

ARO.PWY

FDR: $3.804e-02$

Coefficient: $1.05e-01$

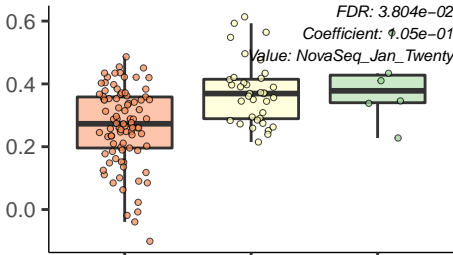
Value: NovaSeq_Jan_Twenty

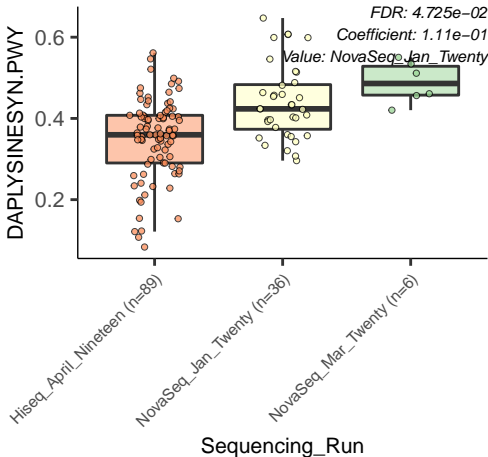
Hiseq_April_Nineteen (n=89)

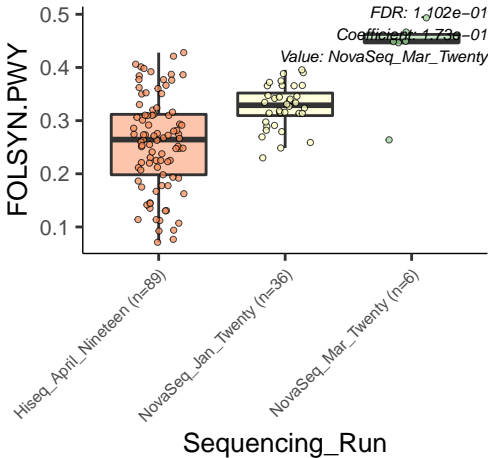
NovaSeq_Jan_Twenty (n=36)

NovaSeq_Mar_Twenty (n=6)

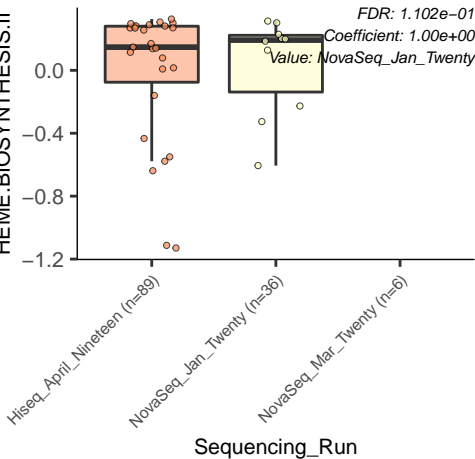
Sequencing_Run



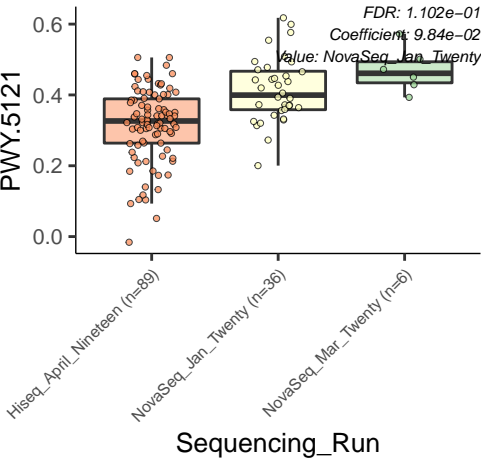




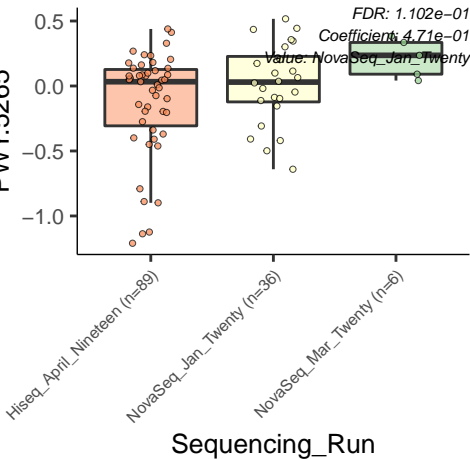
HEME.BIOSYNTHESIS.II

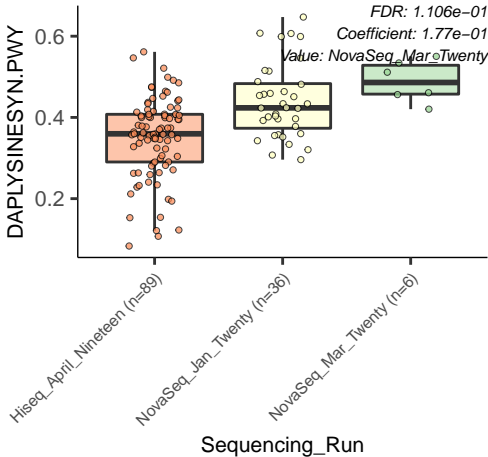


PWY.5121



PWY.5265





NONMEVIPP.PWY

FDR: 1.106e-01

Coefficient: 1.08e-01

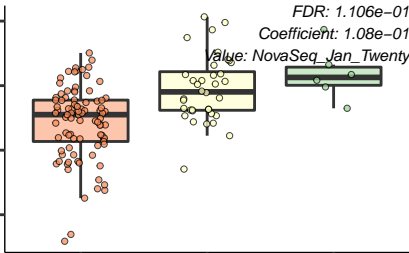
Value: NovaSeq_Jan_Twenty

Hiseq_April_Nineteen (n=89)

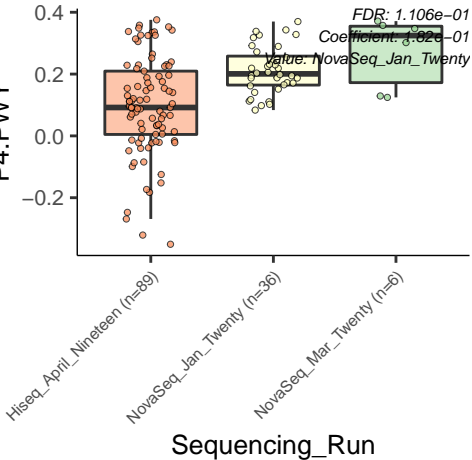
NovaSeq_Jan_Twenty (n=36)

NovaSeq_Mar_Twenty (n=6)

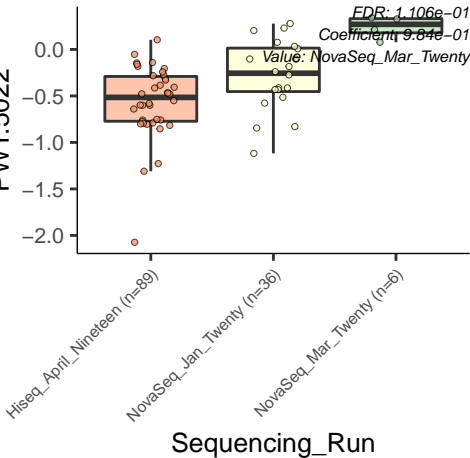
Sequencing_Run



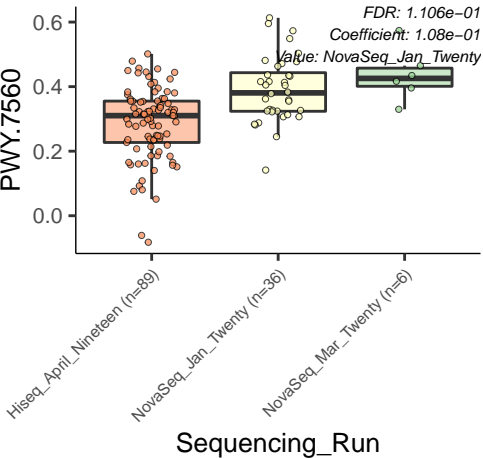
P4.PWY

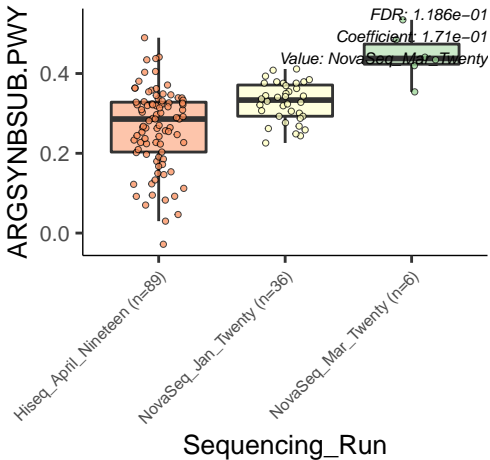


PWY.5022

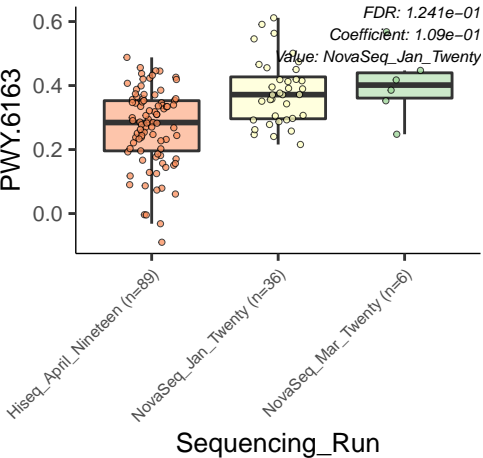


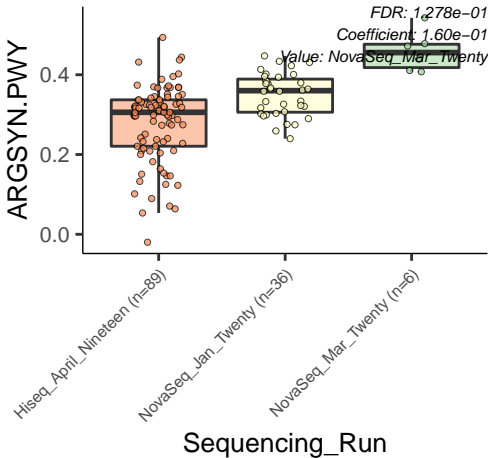
PWY.7560

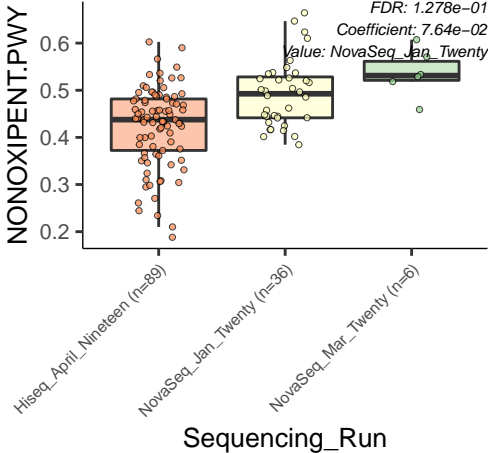




PWY.6163







PWY.7400

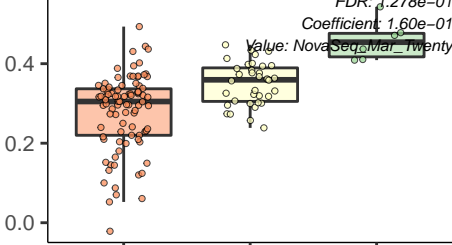
Hiseq_April_Nineteen (n=89)

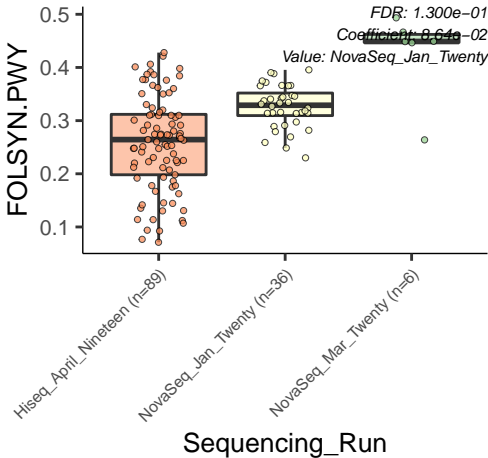
NovaSeq_Jan_Twenty (n=36)

NovaSeq_Mar_Twenty (n=6)

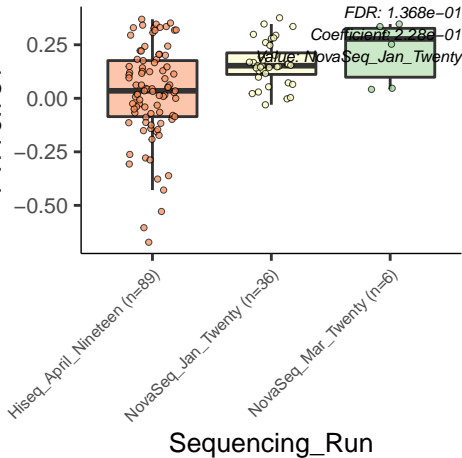
Sequencing_Run

FDR: 1.278e-01
Coefficient: 1.60e-01
Value: NovaSeq_Mar_Twenty

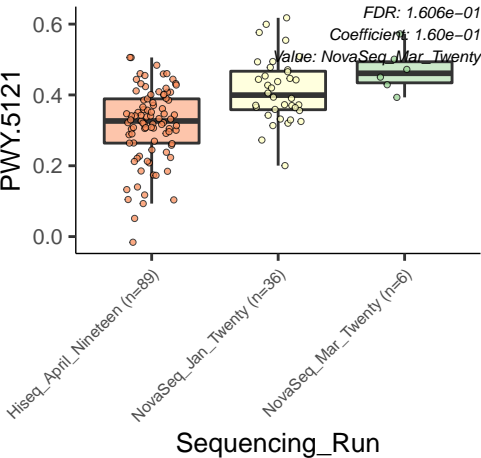




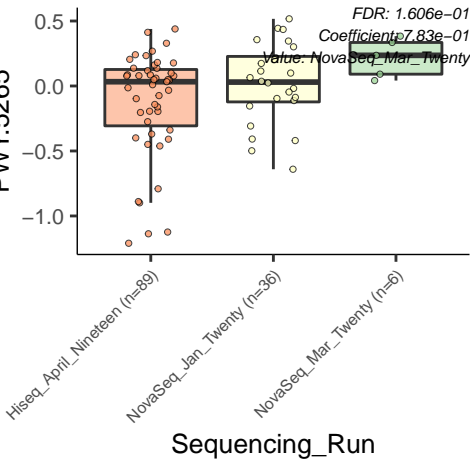
PWY0.781



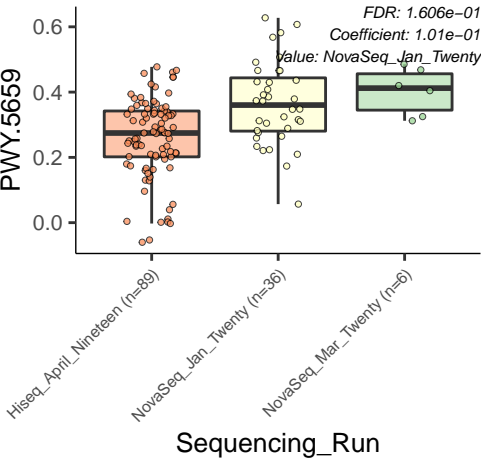
PWY.5121



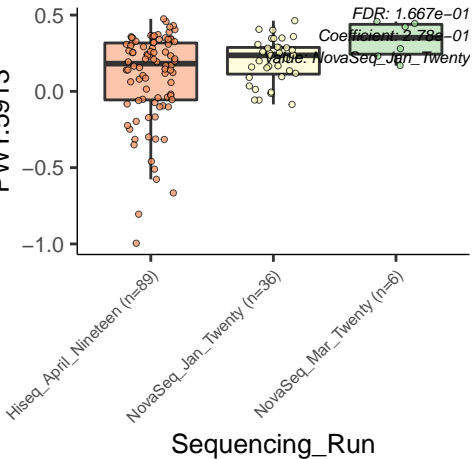
PWY.5265



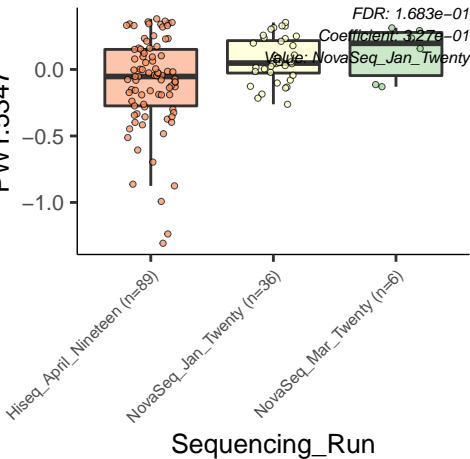
PWY.5659



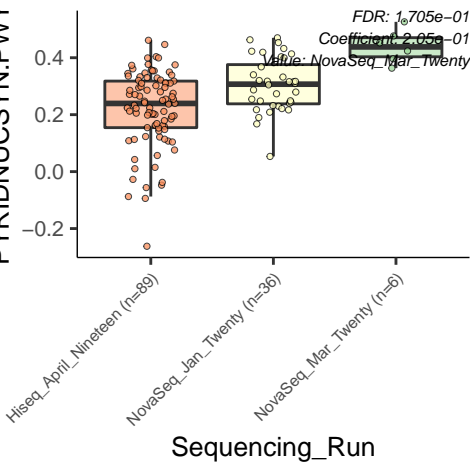
PWY.5913

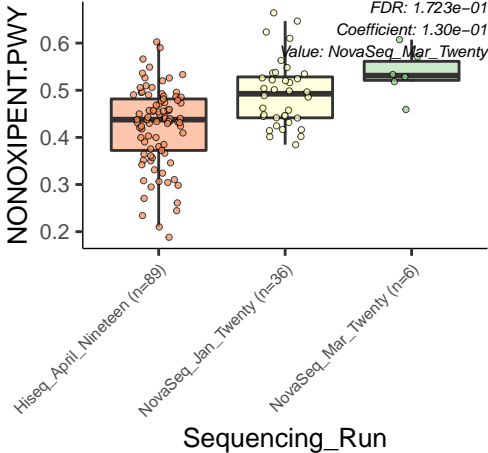


PWY.5347

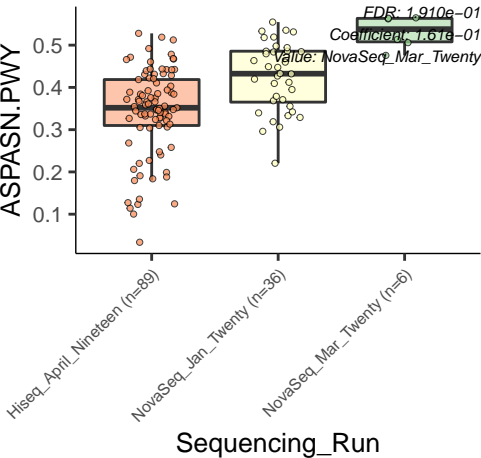


PYRIDNUCSYN.PWY

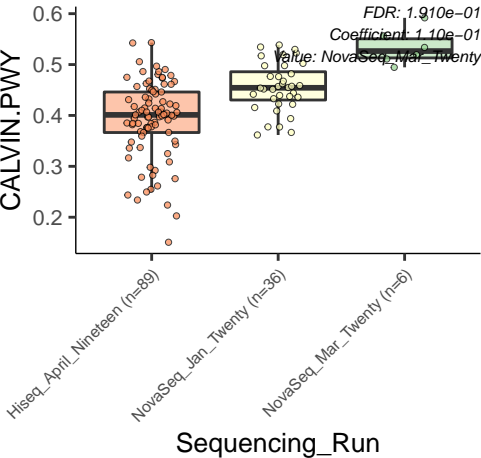




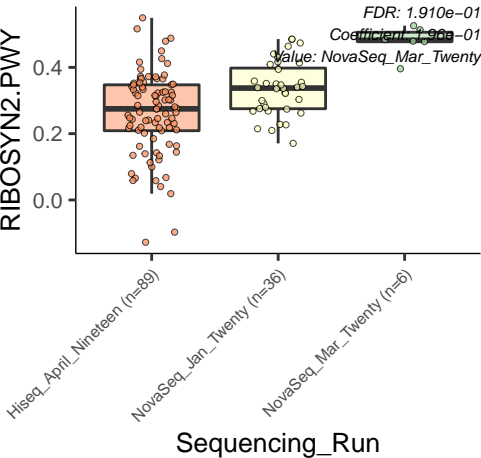
ASPASN.PWY

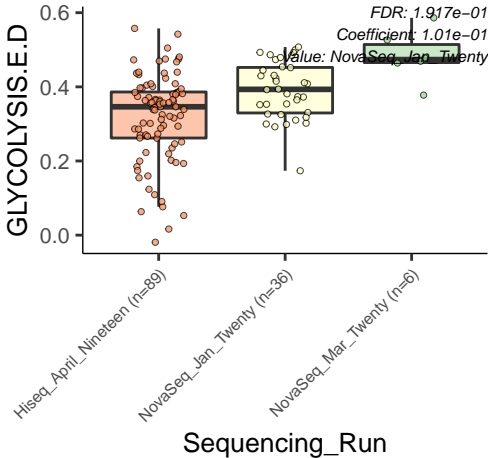


CALVIN.PWY



RIBOSYN2.PWY





PWY.5103

Hiseq_April_Nineteen (n=89)

NovaSeq_Jan_Twenty (n=36)

NovaSeq_Mar_Twenty (n=6)

Sequencing_Run

FDR: 1.917e-01

Coefficient: 1.54e-01

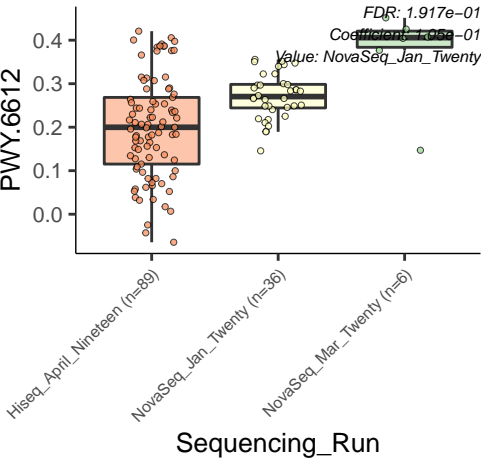
Value: NovaSeq_Mar_Twenty

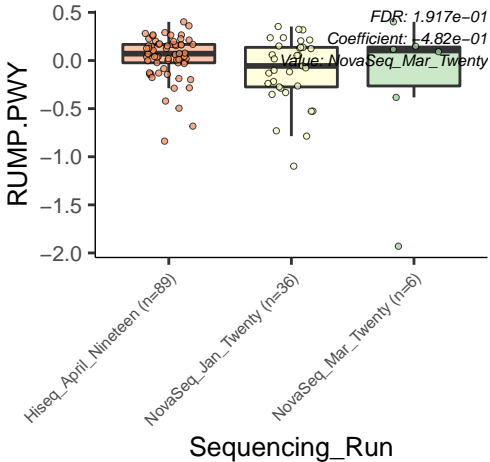
0.4

0.2

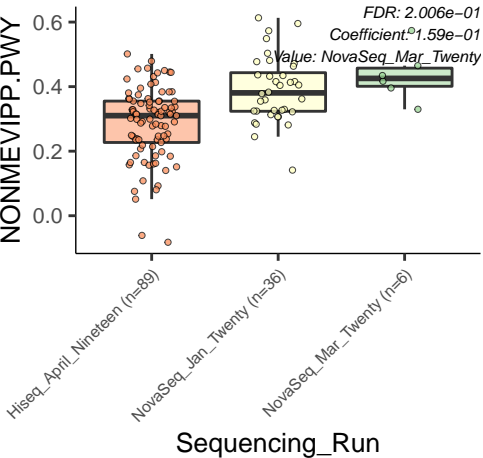
0.0

PWY.6612





NONMEVIPP.PWY



PWY.5103

Hiseq_April_Nineteen (n=89)

NovaSeq_Jan_Twenty (n=36)

NovaSeq_Mar_Twenty (n=6)

Sequencing_Run

FDR: 2.006e-01

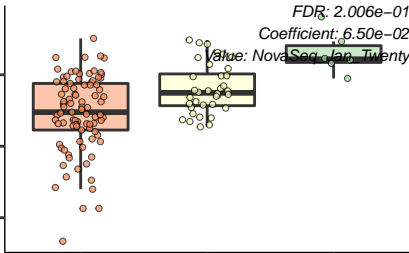
Coefficient: 6.50e-02

Value: NovaSeq_Jan_Twenty

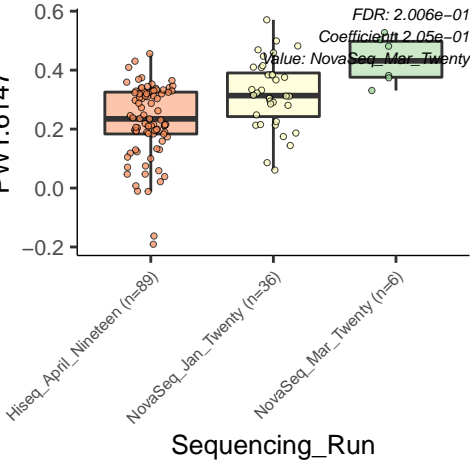
0.4

0.2

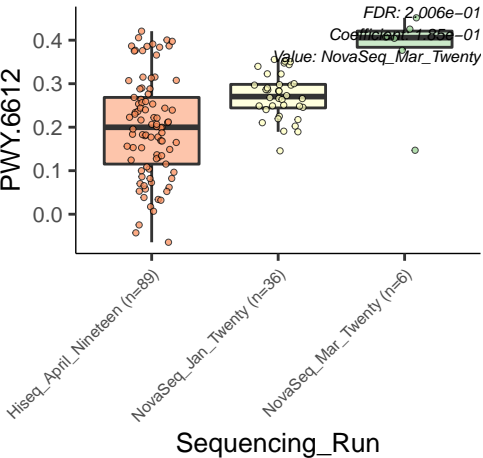
0.0



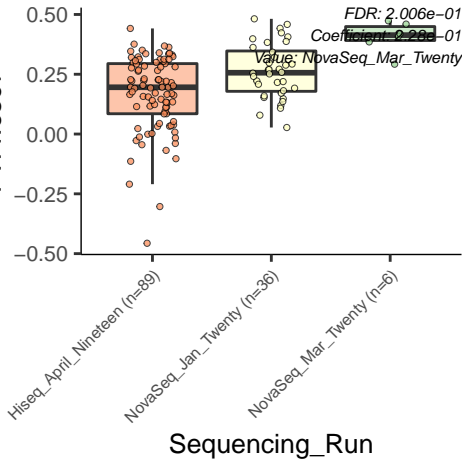
PWY.6147



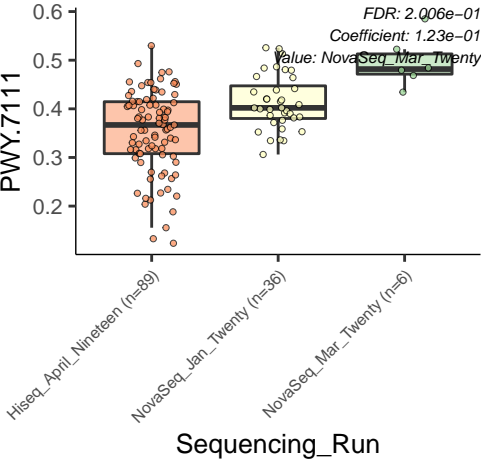
PWY.6612



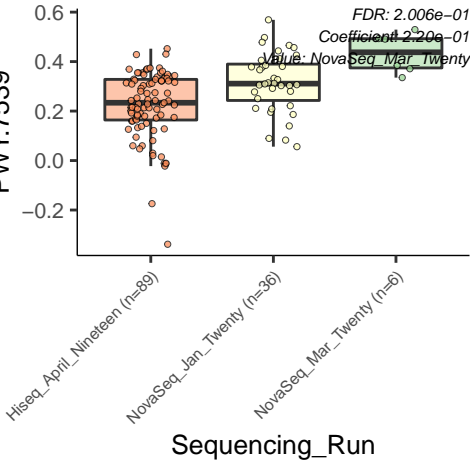
PWY.6897



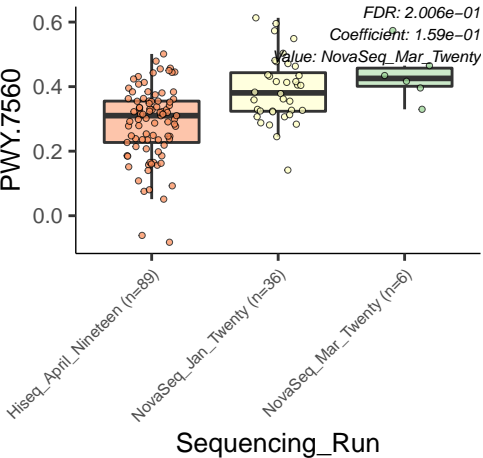
PWY.7111



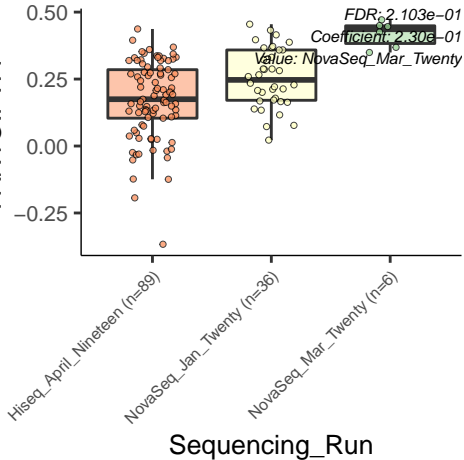
PWY.7539



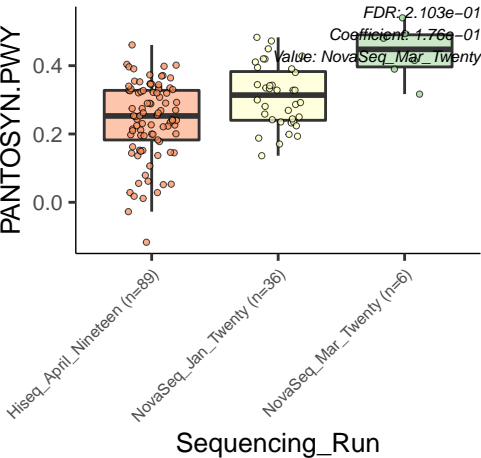
PWY.7560

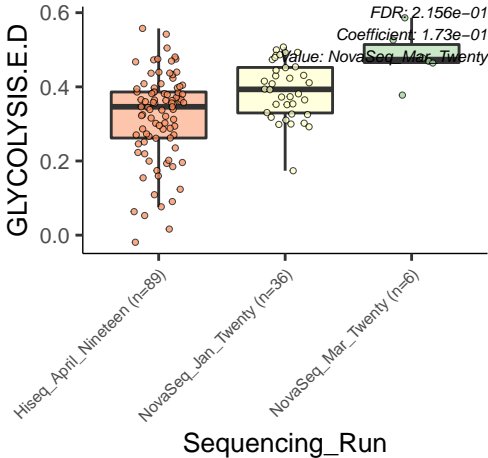


PANTO.PWY

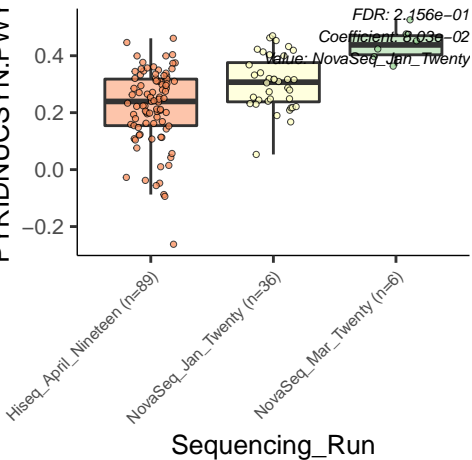


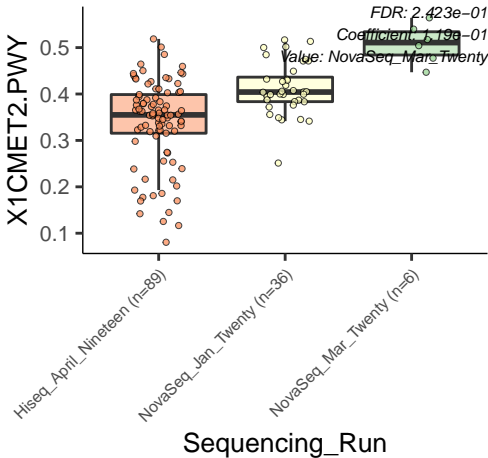
PANTOSYN.PWY

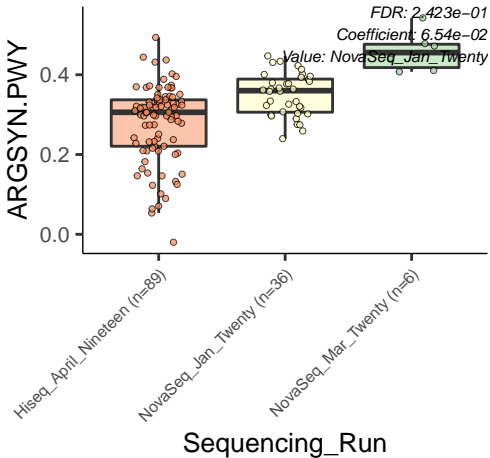


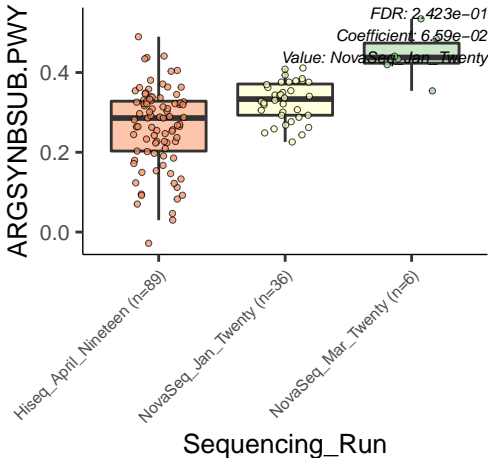


PYRIDNUCSYN.PWY

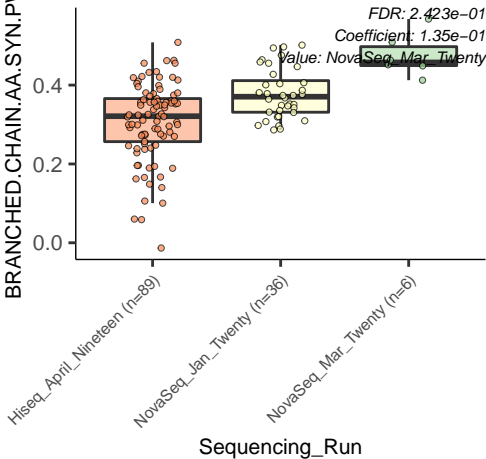




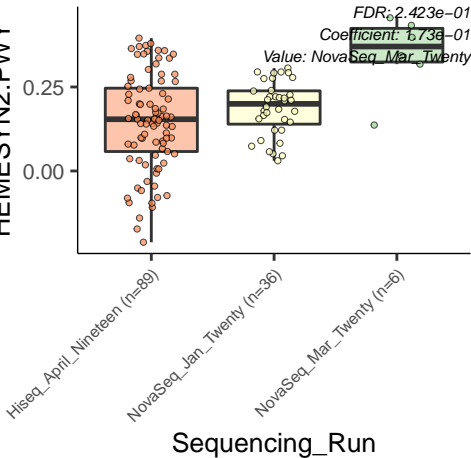




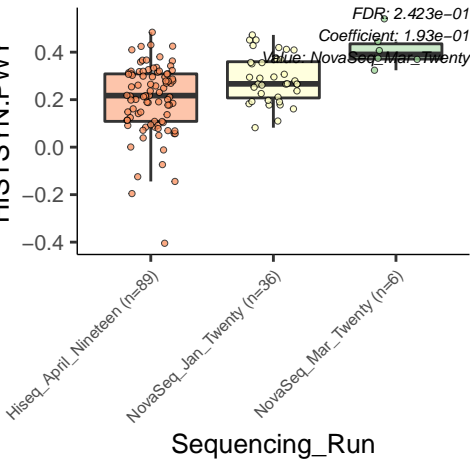
BRANCHED.CHAIN.AA.SYN.PV



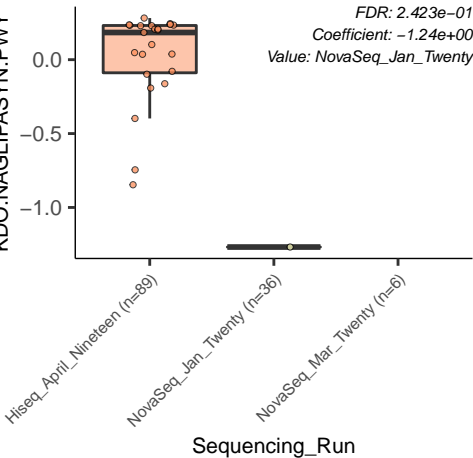
HEMESYN2.PWY



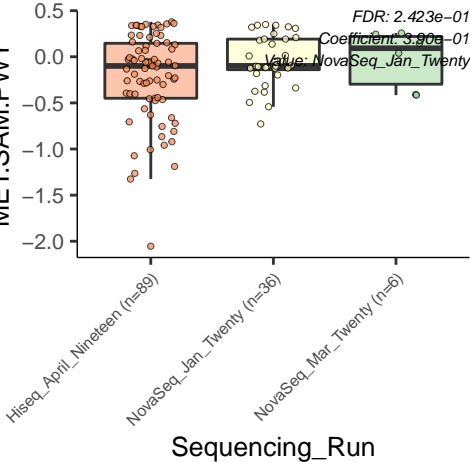
HISTSYN.PWY



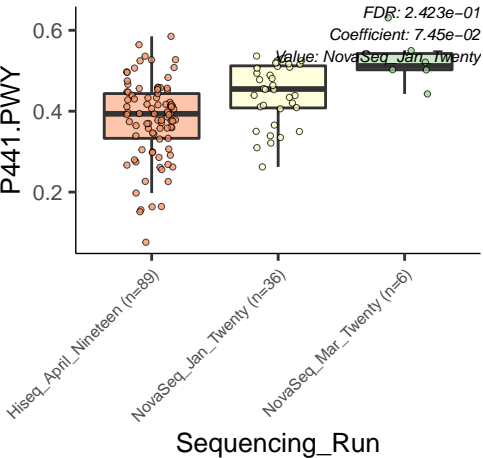
KDO.NAGLIPASYN.PWY



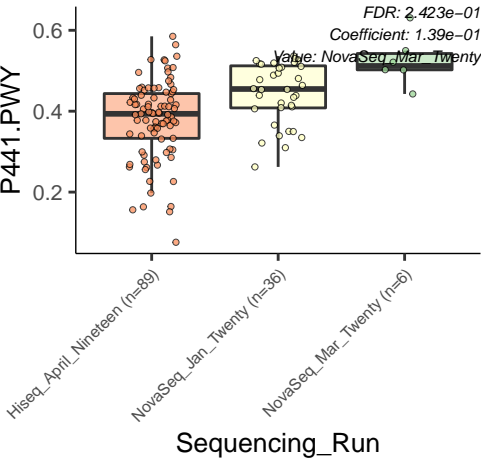
MET.SAM.PWY



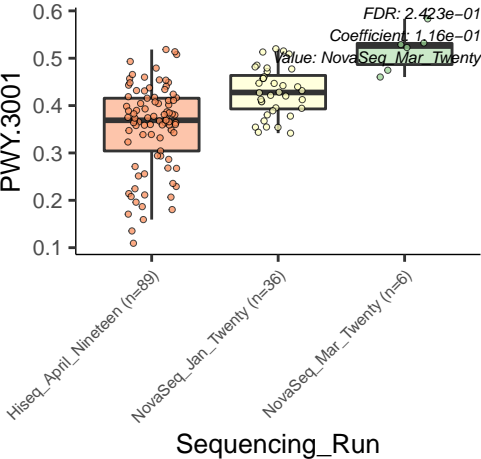
P441.PWY

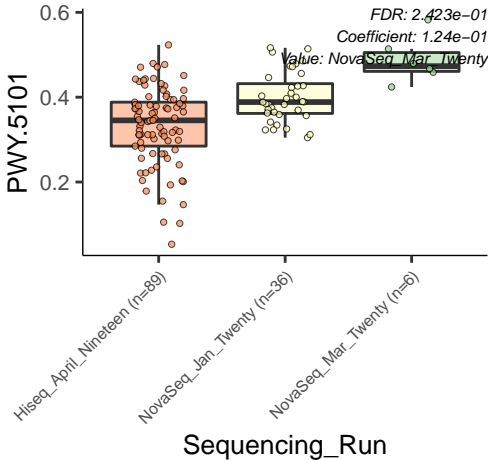


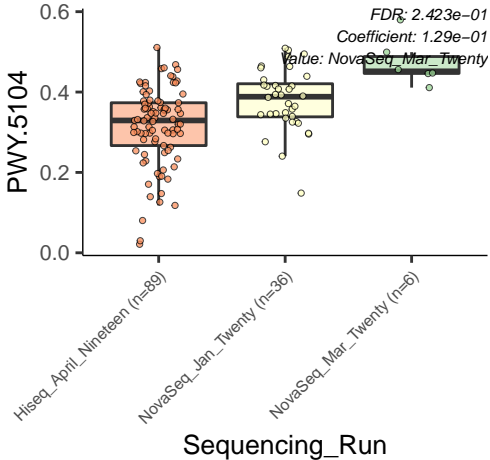
P441.PWY



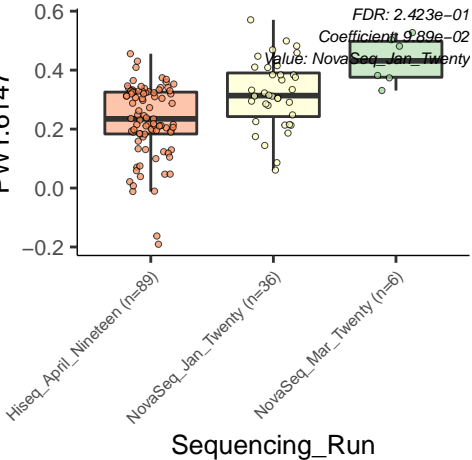
PWY.3001



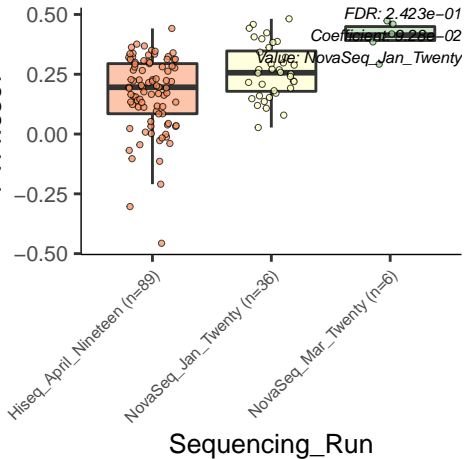




PWY.6147



PWY.6897



PWY.7400

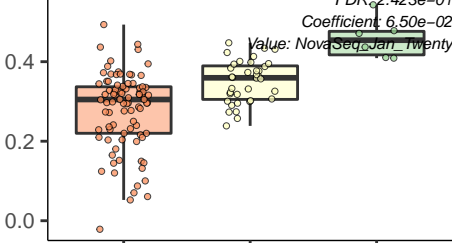
Hiseq_April_Nineteen (n=89)

NovaSeq_Jan_Twenty (n=36)

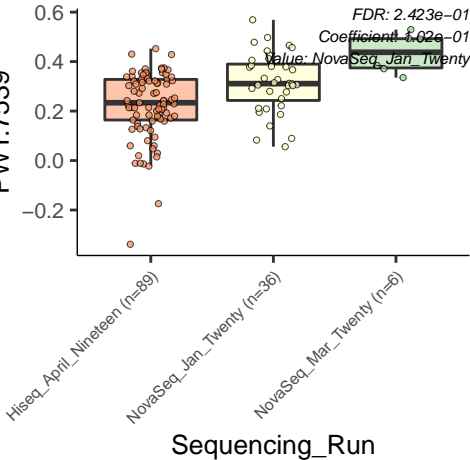
NovaSeq_Mar_Twenty (n=6)

Sequencing_Run

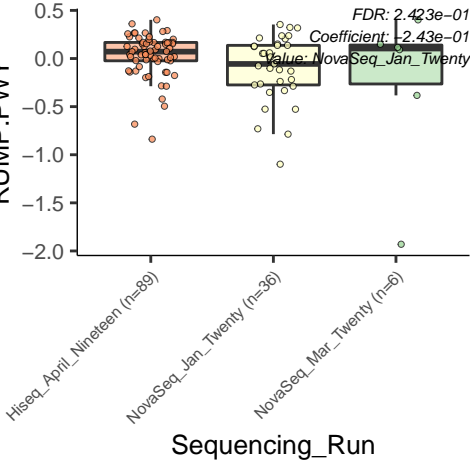
FDR: 2.423e-01
Coefficient: 6.50e-02
Value: NovaSeq_Jan_Twenty

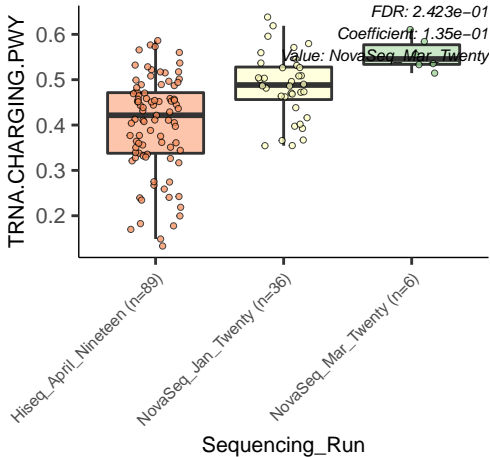


PWY.7539

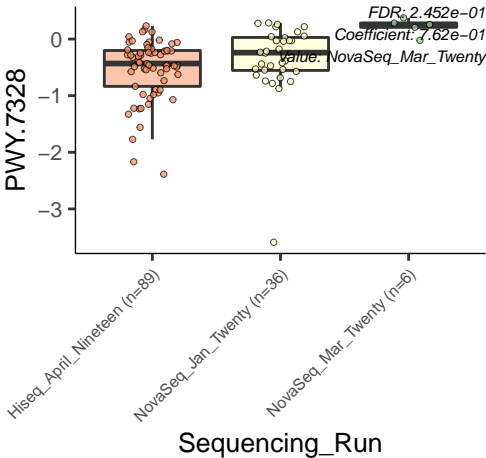


RUMP.PWY

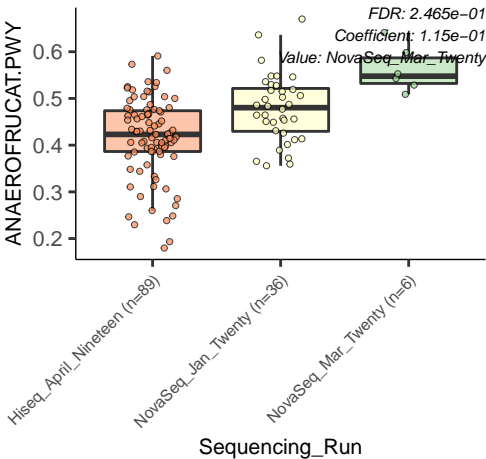




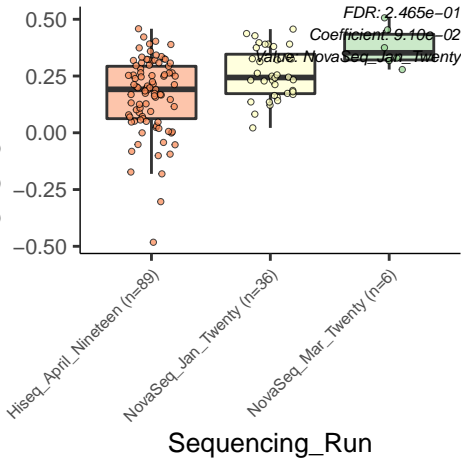
PWY.7328



ANAEROFRUCAT.PWY



GLUTURN.PWY



HISTSYN.PWY

FDR: $2.465e-01$

Coefficient: $8.07e-02$

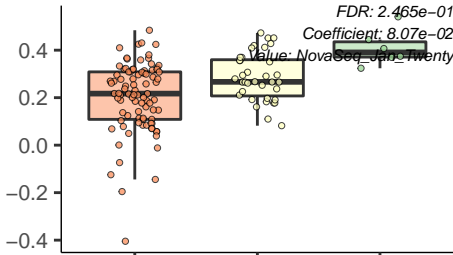
Value: NovaSeq_Jan_Twenty

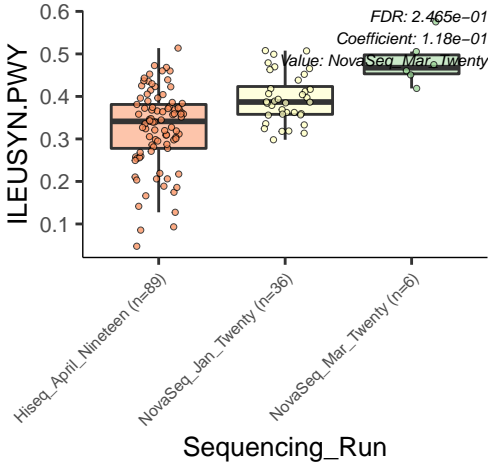
Hiseq_April_Nineteen (n=89)

NovaSeq_Jan_Twenty (n=36)

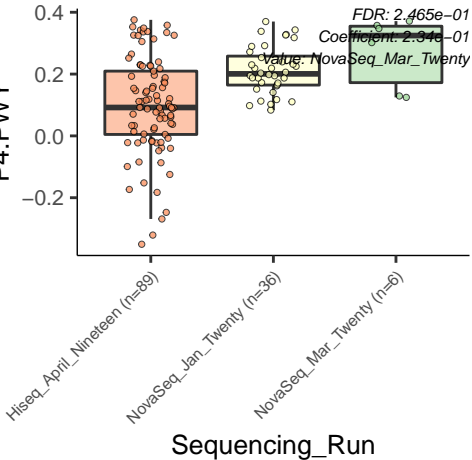
NovaSeq_Mar_Twenty (n=6)

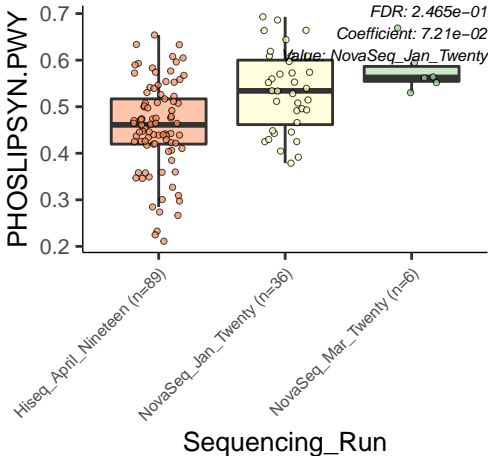
Sequencing_Run

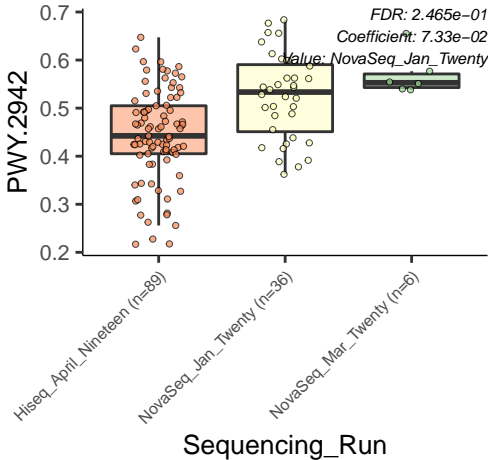




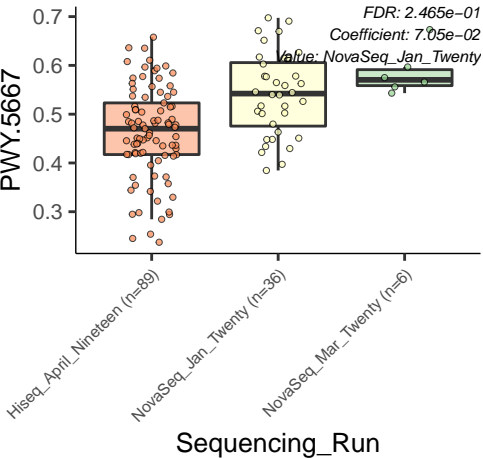
P4.PWY



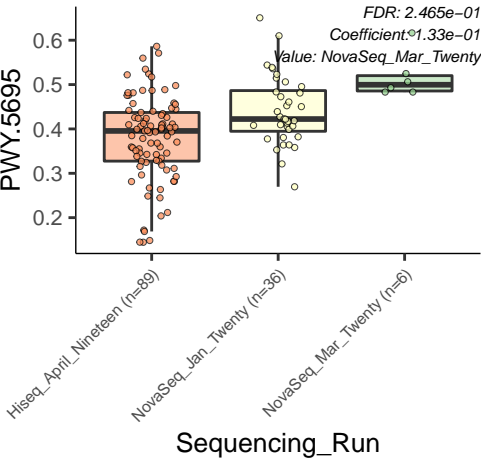




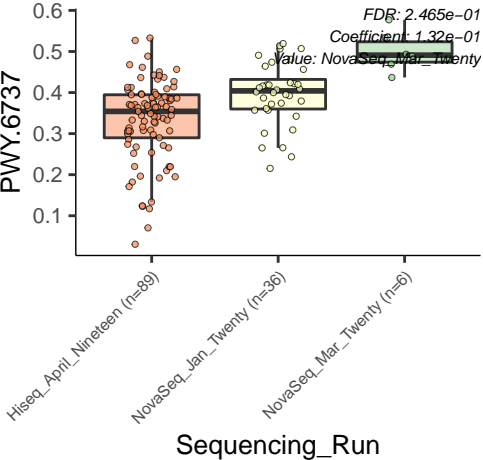
PWY.5667



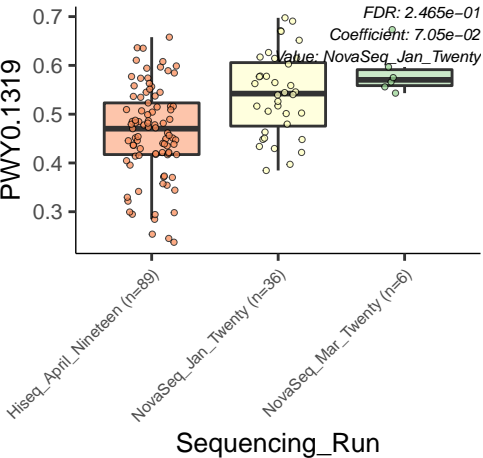
PWY.5695



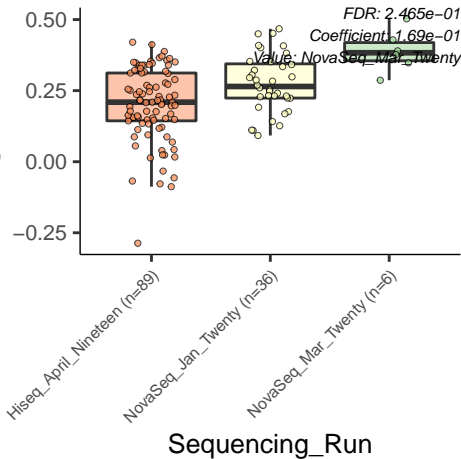
PWY.6737

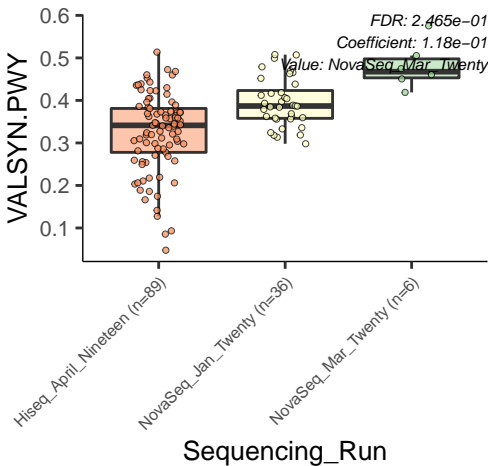


PWY0.1319

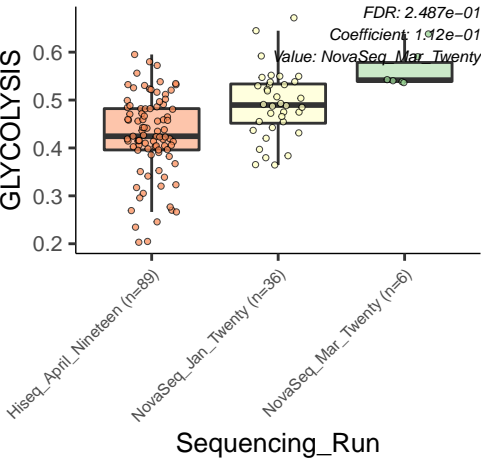


TRPSYN.PWY

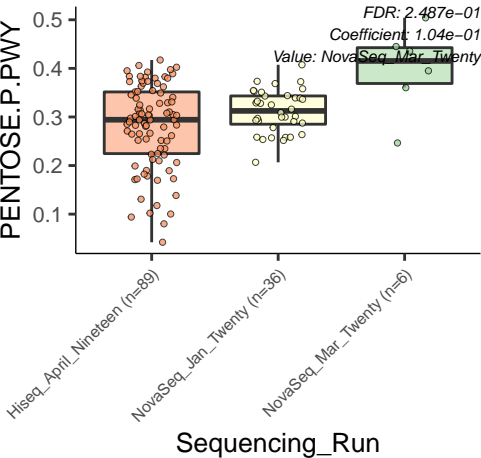




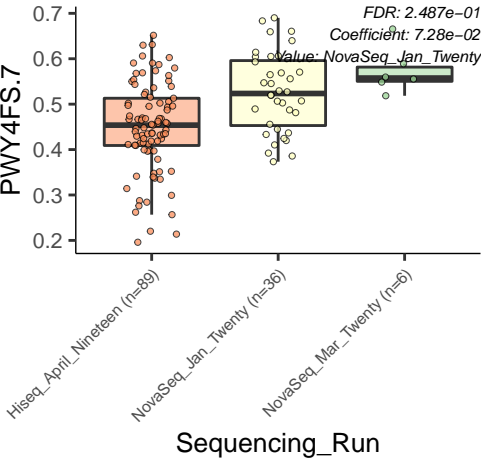
GLYCOLYSIS



PENTOSE.P.PWY



PWY4FS.7



PWY4FS.8

