

Table SI. YTHDF2 siRNA sequences.

Name	Sequence (5'-3')	Molecular weight, Da	Temperature, °C
YTHDF2 si-1 sense	AAGGACGUUCCCAAUAGCCAATT	7848.0	58.40
YTHDF2 si-1 antisense	UUGGCUAUUGGGAACGUCCUUTT	7275.4	58.40
YTHDF2 si-2 sense	GCUACUCUGAGGACGAUAUUCTT	7818.9	58.40
YTHDF2 si-2 antisense	GAAUAUCGUCCUCAGAGUAGCTT	7304.5	58.40
YTHDF2 si-3 sense	CCUAGGUAGCACUCCAUUUCUTT	7715.8	58.40
YTHDF2 si-3 antisense	AGAAAUGGAGUGCUACCUAGGTT	7407.6	58.40
NC sense	UUCUCCGAACGUGUCACGUdTdT	1.	2.
NC antisense	ACGUGACACGUUCGGAGAAdTdT	3.	4.

YTHDF2, YTH domain-containing family protein 2; si-/siRNA, small interfering RNA; NC, negative control.

Table SII. Two-way ordered categorical variables with different attributes between different m1A clusters and different risk stratifications.

ISS stage	Cluster A (n=248)	Cluster B (n=190)	Cluster C (n=121)
ISS stage I (n=315)	155	94	66
ISS stage II (n=177)	71	69	37
ISS stage III (n=67)	22	27	18

Pearson's χ^2 test: $\chi^2=9.0291$; degrees of freedom=4; P-value=0.06038. ISS, International Staging System.