A
Study of origin
PABPC1L 10%
ALYREF 0.8%
PPIH 0.2%
RAE1 9%
XPO5 1%
XPOT 0%
OAS3 0%
Genetic alteration
Amplification
No alteration
Study of origin
COAD (TCGA, Nature2012)
COAD (TCGA, Provisional)
B
84
2512
-log10 pvalue (BH-adjusted)
△PSI(high vs low)
C
siNC
siPABPC1L
Relative mRNA/GAPDH expression (Fold change)
PABPC1L
NOXA1-L
NOXA1-s
SMARCD3-L
SMARCD3-s
METTL3-L
METTL3-s
ACCS-L
ACCS-s
EZH1-L
EZH1-s
D
Ratio (aberrant spliced /normal)
NOXA1
SMARCD3
METTL3
CD46
ACCS
EZH1
NS
E
PABPC1L
MYC
Relative mRNA/RPL9 expression (fold change)
siNC
siPABPC1L
siNC
siMYC