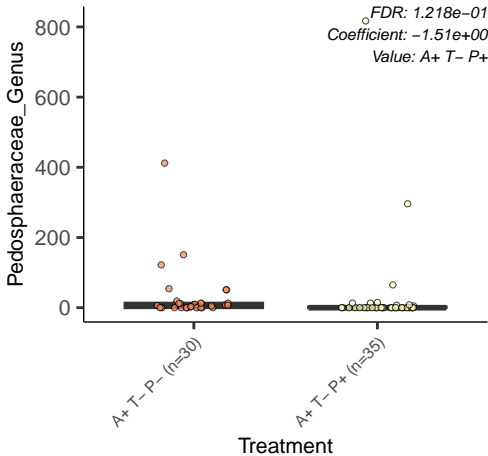
FDR: 1.139e-01
Coefficient: -1.43e+00
Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





Pseudoxanthomonas

FDR: 1.218e-01

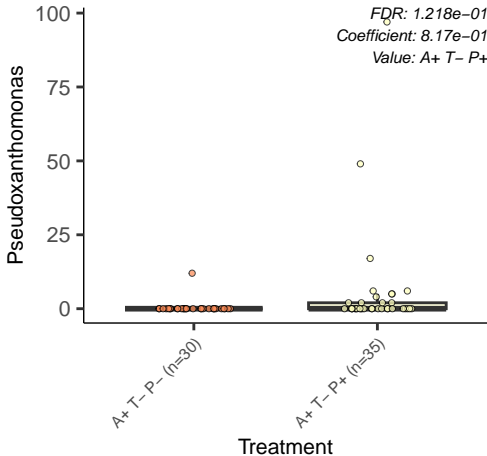
Coefficient: 8.17e-01

Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





FDR: 1.218e-01

Coefficient: 8.56e-01

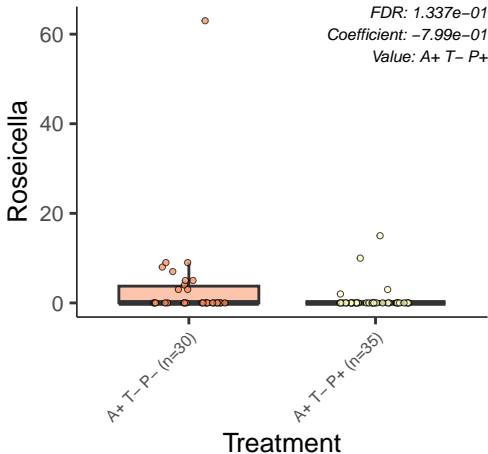
Value: A+ T- P+



A+T-P-(n=30)

A+ T-P+ (n=35)

Treatment



Fluviicola

FDR: 1.388e-01
Coefficient: -2.00e+00
Value: A+ T- P+

1000

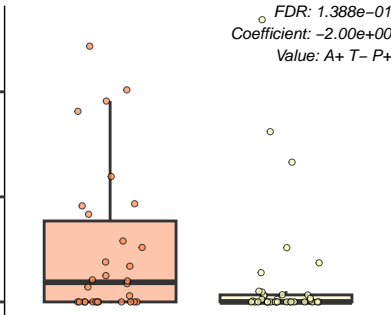
500

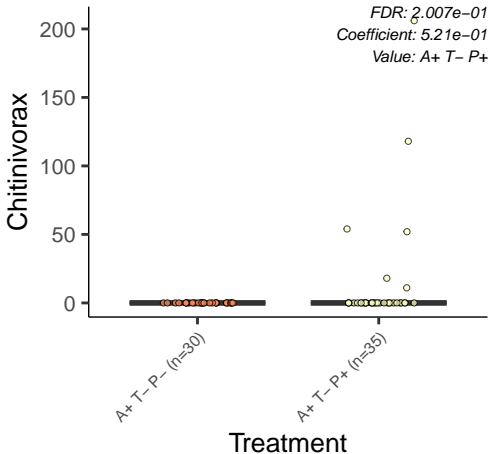
0

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment





Paucibacter

FDR: 2.336e-01

Coefficient: -8.50e-01

Value: A+ T- P+

A+ T- P- (n=30)

A+ T- P+ (n=35)

Treatment

