

Table S3 Morpholinos used.

Gene	MO phenotype	MO type	Ensembl transcript ID (MO predicted result)	MO sequence (5' to 3')
<i>ttl11</i>	none/mild	Splice, 1st EXON-intron	Retention of 14.2 kb intron (ENSDART00000085684)	AGTGGGTCCTGAGCTCTTACCTCTT
<i>irs2</i>	hPC, acute hypoxia-sensitive	ATG	Translational inhibition (ENSDART00000053924)	CCCCTTTAAGAGGCGGACTTGCCAT
<i>btr01</i>	hPC	Splice, 1st EXON-intron	Retention of 1.2 kb intron (ENSDART00000073689)	AACCGTATAAATGTGCTTACCTTCC
<i>camk2g2</i>	hPC	Splice, 1st EXON-intron	Retention of 10.8 kb intron (ENSDART00000078652)	GAACAGGACACTGAGACTCACCTAT
<i>crtc3</i>	hPC	Splice, 1st EXON-intron	Retention of 15.7 kb intron (ENSDART00000073903)	TCCTAATTTGGCTGAGCTTACCCTT
<i>ncam2</i>	hPC	Splice, 1st EXON-intron	Retention of 319.1 kb intron (ENSDART00000100681)	TCACGATCTCGCATAAATACCTTGA
<i>pacsin3</i>	developmental	ATG	Translational inhibition (ENSDART00000098296)	TGCAGATCACCGTTGGAAGACATT
<i>mical2b</i>	developmental	ATG	Translational inhibition (ENSDART00000139013)	TCCGTTCTTCTCCGTCTCCCCAT
<i>inhbb</i>	developmental	Splice, 1st intron-EXON	Skip 2 nd of 2 exons (ENSDART00000059762)	GCCGTCTGTGGAGAAAACACACACA
<i>opn5</i>	developmental	Splice, 3rd EXON-intron	Retention of 6.4 kb intron (ENSDART00000033404)	TGAAAACCTGAATACTGCACCTTCAC