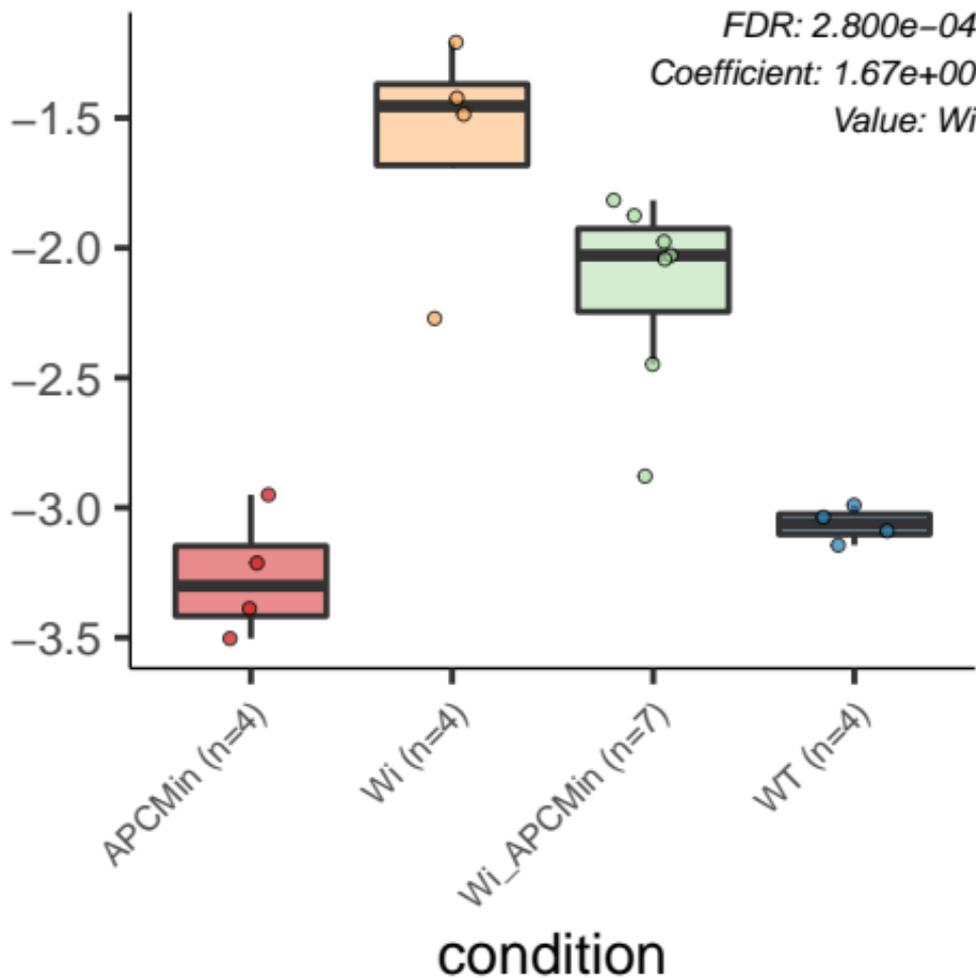
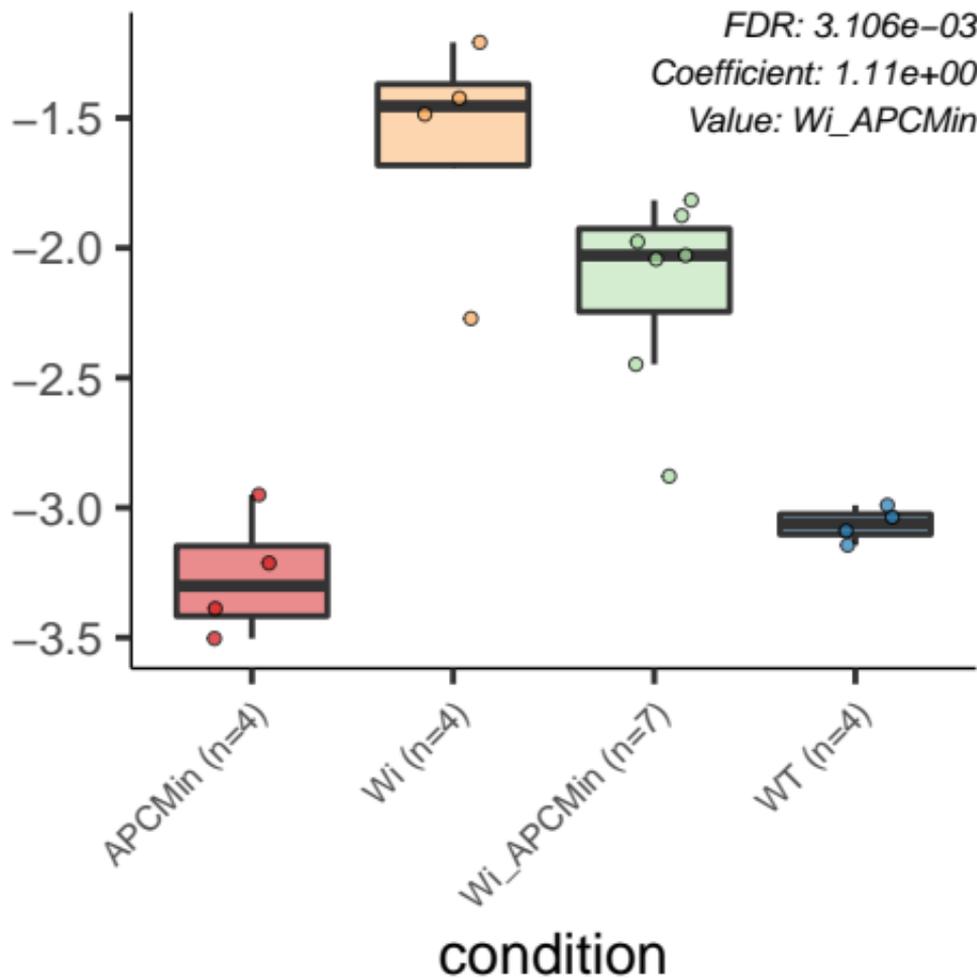


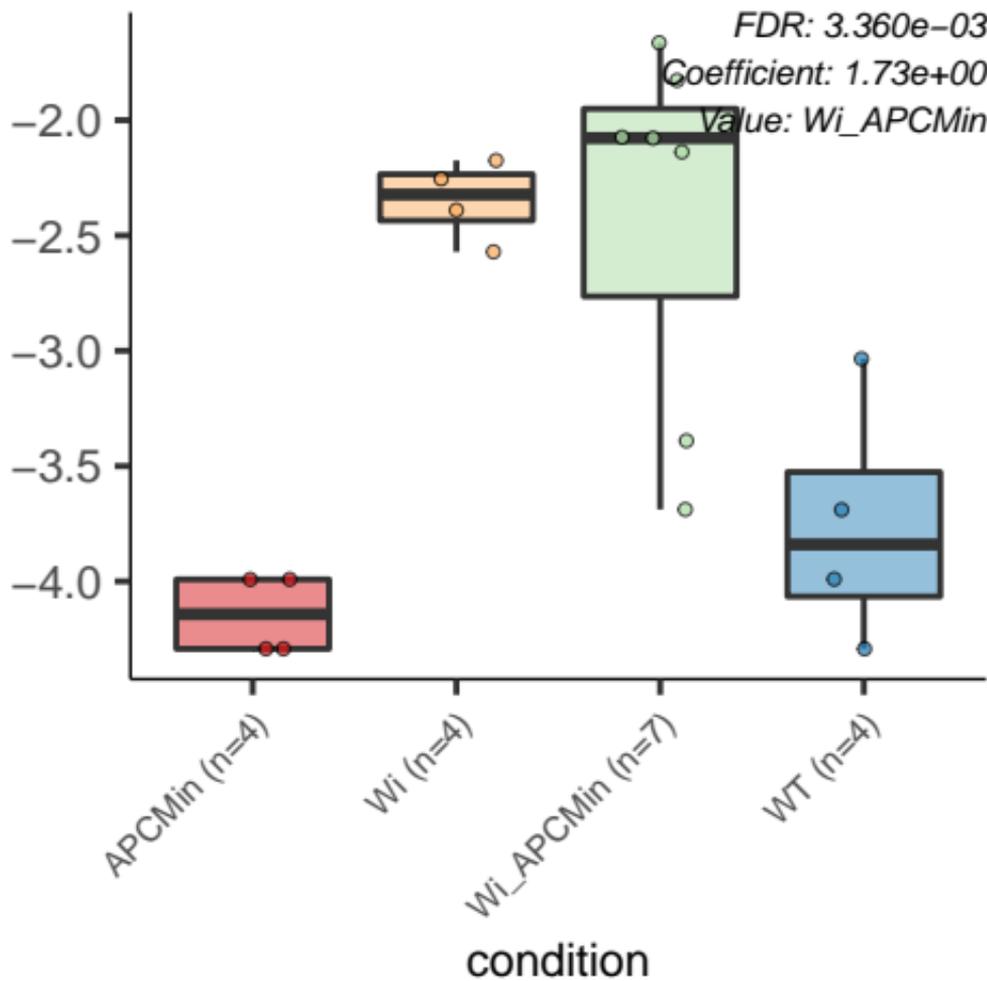
Eubacteriaceae



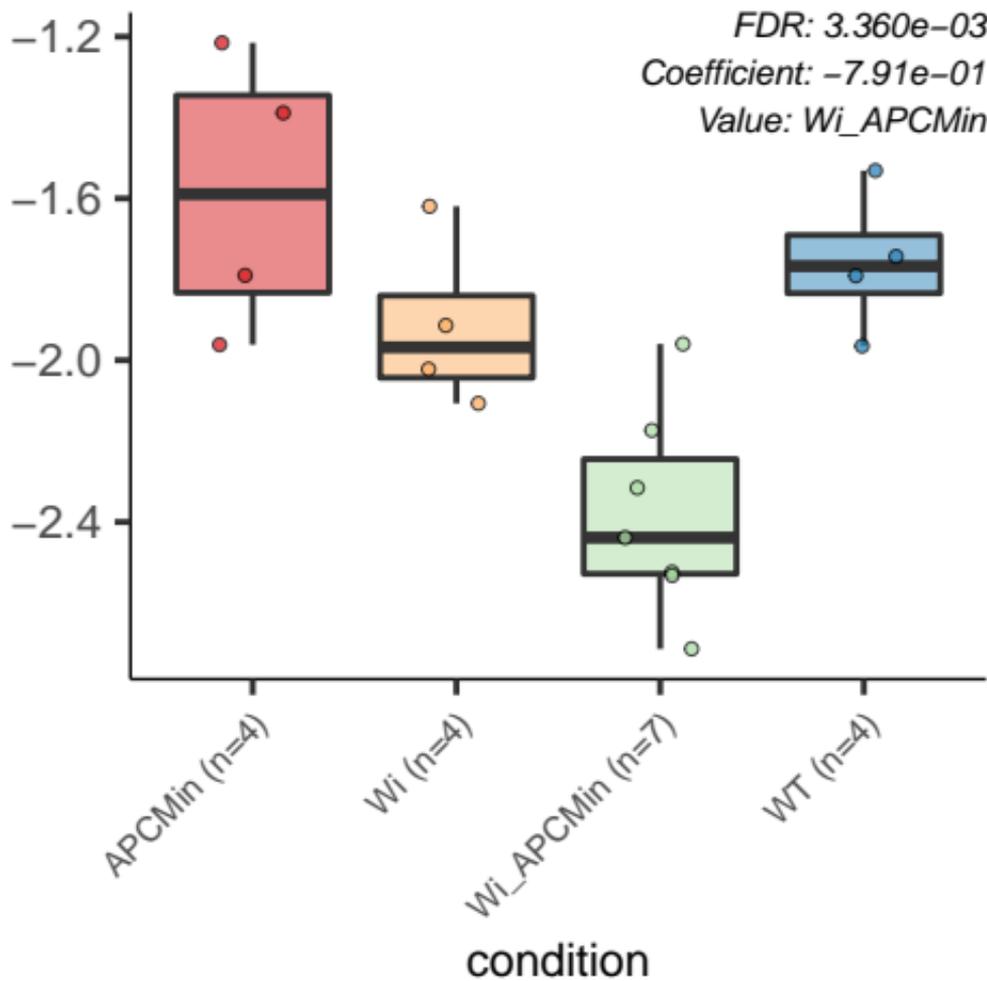
Eubacteriaceae



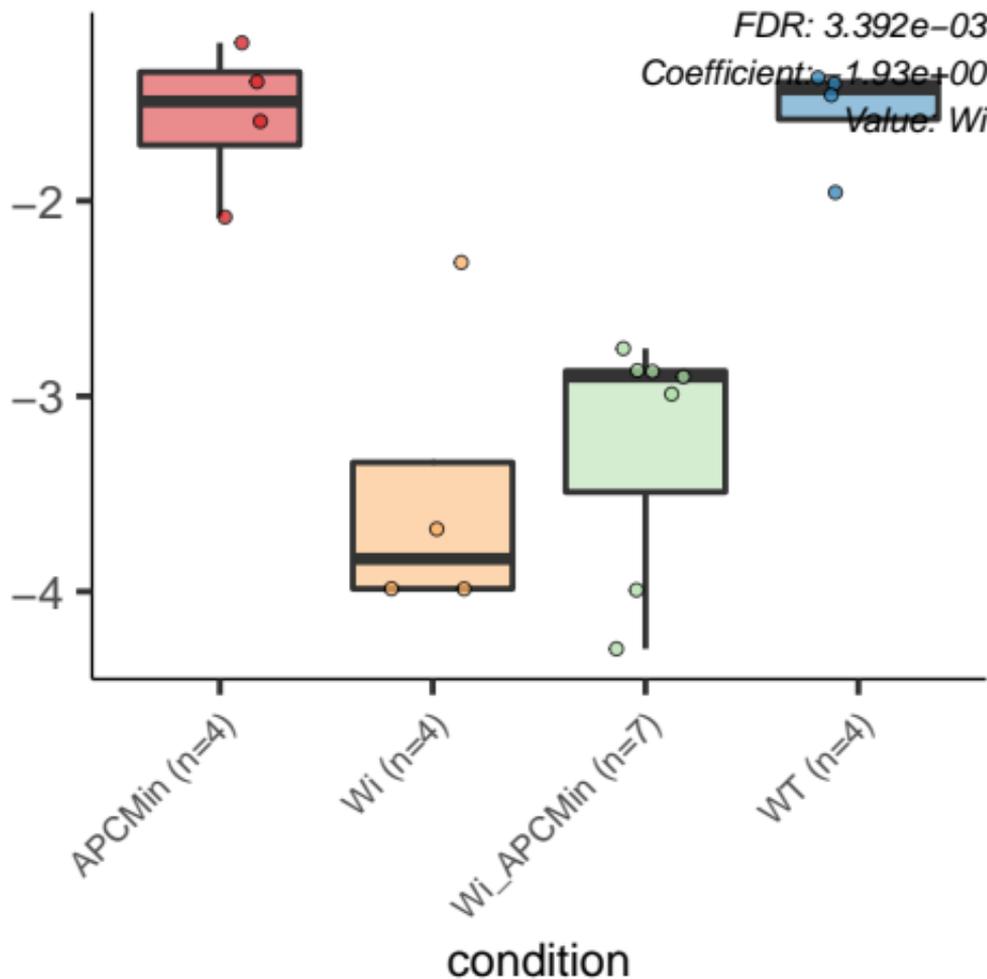
Acholeplasmataceae



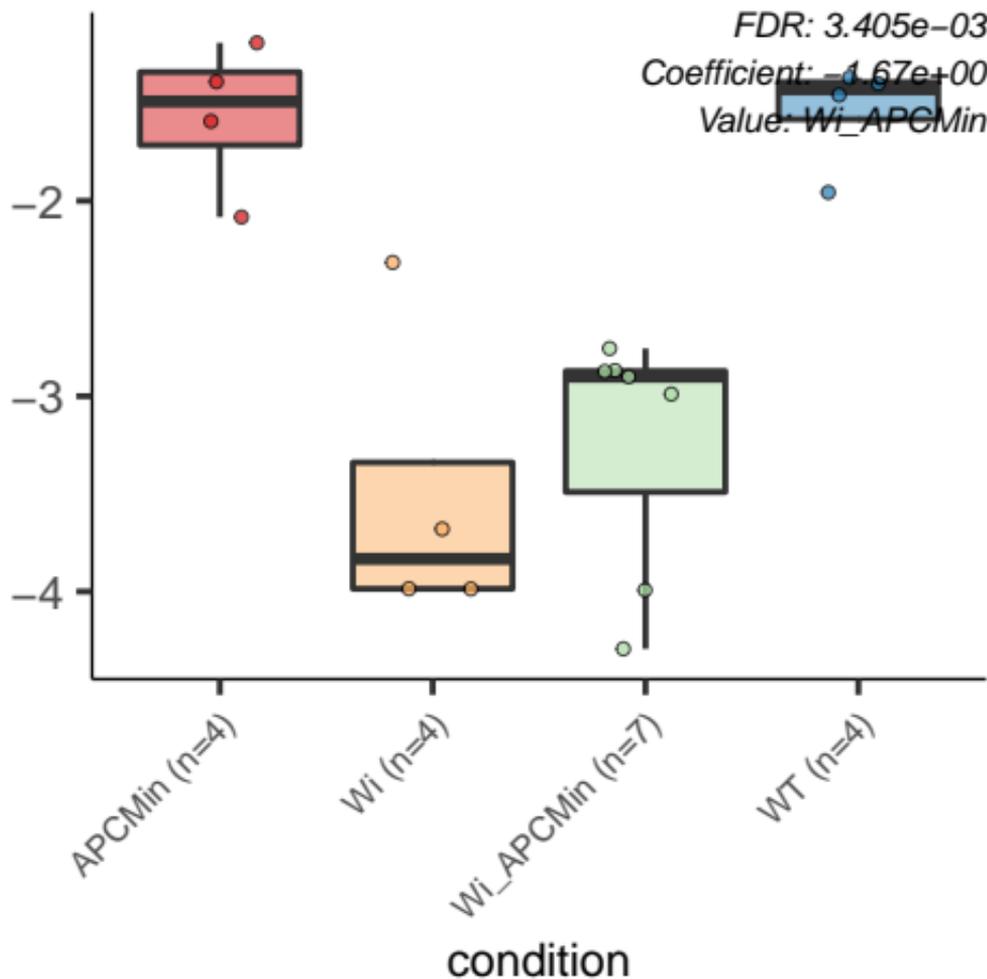
Lactobacillaceae



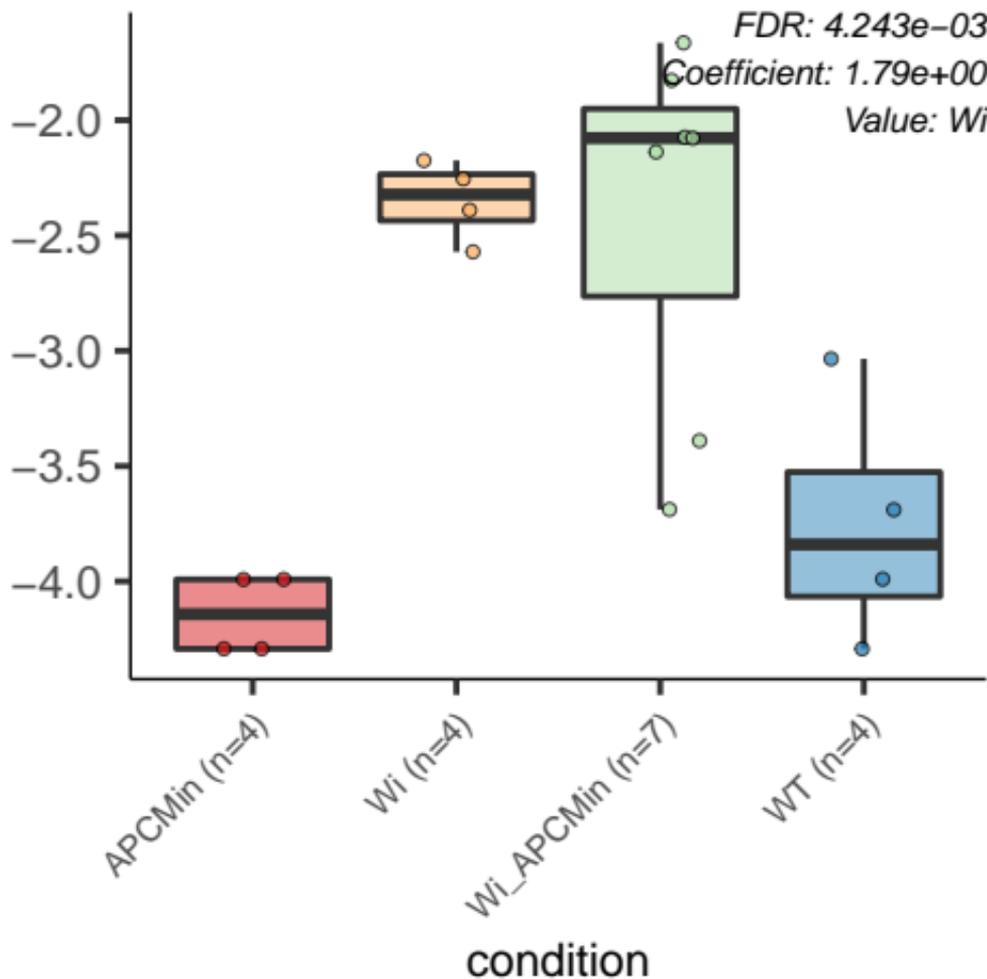
Deferribacteraceae

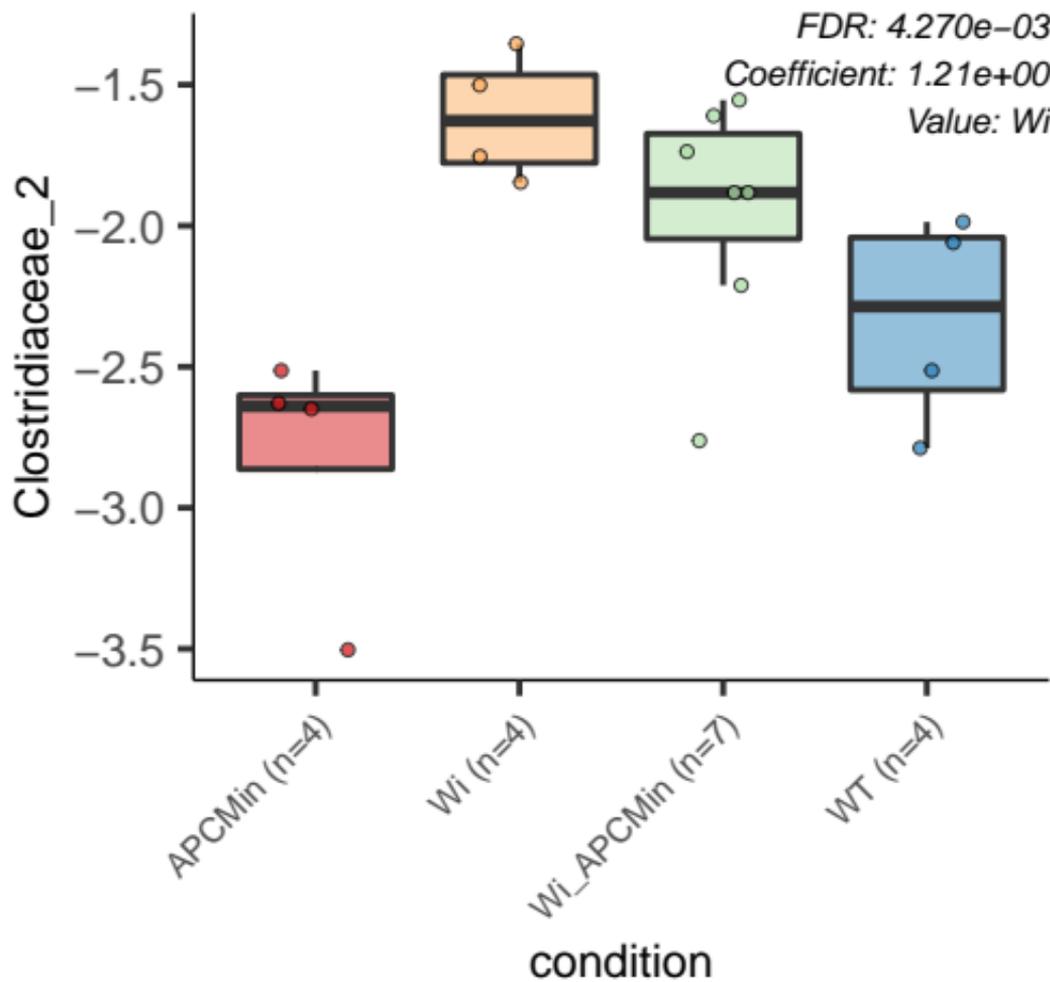


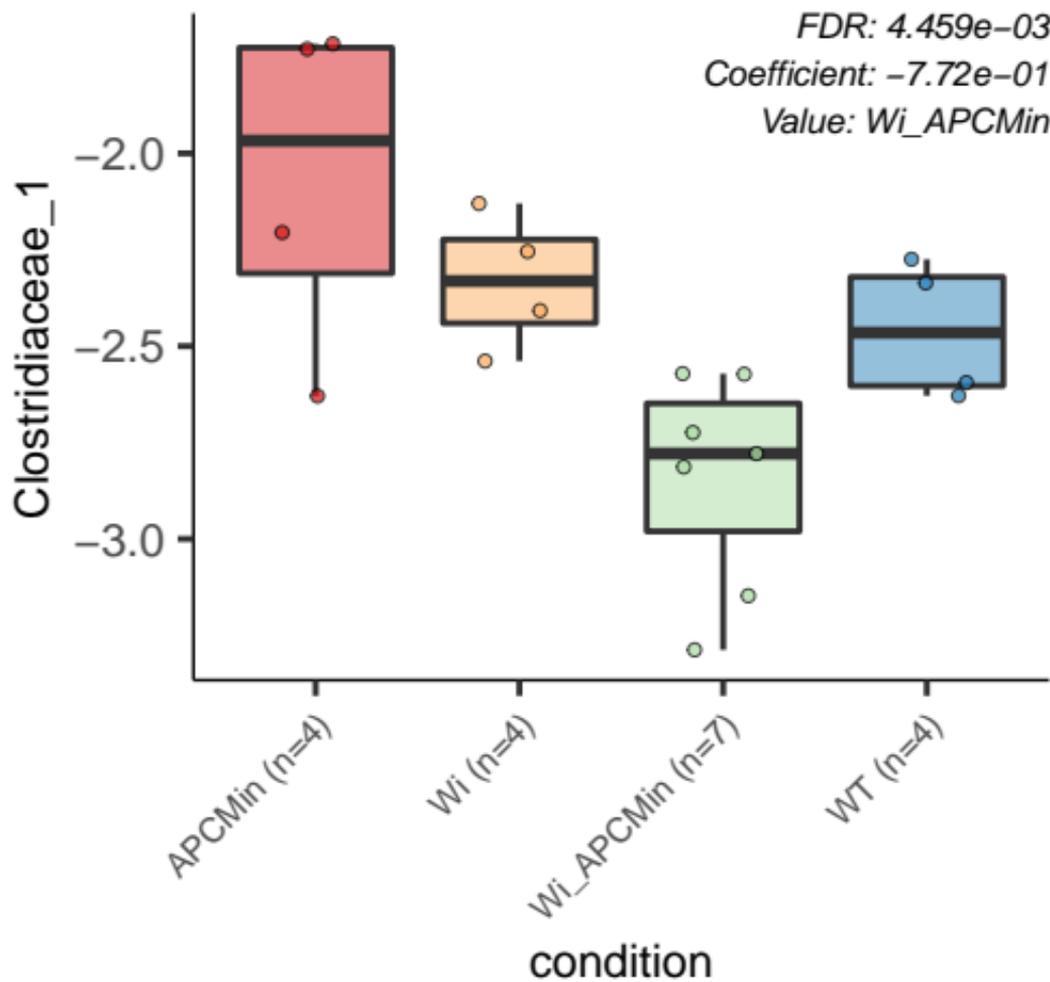
Deferribacteraceae



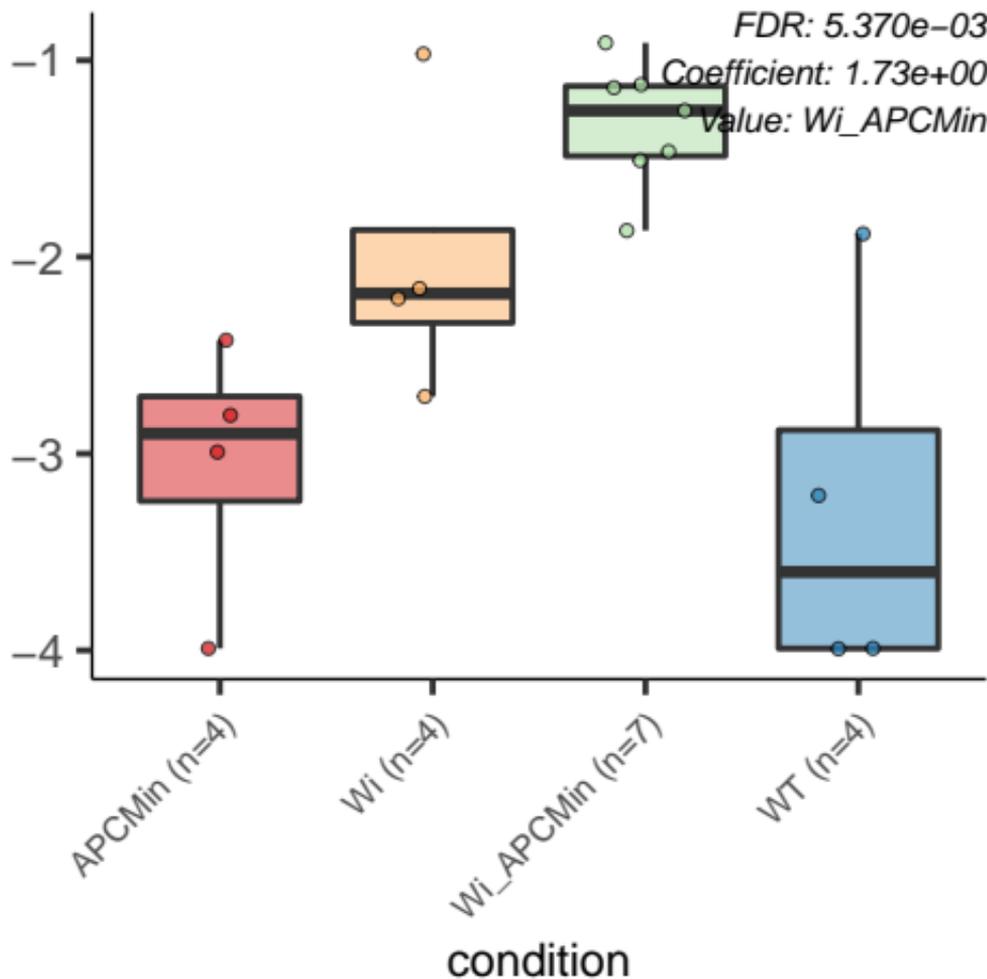
Acholeplasmataceae



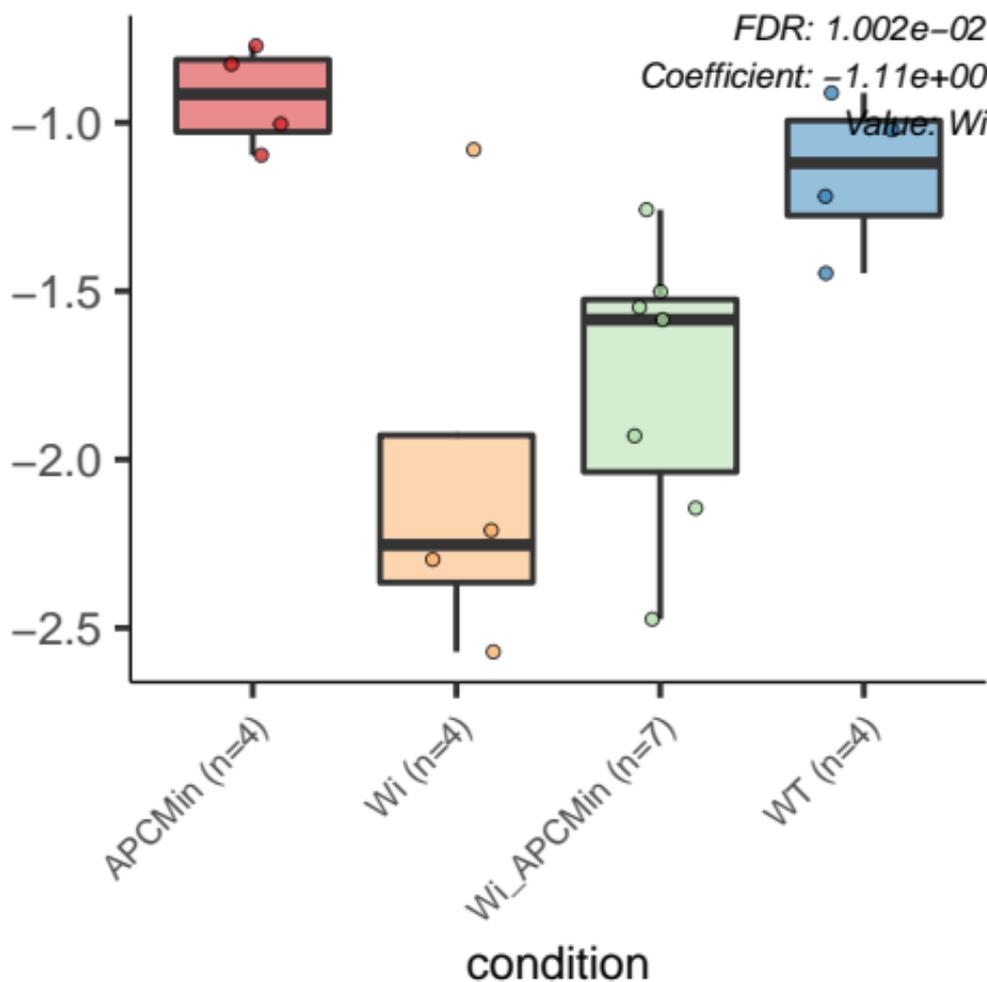




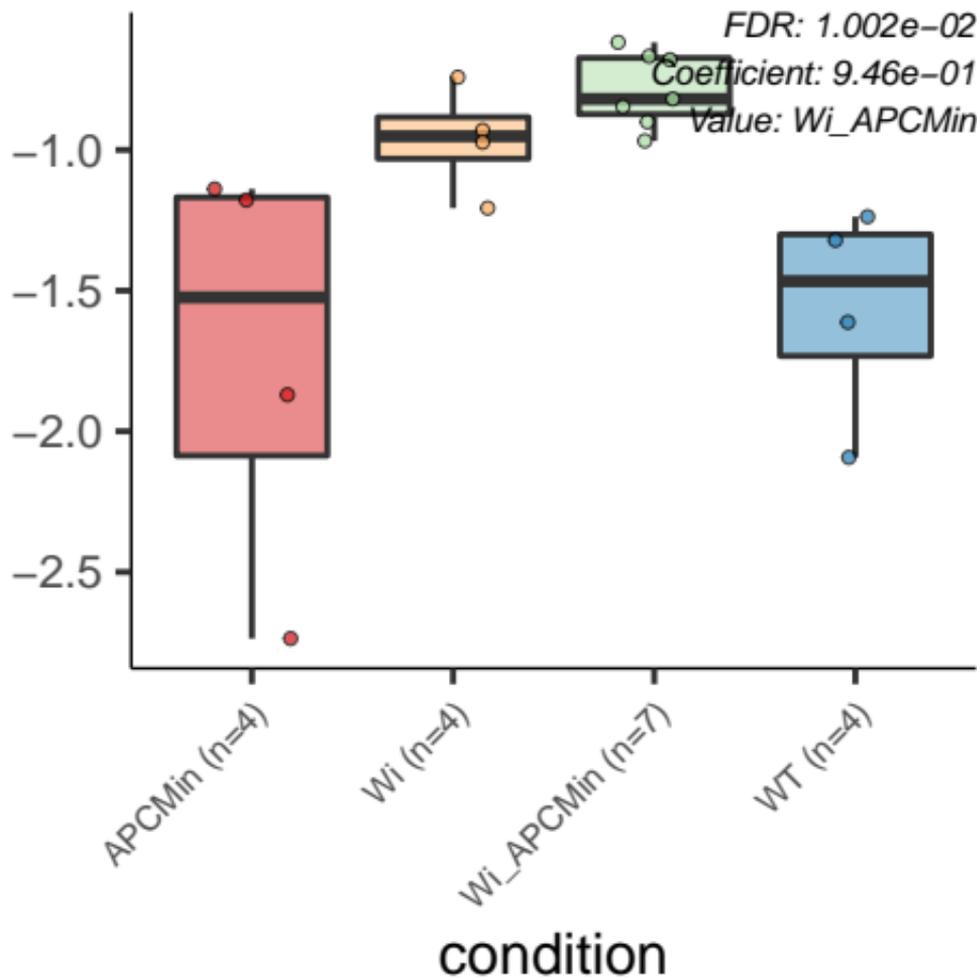
Verrucomicrobiaceae



Helicobacteraceae



Prevotellaceae



Coriobacteriaceae

-2.0
-2.5
-3.0

FDR: 1.340e-02
Coefficient: -5.58e-01
Value: W_i_APCMin

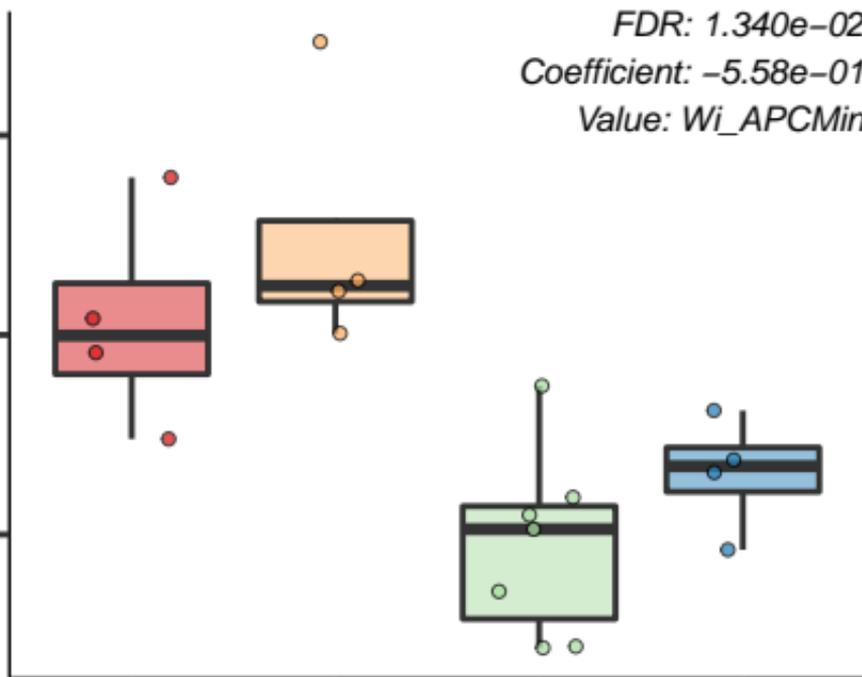
APCMin (n=4)

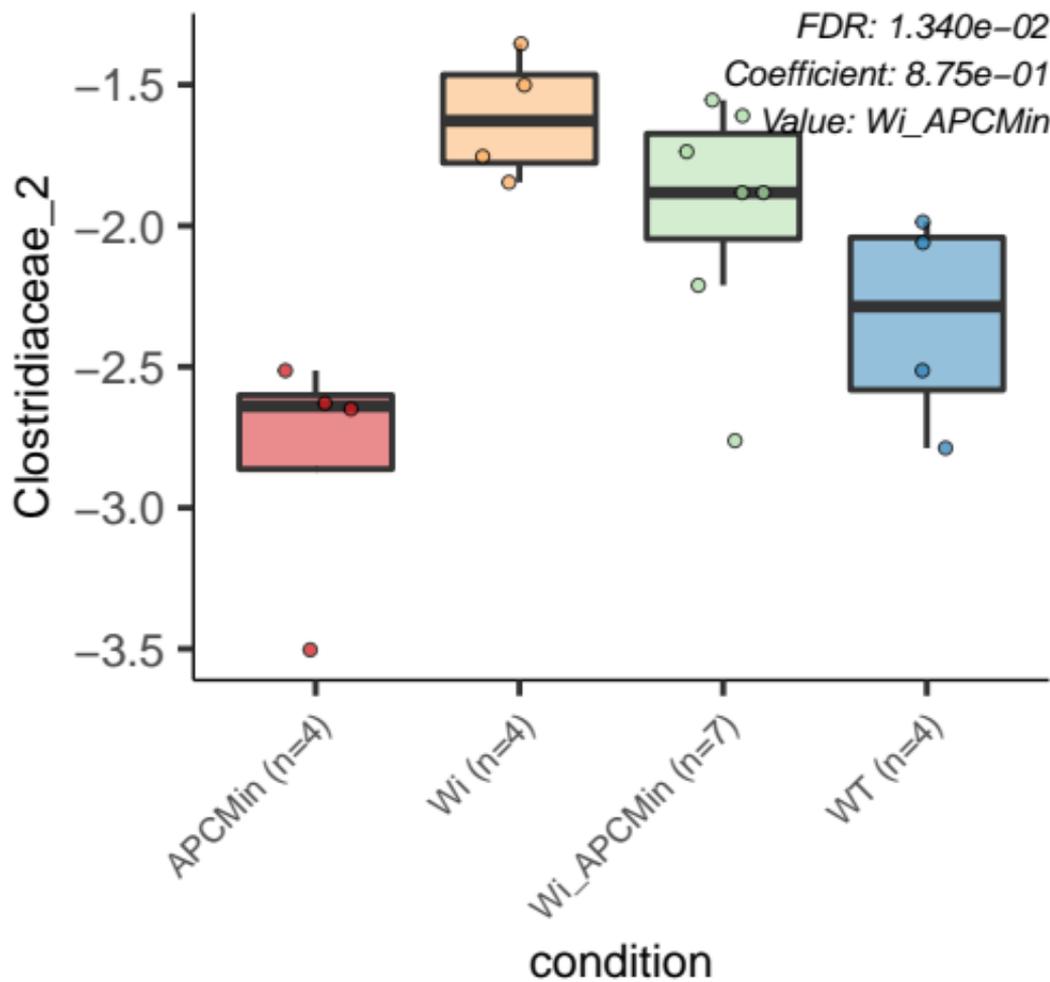
Wi (n=4)

Wi_APCMin (n=7)

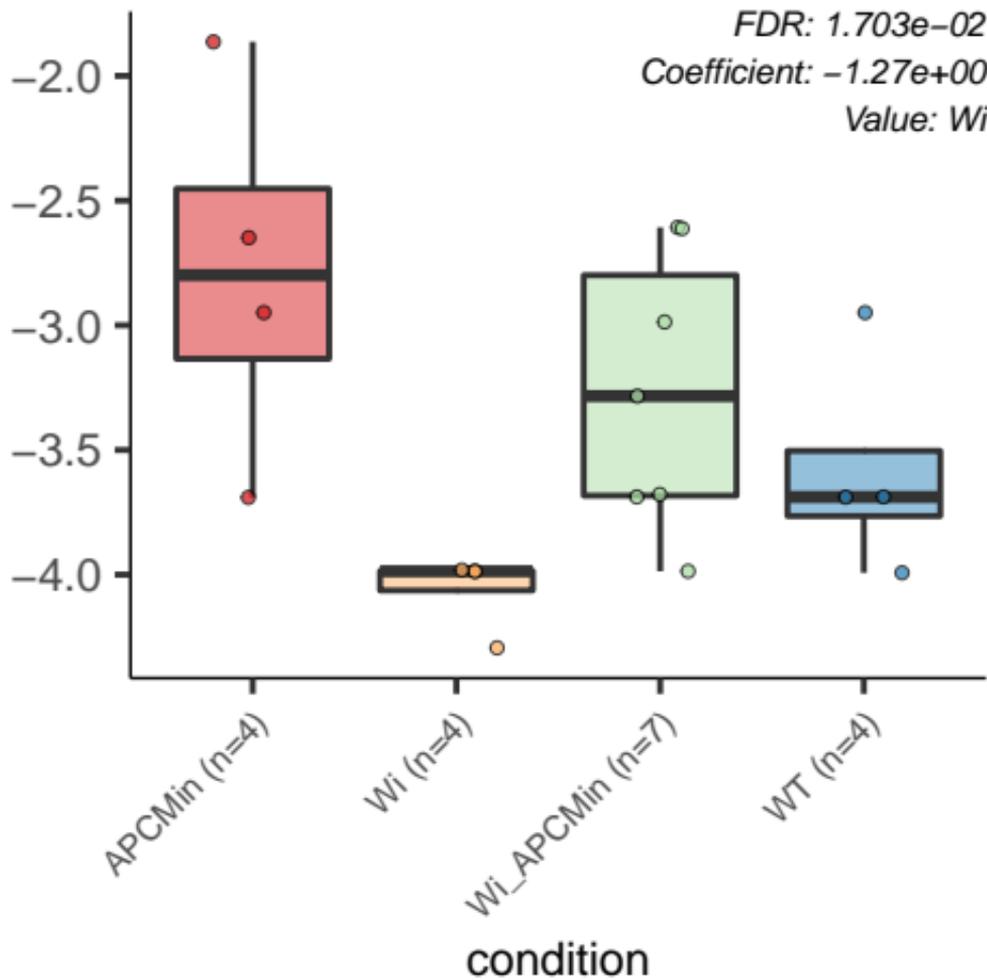
WT (n=4)

condition

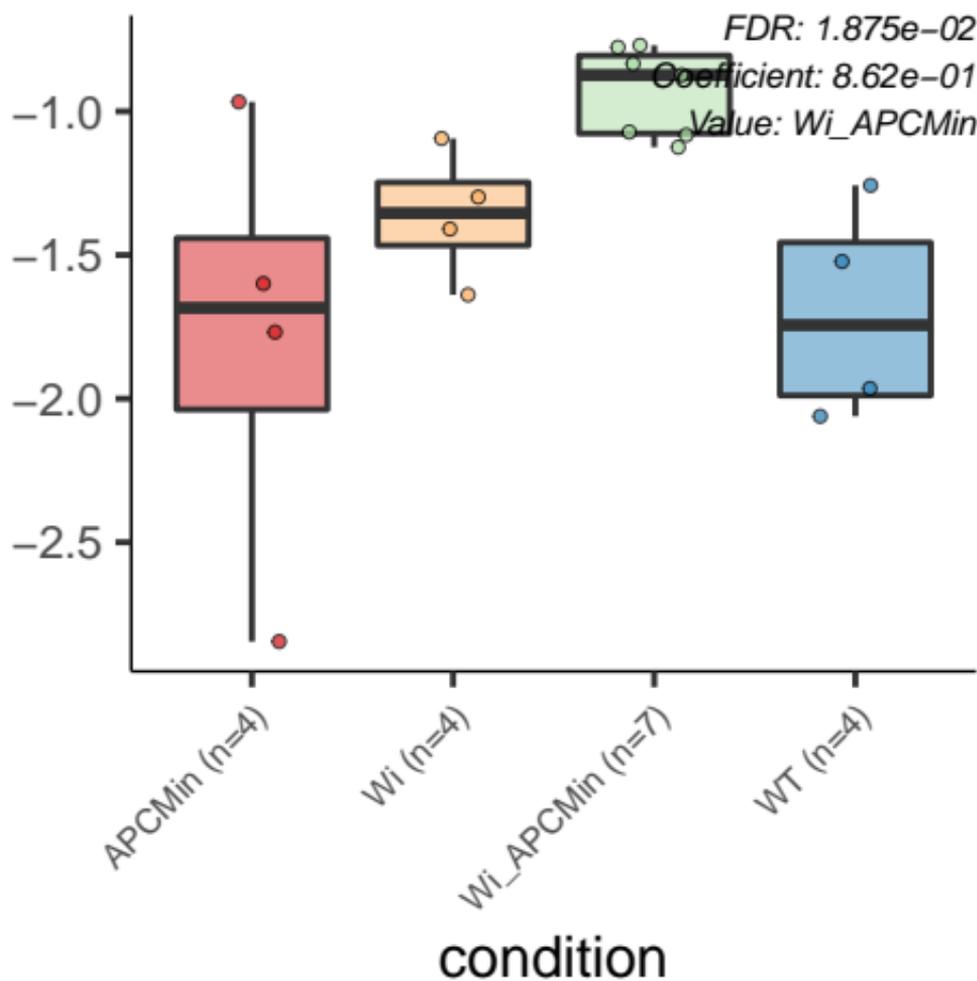




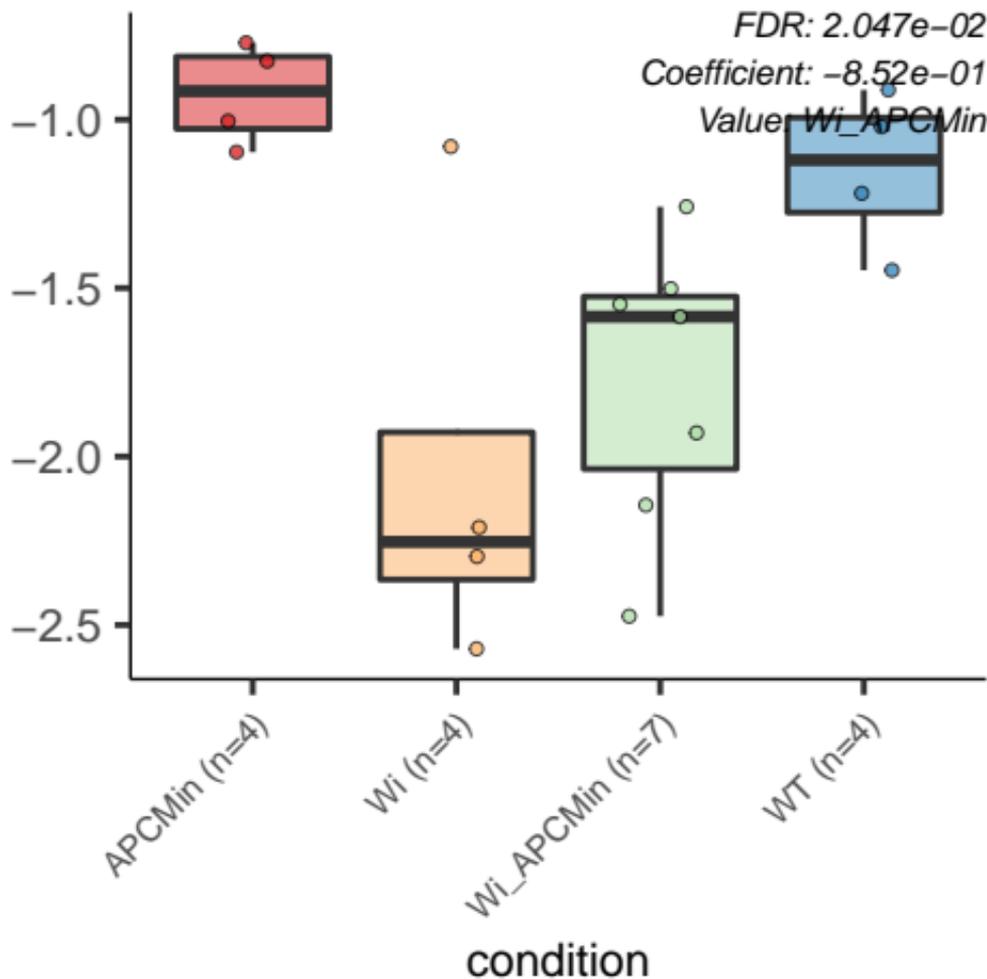
Enterobacteriaceae



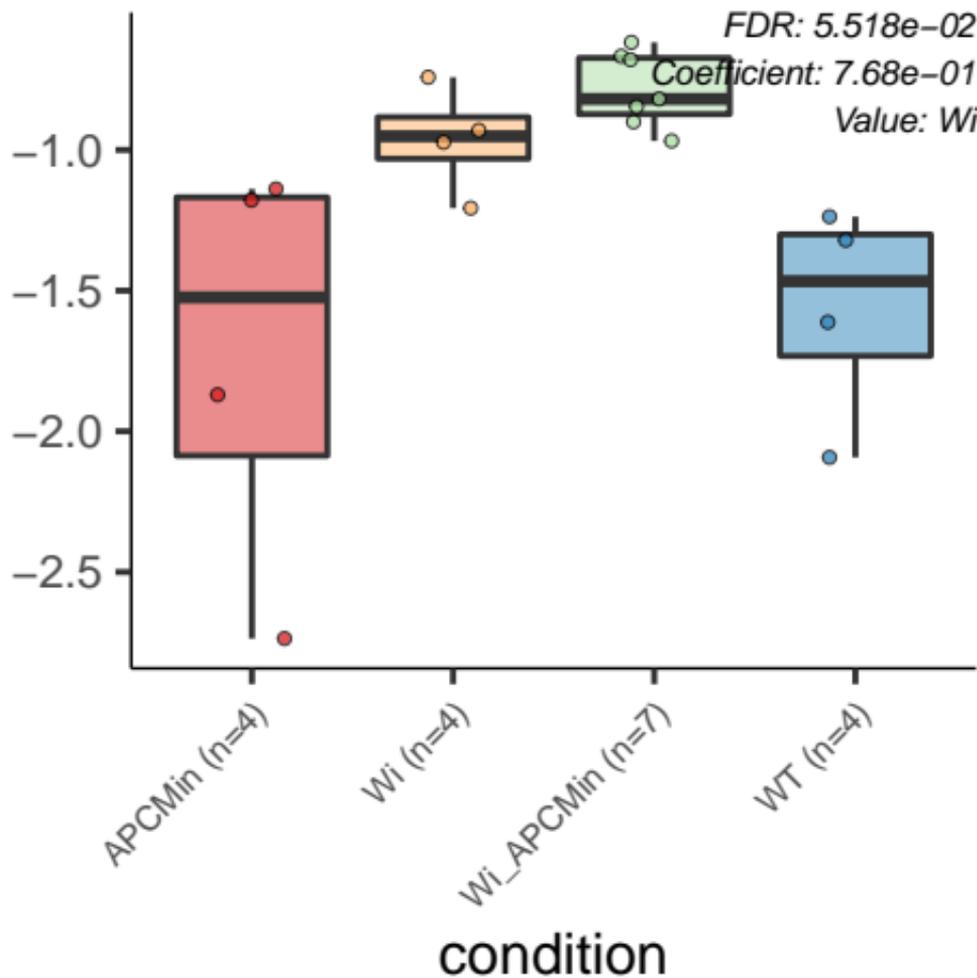
Bacteroidaceae



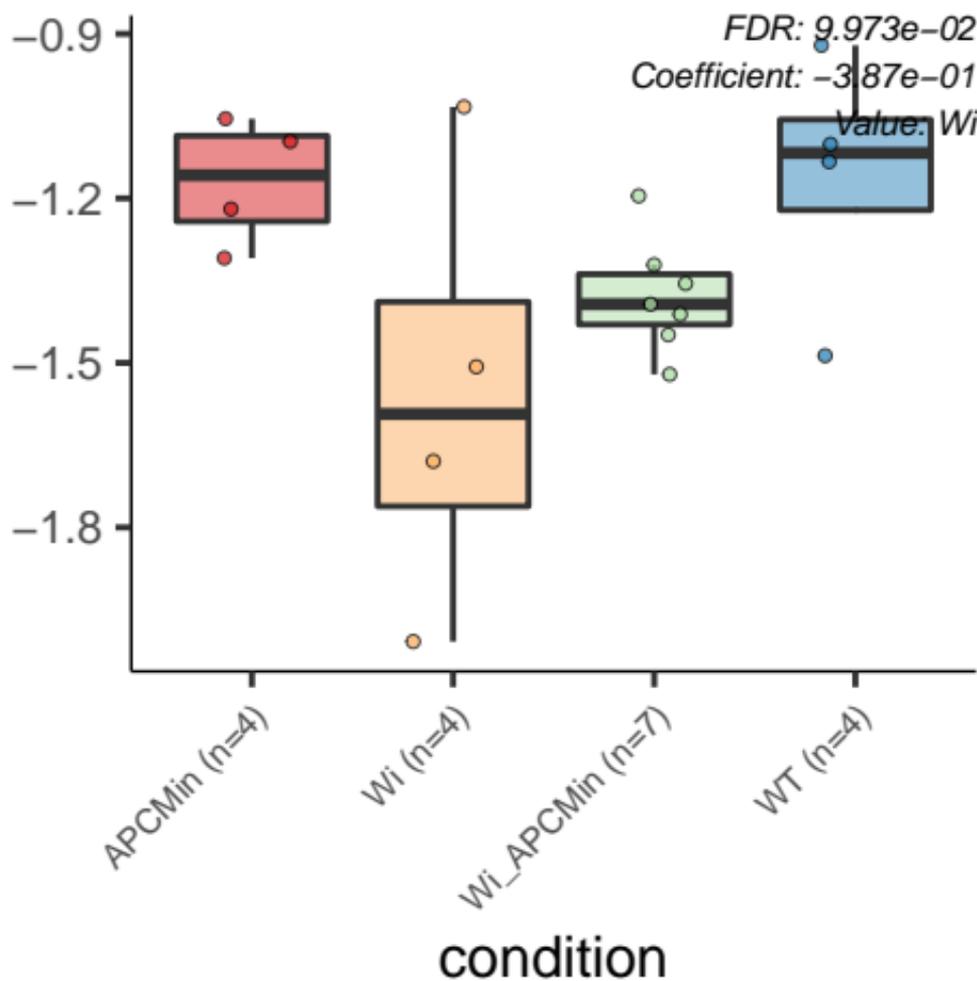
Helicobacteraceae



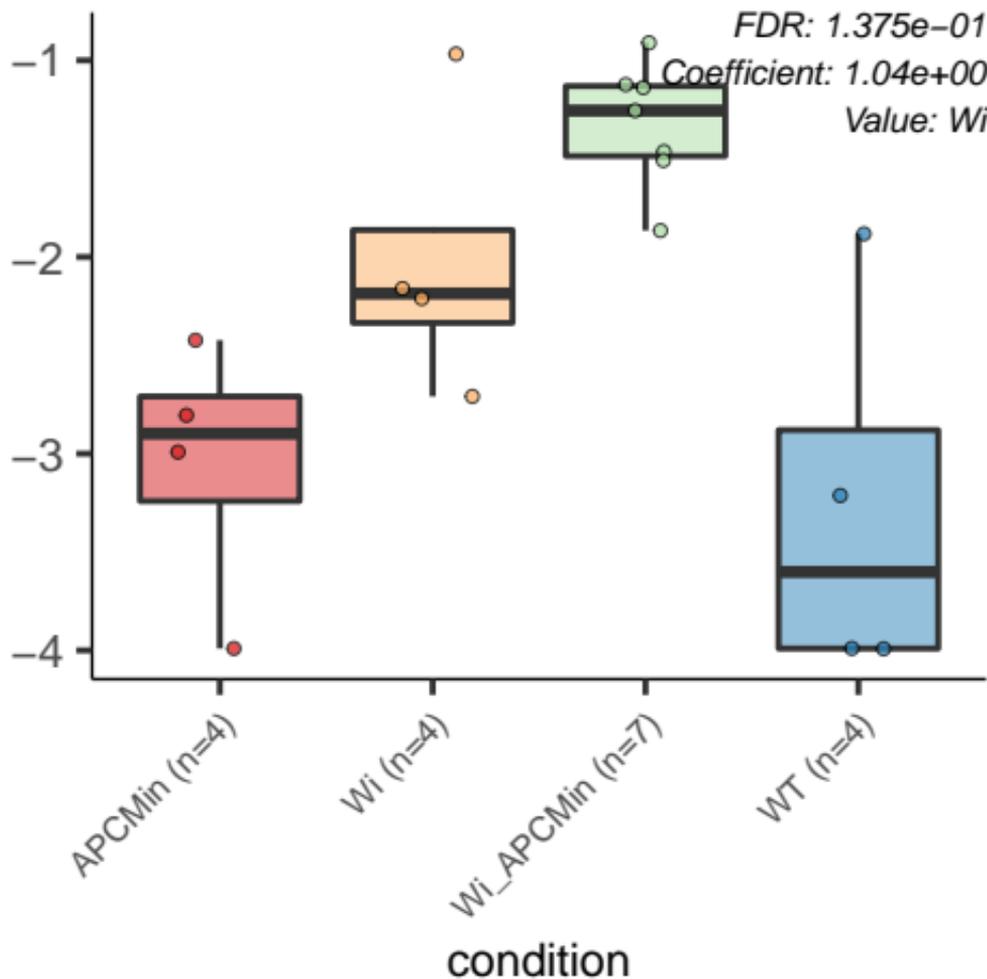
Prevotellaceae



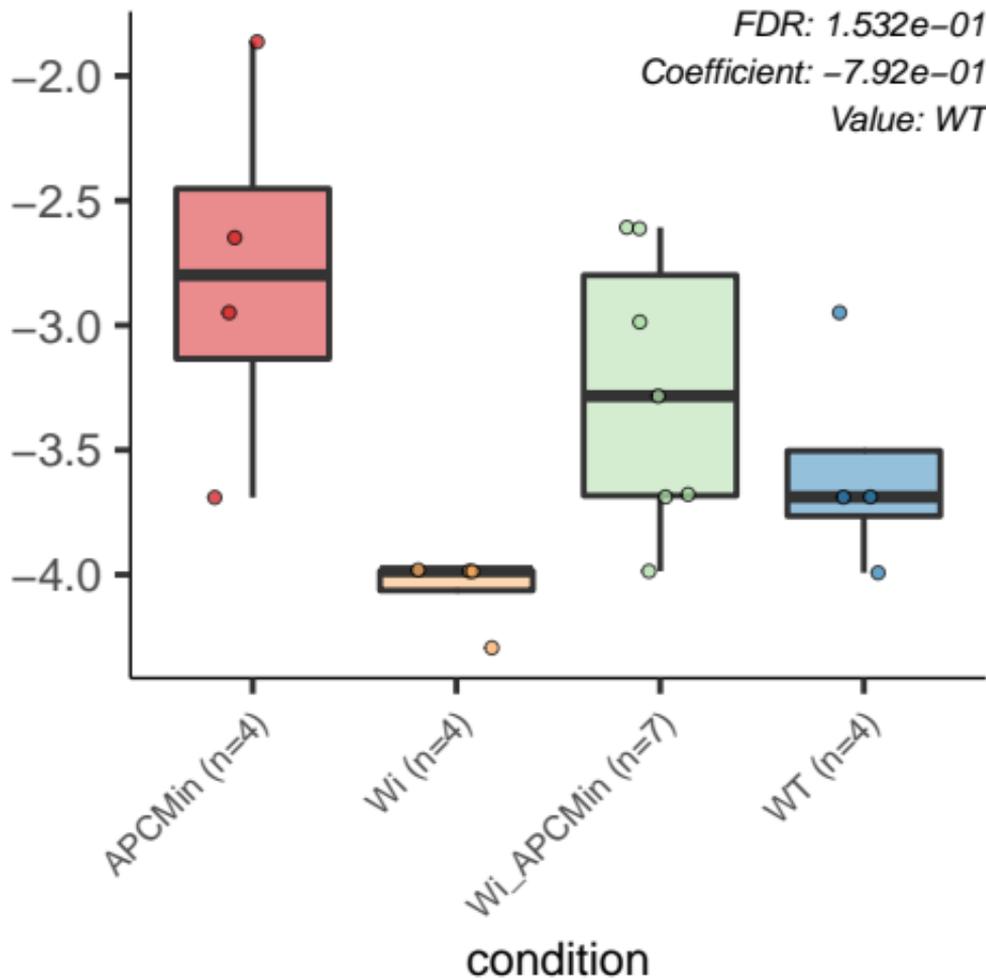
Rikenellaceae



Verrucomicrobiaceae



Enterobacteriaceae



Coriobacteriaceae

-2.0
-2.5
-3.0

FDR: 1.532e-01
Coefficient: -3.79e-01
Value: WT

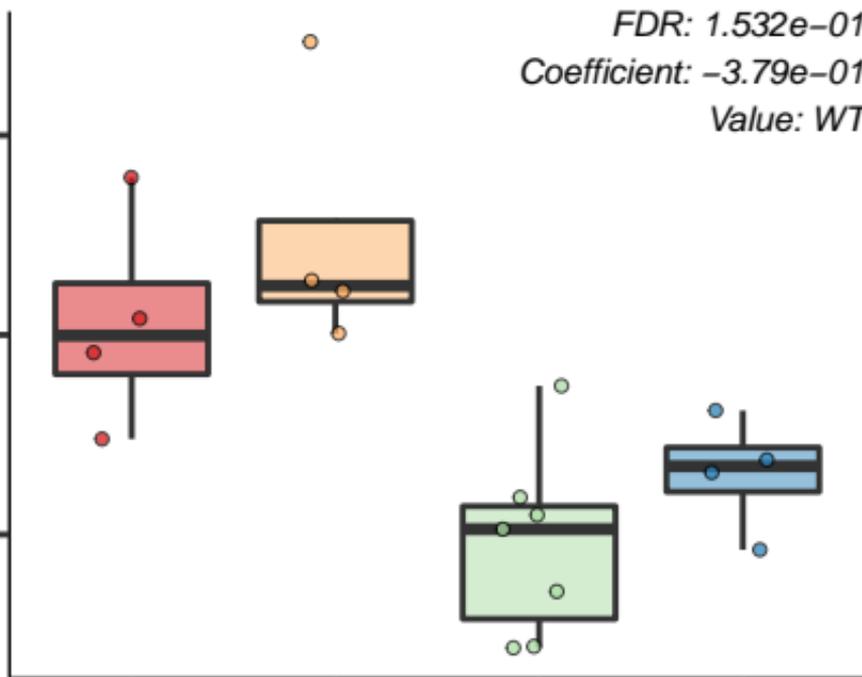
APCMin (n=4)

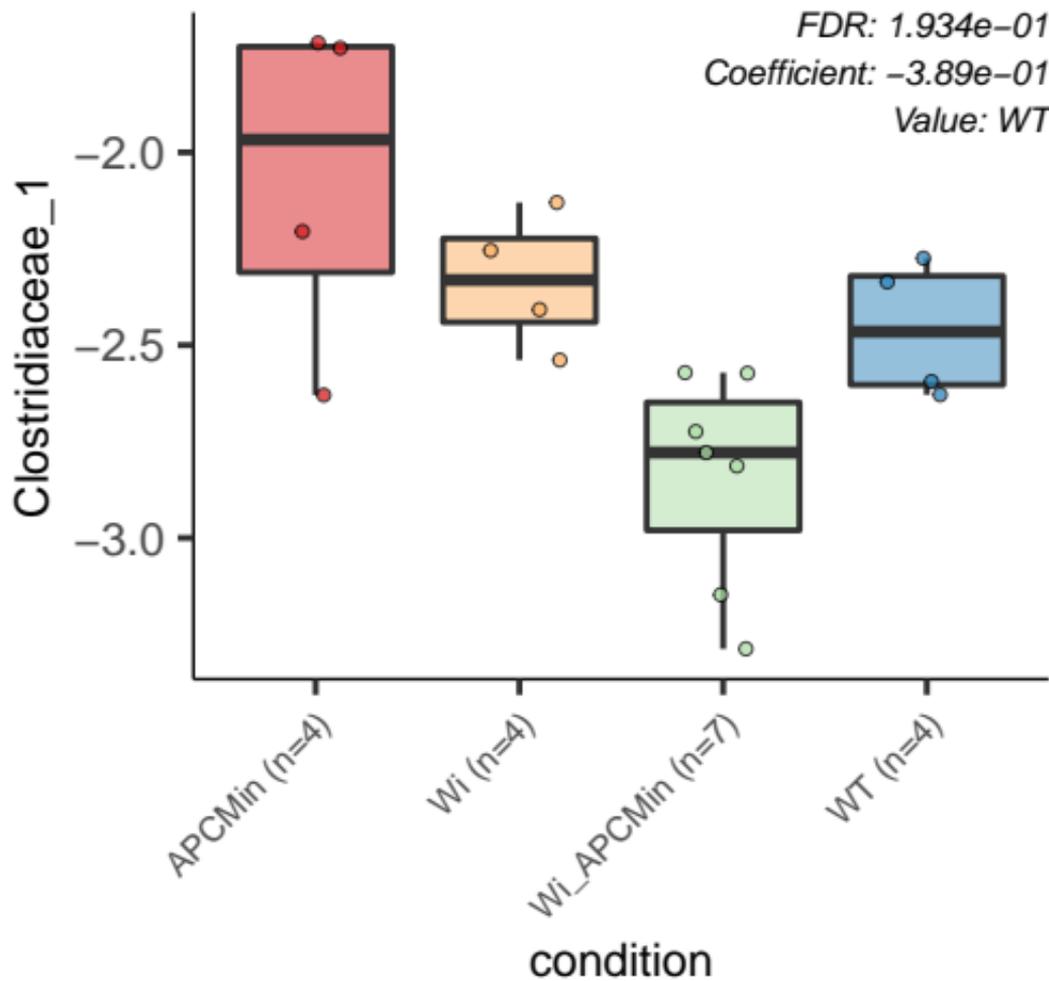
Wi (n=4)

Wi_APCMin (n=7)

WT (n=4)

condition





Cyclobacteriaceae

-2.0
-2.5
-3.0

APCMin (n=4)

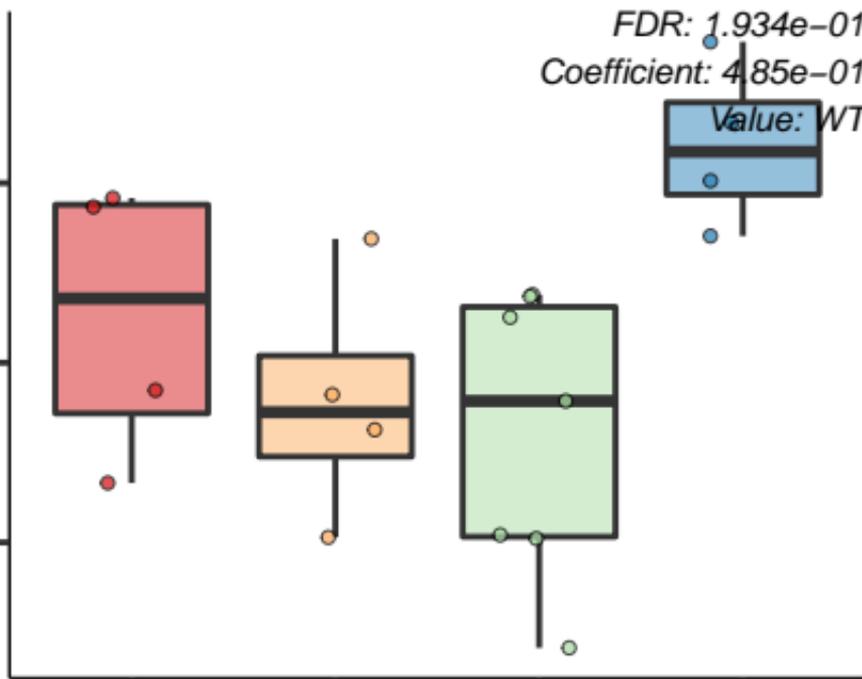
Wi (n=4)

Wi_APCMin (n=7)

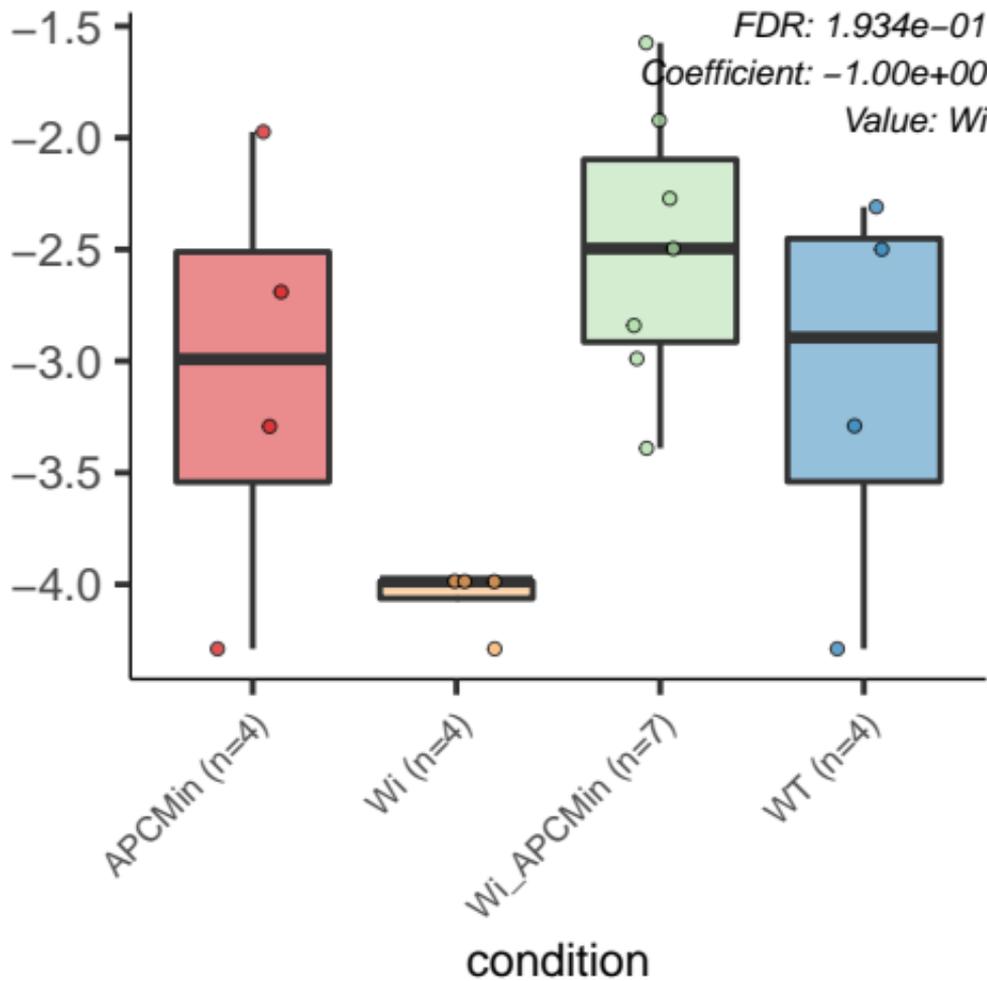
WT (n=4)

FDR: $1.934e-01$
Coefficient: $4.85e-01$
Value: WT

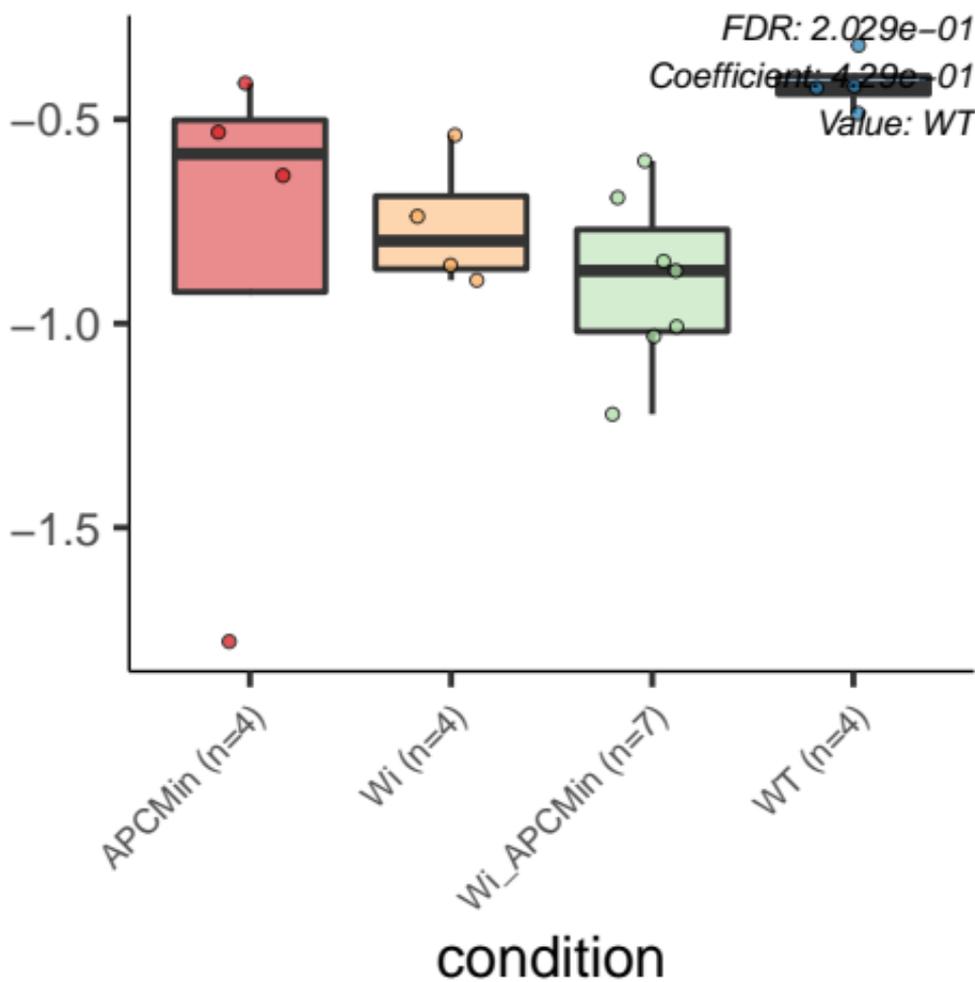
condition



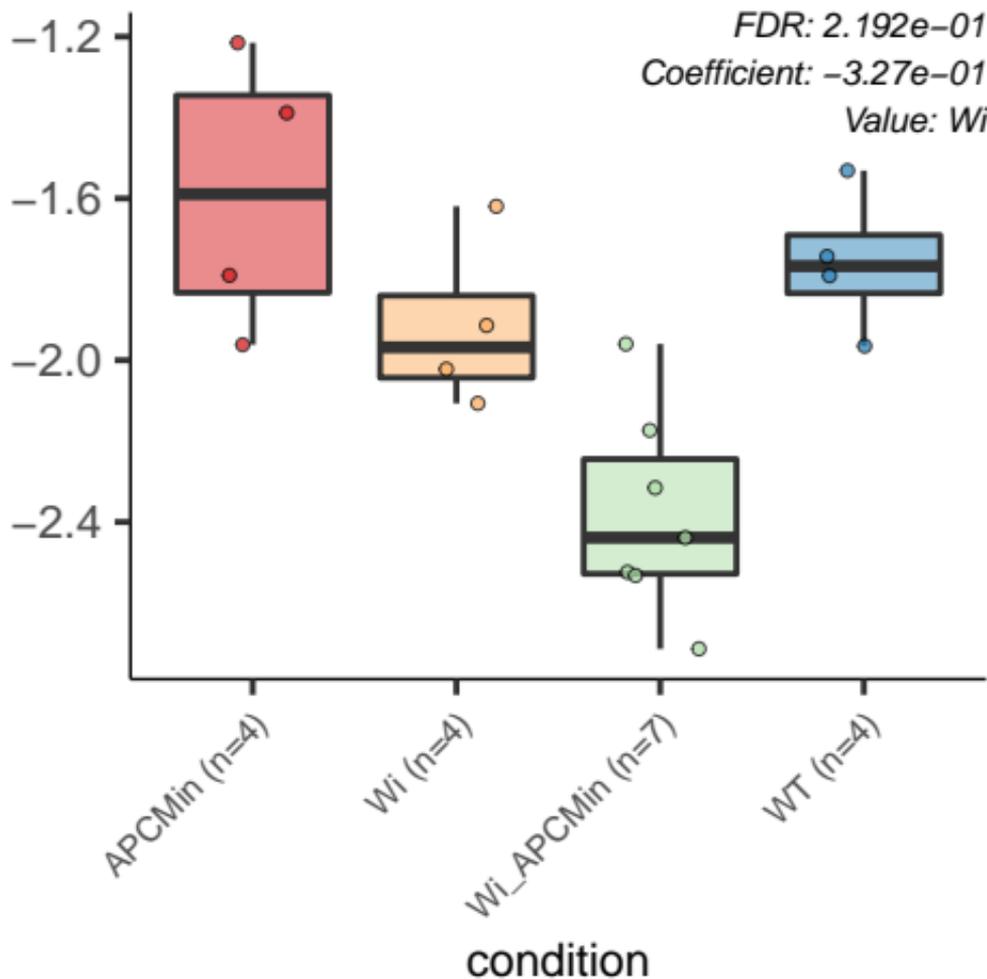
Anaeroplasmataceae

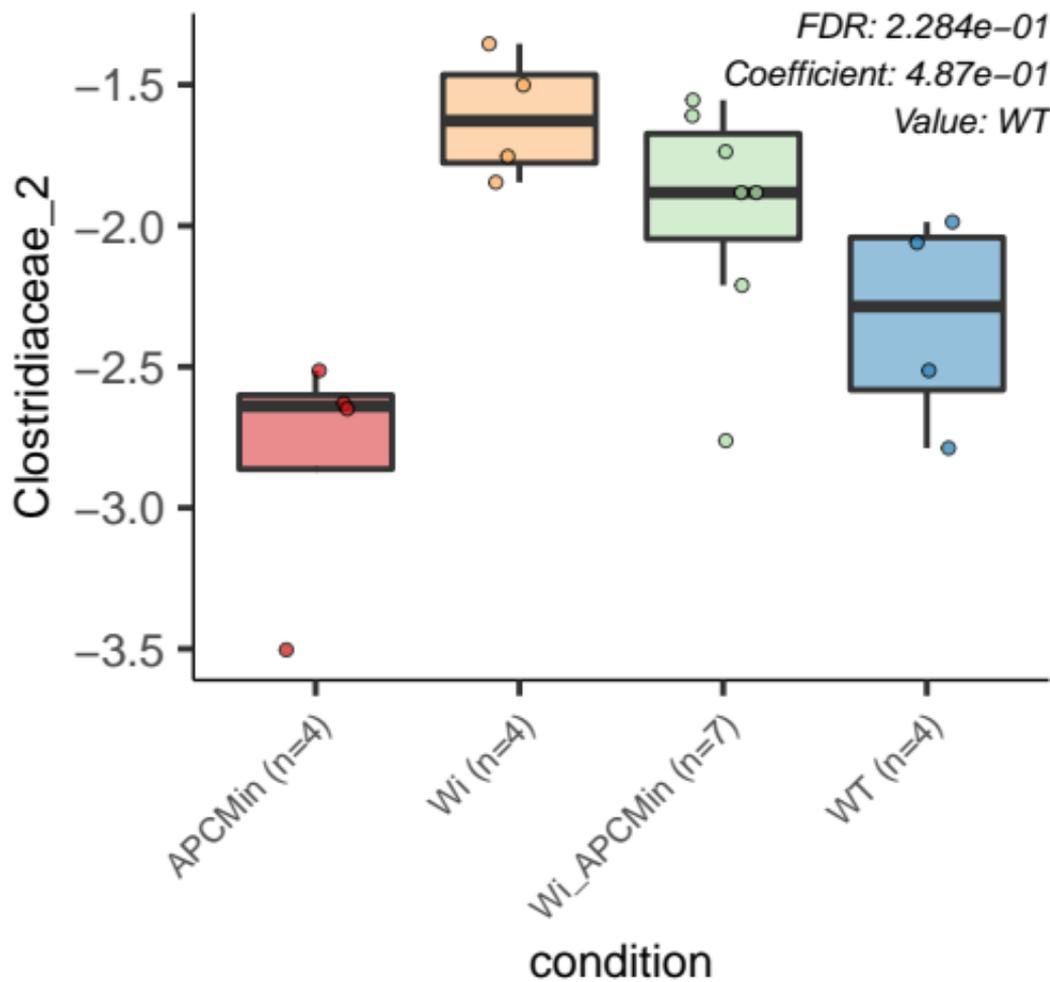


Lachnospiraceae



Lactobacillaceae





Desulfovibrionaceae

-1.6
-2.0
-2.4

APCMin (n=4)

Wi (n=4)

Wi_APCMin (n=7)

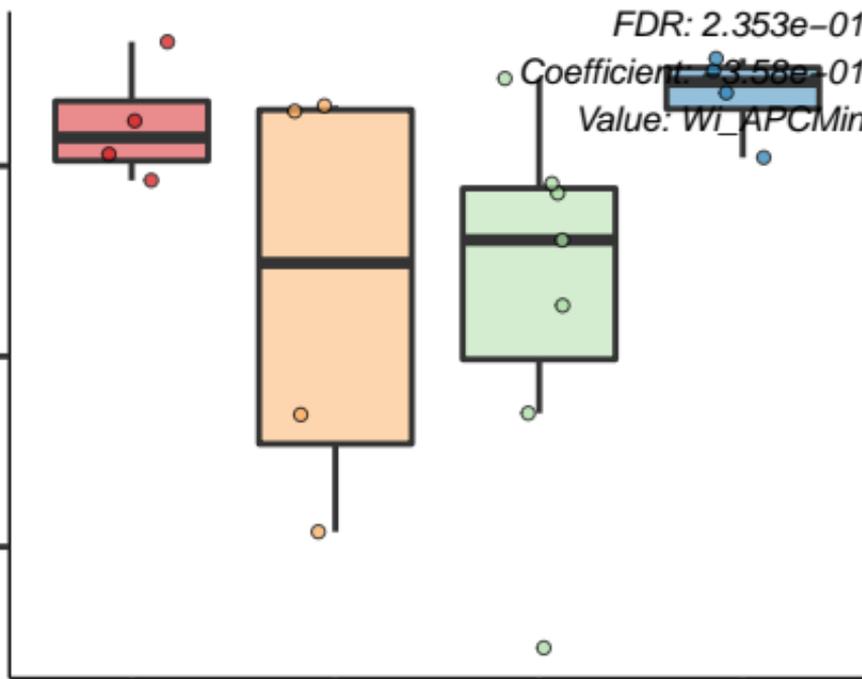
WT (n=4)

condition

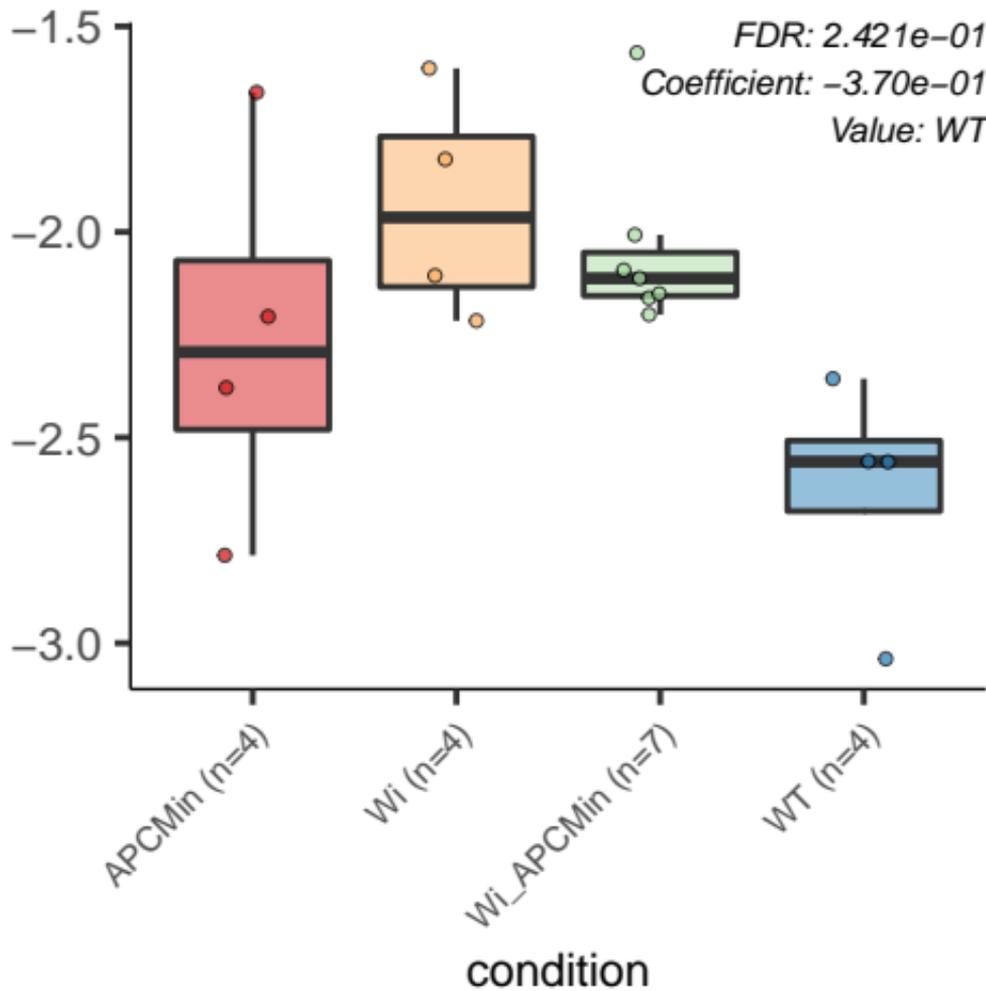
FDR: $2.353e-01$

Coefficient: $-3.58e-01$

Value: *Wi_APCMin*



Erysipelotrichaceae



Porphyromonadaceae

