

Supplementary figures and legend

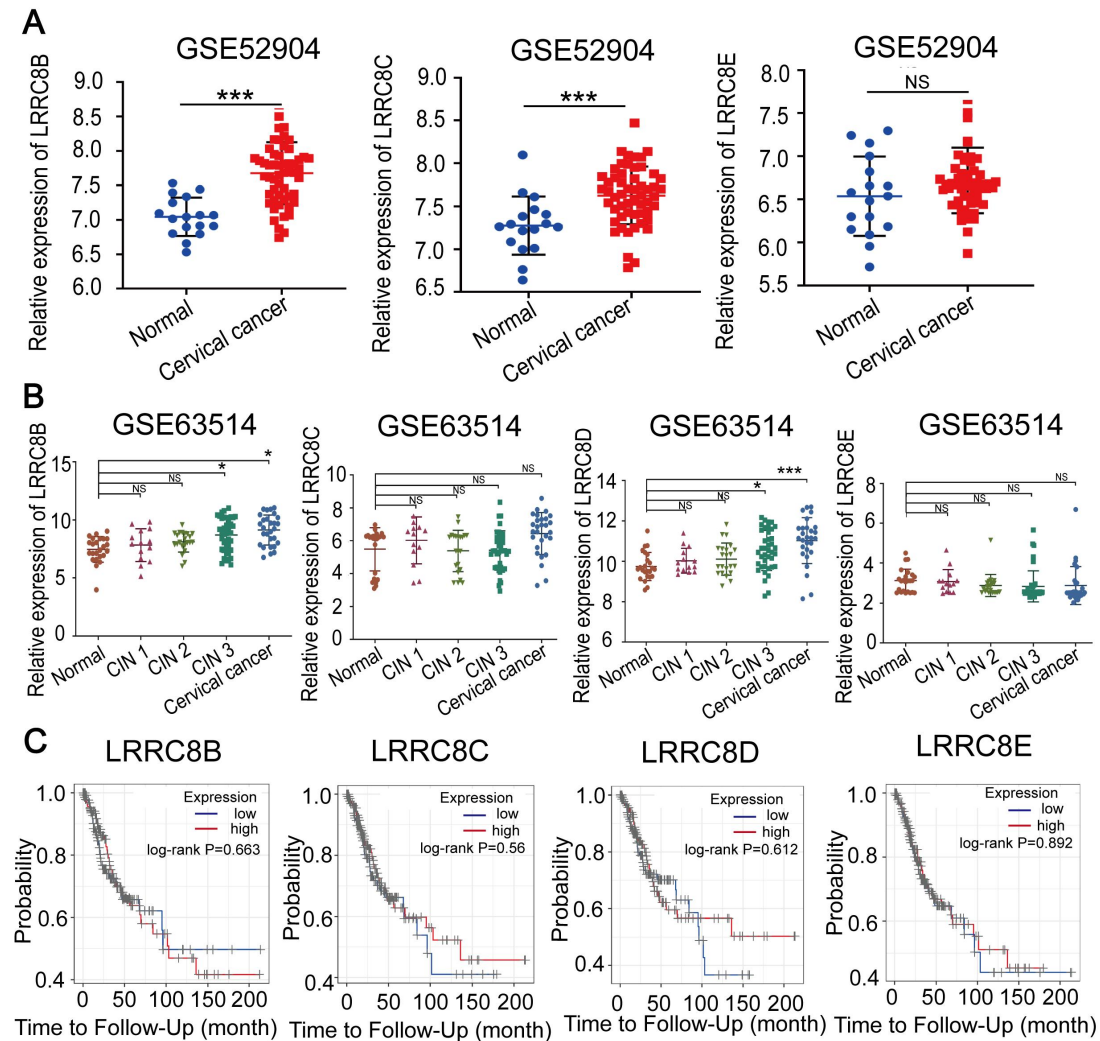


Fig S1. (A and B) Relative RNA levels of LRRC8B, LRRC8C and LRRC8E in CC tissues and normal cervical epithelium in GEO datasets (GSE52904 and GSE63514). **(C)** Kaplan-Meier analysis of CC patients' RFS based on LRRC8B, LRRC8C, LRRC8D, LRRC8E expression by TIMER website. * $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, NS, not significant.

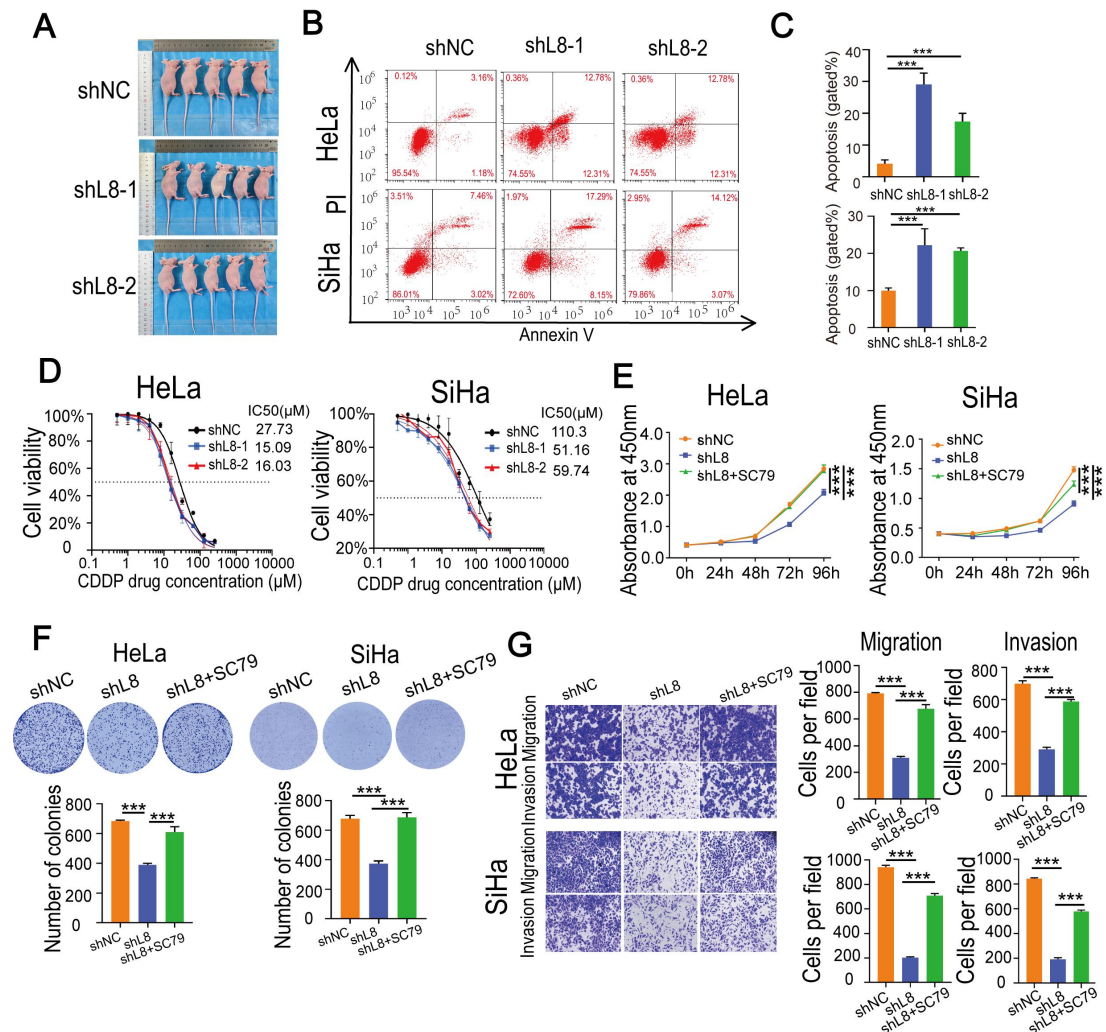


Fig S2. (A) Images of xenograft tumors derived from subcutaneous implantation of LRRC8A knockdown and control cells on nude mice. **(B)** The effect of LRRC8A knockdown on apoptosis of cervical cancer cells. **(C)** Statistical analysis percentage of early-and late-apoptotic cells in LRRC8A knockdown cells. **(D)** Cell viability of control and LRRC8A knockdown cells after treatment with increasing concentrations of cisplatin for 24 h. n = 3. **(E)** The AKT phosphorylation agonist rescues CC cells' proliferation suppressed by LRRC8A depletion. **(F)** SC79 treatment restores the colony formation of CC cells suppressed by LRRC8A knockdown. **(G)** Cell migration and invasion analysis of LRRC8A knockdown in HeLa and SiHa cells with or without SC79 treatment. Scale bar, 200 μ m. Data are presented as means \pm S.D: ***P < 0.001.

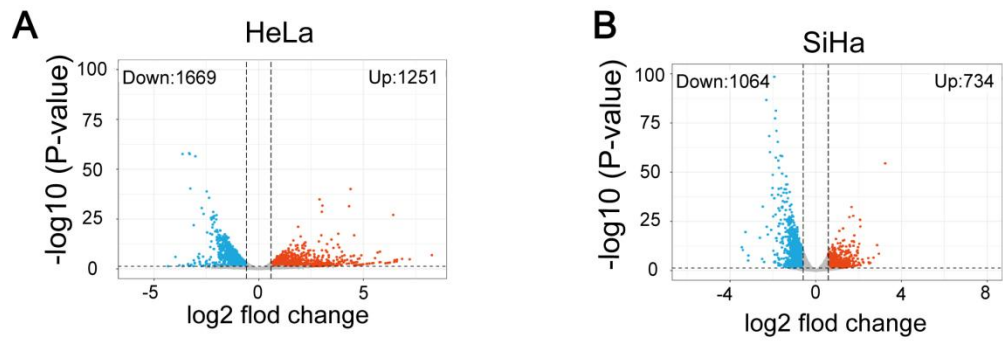


Fig S3. (A) Volcano plot of differentially expressed genes from RNA-seq analysis of NSUN2 knockdown in HeLa cells. **(B)** Volcano plot of differentially expressed genes from RNA-seq analysis of NSUN2 knockdown in SiHa cells.

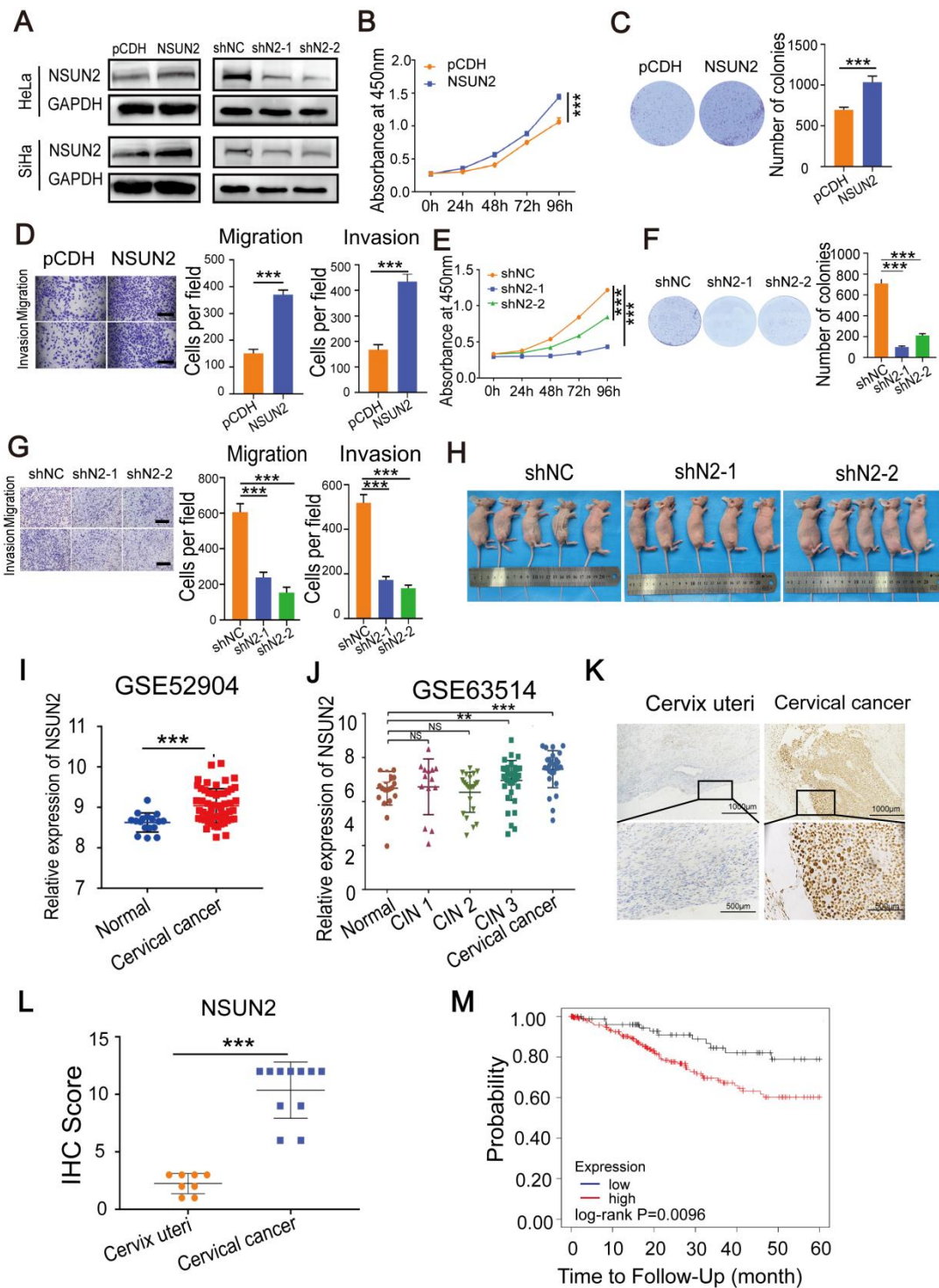


Fig S4. (A) Protein expression of NSUN2 overexpression and knockdown in HeLa and SiHa cells. **(B)** CCK-8 assays showed the proliferation of SiHa cells upon NSUN2 overexpression. **(C)** Cell colony formation of SiHa cells with NSUN2 overexpression. **(D)** Transwell cell migration and invasion analysis of NSUN2 overexpression and control cells. Scale bar, 200 μ m. **(E)** The effect of NSUN2 knockdown on cell growth was determined by CCK-8 assays. **(F)**

Colony formation assays were performed in NSUN2 knockdown and control cells. **(G)** Migration and invasion analysis of SiHa cells upon NSUN2 knockdown. Scale bar, 200 μm . **(H)** Images of xenograft tumors derived from subcutaneous implantation of NSUN2 knockdown and control cells on nude mice. **(I and J)** Relative RNA levels of NSUN2 in CC tissues and normal cervical epithelium in GEO datasets (GSE52904 and GSE63514). **(K)** Representative IHC staining images of NSUN2 protein levels in CC tissues and cervical epithelium tissues. Scale bar, upper: 100 μm , lower: 500 μm . **(L)** The quantitative analysis of NSUN2 protein expression in CC tissues and cervical epithelium tissues assessed by IHC staining. **(M)** Kaplan-Meier analysis of CC patients' RFS based on NSUN2 expression. Data are presented as means \pm S.D: ***P < 0.001.

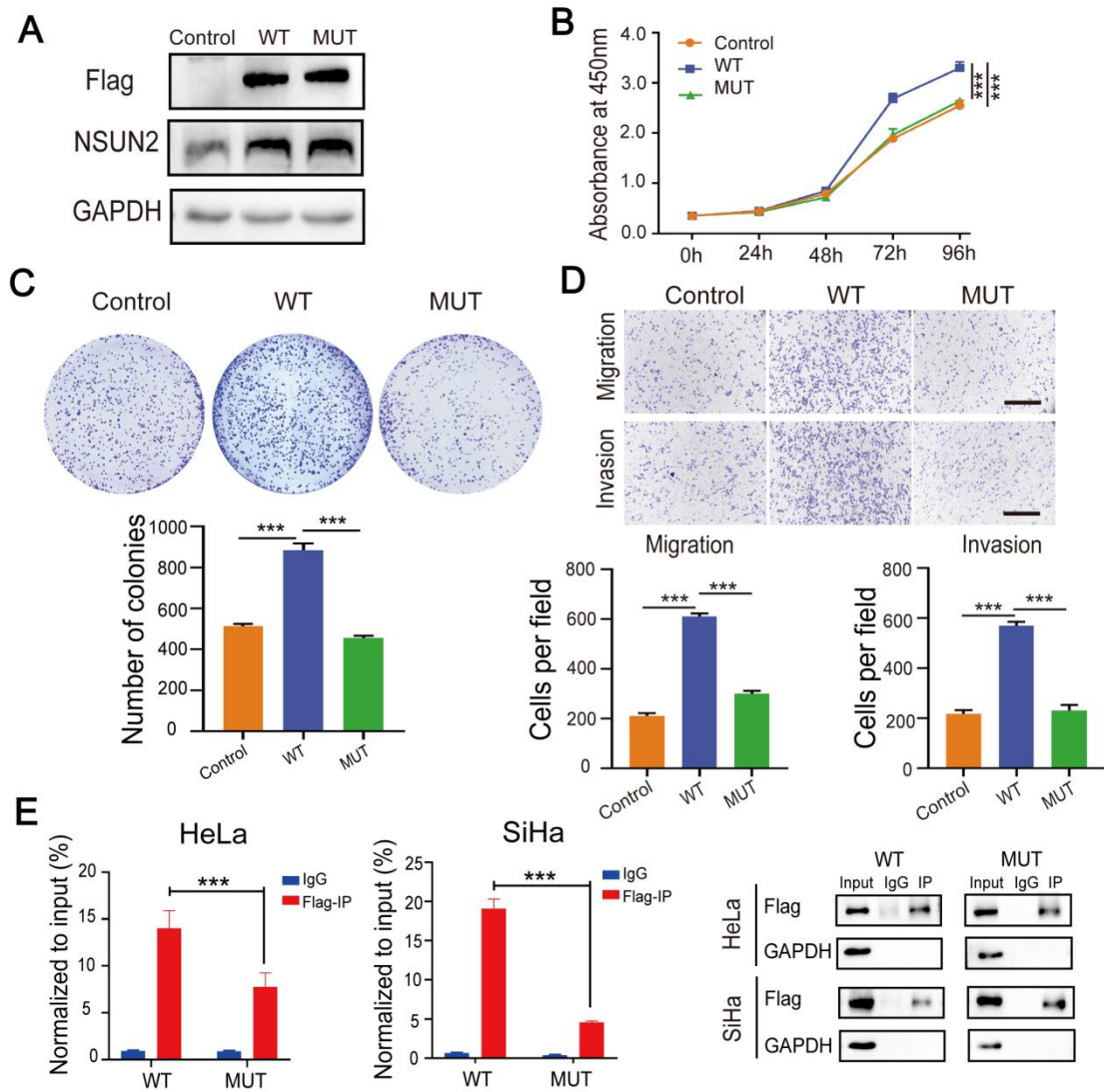


Fig S5. (A) Western blotting assays showed the protein levels of NSUN2 in NSUN2-WT, NSUN2-MUT and control HeLa cells. **(B)** CCK-8 assays of HeLa cells with forced expression of wild-type or catalytic-mutant NSUN2. **(C)** Colony formation assays were performed in CC cells with forced expression of wild-type or mutated NSUN2. **(D)** Transwell cell migration and invasion assay of HeLa cells with forced expression of wild-type or mutated NSUN2. Scale bar, 200 μ m. **(E)** RIP-PCR assays detecting the interactions between wild-type or mutated NSUN2 and LRRRC8A mRNA. IgG was used as an internal control. GAPDH was used as the negative control in western blot assays. Data are presented as means \pm S.D: ***P < 0.001.

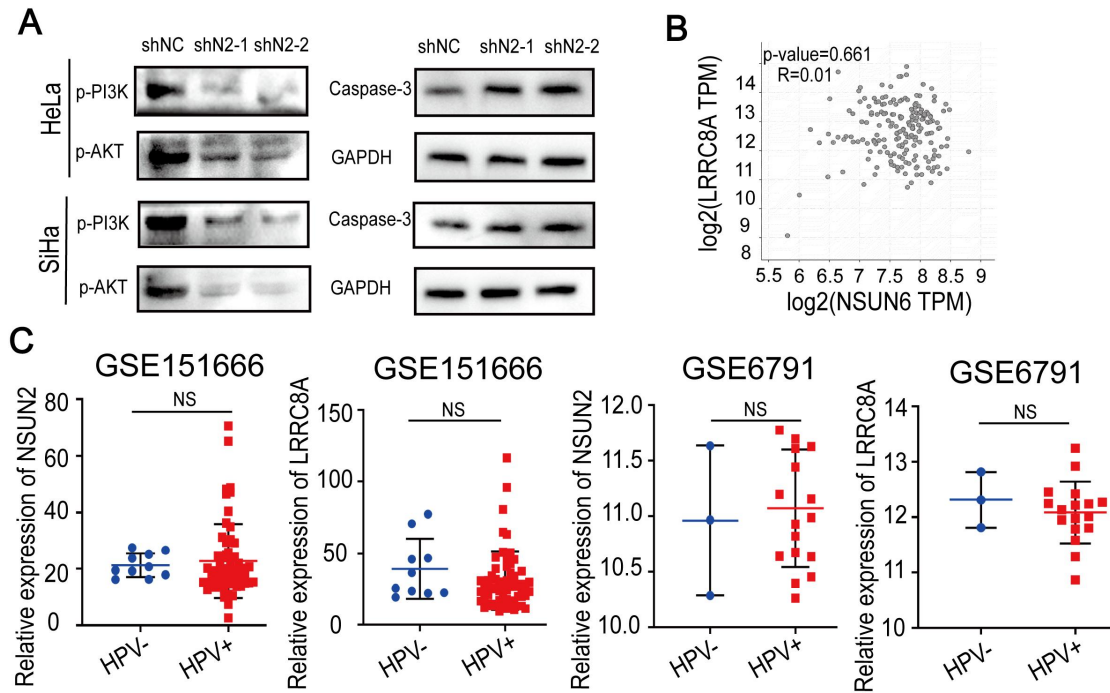


Fig S6. (A) Western blot assays showed protein levels of p-AKT, p-PI3K and Caspase-3 in HeLa and SiHa cells upon NSUN2 knockdown. **(B)** Spearman's correlation analysis between mRNA levels of NSUN6 and LRRC8A in cervical cancer. **(C)** Relative RNA levels of NSUN2 and LRRC8A in HPV negative and HPV positive CC tissues in GEO datasets. NS, not significant.

Supplementary table 1: Sequences of primers and oligos used in this study

	Forword	Reverse
NSUN2	CAAGCTGTTTCGAGCACTA CTAC	CTCCCTGAGAGCGTCCAT GA
GAPDH	CTGGGCTACACTGAGCA CC	AAGTGGTCGTTGAGGGCA ATG
LRR8A	TGTGTGCTATGAGAACCG ACT	GTGCGCGGGAATTTGAAC C
YBX1	GGGGACAAGAAGGTCAT CGC	CGAAGGTACTTCCTGGGG TTA
18S	CGATAACGAACGAGACTC TGCC	CGGACATCTAAGGGCATC ACA
shLRR8A-1	CCGGCCGCAACAAGATC GAGAAGATCTC GAGATCTTCTCGATCTTG TTGCGGTTTTTG	AATTCAAAAACCGCAACAA GATCGAGAAG ATCTCGAGATCTTCTCGAT CTTGTTGCGG
shLRR8A-2	CCGGCATCTGCTACACCG TCTACTACTCGAGTAGTA GACGGTGTAGCAGATGTT TTTG	AATTCAAAAACATCTGCTAC ACCGTCTACTACTCGAGTA GTAGACGGTGTAGCAGATG
shNSUN2-1	CCGGGCGTGTTAGAAATC ACTTGTCTCGAGAACAA GTGATTTCTAACACGCTT TTTG	AATTCAAAAAGCGTGTTAG AAATCACTTGTCTCGAGA ACAAGTGATTTCTAACACG C
shNSUN2-2	CCGGGCACACAAATTTAA GTCGAAACTCGAGTTTCG ACTTAAATTTGTGTGCTTT TTG	AATTCAAAAAGCACACAAA TTTAAGTCGAAACTCGAGT TTCGACTTAAATTTGTGTG C
shYBX1	CCGGCCAGTTCAAGGCA GTAAATATCTCGAGATAT TACTGCCTTGAAGTGGT TTTTG	AATTCAAAAACAGTTCAA GGCAGTAAATATCTCGAGA TATTTACTGCCTTGAAGT G