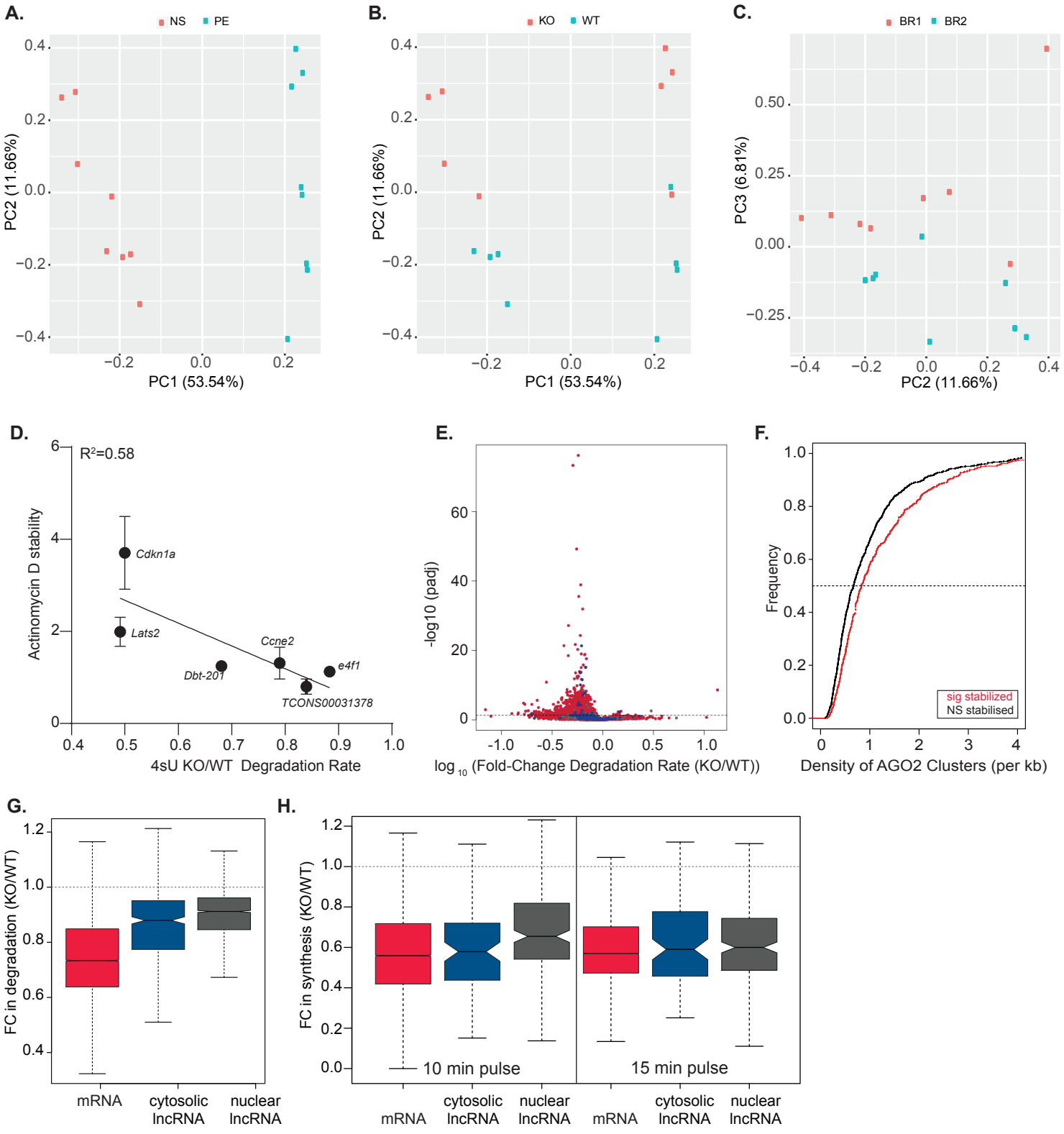
Figure S2


A.
NS
PE
PC2 (11.66%)
PC1 (53.54%)
B.
KO
WT
PC2 (11.66%)
PC1 (53.54%)
C.
BR1
BR2
PC3 (6.81%)
PC2 (11.66%)
D.
R²=0.58
Actinomycin D stability
Cdkn1a
Lats2
Ccne2
Dbt-201
e4f1
TCONS00031378
4sU KO/WT Degradation Rate
E.
-log10 (padj)
log 10 (Fold-Change Degradation Rate (KO/WT))
F.
Frequency
sig stabilized
NS stabilised
Density of AGO2 Clusters (per kb)
G.
FC in degradation (KO/WT)
mRNA
cytosolic lncRNA
nuclear lncRNA
H.
FC in synthesis (KO/WT)
10 min pulse
15 min pulse
mRNA
cytosolic lncRNA
nuclear lncRNA
mRNA
cytosolic lncRNA
nuclear lncRNA