

Gene	Sequence		Location	Accession No.	
consensus ERRE	TCAAGGTCA				
<i>Acadm</i> , mouse	GAGCC	CGAAGGTCA	CGTTC	-458 to -466	NM_007382
<i>Acadm</i> , rat	GAGCC	CGAAGGTCTG	CGTCC	-430 to -438	NM_016986
<i>Acadm</i> , mouse	CCAAG	TAAAGGTCA	CAGCT	-558 to -550	NM_007382
<i>Acadm</i> , rat	CCGAG	TAAAGGTCA	CAACT	-529 to -521	NM_016986
<i>ACADM</i> , human	CCGGG	TAAAGGTGA	AGGCT	-330 to -338	NM_000016
<i>Aco2</i> , mouse	AACAG	TCAAGGTCA	AACTG	+984 to +976	NM_080633
<i>Aco2</i> , rat	AACAG	TCAAGGTCA	AACTG	+940 to +932	NM_024398
<i>ACO2</i> , human	GGGAG	TCAAGGTCA	AACTG	+961 to +953	NM_001098
<i>Atp5b</i> , mouse	CAAGC	TCAAGGTCA	CGGGG	+197 to +198	NM_016774
<i>Atp5b</i> , rat	CAGGC	TCAAGGTCA	CGGGG	+198 to +190	NM_134364
<i>ATP5B</i> , human	CCGGC	TCAAGGTCA	TGGGG	+281 to +289	NM_001686
<i>Atp5b</i> , mouse	AAGGT	CCAAGGACA	GTCAA	-322 to -318	NM_016774
<i>Atp5b</i> , rat	AAAGT	CCAAGGACA	GTCAA	-318 to -314	NM_134364
<i>ATP5B</i> , human	AAAGC	CCAAGGACA	GGCAA	-253 to -245	NM_001686
<i>Cycs</i> , mouse	CTAGC	CCAAGGACA	CGCCG	-4 to -12	NM_007808
<i>Cycs</i> , rat	CTAGC	CCAAGGACA	CGCCG	-4 to -12	NM_012839
<i>CYCS</i> , human	TAAGT	CCAAGGACA	CGCCG	-5 to -13	NM_018947
<i>Cpt1b</i> , mouse	TAGGG	AAAAGGTCA	CCAGG	-754 to -762	NM_009948
<i>Cpt1b</i> , rat	TAGGG	AAAAGGTCA	CCAGG	-293 to -301	NM_013200
<i>CPT1B</i> , human	TAGGG	AAAAGGTCA	CCGAT	-233 to -241	NM_004377
<i>Idh3a</i> , mouse	CGGCG	TCAAGGTCA	AGGCG	+220 to +228	NM_029573
<i>Idh3a</i> , rat	CGGCG	TCAAGGTCA	AGGCG	+244 to +252	NM_053638
<i>IDH3A</i> , human	CGGCG	TCAAGGTCA	AGGCG	+284 to +292	NM_005530