

X87_Alistipes.

FDR: 3.600e-04

Coefficient: 2.87e+00

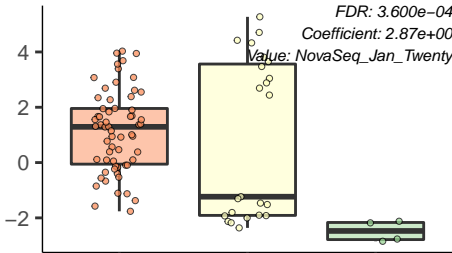
Value: NovaSeq_Jan_Twenty

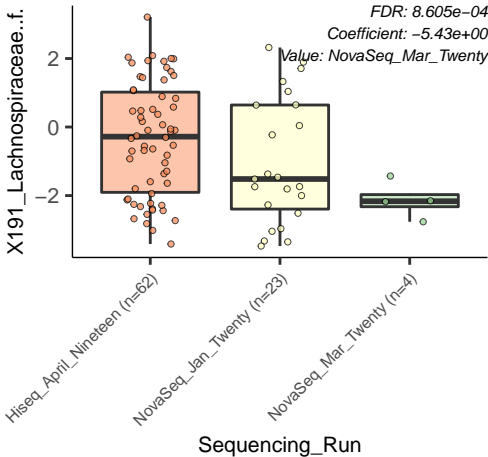
Hiseq_April_Nineteen (n=62)

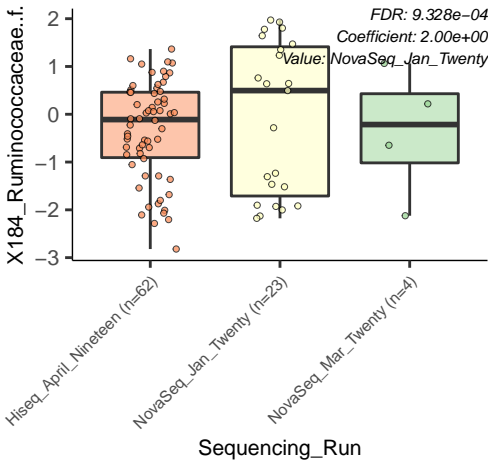
NovaSeq_Jan_Twenty (n=23)

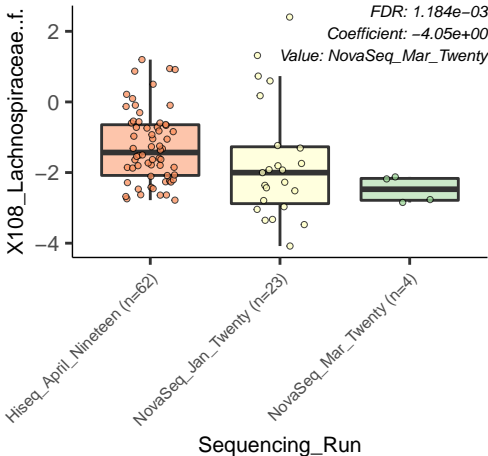
NovaSeq_Mar_Twenty (n=4)

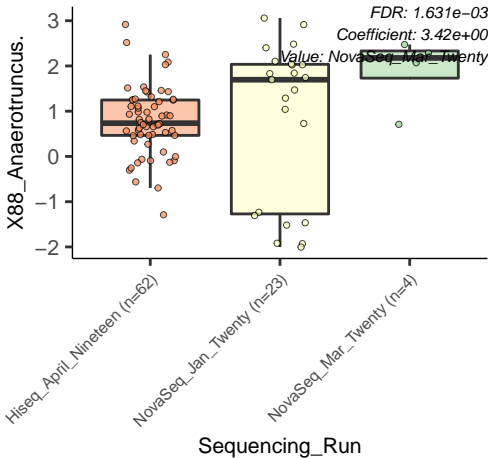
Sequencing_Run

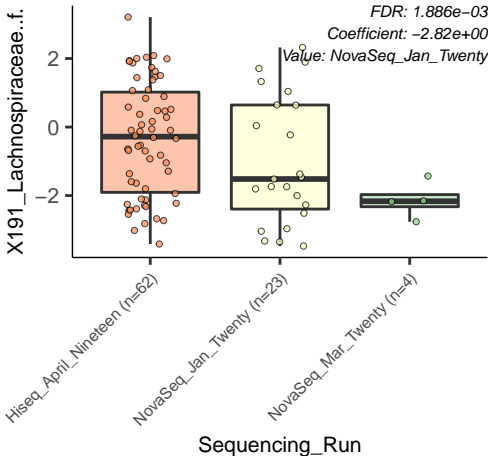


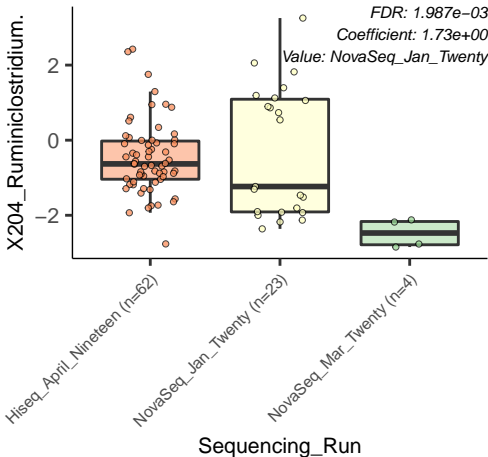


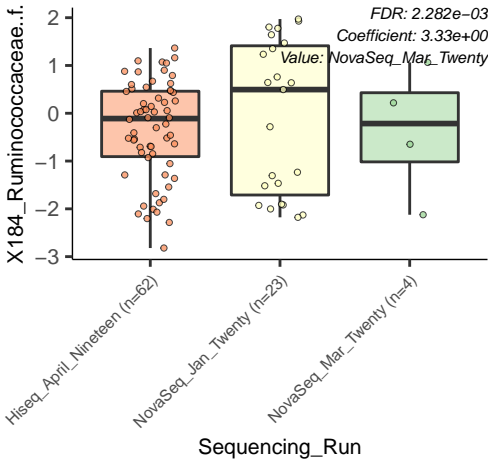


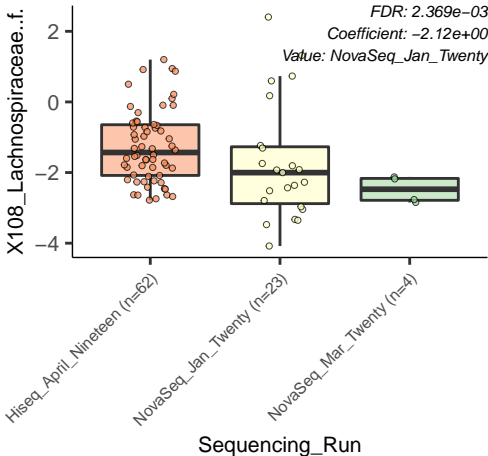


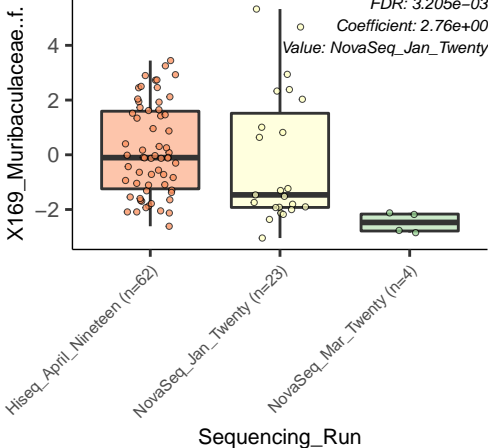


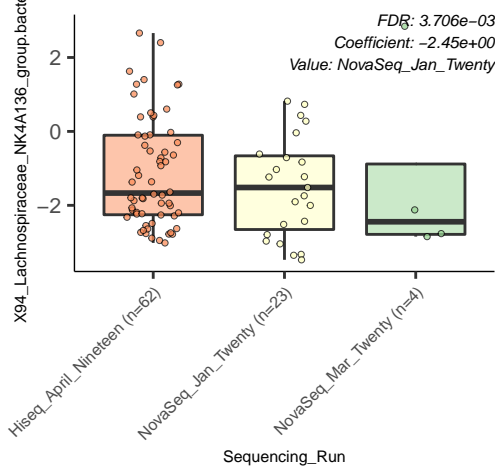


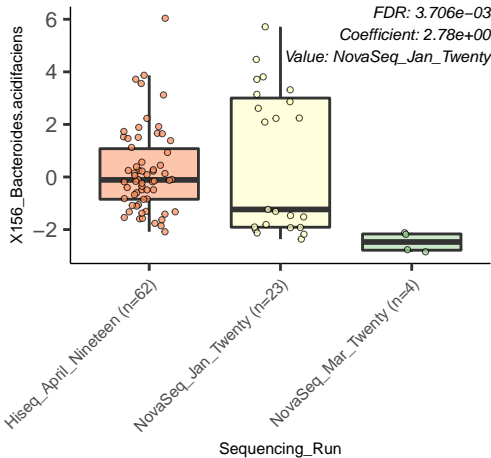


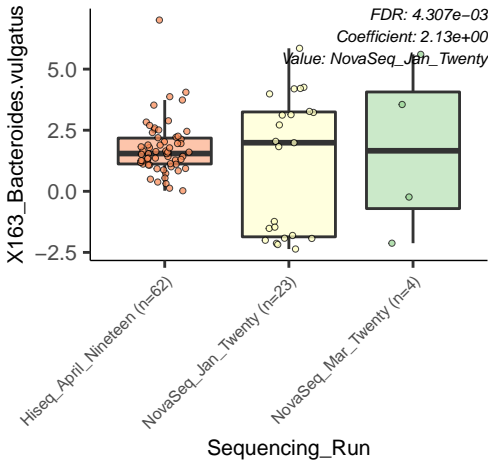


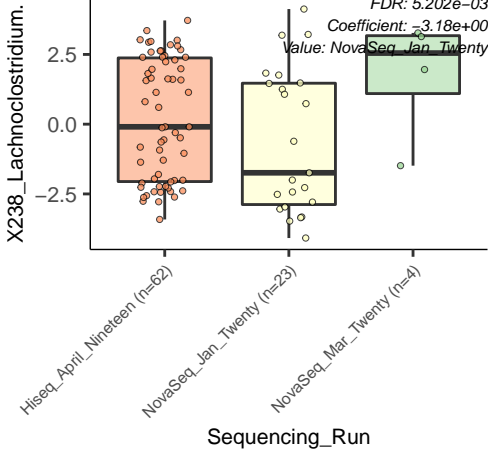




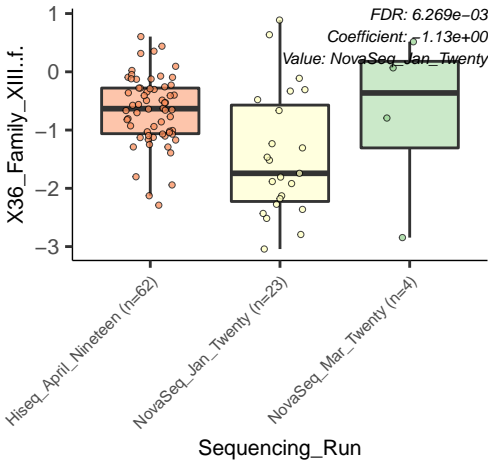


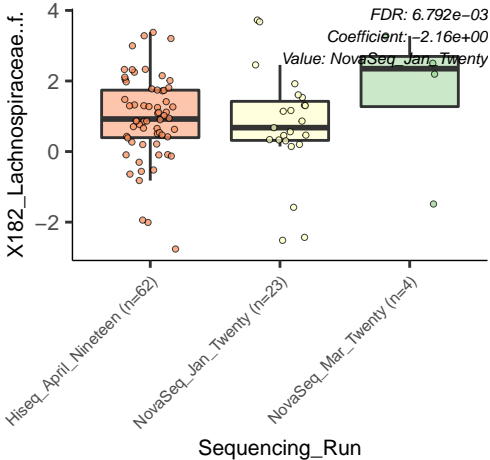


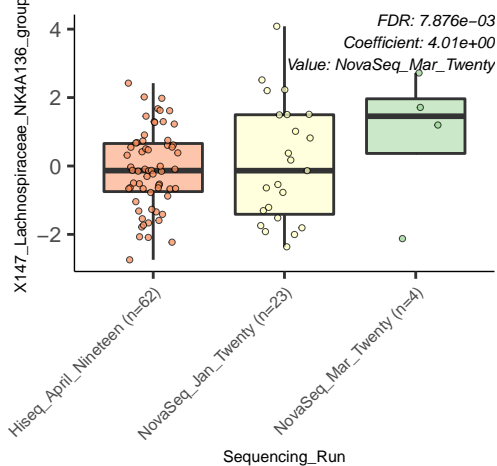


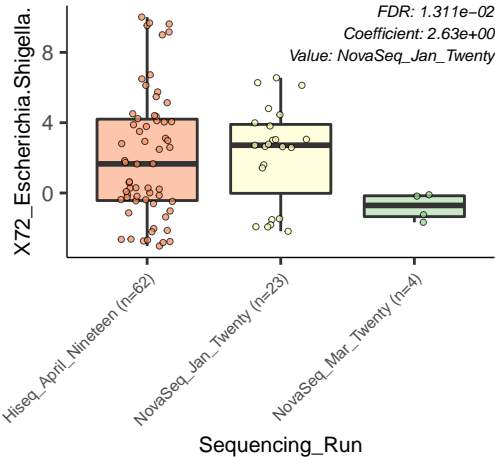


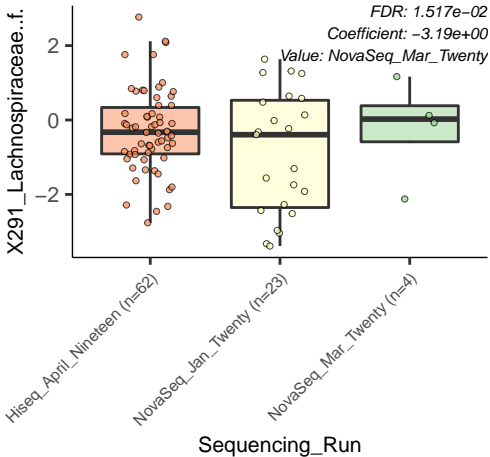
X36_Family_XIII.f.

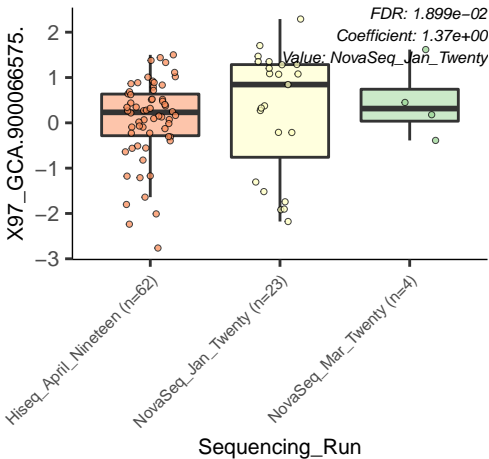


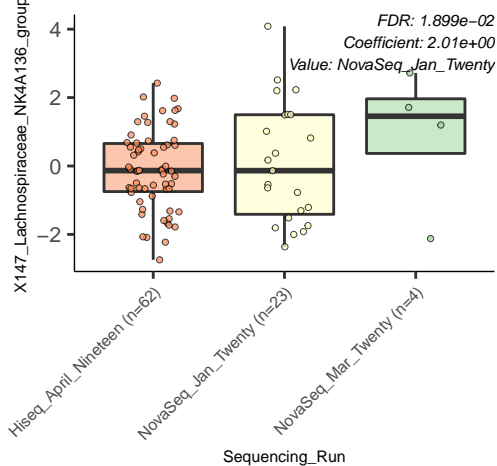


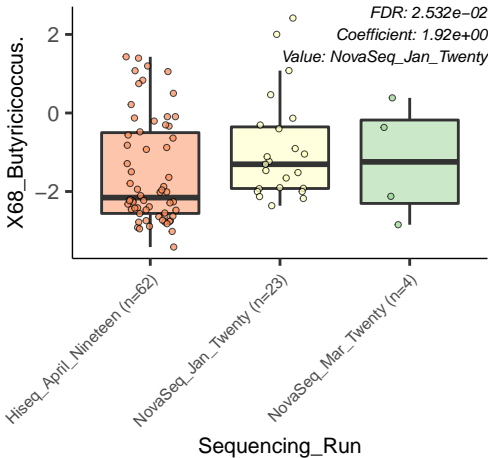


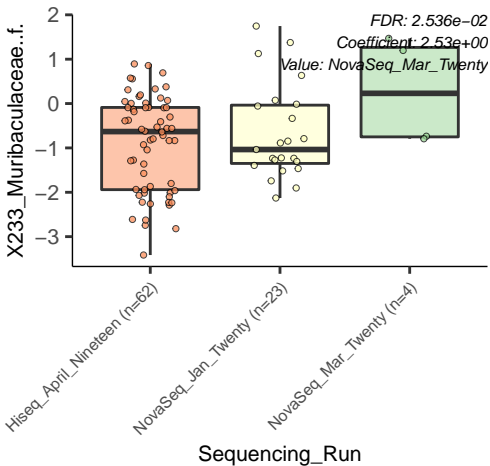


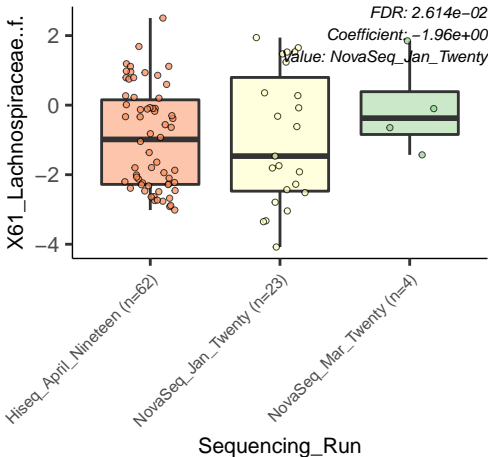


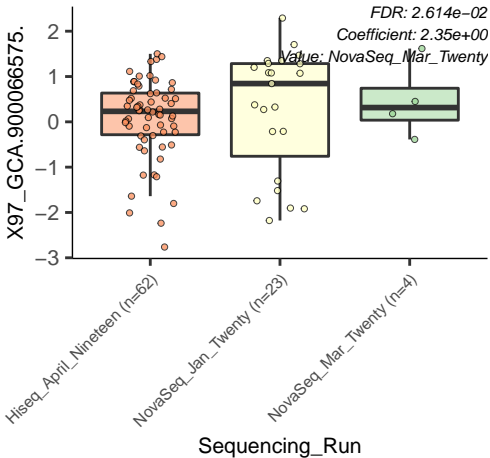


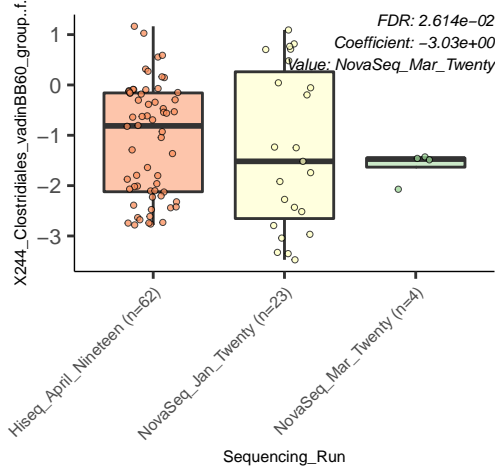


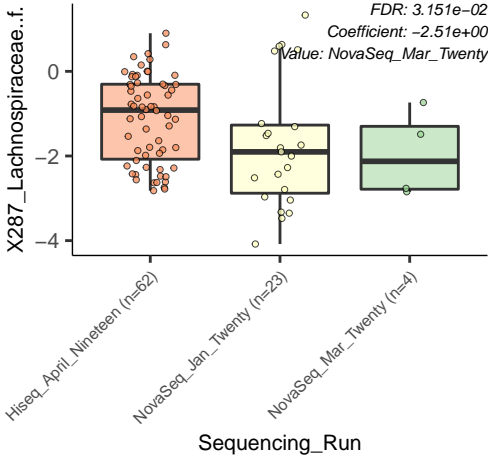


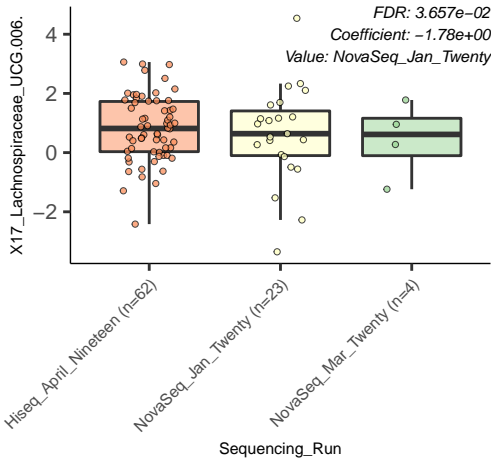


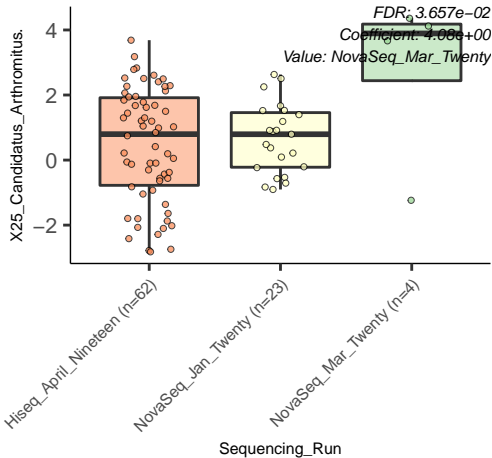




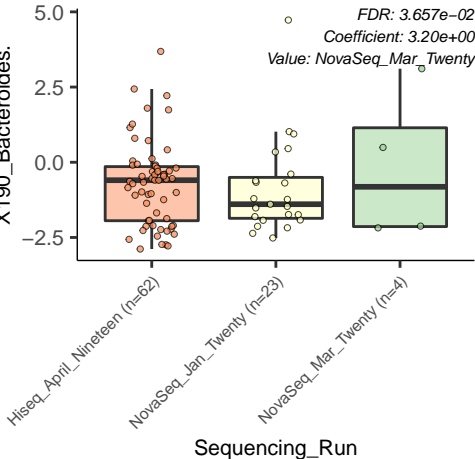


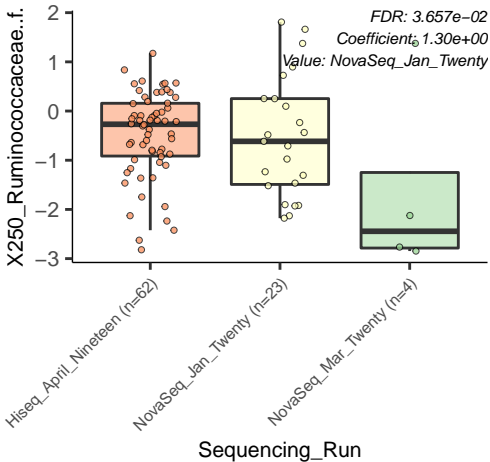


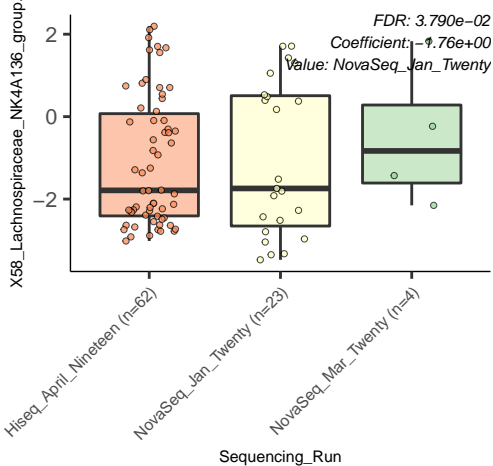


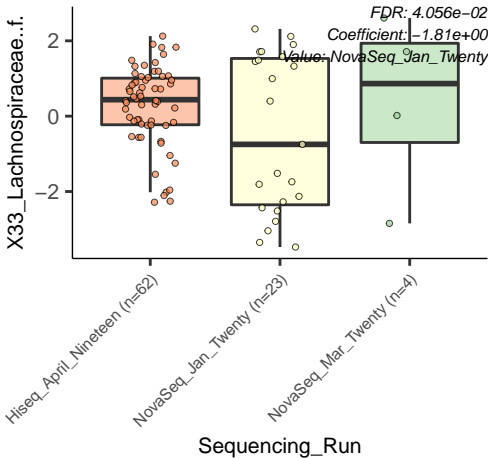


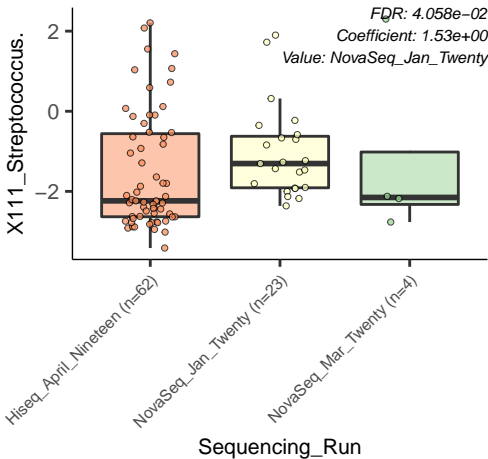
X190_Bacteroides.

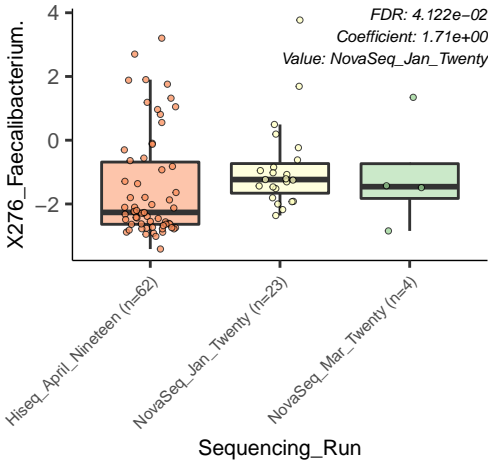


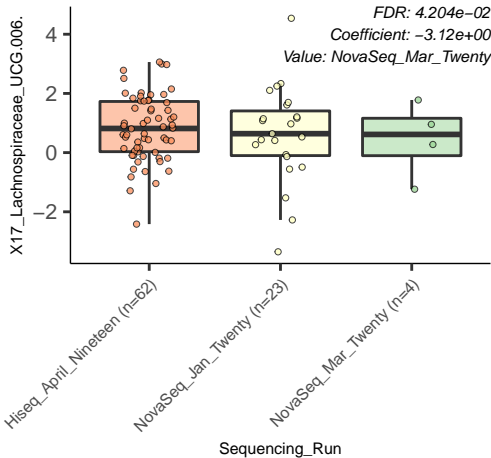




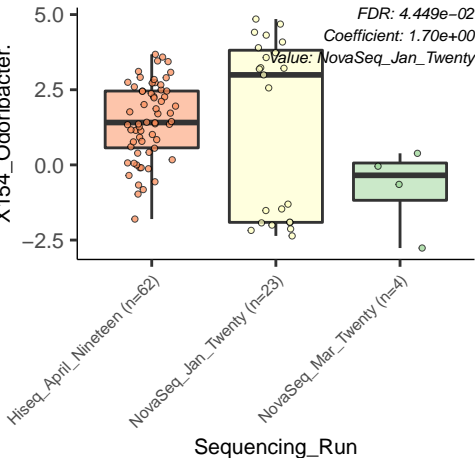


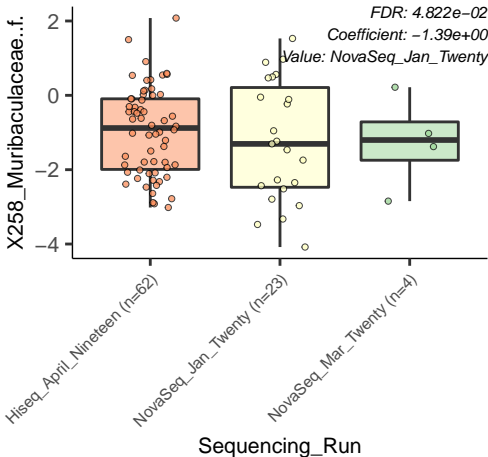




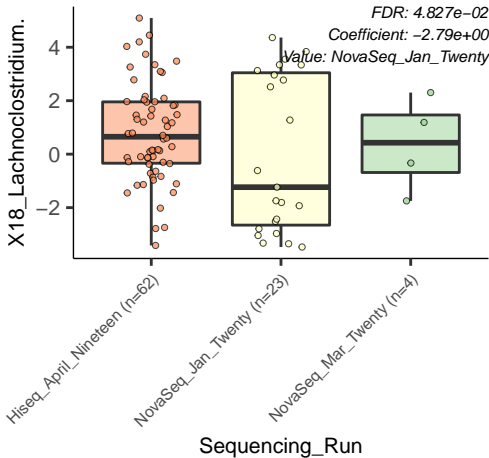


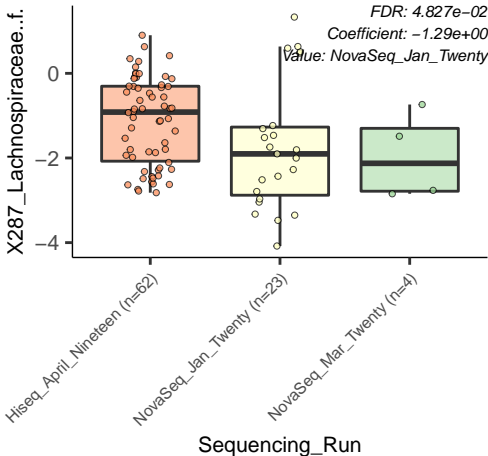
X154_Odoribacter.

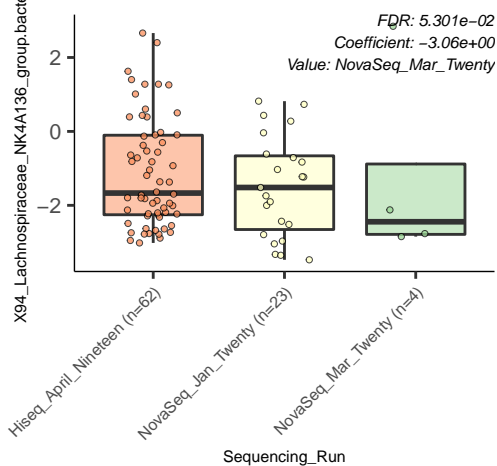


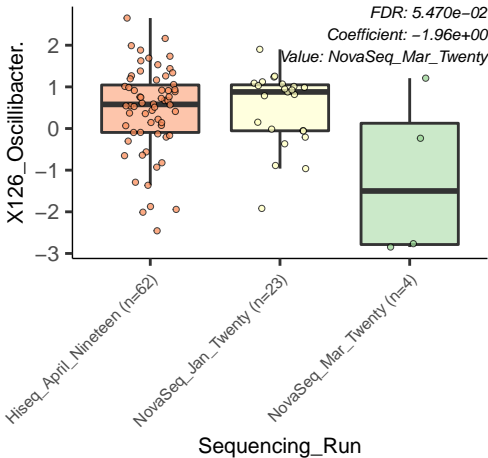


X18_Lachnoclostridium.

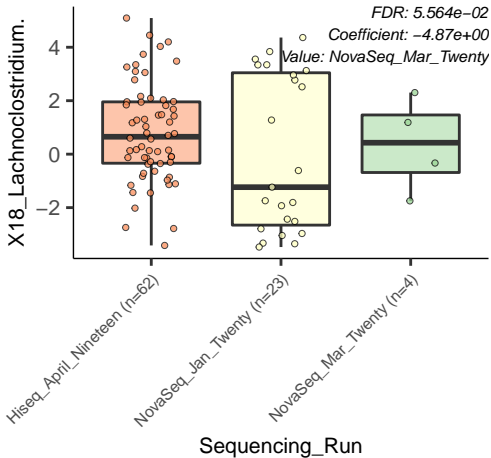




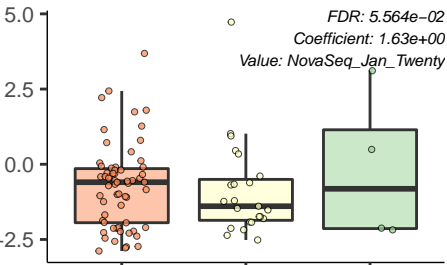




X18_Lachnoclostridium.



X190_Bacteroides.



Hiseq_April_Nineteen (n=62)

NovaSeq_Jan_Twenty (n=23)

NovaSeq_Mar_Twenty (n=4)

Sequencing_Run

X219_Blautia.

FDR: 5.584e-02

Coefficient: 2.32e+00

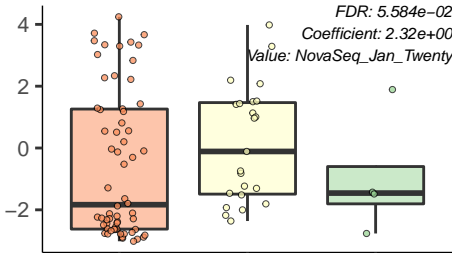
Value: NovaSeq_Jan_Twenty

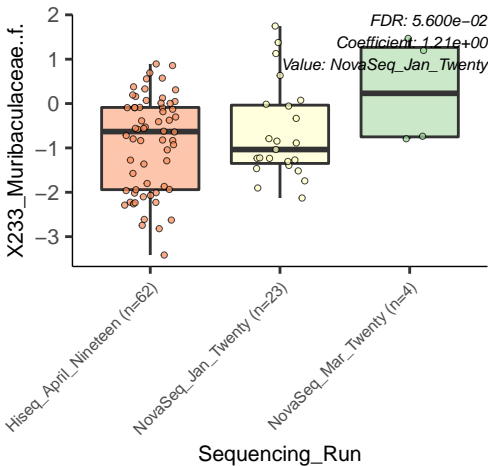
Hiseq_April_Nineteen (n=62)

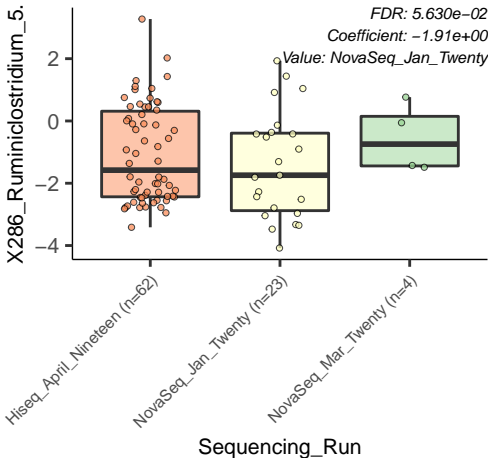
NovaSeq_Jan_Twenty (n=23)

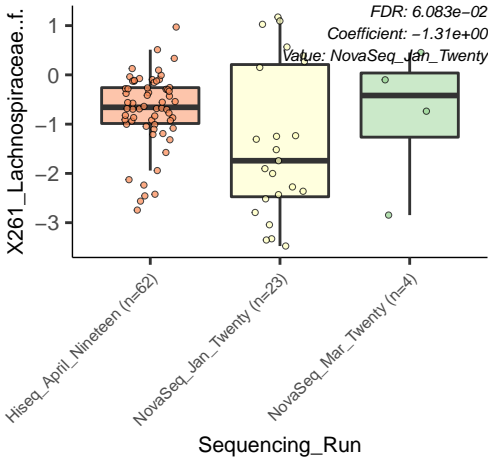
NovaSeq_Mar_Twenty (n=4)

Sequencing_Run









X22_Muribaculaceae.f.

Hiseq_April_Nineteen (n=62)

NovaSeq_Jan_Twenty (n=23)

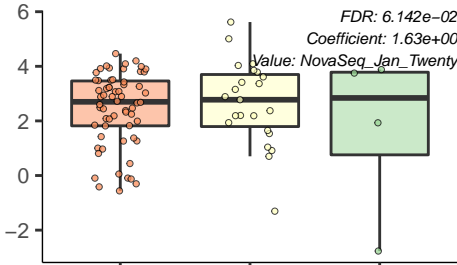
NovaSeq_Mar_Twenty (n=4)

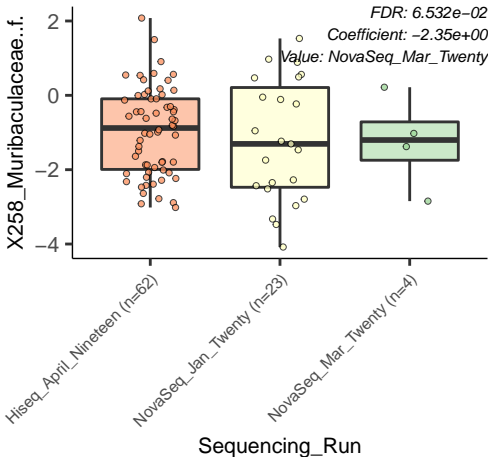
Sequencing_Run

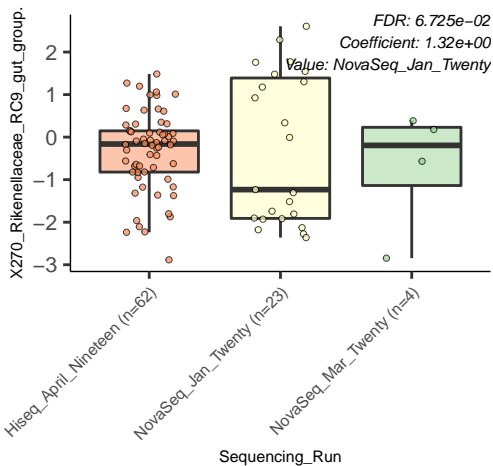
FDR: 6.142e-02

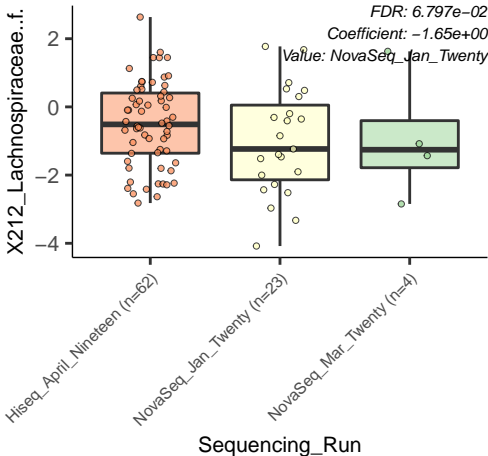
Coefficient: 1.63e+00

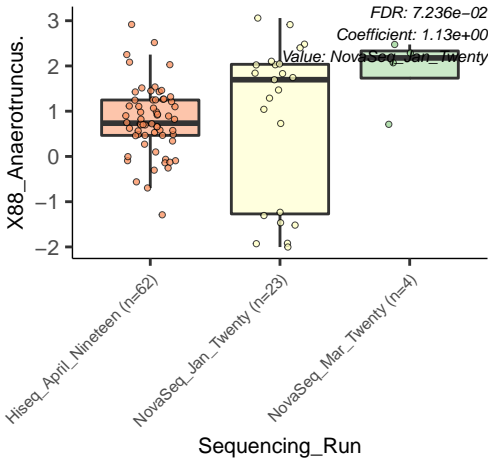
Value: NovaSeq_Jan_Twenty

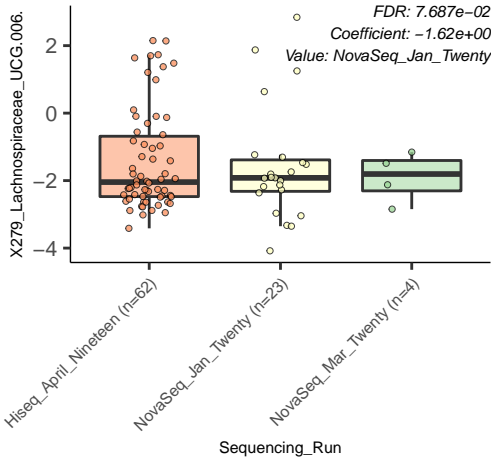


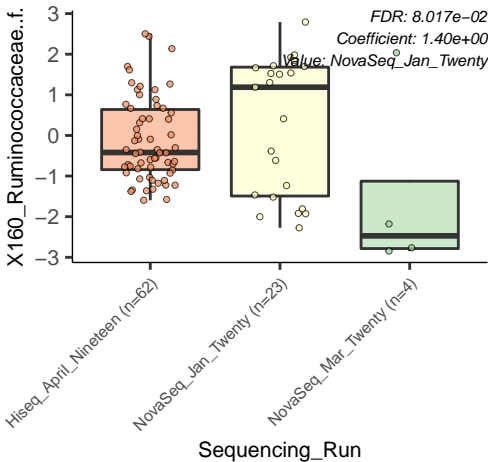


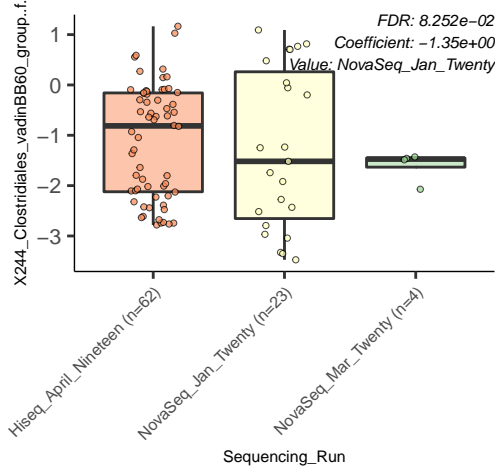


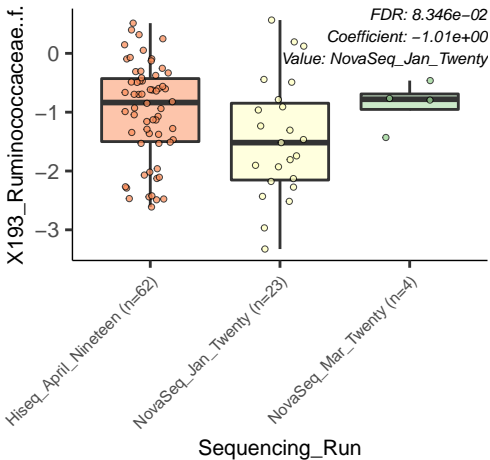


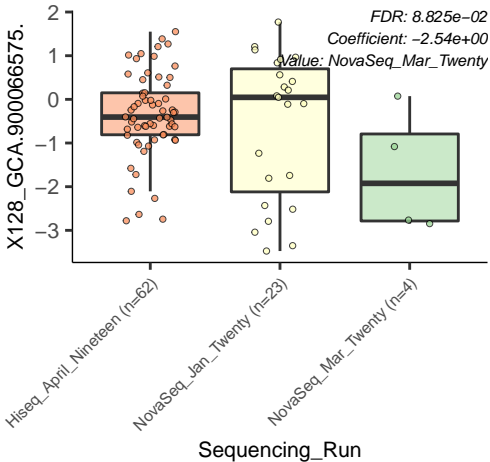


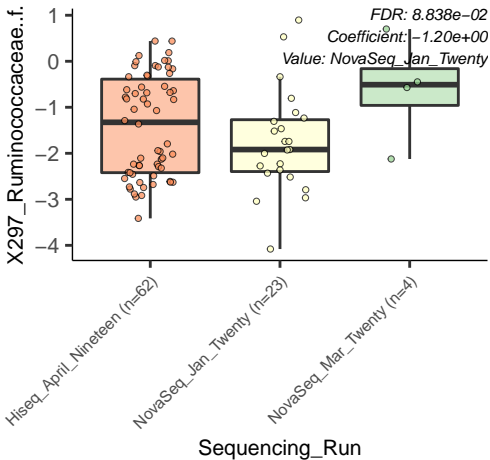


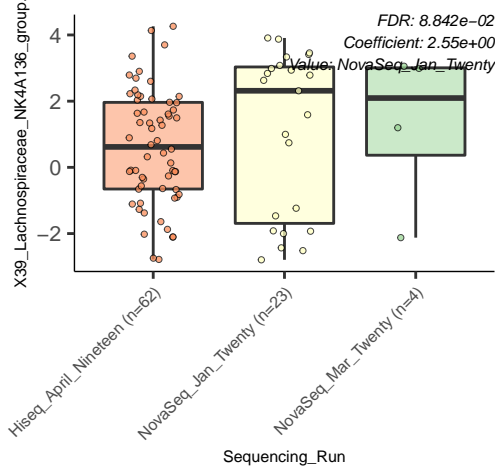


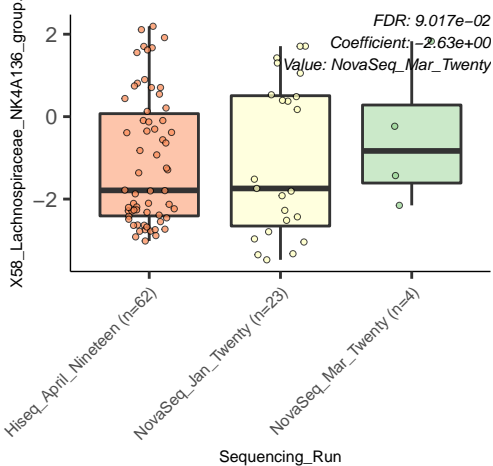


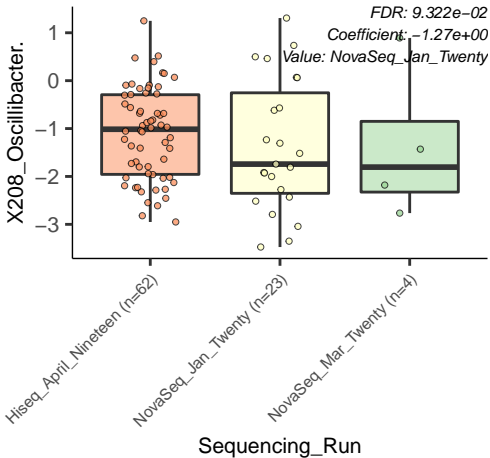


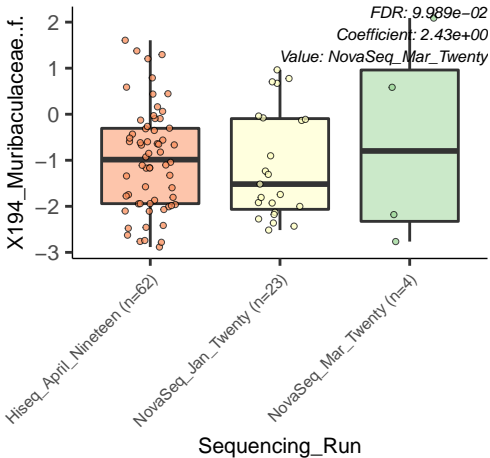


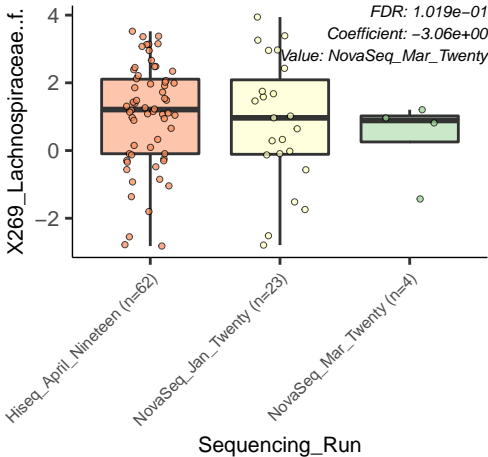












X87_Alistipes.

FDR: 1.023e-01

Coefficient: 2.50e+00

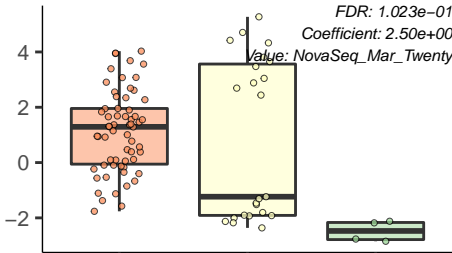
Value: NovaSeq_Mar_Twenty

Hiseq_April_Nineteen (n=62)

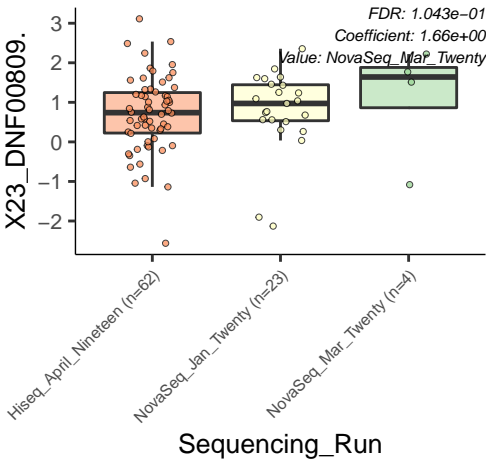
NovaSeq_Jan_Twenty (n=23)

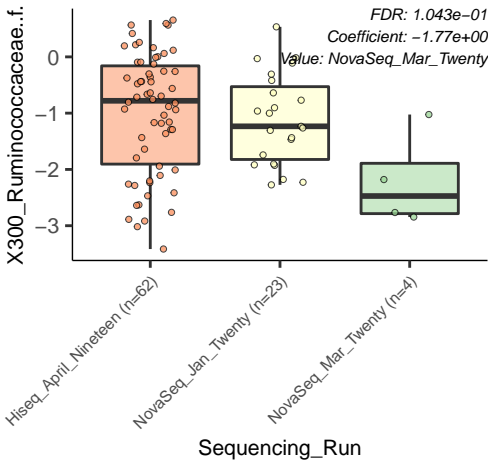
NovaSeq_Mar_Twenty (n=4)

Sequencing_Run

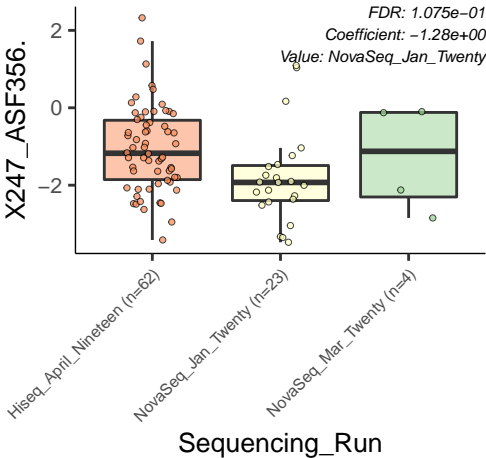


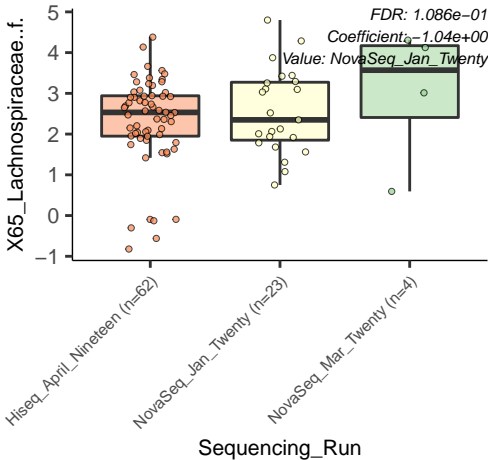
X23_DNF00809.

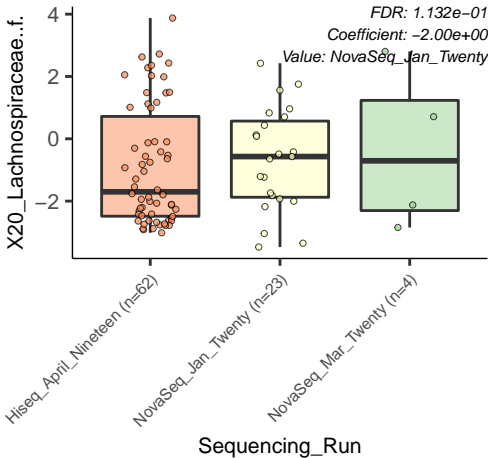


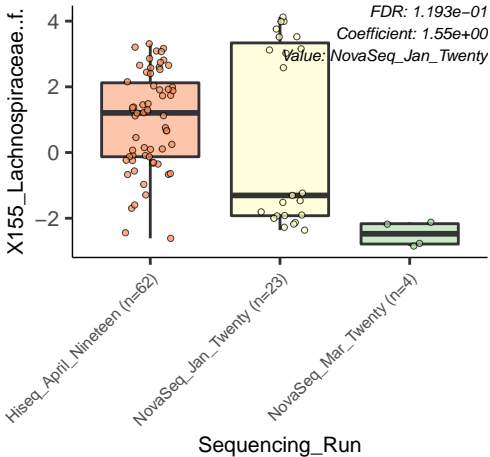


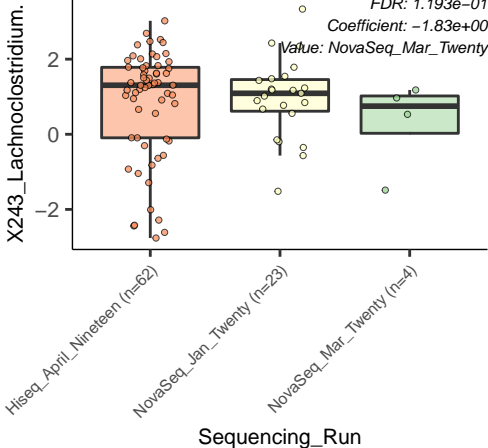
X247_ASF356.

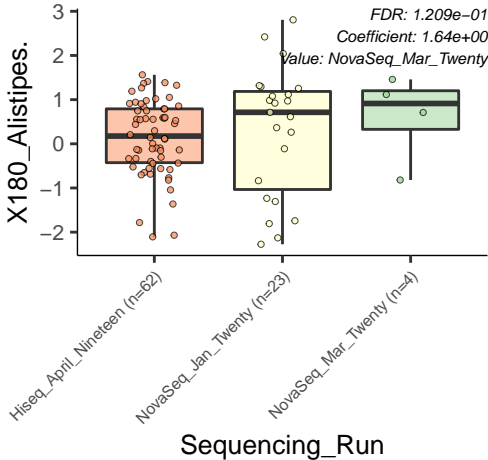


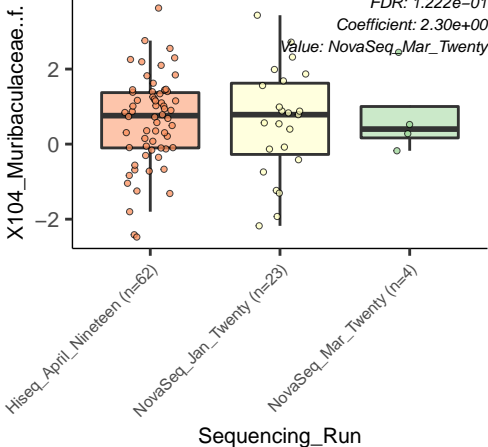


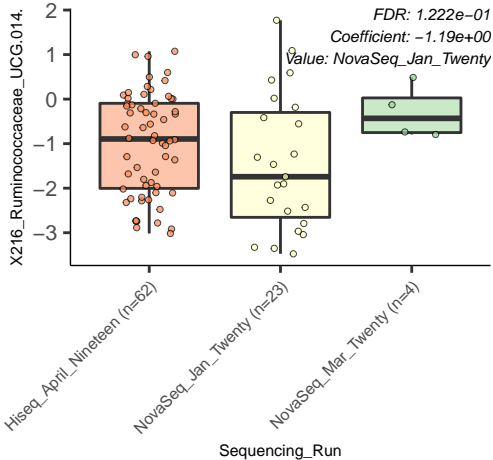


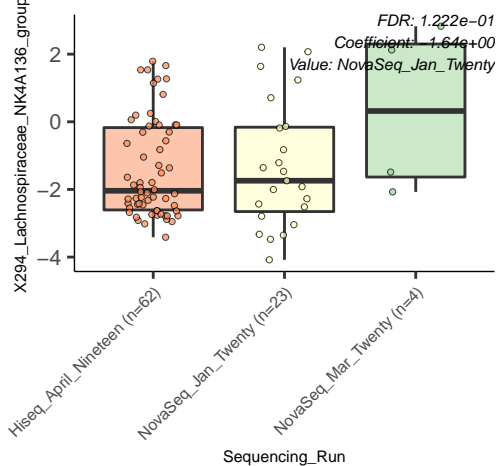


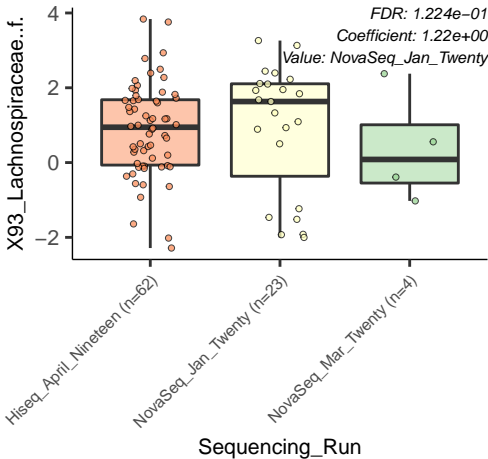


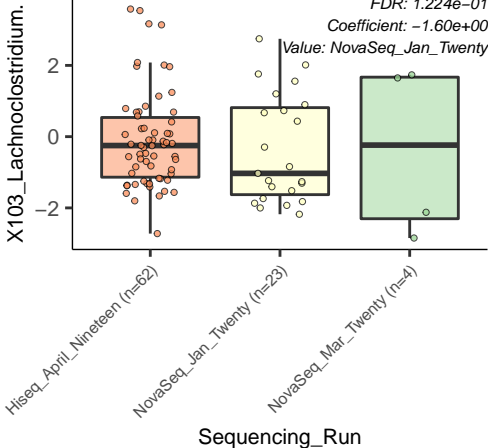


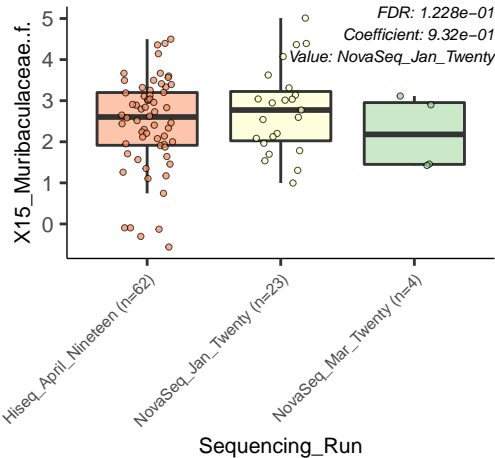


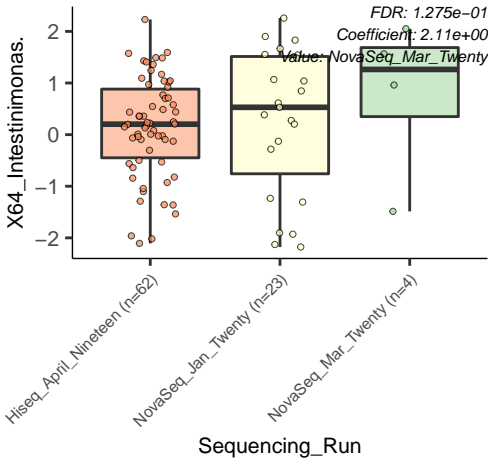




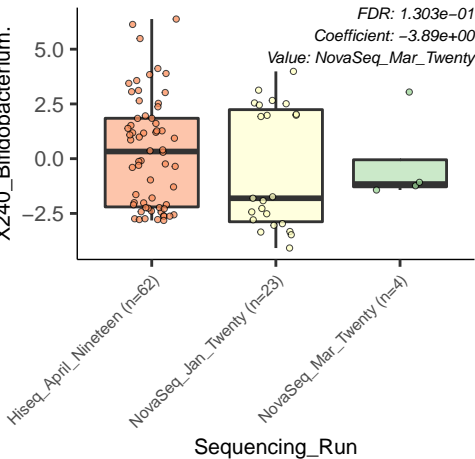




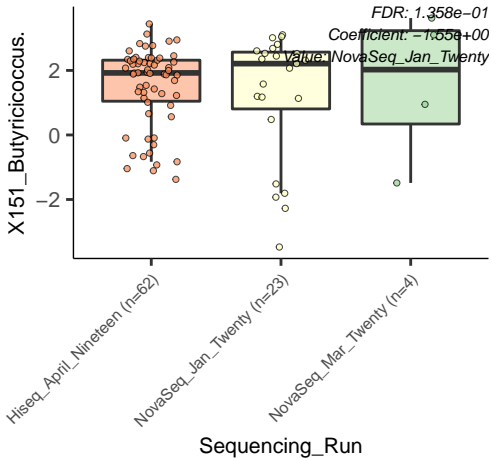


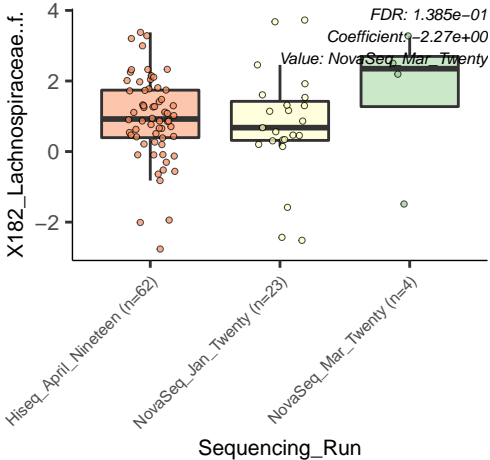


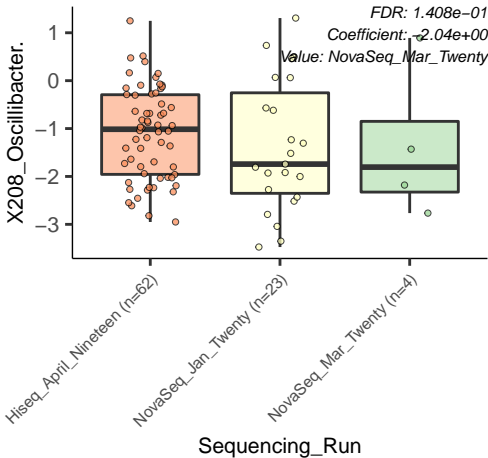
X240_Bifidobacterium.



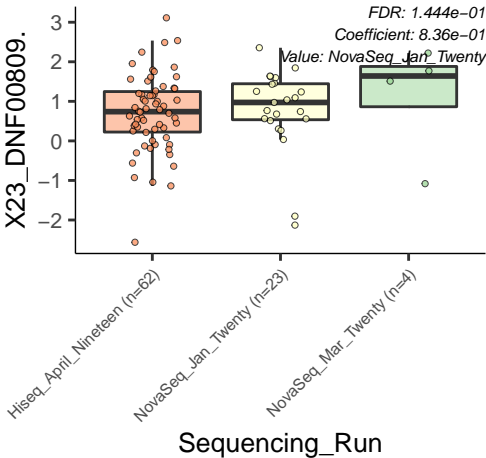
X151_Butyricoccus.

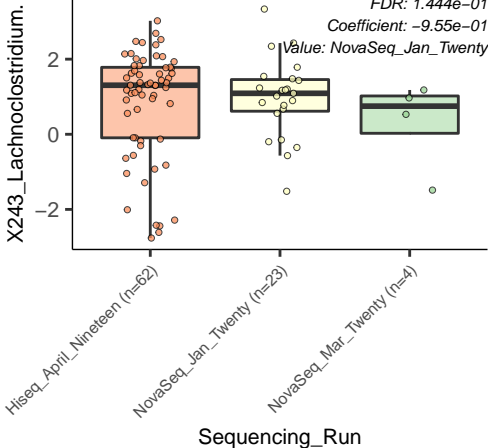


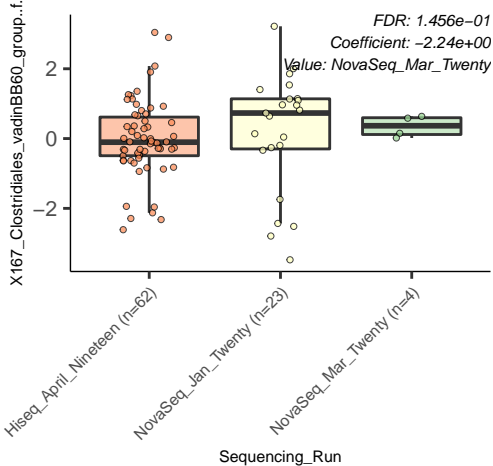




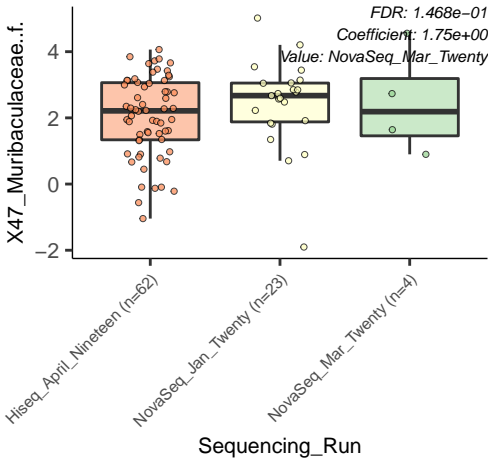
X23_DNF00809.

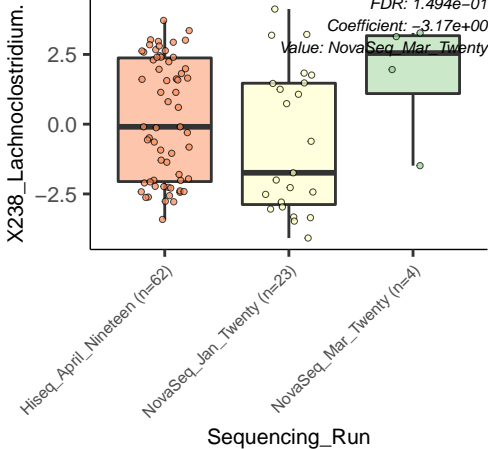




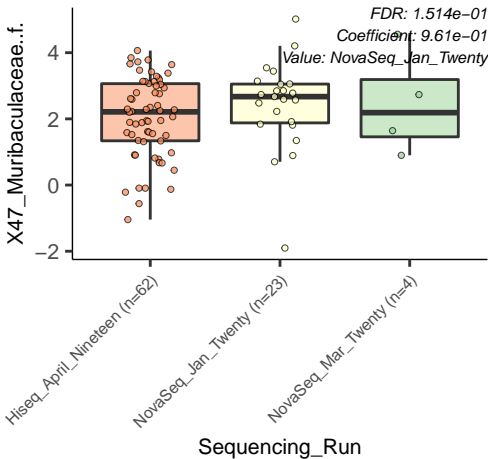


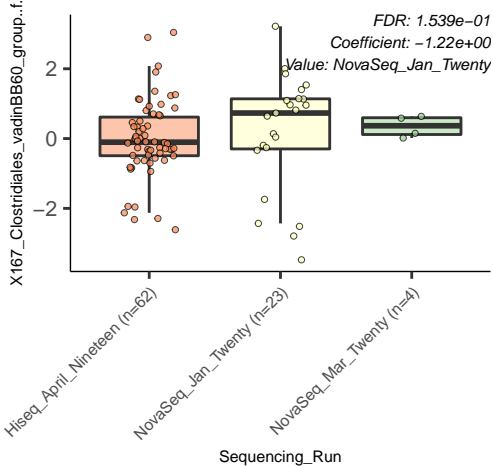
X47_Muribaculaceae..f.

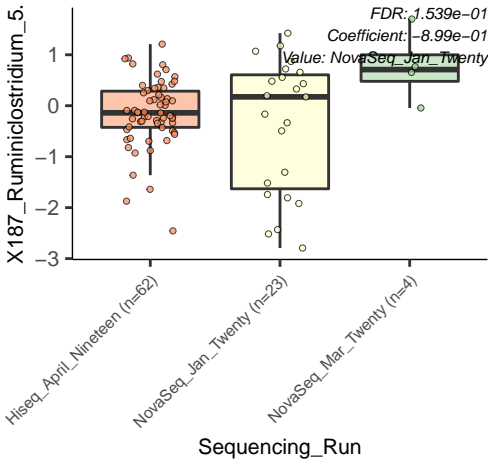


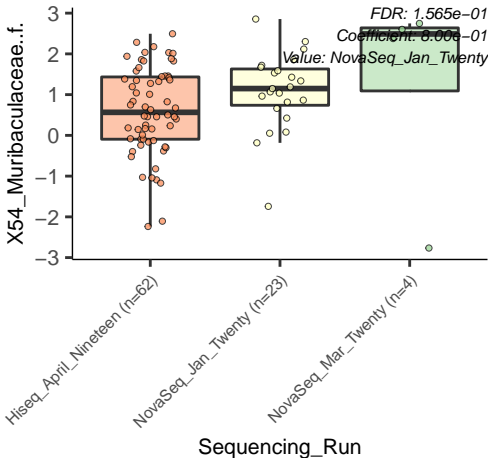


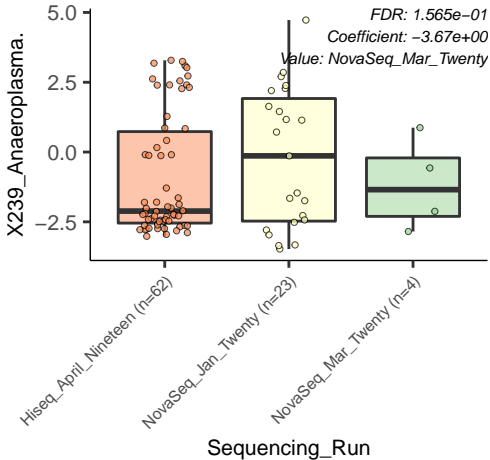
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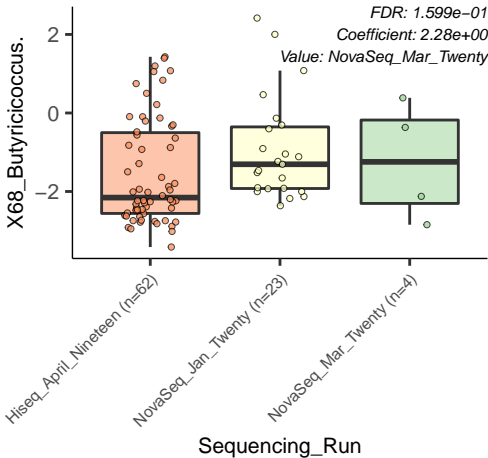


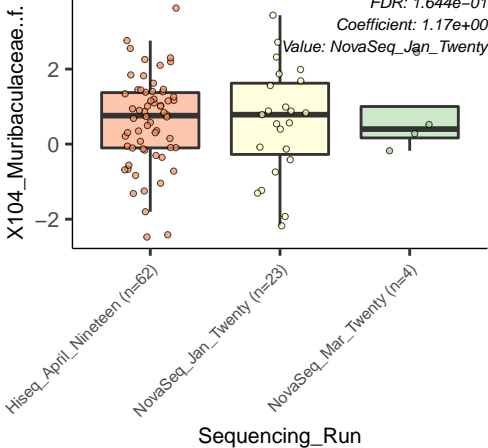


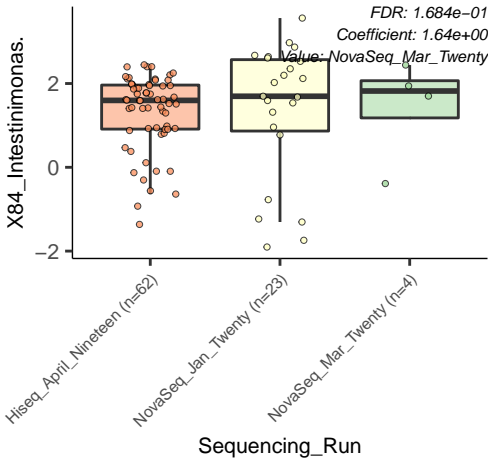


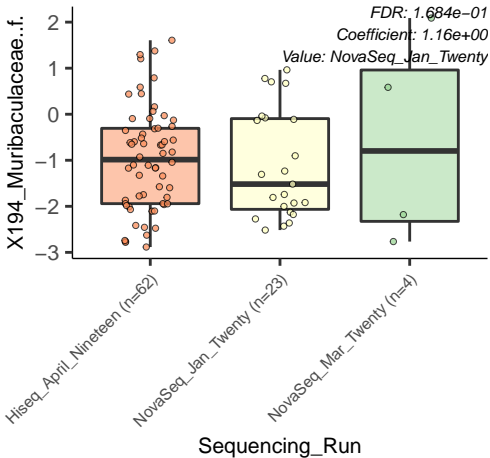


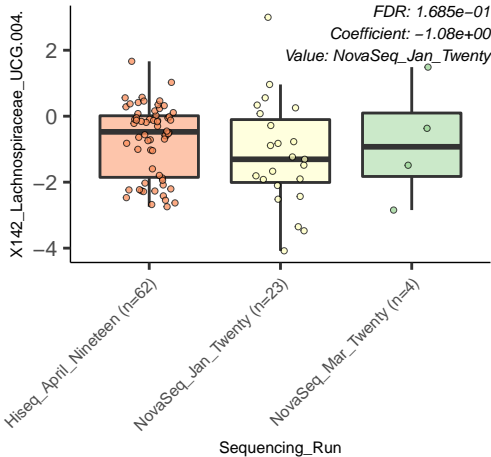


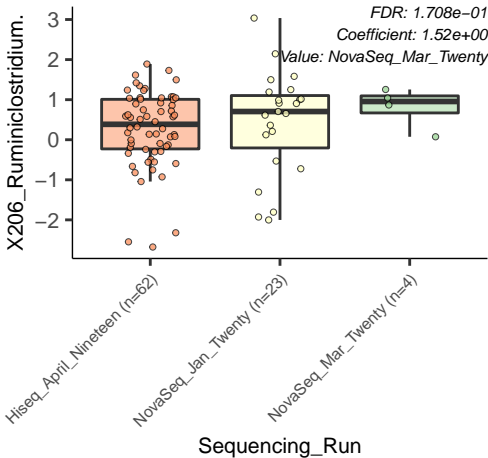




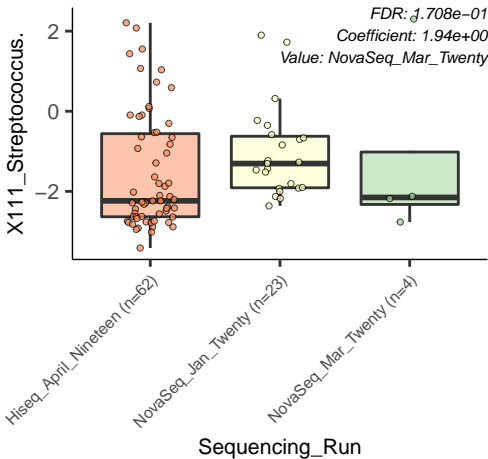


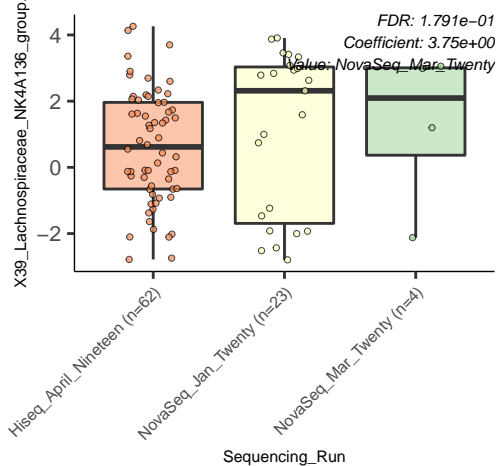


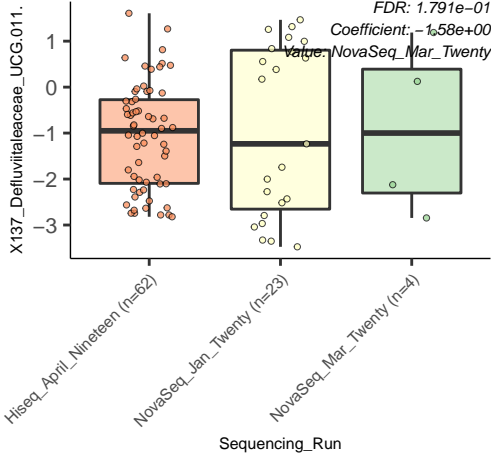


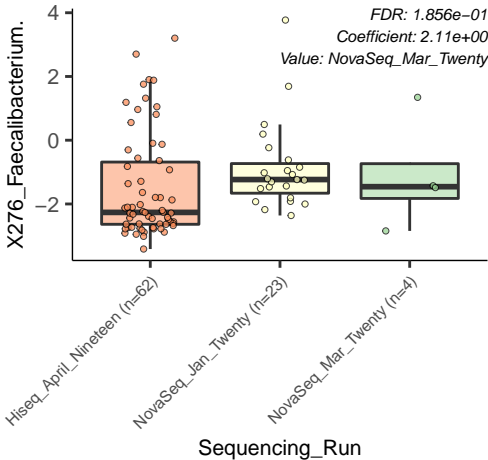


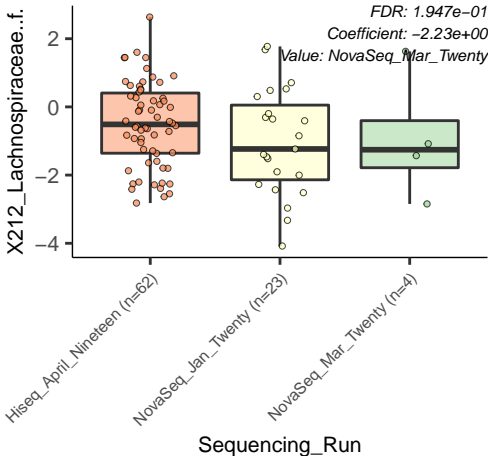
X111_Streptococcus.

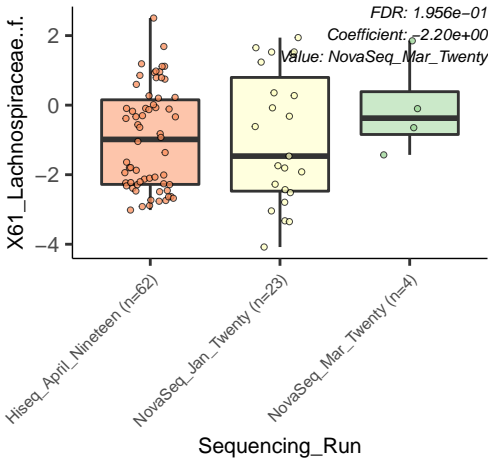


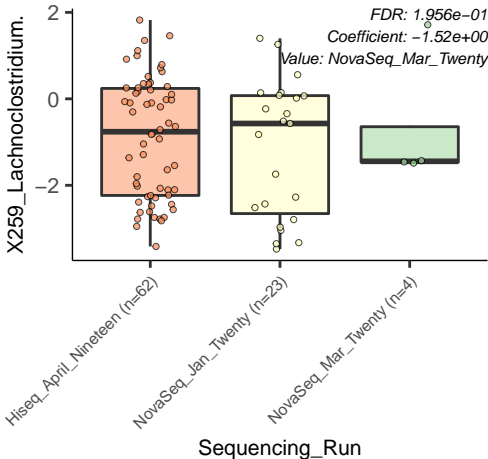


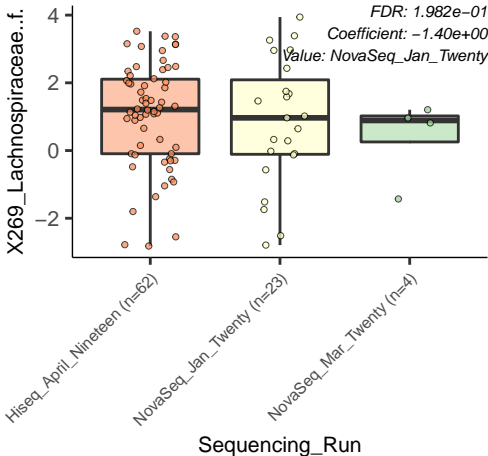


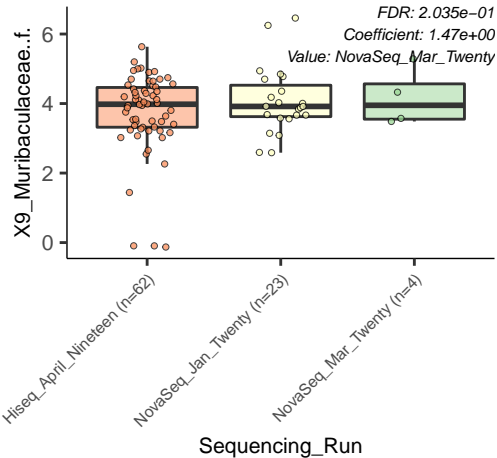


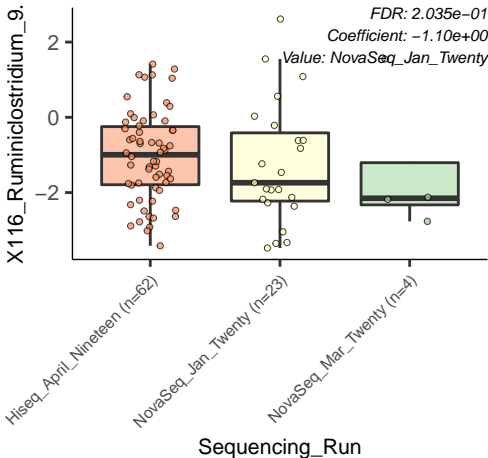


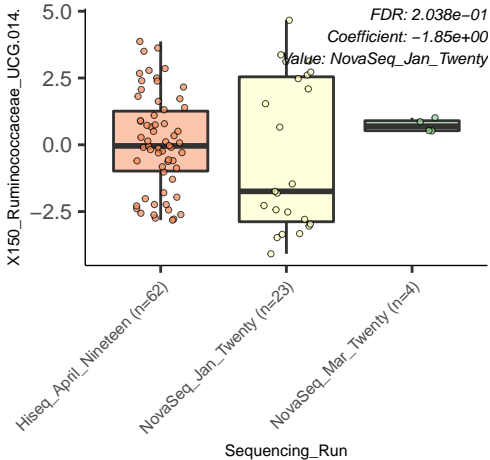


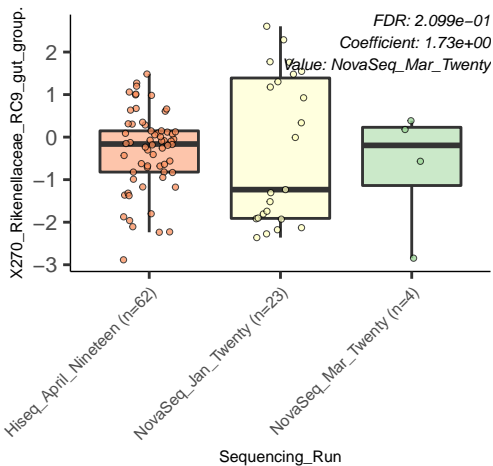


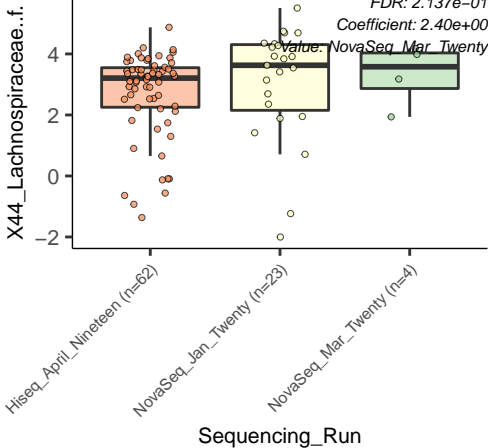


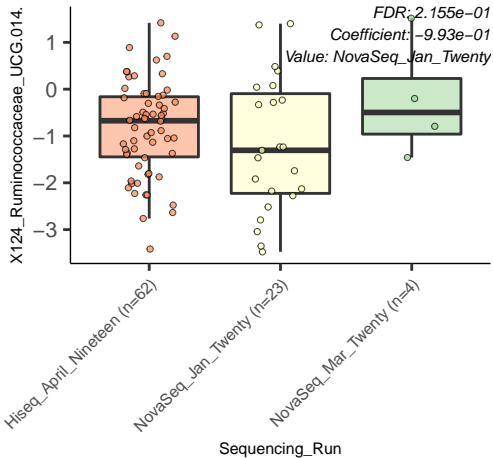


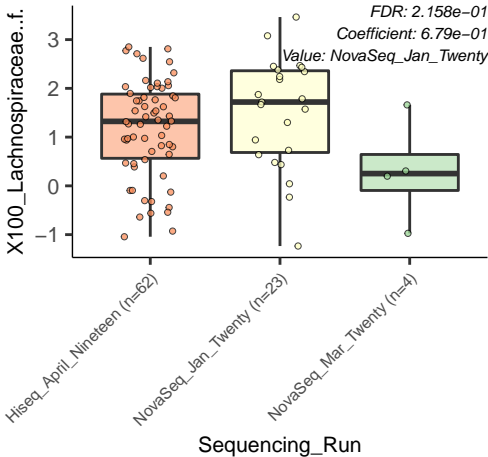


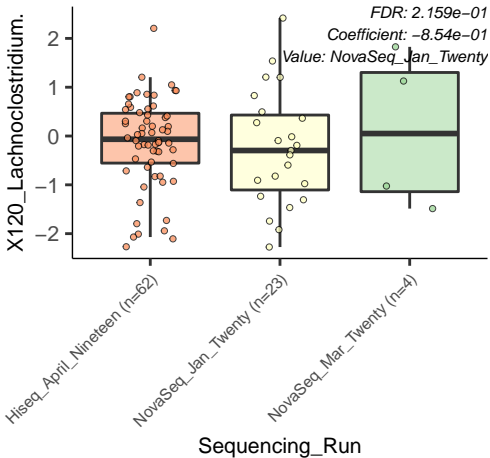


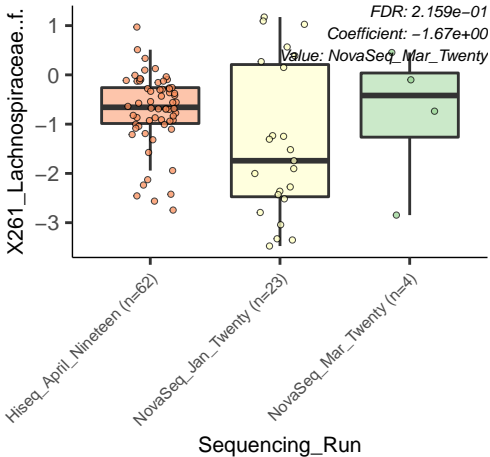


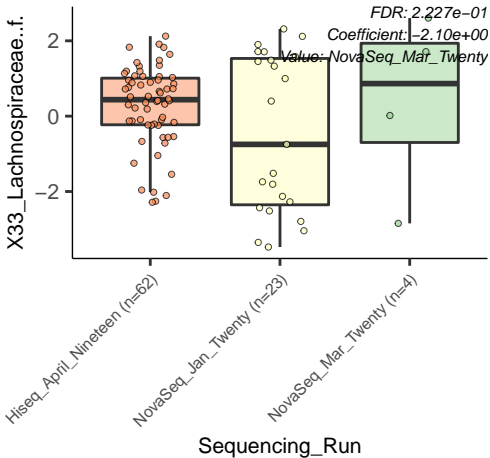


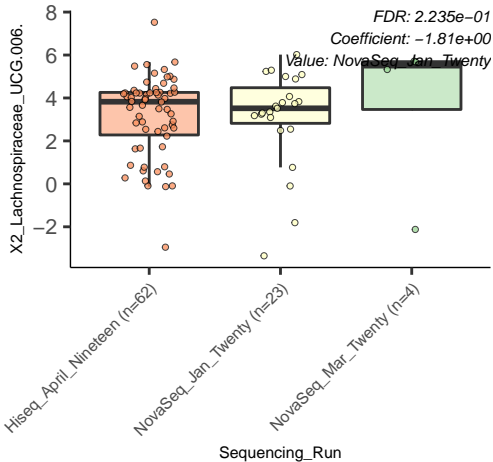


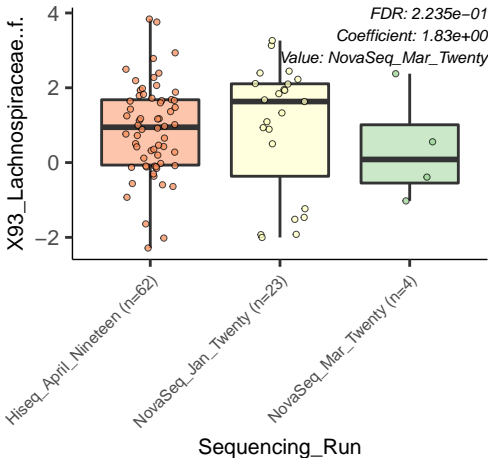












X154_Odoribacter.

