

Figure S1. Cell line screening and flow cytometry. (A) Reverse transcription-quantitative PCR analysis was performed to detect the expression levels of SNHG16, miR-373-3p and TGF- $\beta$ -R2 in RWPE1, DU-145, PC-3, 22Rv-1 and LNCaP cell lines. (B) SNHG16 regulated DU-145 cell apoptosis. (C) miR-373-3p inhibitor partially reversed sh-SNHG16-induced apoptosis of DU-145 cells. (D) TGF- $\beta$ -R2 knockdown partially reversed the antiapoptotic effect of miR-373-3p inhibitor. \*P<0.05 vs. RWPE1. SNHG16, small nucleolar RNA host gene 16; miR, microRNA; TGF- $\beta$ -R2, transforming growth factor- $\beta$  receptor type 2; sh, short hairpin; lncRNA, long non-coding RNA; PI, propidium iodide; NC, negative control.

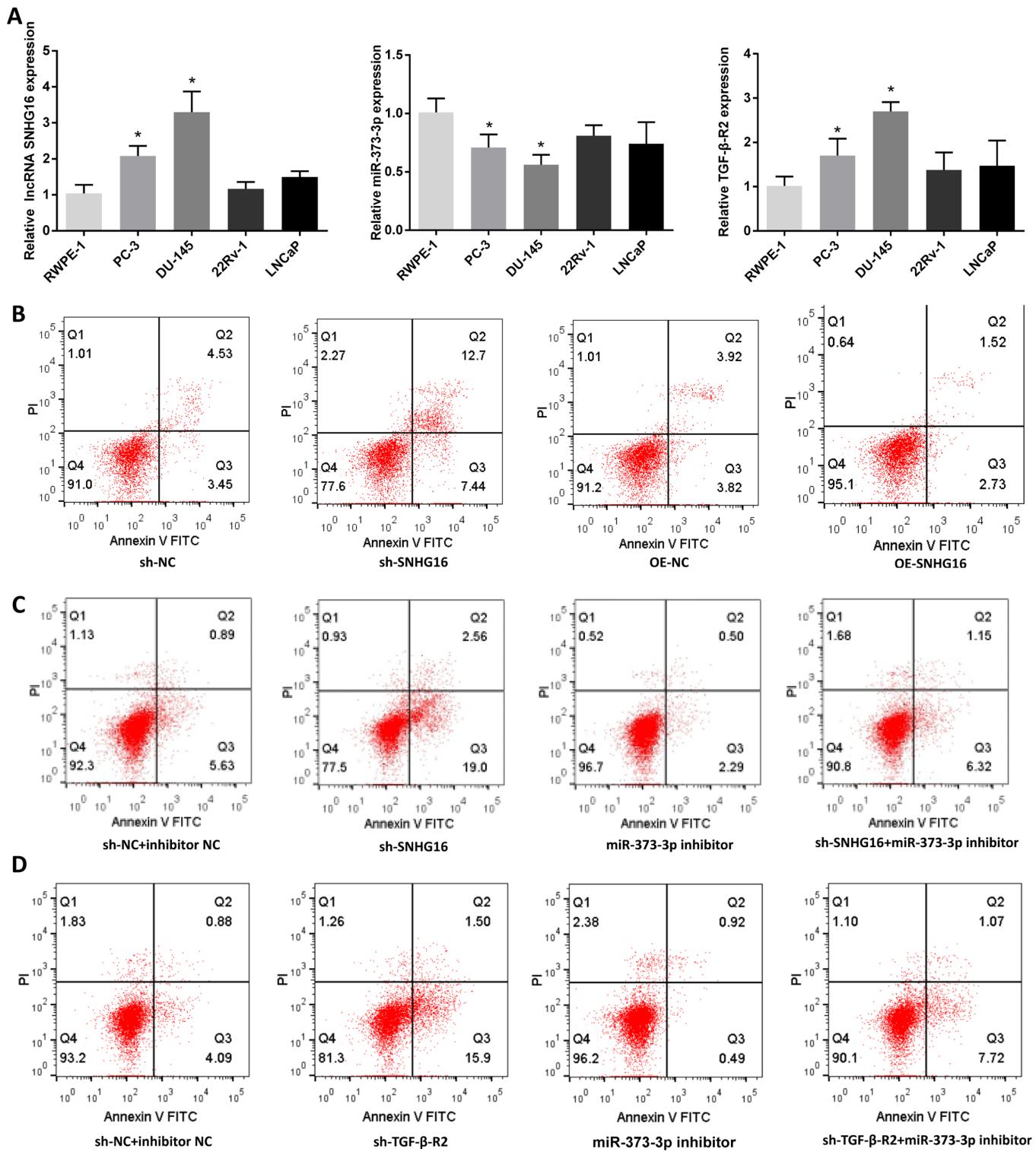


Table SI. Luciferase reporter constructs of SNHG16 and TGF- $\beta$ -R2 3'-UTR wild-type and mutated form.

Gene	Sequence
SNHG16 3'-UTR wild-type	5'-tcgatggactgcaaggattgtaaactgaaatgaacatgattatactctgttgcagagccctaagaggaaactgatgccatgagttcagaga gtaatgcttaaccccagttcacacaggatccgttgcctgttttagttaccactacagtgtttgtatctgtaatgggttgcaccc acaaccattgttttagtcaatcaatgaag gattgataaaaggttctcggtgtccgcagagtgcctccaggaaacagatctt gcatagaatatcagtggttcccttttgttcaaataatggtcagaaaatccccatgt tgactccaccaaggaaatcagct-3'
SNHG16 3'-UTR mutant	5'-tcgatggactgcaaggattgtaaactgaaatgaacatgattatactctgttgcagagccctaagaggaaactgatgccatgagttcagaga gtaatgcttaaccccagttcacacaggatccgttgcctgttttagttaccactacagtgtttgtatctgtaatgggttgcaccc caaccattgttttaACGCTTActtcaaataatgaaggatgtataaaaggttctcggtgtccgcagagtgcctccaggaaacagatc tttgcataatcagtggttcccttttgttcaaataatggtcagaaaatccccatgttgactccaccaaggaaatcagct-3'
TGF- $\beta$ -R2 3'-UTR wild-type	5'-ttcttattcaaaaaaaaaagccaaggaaataacattctgttagttccaaaaactgacttttcaactactatacataaaaggaaagttttattc tttatggAACACTTCAGCTGTACTCATGTTAAATAGGAATGTGAATGCTATACTCTTTATCAAAAGTCTAAGCACTTATTCTATGC ATGGTTGCTTTACATAAATAAAATGTTATTAGATGAATAAAGCAAATACTCAGGTGAGCATCCTGCCTGTCCCATTCCTAGTAGCTA-3'
TGF- $\beta$ -R2 3'-UTR mutant	5'-ttcttattcaaaaaaaaaagccaaggaaataacattctgttagttccaaaaactgacttttcaactactatacataaaaggaaagttttattcttt atggAACACTTCAGCTGTACTCATGTTAAATAGGAATGTGAATGCTATACTCTTTATCAAAAGTCTCA CAGAU CUATTTTATTCTATG CATTGTTGCTTTACATAAATAAAATGTTATTAGATGAATAAAGCAAATACTCAGGTGAGCATCCTGCCTGTCCCATTCCTAGTAGCTA-3'

UTR, untranslated region; SNHG16, small nucleolar RNA host gene 16; TGF- $\beta$ -R2, transforming growth factor- $\beta$  receptor type 2.