

## **Supplementary Data**

### **UASR1 Exon 1**

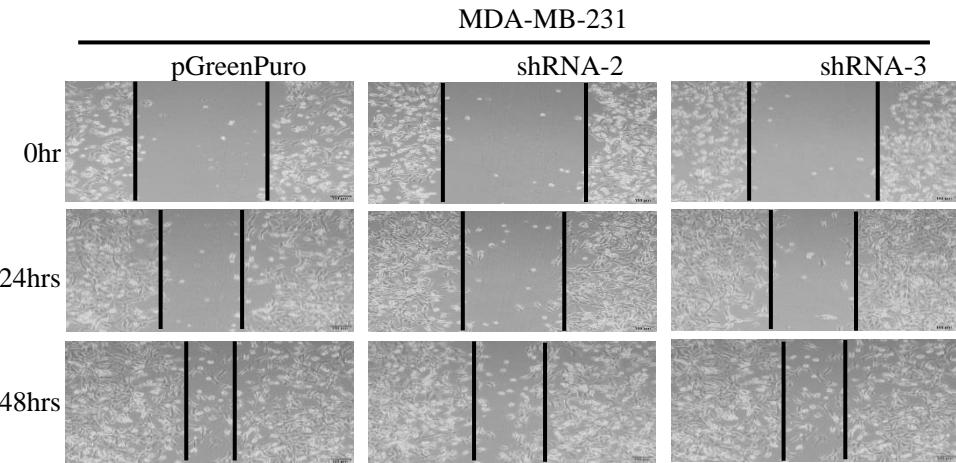
UASR1	1	CCCTCCTCAAACACACATCCATCCTCCGGcacacacccagtc 	catgcctcgccccacac
UNC5B-Int1	5164	CCCTCCTCAAACACACATCCATCCTCCGGCACACACCAGTCC 	CATGCCTCGCCCCACAC
UASR1	61	acacctgacaccccagtgcaacaccacaccaccc 	ccctccctccgtcaaactccacctc
UNC5B-Int1	5104	ACACCTGACACCCCAGTGCAACACCACACCACCC 	CTCCCTCCGTGAAACTCCCACCTC
UASR1	121	CGCCCACCTGCTTAATACACATTCTCACCCCCCACACACTC 	CTTAATACATACTCTCACA
UNC5B-Int1	5044	CGCCCACCTGCTTAATACACATTCTCACCCCCCACACACTC 	CTTAATACATACTCTCACA
UASR1	181	CCCACAAGCCTGCCTTCTTGGAGAAGTGAGCCGAGCGTGC 	AGCGCCGCGAAGG
UNC5B-Int1	4984	CCCACAAGCCTGCCTTCTTGGAGAAGTGAGCCGAGCGTGC 	AGCGCCGCGAAGG

### **UASR1 Exon 2**

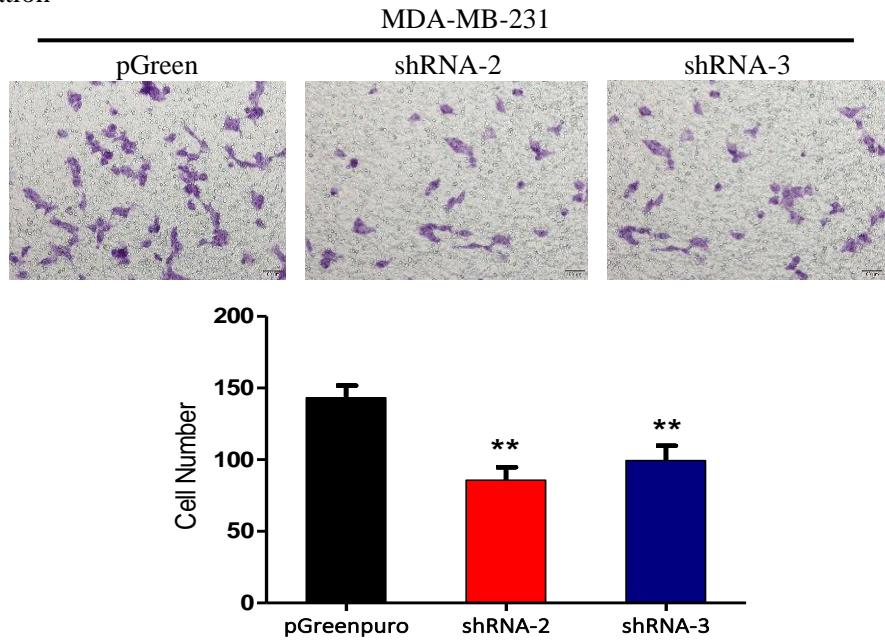
UASR1	235	GCATCCCCGAAGACCGGGAGGAACGCCGCGGGGACCTGT 	GGCTTAGCGCGCTCCGCCCGG
UNC5B-Int1	4572	GCATCCCCGAAGACCGGGAGGAACGCCGCGGGGACCTGT 	GGCTTAGCGCGCTCCGCCCGG
UASR1	295	GCTTGTCTGCCCGCGGGGGCGCAGCGGCTGAGGCG 	GCTCCGGCCGGAGTTCCAATCAAG
UNC5B-Int1	4512	GCTTGTCTGCCCGCGGGGGCGCAGCGGCTGAGGCG 	GCTCCGGCCGGAGTTCCAATCAAG
UASR1	355	CGCCACCCAACTCCCAGTCGGGGCCGAGGCCAGGCC 	GGGATGCCAGCTCCCCAAAA
UNC5B-Int1	4452	CGCCACCCAACTCCCAGTCGGGGCCGAGGCCAGGCC 	GGGATGCCAGCTCCCCAAAA
UASR1	415	AGATCCTGCCTCAGGGAAATGCATGGAGCCGGAAAAG 	CCCGCGGCCGGCGGA
UNC5B-Int1	4392	AGATCCTGCCTCAGGGAAATGCATGGAGCCGGAAAAG 	CCCGCGGCCGGCGGA
UASR1	475	TCGCAGACCTAACGGGGCGGGAGGTGGCG 	CCCCAGTCCAACCTCTTGAGCCAACCCAG
UNC5B-Int1	4332	TCGCAGACCTAACGGGGCGGGAGGTGGCG 	CCCCAGTCCAACCTCTTGAGCCAACCCAG
UASR1	535	TGGGTGGGAAGTGCCTTACCTAGGCCTCCGCAAAGTGT 	TCTCTCCTGTATTATTCT
UNC5B-Int1	4272	TGGGTGGGAAGTGCCTTACCTAGGCCTCCGCAAAGTGT 	TCTCTCCTGTATTATTCT
UASR1	595	AATTACGGTATTTAACCTTaaaaaaaataaga 	aaacagaaaagcacagaa
UNC5B-Int1	4212	AATTACGGTATTTAACCTTAAAAAAATAAGAACAG 	AAAAGCACAGAA

**Supplementary Figure S1. Structure of UASR1 gene.** Data show the blast between UASR1 cDNA and Intron 1 DNA sequence of UNC5B gene. UASR1 gene consists of two exons with 647bp in length and one intron of 358bp. UNC5B-Int1 indicates intron 1 of UNC5B gene, which is composed of 66,758bp.

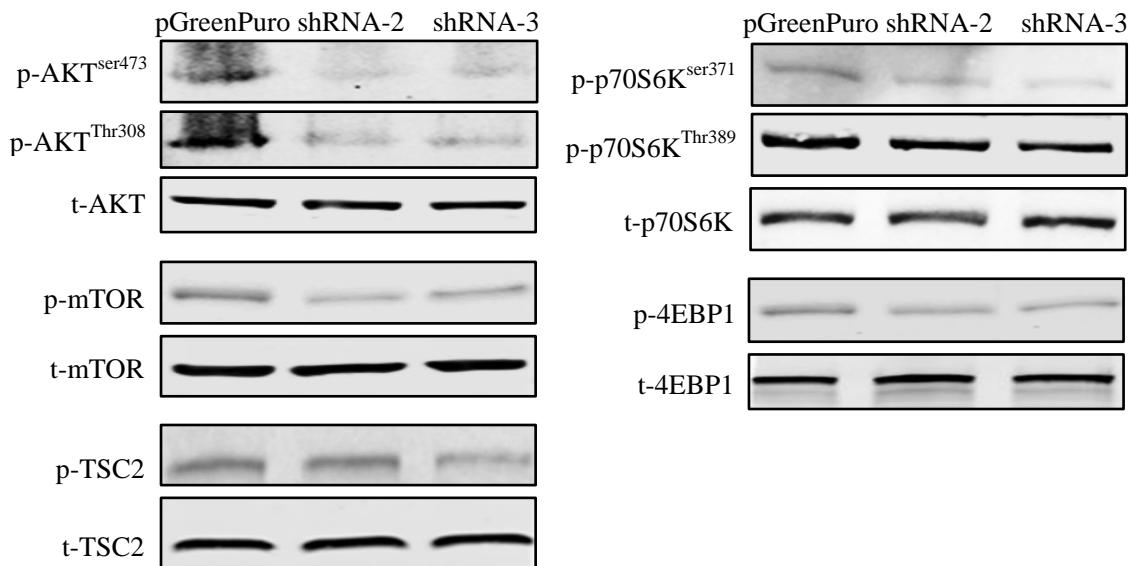
### A) Wound healing



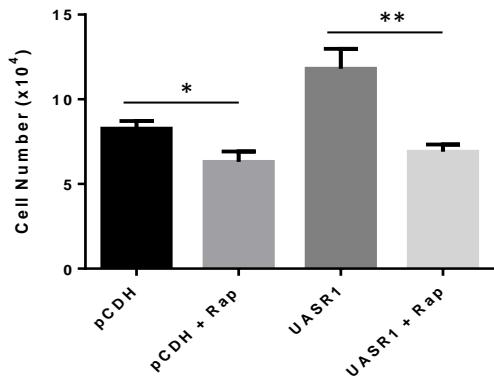
### B) Migration



### C) Western blot



**Supplementary Figure S2. Effects of UASR1 silencing in MDA-MB-231 cells.** UASR1 shRNAs were delivered by lentiviral vectors into MDA-MB-231 cells; wound healing, migration assays and Western blot were conducted as described in Materials and Methods. (A) Wound healing. (B) Migration. Bar chart indicates cell number migrated. Data indicate mean  $\pm$  SD from three independent experiments. \*\* p<0.01 compared to control. (C) Western blot, showing decrease of pAKT, p-mTOR, p-TSC2, p-p70S6K (Ser371 and Thr389), and p-4EBP1 proteins, i.e., inhibition of the AKT/mTOR pathway by UASR1 silencing.



**Supplementary Figure S3. Rapamycin inhibits cell proliferation.** MCF7 cells ( $3 \times 10^4$ /well) were spread into 12-well plates. After incubation overnight, cells were fed with fresh medium containing rapamycin (20 nM) for 24 hours and then collected by trypsinization. After trypan blue staining, viable cells were counted using a Vi-cell counter. Rap indicates rapamycin. \*  $p < 0.05$  and \*\*  $p < 0.001$ .