

Alloprevotella

FDR: 7.814e-05

Coefficient: 1.79e+00

Value: Wi_APCMin

-2

-3

-4

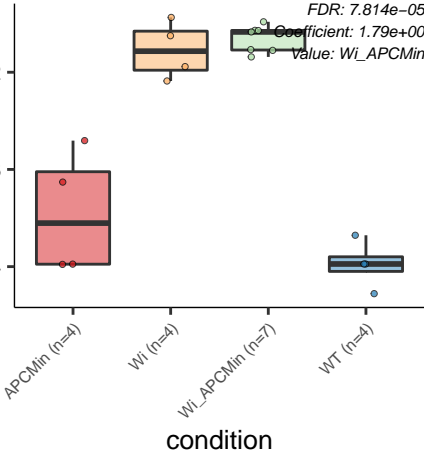
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

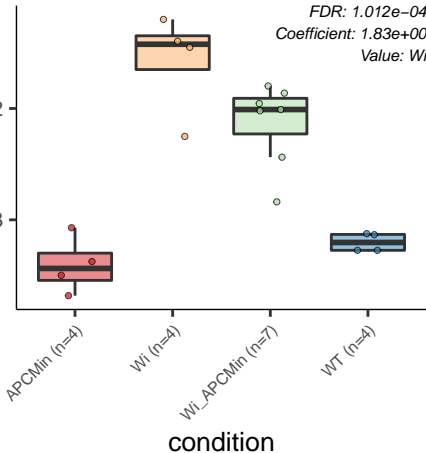
WT (n=4)

condition



Eubacterium

FDR: 1.012e-04
Coefficient: 1.83e+00
Value: Wi



Alloprevotella

FDR: 2.522e-04

Coefficient: 1.67e+00

Value: Wi

APCMin (n=4)

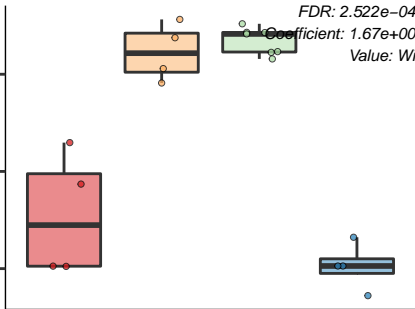
Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

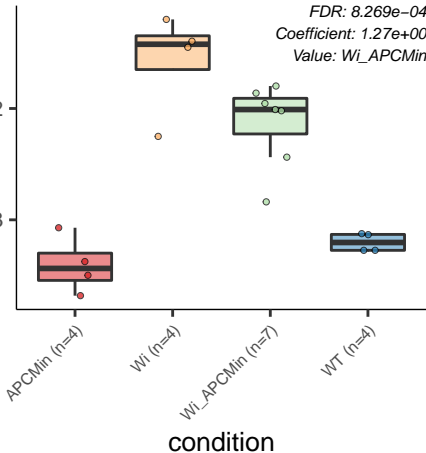
condition

-2
-3
-4

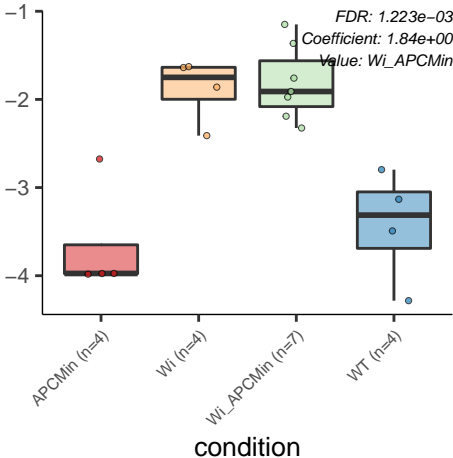


Eubacterium

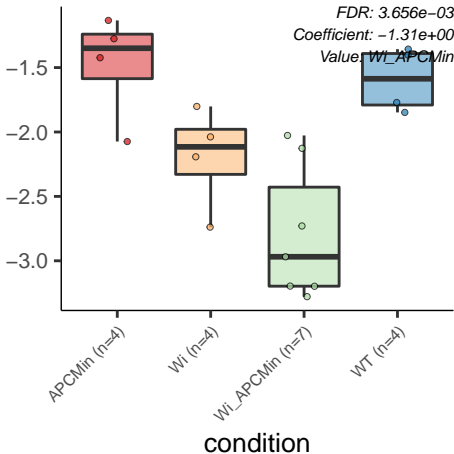
FDR: 8.269e-04
Coefficient: 1.27e+00
Value: Wi_APCMin



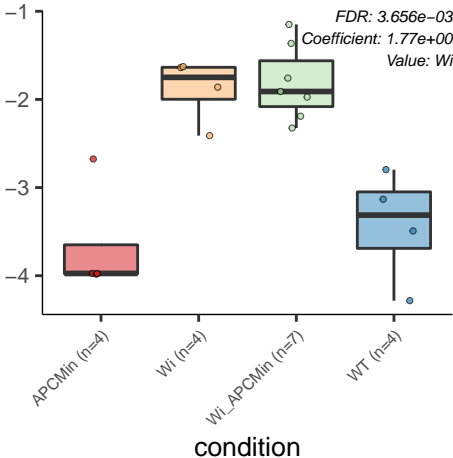
Paraprevotella



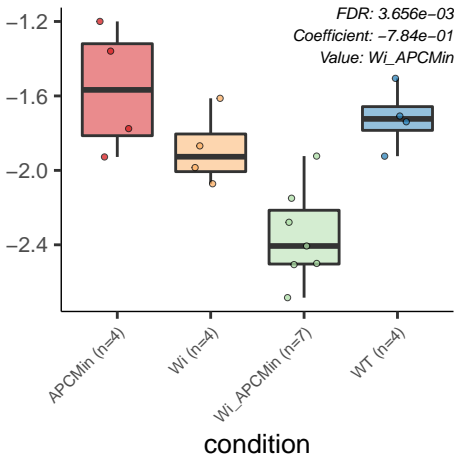
Rikenella



Paraprevotella



Lactobacillus



Acholeplasma

FDR: 3.656e-03

Coefficient: 1.75e+00

Value: W_i_APCMin

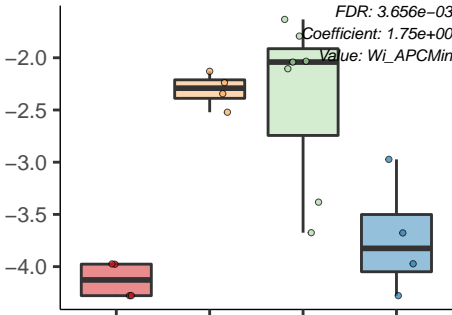
APCMin (n=4)

Wi (n=4)

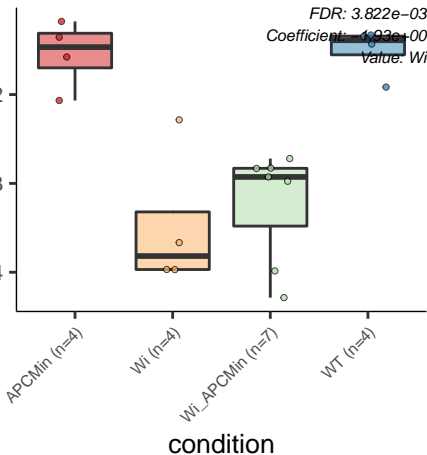
Wi_APcMin (n=7)

WT (n=4)

condition



Mucispirillum

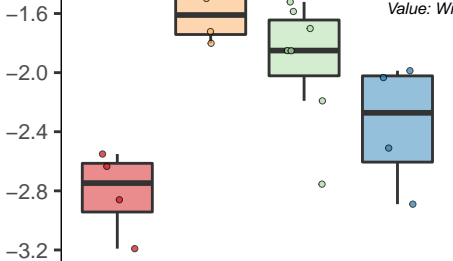


Alkaliphilus

FDR: 3.822e-03

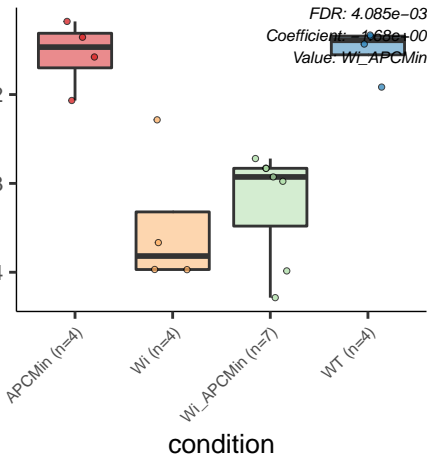
Coefficient: 1.22e+00

Value: Wi



condition

Mucispirillum



Acholeplasma

FDR: 5.279e-03

Coefficient: 1.82e+00

Value: Wi

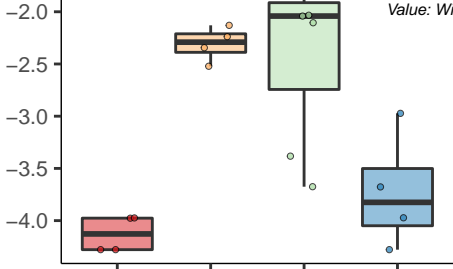
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

condition

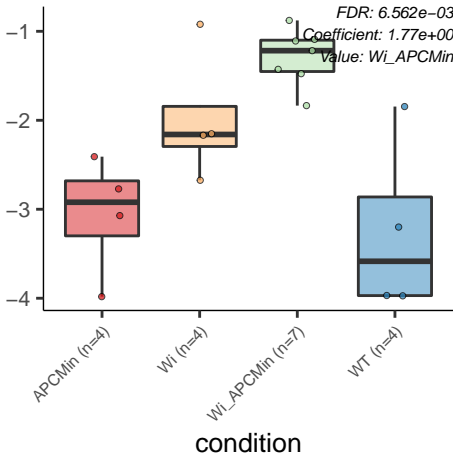


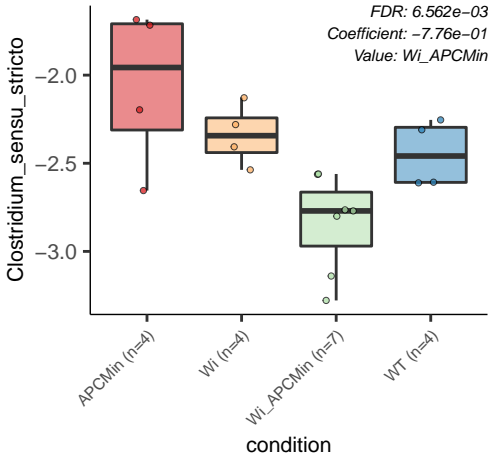
Akkermansia

FDR: 6.562e-03

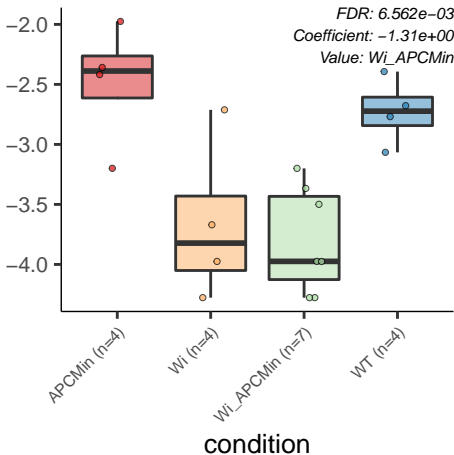
Coefficient: 1.77e+00

Value: Wi_APCMin

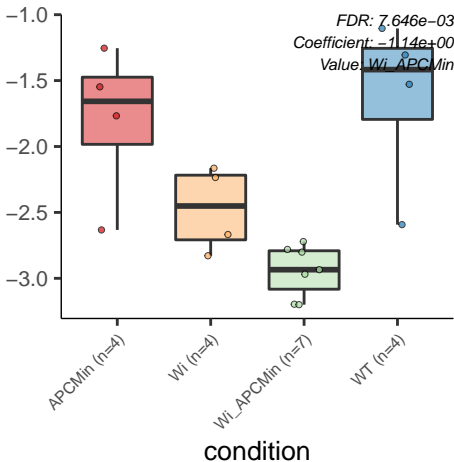




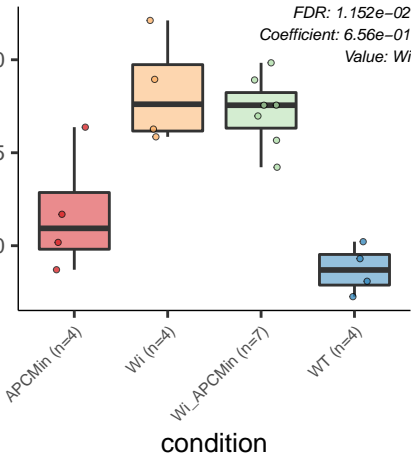
Ureaplasma



Stomatobaculum



Porphyromonas

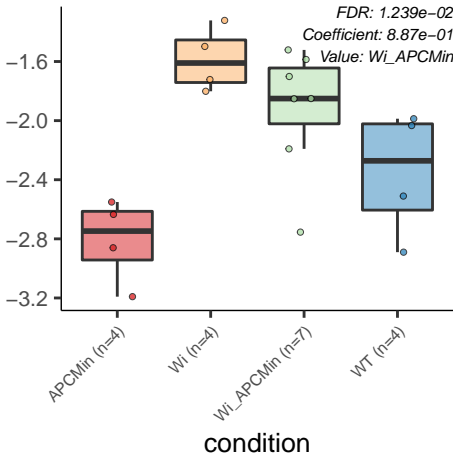


Alkaliphilus

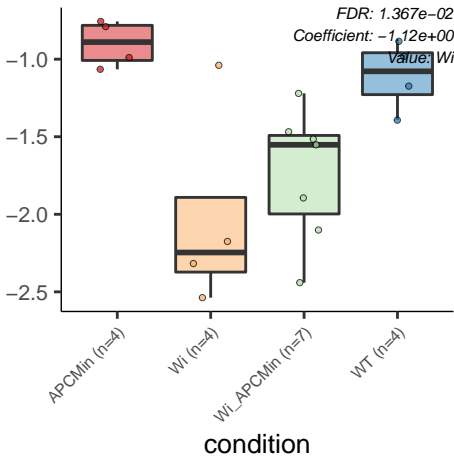
FDR: 1.239e-02

Coefficient: 8.87e-01

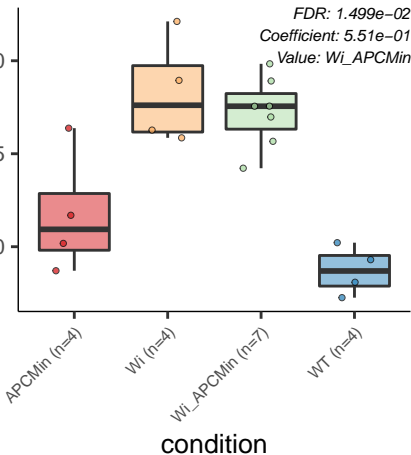
Value: W_i_APCMin



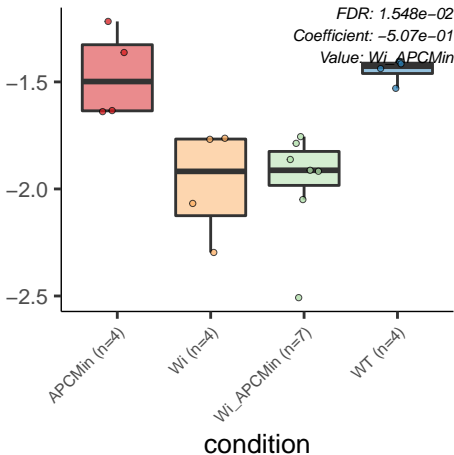
Helicobacter



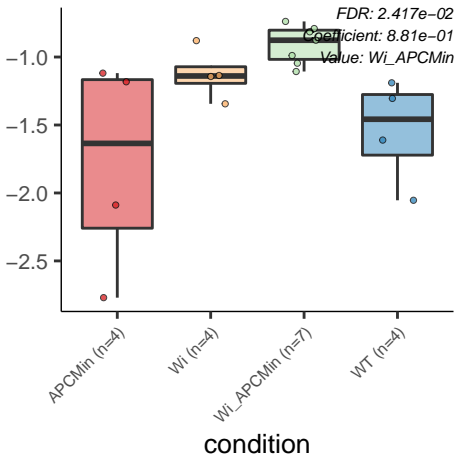
Porphyromonas



Odoribacter



Prevotella



Ureaplasma

FDR: $2.417\text{e-}02$
Coefficient: $-1.17\text{e}+00$
Value: W_i

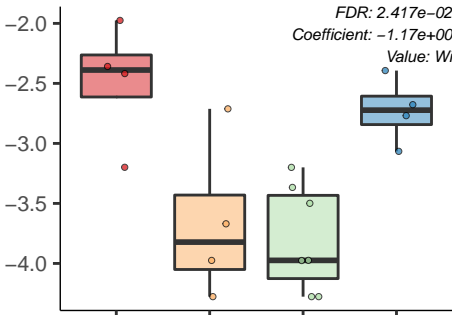
APCMin (n=4)

Wi (n=4)

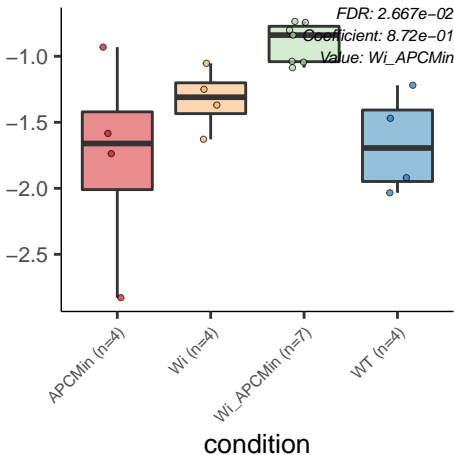
Wi_APCMin (n=7)

WT (n=4)

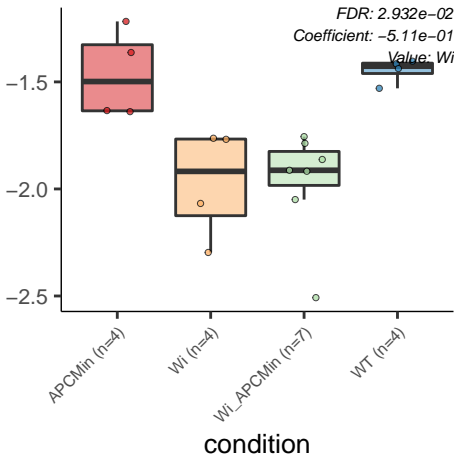
condition



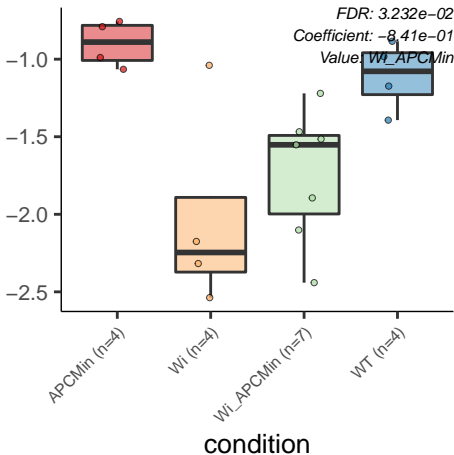
Bacteroides



Odoribacter



Helicobacter



Escherichia.Shigella

FDR: 4.176e-02

Coefficient: -1.35e+00

Value: Wi

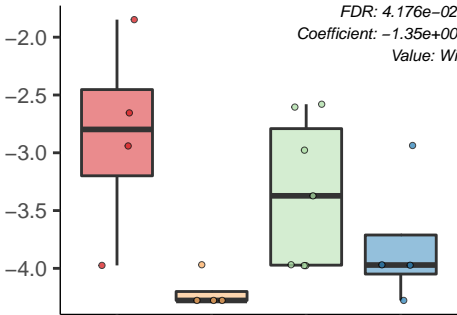
APCMin (n=4)

Wi (n=4)

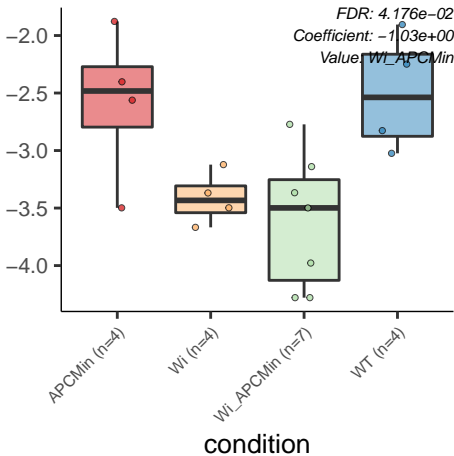
Wi_APCMin (n=7)

WT (n=4)

condition



Syntrophococcus



Ruminococcus

FDR: $6.947e-02$
Coefficient: $1.05e+00$
Value: Wi

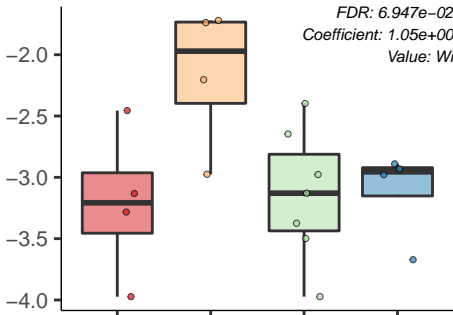
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

condition



Coprobacillus

FDR: 1.116e-01

Coefficient: 9.12e-01

Value: Wi

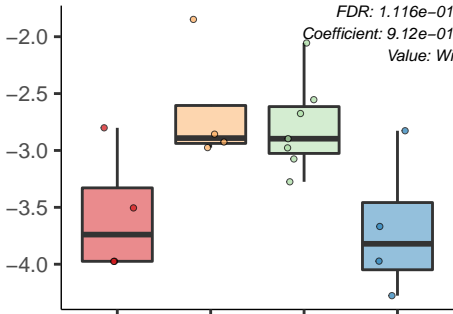
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

condition



Lachnospiraceae_incertae_sedis

-2.0
-2.5

APCMin (n=4)

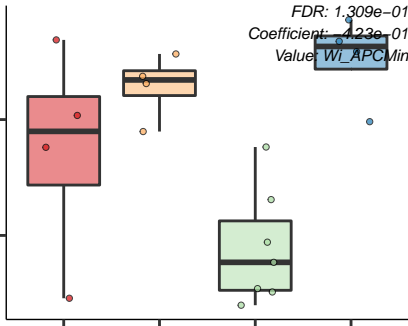
Wi (n=4)

Wi_APcMin (n=7)

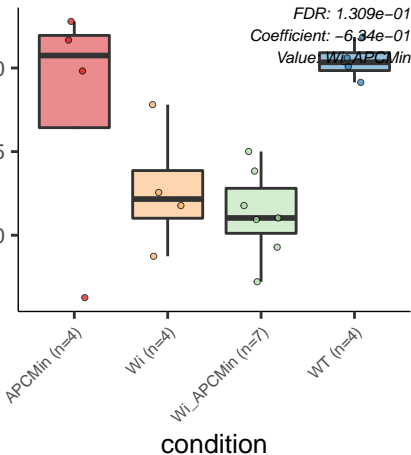
WT (n=4)

condition

FDR: 1.309e-01
Coefficient: -4.23e-01
Value: Wi_APcMin



Anaerotruncus



Coprobacillus

FDR: 1.309e-01

Coefficient: 7.77e-01

Value: Wi_APCMin

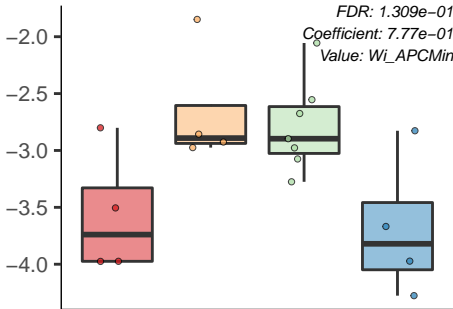
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

condition

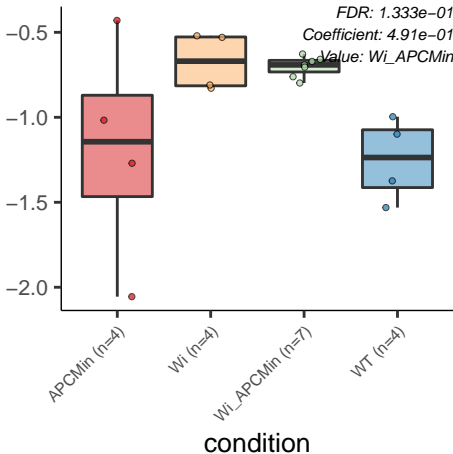


Barnesiella

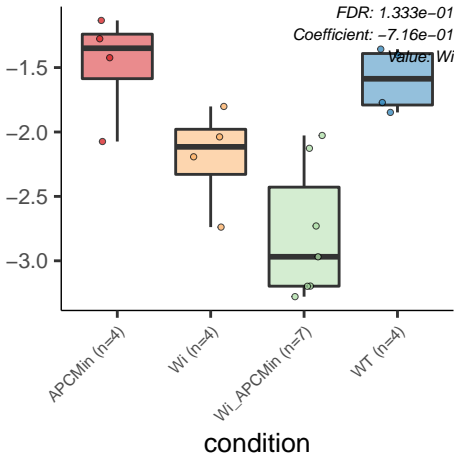
FDR: 1.333e-01

Coefficient: 4.91e-01

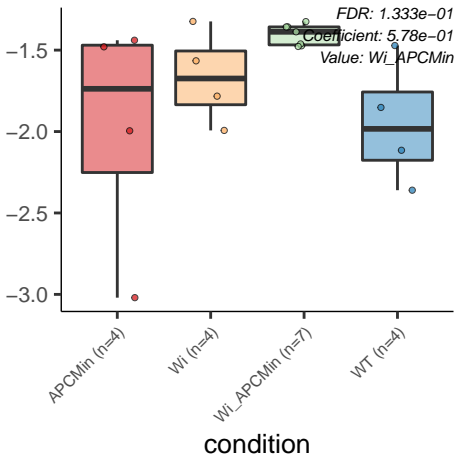
Value: Wi_APCMin



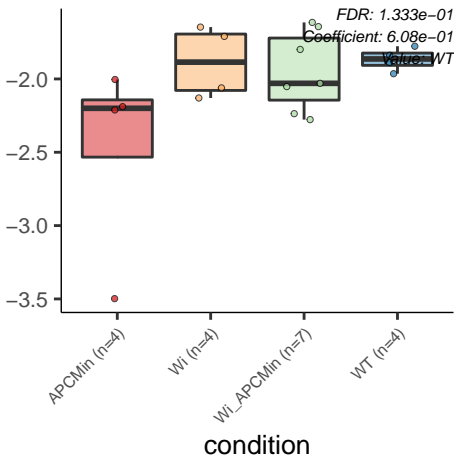
Rikenella



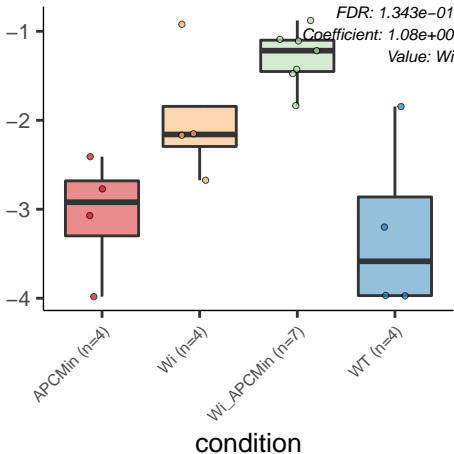
Parabacteroides



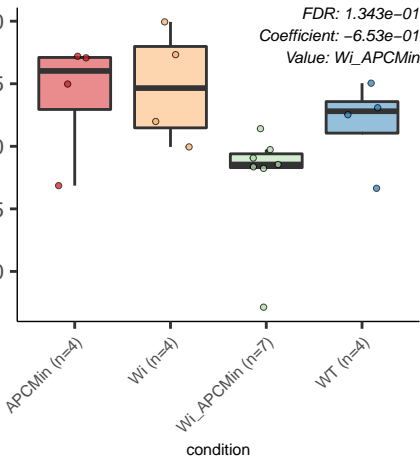
Flavonifractor



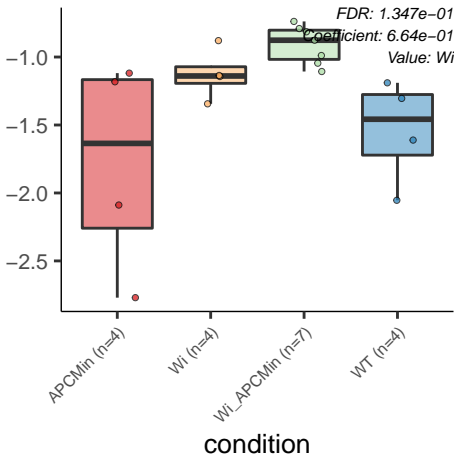
Akkermansia



Saccharibacteria_genera_incertae_sedis



Prevotella

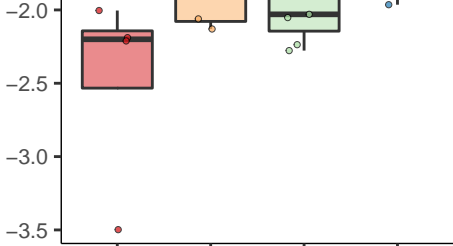


Flavonifractor

FDR: 1.357e-01

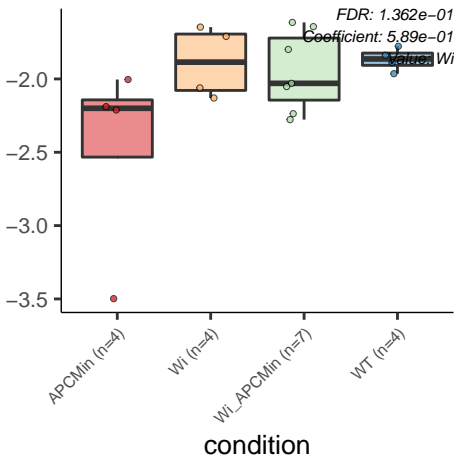
Coefficient: 5.25e-01

Value: W_i APCMin



condition

Flavonifractor

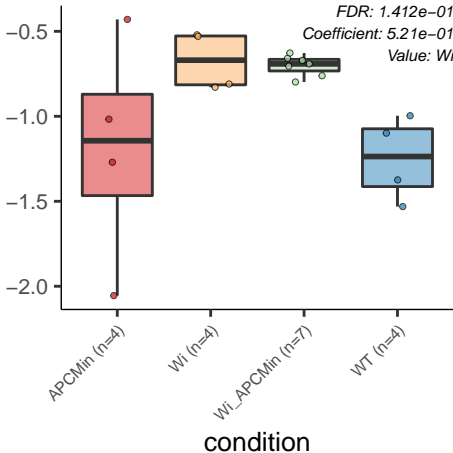


Barnesiella

FDR: 1.412e-01

Coefficient: 5.21e-01

Value: Wi



Alloprevotella

FDR: 1.465e-01

Coefficient: -5.29e-01

Value: WT

-2

-3

-4

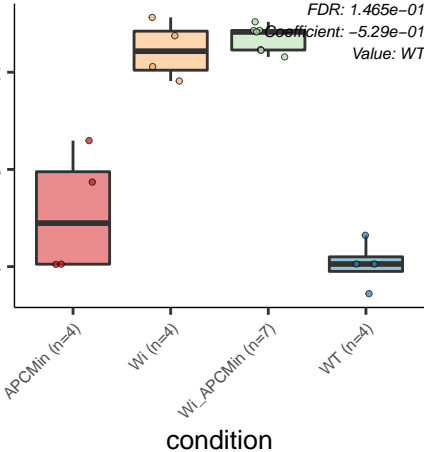
APCMin (n=4)

Wi (n=4)

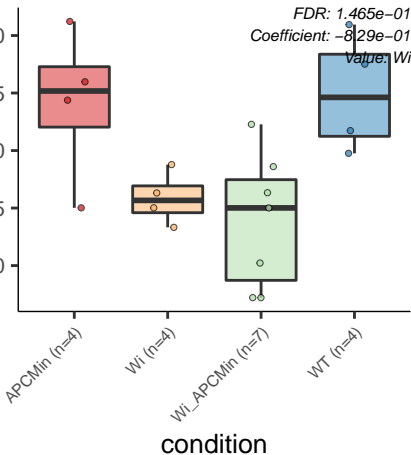
Wi_APCMin (n=7)

WT (n=4)

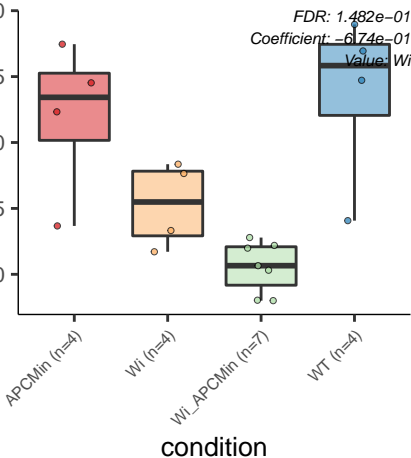
condition



Syntrophococcus



Stomatobaculum



Escherichia.Shigella

FDR: 1.586e-01

Coefficient: -9.35e-01

Value: WT

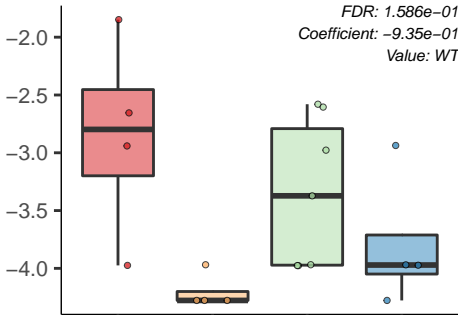
APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

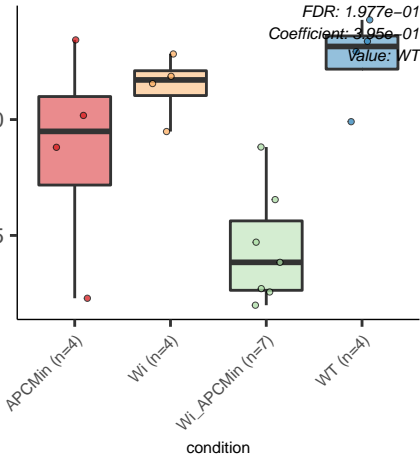
WT (n=4)

condition



Lachnospiraceae_incertae_sedis

FDR: 1.977×10^{-1}
Coefficient: 3.95×10^{-1}
Value: WT



Anaeroplasm

FDR: 2.001e-01

Coefficient: -1.01e+00

Value: W_i

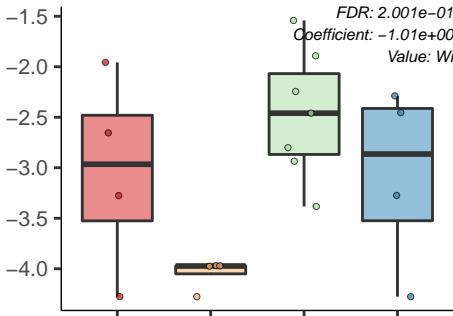
APCMin (n=4)

W_i (n=4)

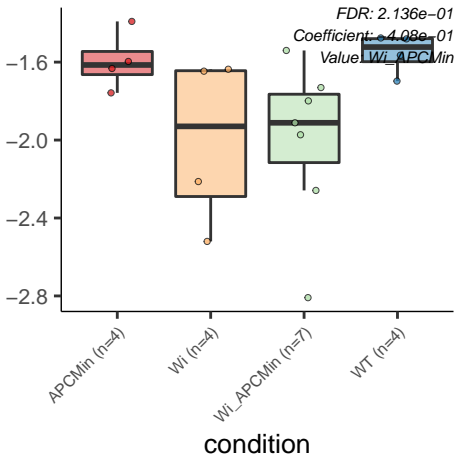
W_i _APCMin (n=7)

WT (n=4)

condition



Desulfovibrio



Turcibacter

FDR: 2.245e-01
Coefficient: -9.37e-01
Value: WT

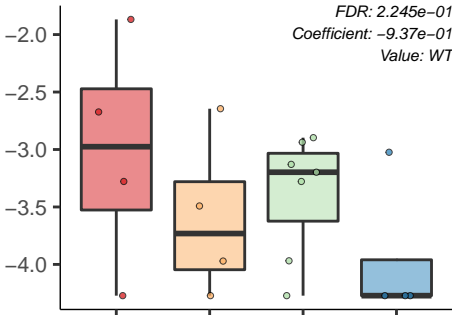
APCMin (n=4)

Wi (n=4)

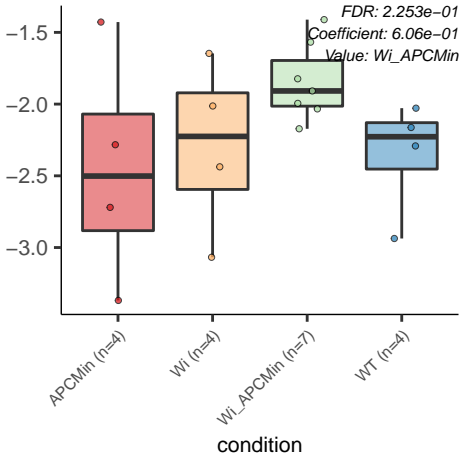
Wi_APCMin (n=7)

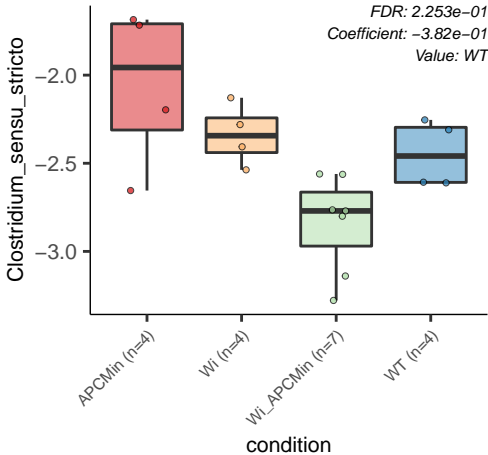
WT (n=4)

condition

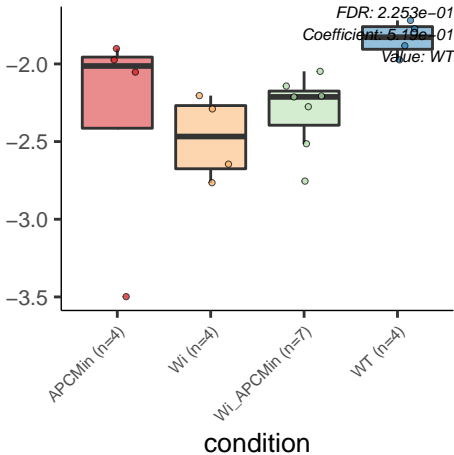


Falsiporphyrromonas

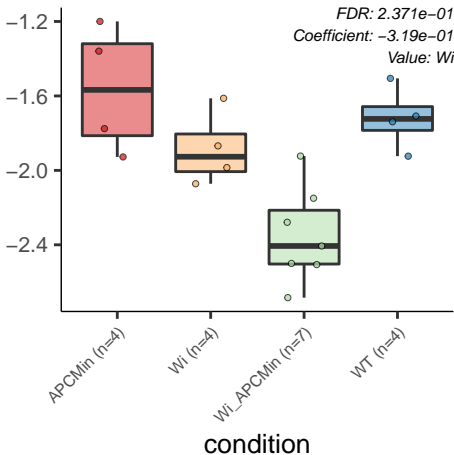




Oscillibacter



Lactobacillus

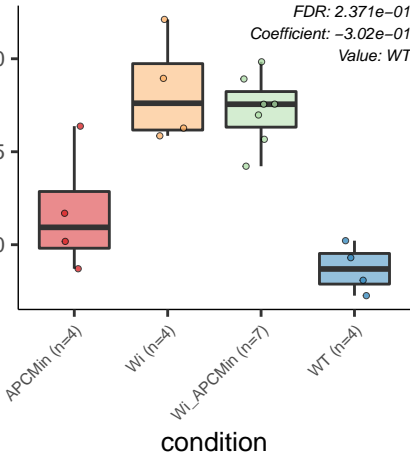


Porphyromonas

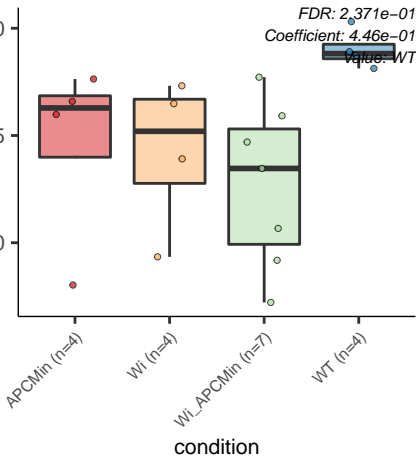
FDR: 2.371e-01

Coefficient: -3.02e-01

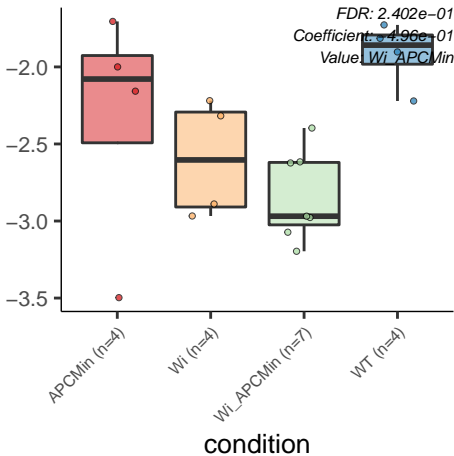
Value: WT



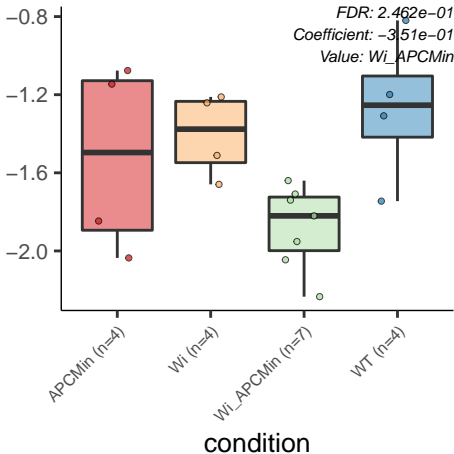
Clostridium_XIVb



Acetatifactor



Clostridium_IV



Alkaliphilus

FDR: 2.462e-01

Coefficient: 4.54e-01

Value: WT

