a.
CHEQ-seq 5′ - H3K27ac enhancers
Log2 FC
Enhancers
Enhancer Phenotype
MES
MEL
COL5A1
SERPINE1
Others
Adjusted p–value
<0.05
>=0.05
b.
Pearson's r = 0.8292
Spearman's rho = 0.7437
CHEQ–seq Intron Log2 FC
CHEQ–seq 5' Log2 FC
Enhancer phenotype
MES
MEL
c.
STARR-seq - ATAC-seq enhancers
Log2 FC
Enhancers
Enhancer Phenotype
MES
MEL
COL5A1
SERPINE1
Others
Adjusted p–value
<0.05
>=0.05
d.
Pearson's r = 0.6569
Spearman's rho = 0.6816
STARR–seq Log2 FC
CHEQ–seq Log2 FC
Enhancer phenotype
MES
MEL
e.
Pearson's r = 0.5205
Spearman's r = 0.4418
CHEQ–seq ATAC Log2 FC
CHEQ–seq Intron H3K27ac Log2 FC
Enhancer phenotype
MES
MEL