Supplemantary Table S1. Sequence of the oligonucleotides used in the transfection experiments

Oligonucleotide	Sequence
miR-133a-mimic	5'-UUUGGUCCCCUUCAACCAGCUGUU-3'
	5'-CAGCUGGUUGAAGGGGACCUUAUU-3'
DS control mimic	5'-UUCUCCGAACGUGUCACGUTT-3'
	5'-ACGUGACACGUUCGGAGAATT-3'
miR-133a-decoy	5'-CAGCUGGUUGAAGGGGACCAAAT-3'
SS control-decoy	5'-CAGUAUUUUGUGUAGUACAA-3'

DS= double strand; SS= single strand. To increase stability, the RNA decoy nucleotides were modified by methylation in the 2' position of the ribose

Supplementary Table S2. Sequence of the primers used in the rtPCR experiments

Gene	Sequence	Accession number
MitoK	F 5'-GGAGCCTAATAAGCTTAGTCTCTTCA-3'	NM_001014098.1
	R 5'-CGAAAGGTTCTCCATACCAGTC-3'	
MitoSur	F 5'-CTCAAACATCGCTTTCAACTGT-3'	NM_001007796.1
	R 5'-ACATGAGGTCTCCCCCTTTC-3'	
Dio3	F 5'-TTTTGGTACTTTTTGGCTTGC-3'	NM_017210.4
	R 5'-CTTGTGCGTAGTCGAGGATG-3'	
Hprt	F 5'-CCCAGCGTCGTGATTAGTGATG-3'	NM_012583.2
	R 5'- ACTCTTCATTCAGGCCCTTG-3'	
Hmbs	F 5' TCAGSTGGCTCAGATAGCATGCA-3'	NM_013168.2
	R 5' TGGACCATCTTCTTGCTGAACA-3'	
miRNA	**Sequennce**	**Accession number**
133a-3p	5'-TTTGGTCCCCTTCAACAAGCTG-3'	MIMAT0000839
SnRNA-U1	5'-CGACTGCATAATTTGTGGTAG-3'	
SnRNA-U6	5'-CGCAAGGATGACACGCAAATT-3'	

Figure S1. Rat mitoK promoter region: in red THRa1 consensus sequence

>NC_051343.1:109717595-109722595 Rattus norvegicus strain BN/NHsdMcwi chromosome 8, mRatBN7.2, whole genome shotgun sequence.

-5000 TGCATCCTTCAGGAATGAGGGCCTCAAAACAATGCCCTTCAGTCCCAACAACCCAGAGAGTTGTCAGGCTTTTTGCAAAACAATGCCAAACTAAACTGTCAGT
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CAACAACAACAATAATACATTCCTTAGCTTCCAAACTTCAAGTCAGTAAAGGAAAACAATGAAAATACATTATCAAGGTCTGCTGTAATATAAATGTCCTCAA
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AGGGAGAAAAAGCAACTGTAGCTCAGGTGGAAGGGTTTACCTGGGATGCACGCCCAGGGCTGAGTAAGGAGGCCTGGCAGCGGTTATGCCTGCAATCCAG
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ATAAAATGCAAAAGGAGTCCAAGTTGAGATAGACCCAGTAAACAGTATTTATTAGTTCCGATCTCAGAGCATGATATAGAGAATCCTTCTGGCATAGCATAT
CTGAATATTCCCCCTTTCGATCCCCACCCACTTTCTGTAGTCT

-1359 C**AGCTCATGGCTG**AAAGGAGTCTGCATTGGCTGTGCCTCTATTATGAATACTATTTACTGAACACACTAG -1289

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GATGGGTGACGTATACAGAGCGCGCCGTTACCCGGGACAGGTTCCTGCCTTGCTGGGCTCAGTACCCTGAGGT

Figure S2. Rat mitoSur (ABCB8) promoter region: in red THRa1 consensus sequence

>NC_051339.1:c10788598-10783598 Rattus norvegicus strain BN/NHsdMcwi chromosome 4, mRatBN7.2, whole genome shotgun sequence

-5000 CTTCCTGCCCAGAGGCTTGGGTGGCCCTCCCCTCACC**CTGGCCTGACCT**TCCTCGACCAATCAGGATGCC -4930

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