

Table S2.Sequences of oligo nucleotides

NAME	SEQUENCES	DESCRIPTION
<i>gRNA</i>		
TOMM7_gRNA3_iPSC	5' - GGAGCCAGTTGCCATTGCGC -3'	gRNA sequencing used for gene editing in iPSCs
tomm7_gRNA3_fish	5' - ATCCCCACTGTGCTCTATCT -3'	gRNA sequencing used for gene editing in zebrafish
<i>ssODN</i>		
TOM7_iPS PS-anti asymmetric ssODN	5' - GCTGTCTCCGCCACCCCTTCCCTCCAGTCACCTTCCGGTTCTTC TCCCCTGACCCAGGTAATCACAAAGAAGGATAAAGCCCCAACGGA TGCGAAACTGGCTCCCTTGAAAGAGCTGCTGAAGTCTTGCTTGGC CTCTTGCTCAGCTTACCATGGCGACGGCCGTG -3'	repair template for gene editing in iPSCs
<i>PCR</i>		
TOMM7_exon1_F	5' - CTTTCGGATTCCCCGACG -3'	confirmation of TOM7 c.86C>T in patient's genomic DNA
TOMM7_exon1_R	5' - AAAATGGGGCTGCTGCTC -3'	
T7_cDNA_Flp-In genotyping_F	5' - ATGGTGAAGCTGAGCAAGA -3'	amplification of Flp-In sequences
lacZ_Flp_In_genotyping_R	5' - CCTTCCTGTAGCCAGCTTCATCAA -3'	
TOMM7_ontargetPCR_F	5' - GGAAGAGCAGGGAAAGGTTAAAGGCG -3'	amplification of CRISPR-targeted regions from genomic DNA in iPSCs
TOMM7_ontargetPCR_R	5' - CATCAACCCCTCCCTACATCCGAG -3'	
tomm7_ontargetPCR_F	5' - GATGATCGTTACTTCCGTCCACAG -3'	amplification of CRISPR-targeted regions from genomic DNA in zebrafish
tomm7_ontargetPCR_R	5' - GTCATATCCGAGTTCATAGACGC -3'	
<i>cloning</i>		
HindIII_3xmyc TOM7_F	5' - AAATTAAAGCTTGCCGCCATGGAGCAGAAACTCATCTCTGAAGAAG ATCTGGAACAAAAGTTGATTTCAGAAGAAGATCTGGAACAGAAAGCT CATCTCTGAGGAAGATCTGGTGAAGCTGAGC -3'	amplification of myc-tagged TOM7 cDNA sequences
Xhol_STOP_TOM7_R	5' - ATATATCTCGAGCTATTATTATCCCCAAAGTAGGC -3'	
TOMM7 86C>T_Cas9 Cloning_F	5' - acaccGGAGCCAGTTGCCATTGCG -3'	construction of gRNA sequences into eSpCas9(1.1)-T2A-Puro plasmid
TOMM7 86C>T_Cas9 Cloning_R	5' - aaaacGCGAATGGCAAACCTGGCTCC -3'	
<i>qPCR</i>		
qPCR_EGLN1_F	5' - GACCTGATAACGCCACTGTAACG -3'	quantification of EGLN1 expression in iEndo cells
qPCR_EGLN1_R	5' - ACCTCCACTTACCTTGGCATC -3'	
qPCR_CHCHD2_F	5' - CTGCAGTTGGCTCTTGTCT -3'	quantification of CHCHD2 expression in iEndo cells
qPCR_CHCHD2_R	5' - TTCCCTGAGGCTCCTGGTAAG -3'	
qPCR_APLN_F	5' - GATGGGAATGGGCTGGAAGAC -3'	quantification of APLN expression in iEndo cells
qPCR_APLN_R	5' - CCTTCAGTCCTGCTTCAGAAAGG -3'	
qPCR_HK2_F	5' - CAAGTGCAGAAGGTTGACCAGT -3'	quantification of HK2 expression in iEndo cells
qPCR_HK2_R	5' - CTCCGTGTCTGTCCCCATCTG -3'	
qPCR_SLC16A3_F	5' - ACAGCTGGATCTCCCTCAT -3'	quantification of SLC16A3 expression in iEndo cells
qPCR_SLC16A3_R	5' - AAACCCAACCCCGTGATGAC -3'	
qPCR_SLC2A1_F	5' - ACTGTCGTGTCGCTGTTGT -3'	quantification of SLC2A1 expression in iEndo cells
qPCR_SLC2A1_R	5' - AGGACCCACTTCAAAGAAGGC -3'	