

S1 Table. Primers. Mutated sequences are underlined.

Primer	Sequence (5'–3')
<i>Electrophoresis mobility shift assay</i>	
PEPCK-ROREwt-sense	GCAGTAAAATGGGTCAAGGT
PEPCK-ROREwt-antisense	ACCTTGACCCATTTTACTGC
PEPCK-ROREmt-sense	GCAG <u>CGCC</u> ATGGGTCAAGGT
PEPCK-ROREmt-antisense	ACCTTGACCCAT <u>GCGC</u> CTGC
Ikb-RORE-sense	GCAGCGCCATGGGTCAAGGT
Ikb-RORE-antisense	ACCTTGACCCATGGCGCTGC
<i>Luciferase reporter cloning</i>	
proPEPCK-FW	GGTTCGCTAGCTGAGTTTGGTCCGGAGG
proPEPCK-RV	CAGCAAGCTTGTGTTCCAGTGGGAAG
proPEPCK-ROREmt-FW	CAGCAG <u>CGCC</u> ATGGGTCAAGGTTTAGTCAG
proPEPEK-ROREmt-RV	CTGACTAAACCTTGACCCAT <u>GCGC</u> CTGCTG
proPEPCK-C3mt-FW	GAAAAGGGTGTGTGTTT <u>CCGG</u> ACAGCAG
proPEPEK-C3mt-RV	CTGCTGT <u>CCGG</u> AAACACAACACCCTTTC
PGVB2-FW	CAAGTGCAGGTGCCAGAAC
PGVB2-RV	CACCTCGATATGTGCATCTG
<i>qRT-PCR</i>	
rtPEPCK-FW	AAGAAGTGCTTTGCTCTCAG
rtPEPCK-RV	GCCATTTTCTGGGTGATGG
rtRORA-FW	TGTGATCGCAGCGATGAAAG
rtRORA-RV	ACAGGAGTAGGTGGCATTGC
rtACTB-FW	GCACCACACCTTCTACAATG
rtACTB-RV	CTCAAACATGATCTGGGTCATC
rt18SrRNA-FW	CGATAACGAACGAGACTCTGG
rt18SrRNA-RV	TAGGGTAGGCACACGCTGAGC