SUPPLEMENTARY FIGURES AND TABLES



Supplementary Figure S1: HERG1 siRNA silencing inhibits pancreatic cancer progression *in vitro*. (A) Flow cytometery analysis of apoptosis in PANC-1 and CFPAC-1 cells after HERG1 siRNA, NC siRNA transfection or without treatement; (B) Flow cytometery analysis of cell cycle progression in PANC-1 and CFPAC-1 cells after HERG1 siRNA, NC siRNA transfection or without treatement; (C) Migration and invasion assay analysis of PANC-1 and CFPAC-1 cells after HERG1 siRNA, NC siRNA transfection or no treatement. Magnification, ×100.



Supplementary Figure S2: Overexpression of miR-96 inhibits pancreatic cancer cell migration, invasion *in vitro*. Migration and invasion assay analysis of PANC-1 and CFPAC-1 cells after transfection with pre-miR-96 plasmid or empty vector or without treatement. Magnification, ×100.

| Supprementary rapid S1. The mixing potentially targeting the 5 -01K of HEK | IERG1 |
|--|-------|
|--|-------|

| miRNAs Position of HERG1 3'-UTR | miRNA (top) and predicted consequential pairing of target region (bottom) |
|------------------------------------|---|
| hsa-miR-96 | ACACGAUCACGGUUU |
| Position 427-440 | TGTGCT—GTGCTAAG |
| hsa-miR-105 | UCGUAAACU |
| Position 60-68 | GGCATTTGG |
| hsa-miR-152 | AAGACAG |
| Position 419-425 | TTCTGTT |
| hsa-miR-199a-5p | UUGUGACCC |
| Position 382-390 | AACACTGGG |
| hsa-miR-220 | GCCACAC |
| Position 193-199 | TGGTGTG |
| hsa-miR-329 | CCACACA |
| Position 195-201 | GGTGTGT |
| hsa-miR-331-5p | AGGGAC |
| Position 90-95 | TCCCTG |
| hsa-miR-592 | AACUGUGU |
| Position 116-123 | TTGACACA |
| hsa-let-7e | UUCGAU |
| Position 435-440 | AAGCTA |
| hsa-let-7g | GGACAUGU |
| Position 325-333 | TCTGTACA |

Supplementary Table S2. Primer pairs and thermal cycling conditions

| | primers | 5'-CGCACCATTAGCAAGATTCCC-3' 5'-AAGACAGCCGTGTAGATGACCAG-3' |
|------------|----------------------------|---|
| HERG1 | thermal cycling conditions | denaturating phase at 95°C for 5 min, 35 cycles of 1 min at 94°C, 1 min annealing phase at 58°C, and a 1 min extension phase at 72°C, with a final extension phase of 10 min at 72°C |
| beta-actin | primers | 5'-TGCGTGACATTAAGGAGAAGC-3' 5'-GCCAGGGTACATGGTGGTG-3' |
| | thermal cycling conditions | denaturating phase at 95°C for 5 min, 35 cycles of 15 seconds at 95°C, 30 seconds annealing phase at 65°C, and 30 seconds extension phase at 58°C, with a final extension phase of 10 min at 72°C |

Supplementary Table S3. Oligonucleotide sequences of four different HERG1 siRNAs and a negative control siRNA

| Oligo name | oligo sequences 5'to 3' |
|------------------|--|
| HERG1 siRNA-1-F | TGCTGCGATGAGGTCCACCACAGCCAGTTTTGGCCACTGACTG |
| HERG1 siRNA -1-R | CCTGCGATGAGGTCCCACAGCCAGTCAGTCAGTGGCCAAAACTGGCT GTGGTGGACCTCATCGC |
| HERG1 siRNA -2-F | TGCTGATGTGTGGCTGCTCCATGTTGGTTTTGGCCACTGACTG |
| HERG1 siRNA -2-R | CCTGATGTGTGGCTGCCATGTTGGTCAGTCAGTGGCCAAAACCAACA TGGAGCAGCCACACATC |
| HERG1 siRNA -3-F | TGCTGTGCTGTTGTAGGGTTTGCCTAGTTTTGGCCACTGACTG |
| HERG1 siRNA -3-R | CCTGTGCTGTTGTAGTTTGCCTAGTCAGTCAGTGGCCAAAACTAGGC AAACCCTACAACAGCAC |
| HERG1 siRNA -4-F | TGCTGTTTGCAGTGCTGCAGCAGTGAGTTTTGGCCACTGACTG |
| HERG1 siRNA -4-R | CCTGTTTGCAGTGCTAGCAGTGAGTCAGTCAGTGGCCAAAACTCAC TGCTGCAGCACTGCAAAC |
| Negative-F | TgctgAAATGTACTGCGCGTGGAGACGTTTTGGCCACTGACTGAC |
| Negative-R | CetgAAATGTACTGCGTGGAGACGTCAGTCAGTGGCCAAAACGTCTCC ACGCGCAGTACATTTe |

Supplementary Table S4. Plasmid vector screening: silencing efficiency of four different HERG1 siRNAs and a negative control siRNA

| HERG1 | siRNA-1 | siRNA-2 | siRNA-3 | siRNA-4 | NC siRNA |
|---------------------------|----------|----------|-----------|----------|----------|
| ΔCt | 13.4335 | 13.6975 | 13.3029 | 13.49777 | 11.18273 |
| ΔΔCt | 2.7465 | 3.0105 | 2.6159 | 2.810767 | 0 |
| $2-\Delta\Delta Ct$ | 0.149012 | 0.124094 | 0.1631307 | 0.14252 | 1 |
| Interferential efficiency | 0.850988 | 0.875906 | 0.8368693 | 0.85748 | 0 |

Supplementary Table S5. The oligonucleotide sequences of pre-miR-96 forward and reverse PCR primers

| Oligo name | oligo sequences 5'to 3' |
|--------------|-------------------------------|
| pre-miR-96-F | TCTGGGTACCGCACTGGTAGAATTCACTG |
| pre-miR-96-R | CCTTTCTAGACCACGGCACCATTCAGGA |