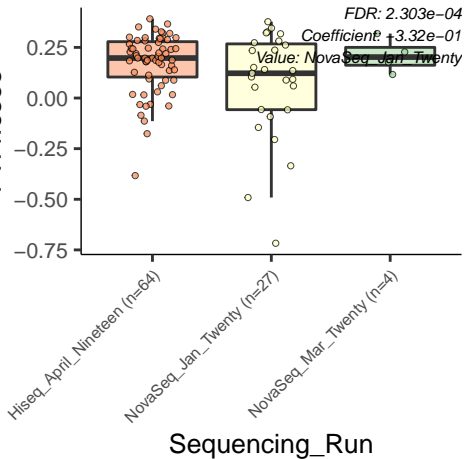
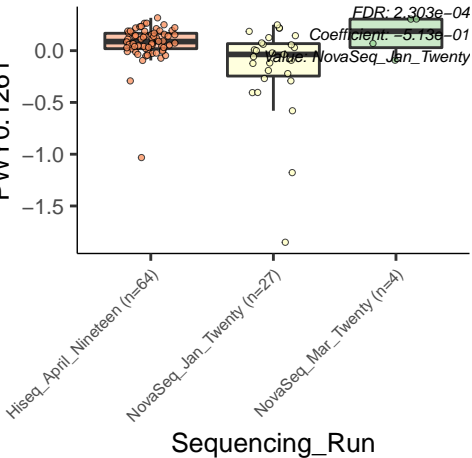
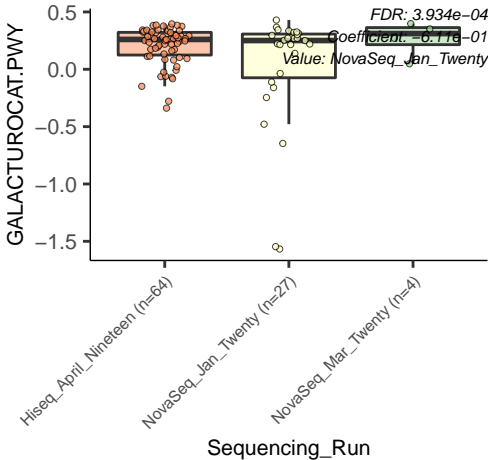


PWY.6353

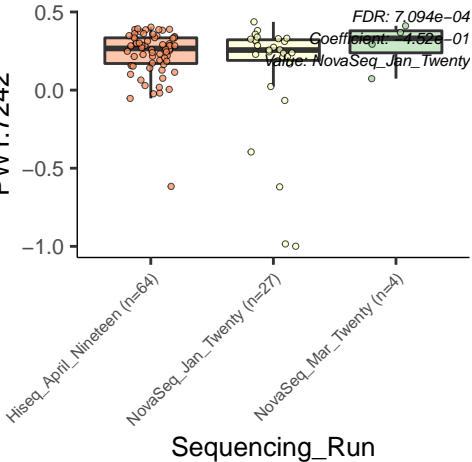


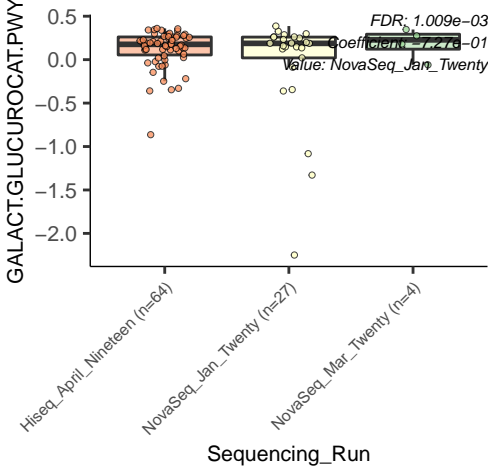
PWY0.1261



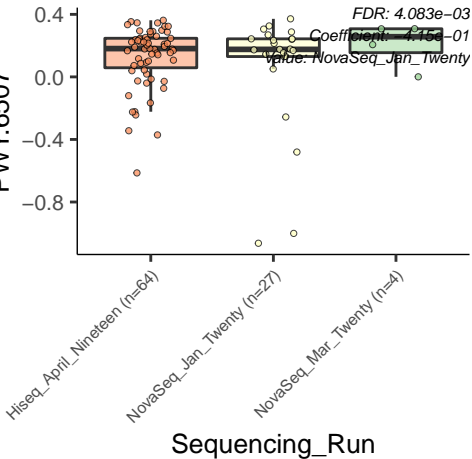


PWY.7242

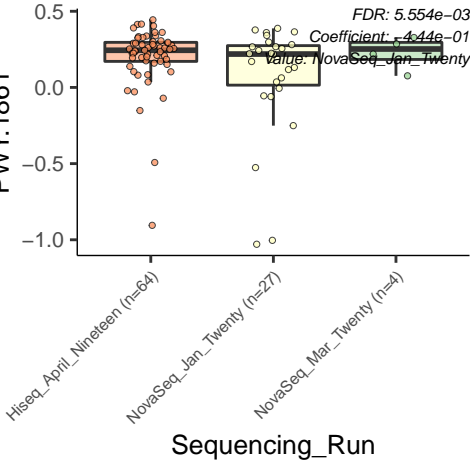




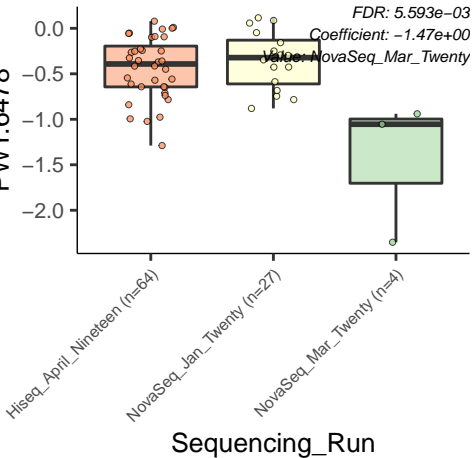
PWY.6507



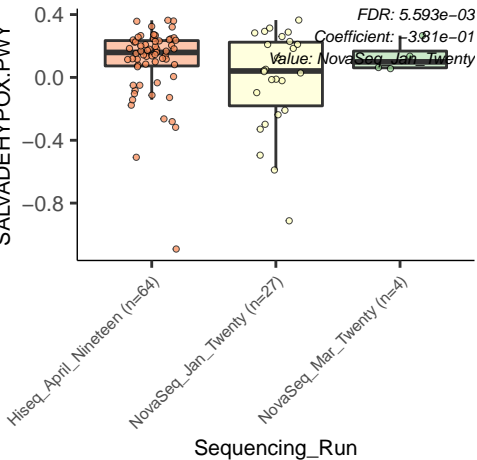
PWY.1861



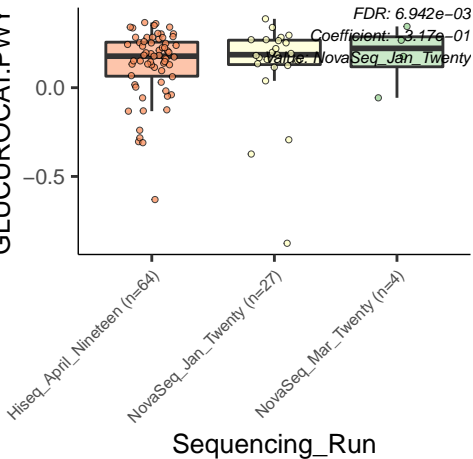
PWY.6478

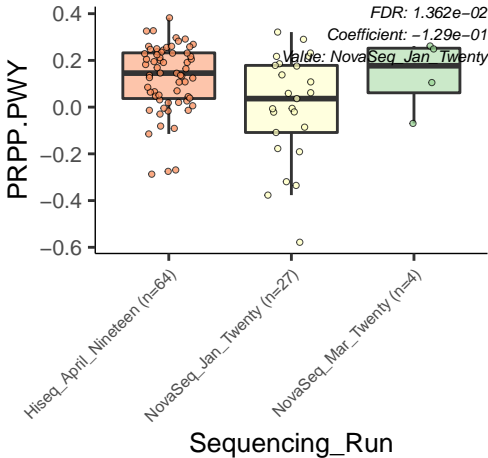


SALVADEHYPOX.PWY

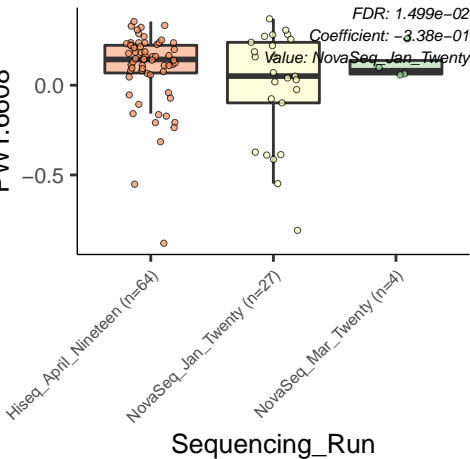


GLUCUROCAT.PWY

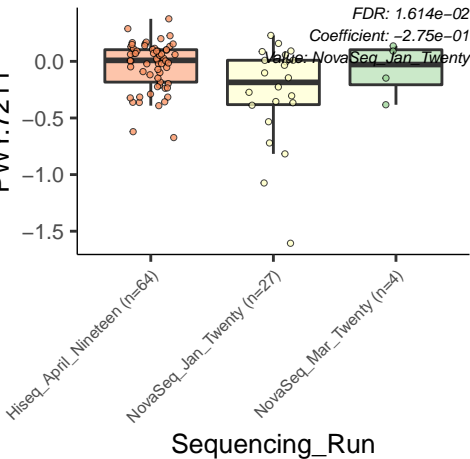




PWY.6608



PWY.7211



P164.PWY

FDR: 1.824e-02

Coefficient: -3.68e-01

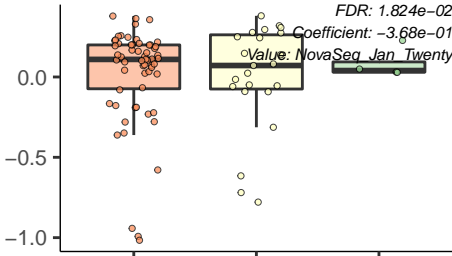
Value: NovaSeq Jan Twenty

Hiseq_April_Nineteen (n=64)

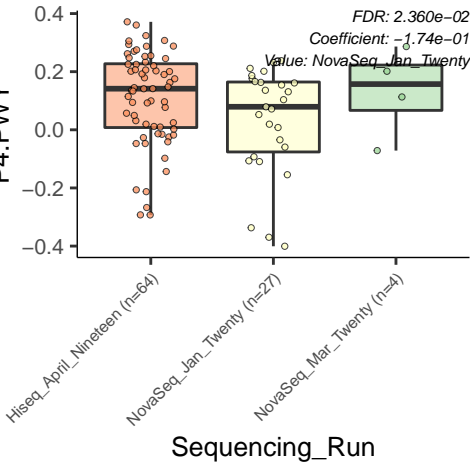
NovaSeq_Jan_Twenty (n=27)

NovaSeq_Mar_Twenty (n=4)

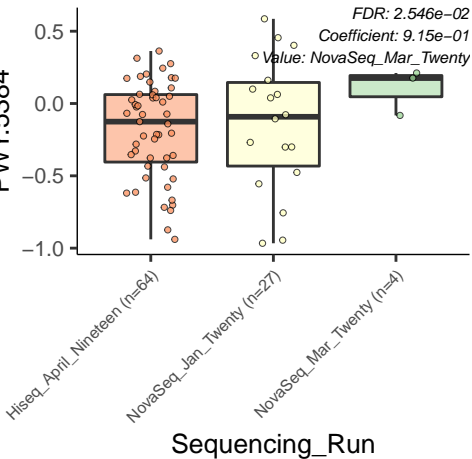
Sequencing_Run



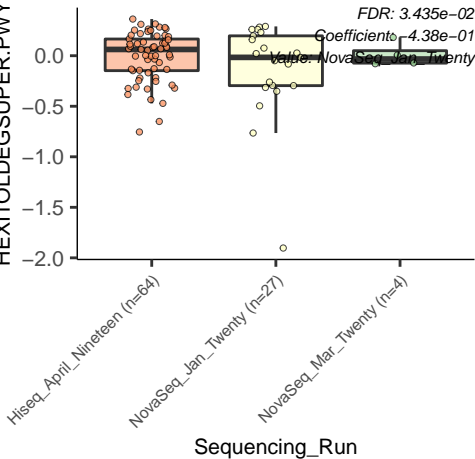
P4.PWY



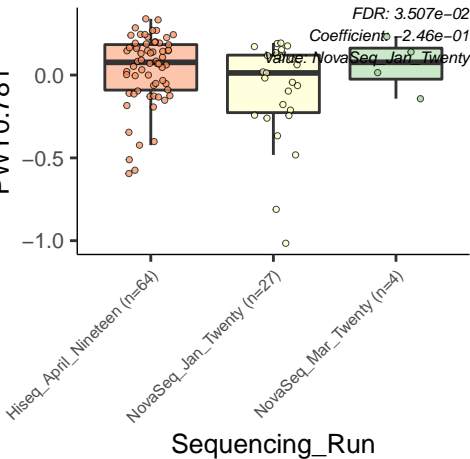
PWY.5384

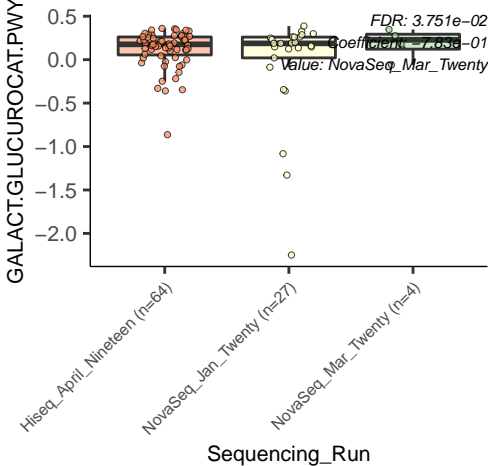


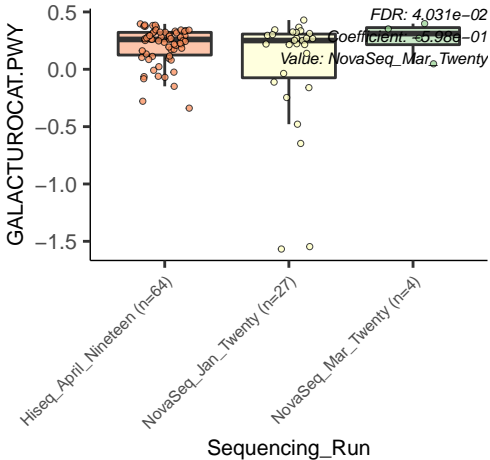
HEXITOLDEGSUPER.PWY



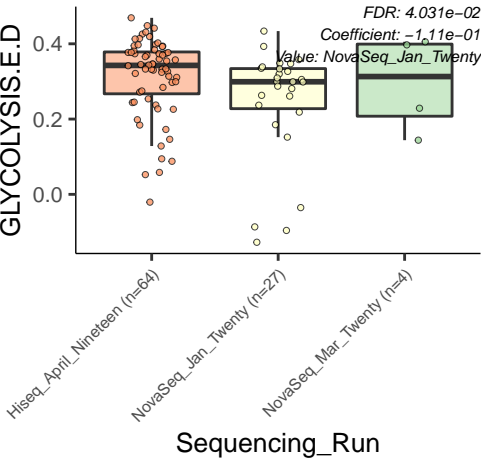
PWY0.781

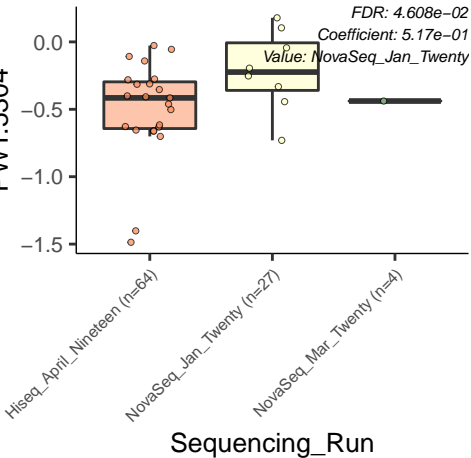




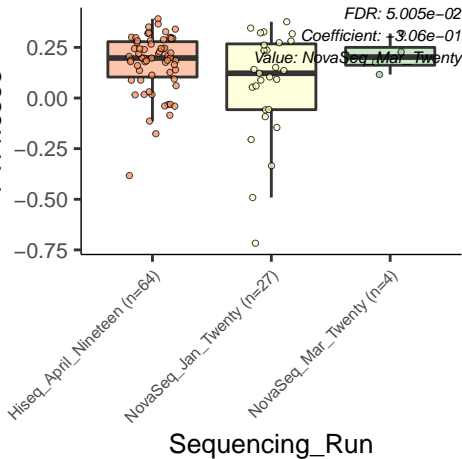


GLYCOLYSIS.E.D

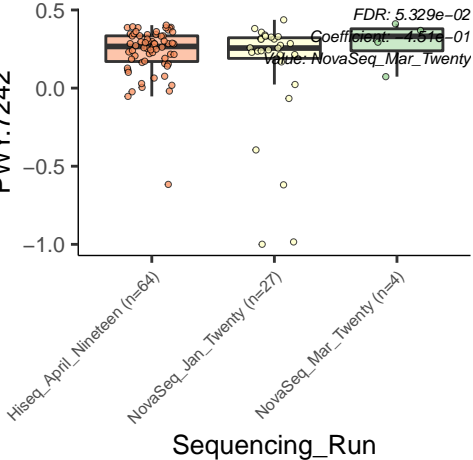




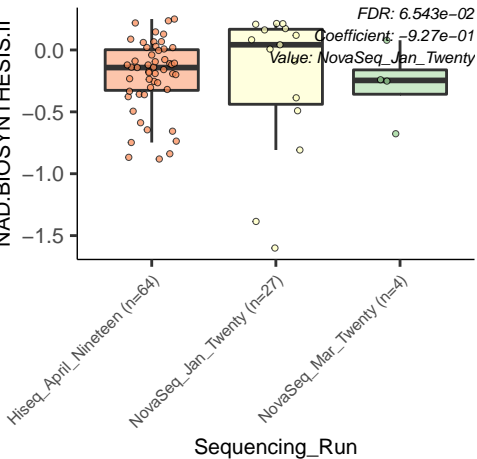
PWY.6353



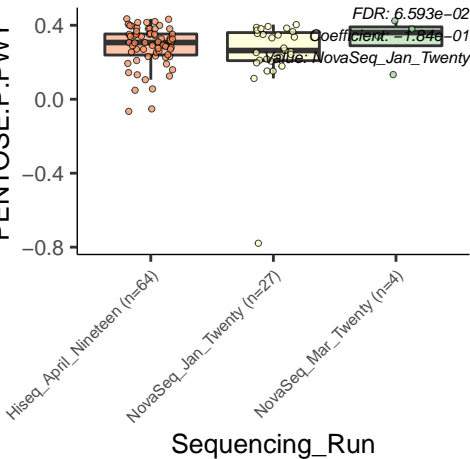
PWY.7242



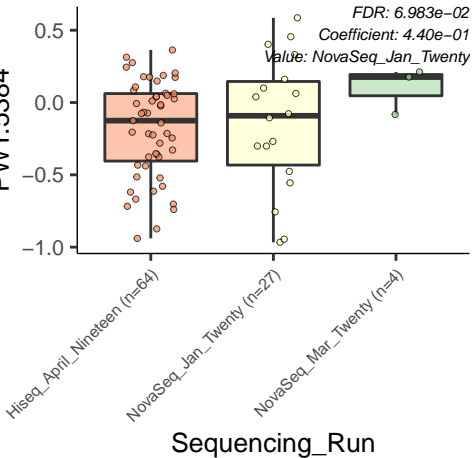
NAD.BIOSYNTHESIS.II

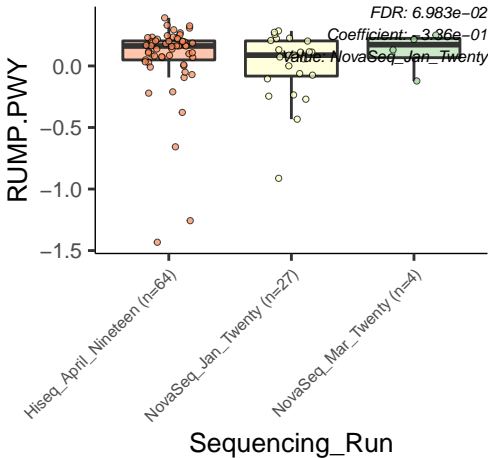


PENTOSE.P.PWY

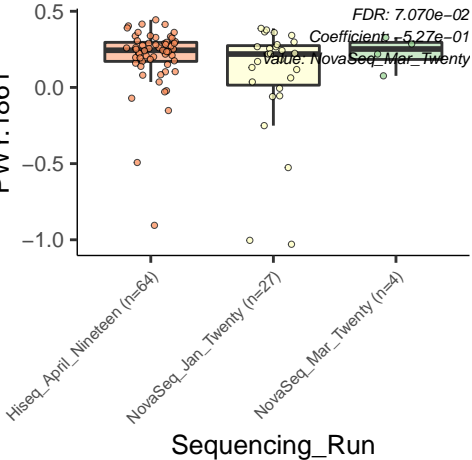


PWY.5384

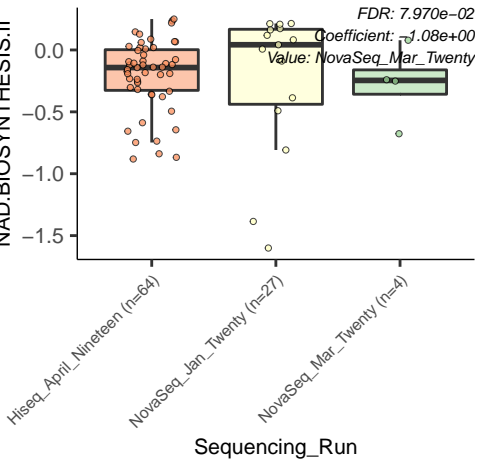




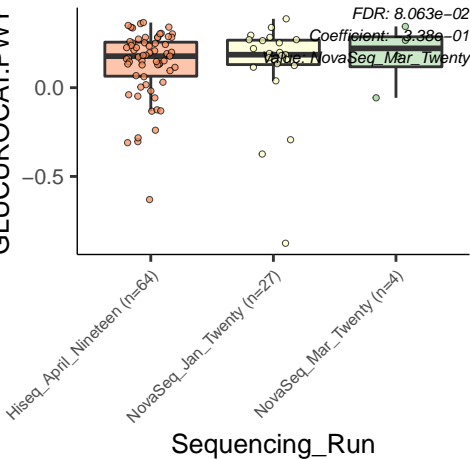
PWY.1861



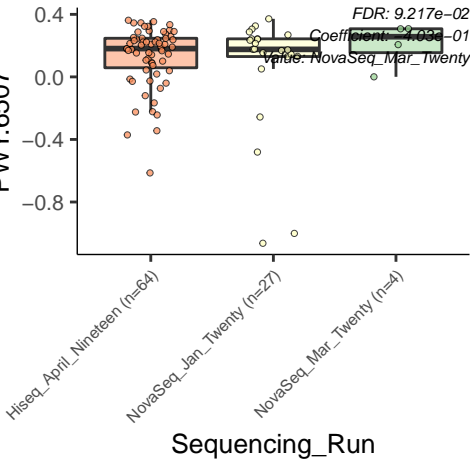
NAD.BIOSYNTHESIS.II



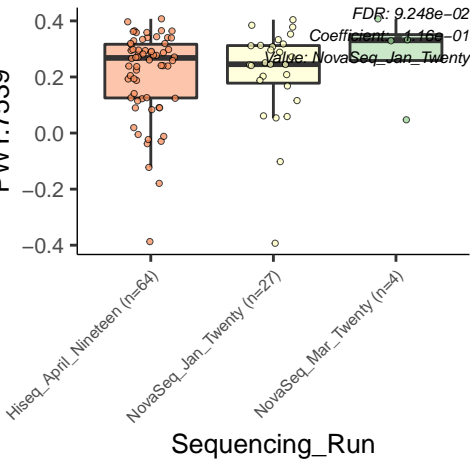
GLUCUROCAT.PWY



PWY.6507



PWY.7539



P441.PWY

FDR: 9.387e-02

Coefficient: -7.94e-02

Value: NovaSeq_Jan_Twenty

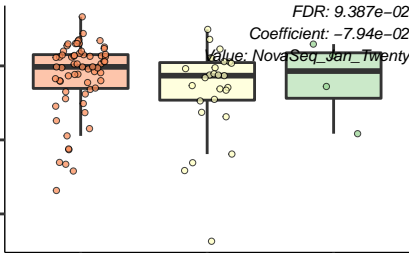
Hiseq_April_Nineteen (n=64)

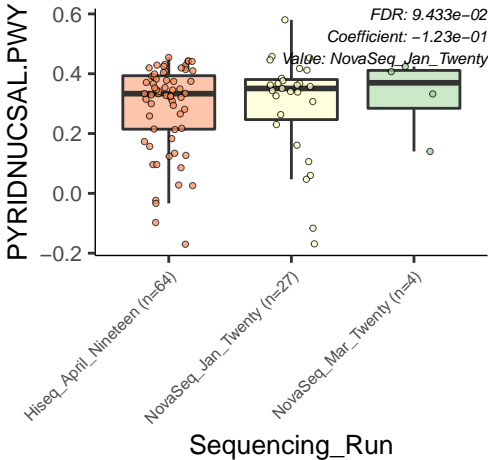
NovaSeq_Jan_Twenty (n=27)

NovaSeq_Mar_Twenty (n=4)

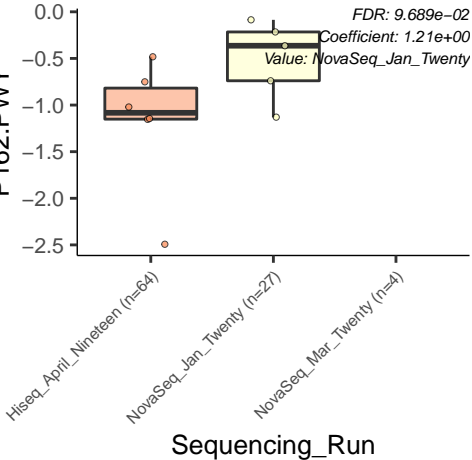
Sequencing_Run

0.0
0.2
0.4

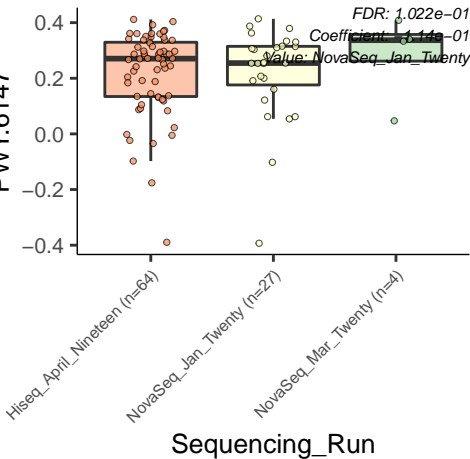




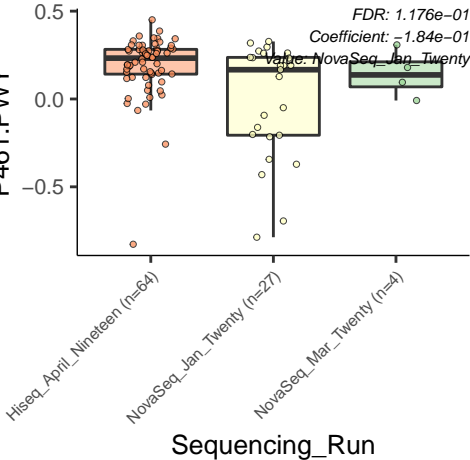
P162.PWY



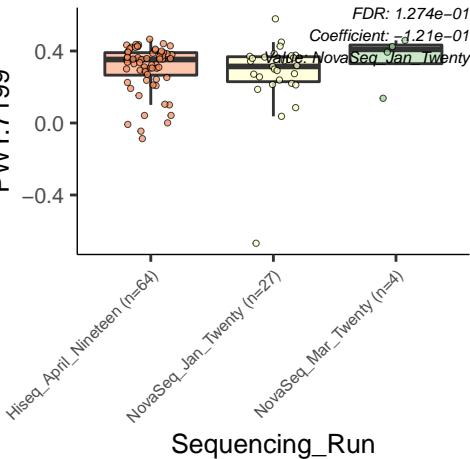
PWY.6147



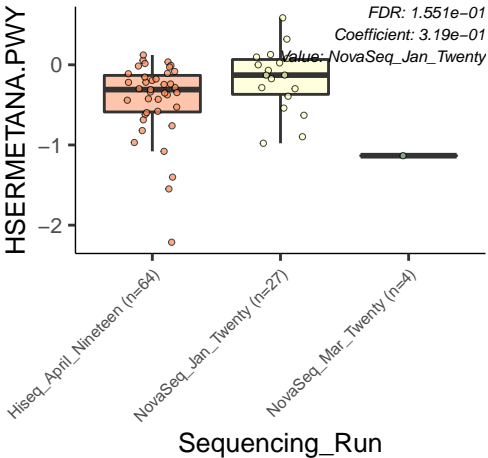
P461.PWY



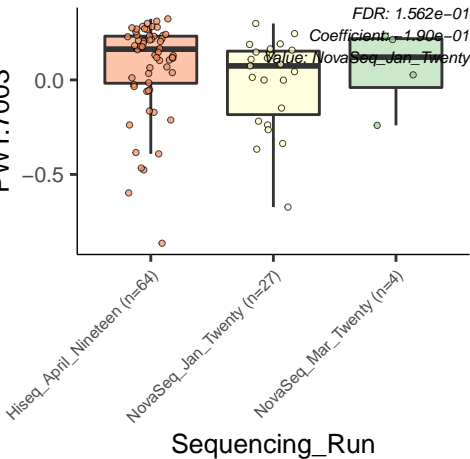
PWY.7199



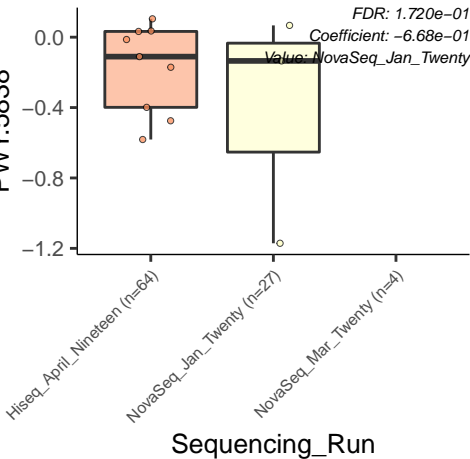
HSERMETANA.PWY



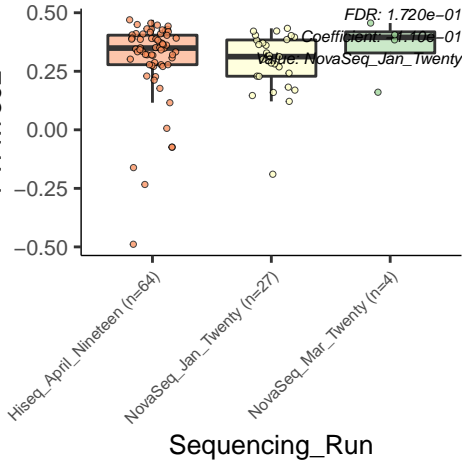
PWY.7003



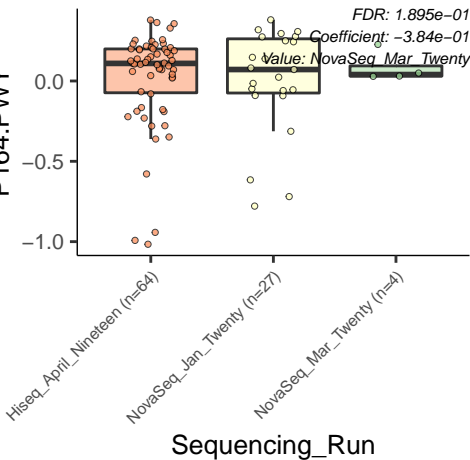
PWY.5838



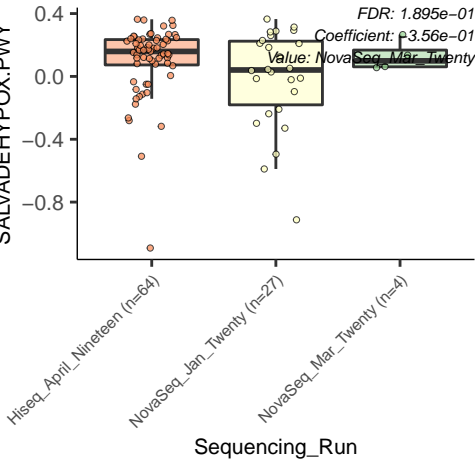
PWY.7392



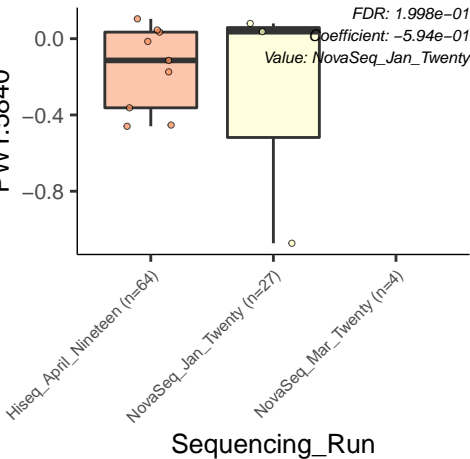
P164.PWY



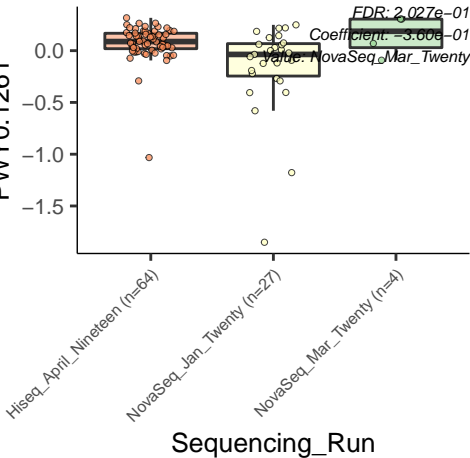
SALVADEHYPOX.PWY



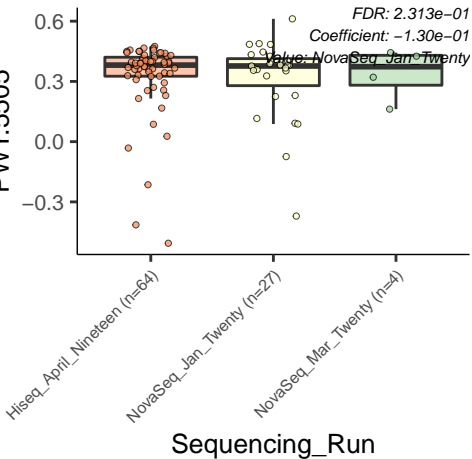
PWY.5840



PWY0.1261



PWY.5505



PWY.7328

