

Supplementary Table S2: Sequences of primers used for quantitative PCR.

Gene name	Forward (5'-3')	Reverse (5'-3')
<i>Acc1</i>	ATGGGCGGAATGGTCTCTTTC	TGGGGACCTTGTCTTCATCAT
<i>Atf4</i>	GGGTTCTGTCTTCCACTCCA	AAGCAGCAGAGTCAGGCTTTC
<i>Atf6</i>	GACTCACCCATCCGAGTTGTG	CTCCCAGTCTTCATCTGGTCC
<i>Cd36</i>	ATGGGCTGTGATCGGAACTG	GTCTTCCCAATAAGCATGTCTC C
<i>Chop</i>	CTGGAAGCCTGGTATGAGGAT	CAGGGTCAAGAGTAGTGAAGG T
<i>Chop (h)</i>	GGAAACAGAGTGGTCATTCCC	CTGCTTGAGCCGTTTATTCTC
<i>Cpt1</i>	CTCCGCCTGAGCCATGAAG	CACCAGTGATGATGCCATTCT
<i>Derl3</i>	AGGAGGGTTCTTTCCGTGG	ACGGCTCCATACATAGACCAG
<i>Derl3 (h)</i>	AGGTCTGGAGGCTCGTCAC	GCAGTAGCGGAACACGAAGAG
<i>Erdj4</i>	AAATAAAAGCCCTGATGCTGA A	CTATTGGCATCCGAGAGTGTTT
<i>Erdj4 (h)</i>	TCTTAGGTGTGCCAAAATCGG	TGTCAGGGTGGTACTTCATGG
<i>Fasn</i>	GGAGGTGGTGATAGCCGGTAT	TGGGTAATCCATAGAGCCCAG
<i>Fatp2</i>	TCCTCCAAGATGTGCGGTACT	TAGGTGAGCGTCTCGTCTCG
<i>Fatp5</i>	CTACGCTGGCTGCATATAGATG	CCACAAAGGTCTCTGGAGGAT
<i>Grp78</i>	ACTTGGGGACCACCTATTCCT	ATCGCCAATCAGACGCTCC
<i>Grp78 (h)</i>	GAAAGAAGGTTACCCATGCAG T	CAGGCCATAAGCAATAGCAGC
<i>Herpud1</i>	AGCAGCCGGACAACTCTAAT	CTTGAAAGTCTGCTGGACA
<i>Herpud1 (h)</i>	TGCTGGTTCTAATCGGGGACA	CCAGGGGAAGAAAGGTTCCG
<i>Hrd1</i>	CCTGGGACAACAAGGCTGTA	AGCTTTCTTGAACTGCCTCATG
<i>Hrd1 (h)</i>	CTTCACCGTTTTTCGGGATGA	CCAGGAGGAACATAAGAGAGA CA

<i>HPS</i>	GATTTACAGCTTCGTCCTGG	TCTTGCCTCCAAGGTCAATG
<i>HPS (h)</i>	ATGGCAAAGGTGTTTCAGTTTCA	ACAATCTGCATACTGCCTCTTG
<i>Hsp40</i>	TTCGACCGCTATGGAGAGGAA	CACCGAAGAACTCAGCAAACA
<i>Hsp90</i>	TGTTGCGGTACTACACATCTGC	GTCCTTGGTCTCACCTGTGATA
<i>Mttp</i>	CTCTTGGCAGTGCTTTTTCTCT	GAGCTTGTATAGCCGCTCATT
<i>Ppara</i>	AGAGCCCCATCTGTCCTCTC	ACTGGTAGTCTGCAAAACCAAA
<i>Scd1</i>	GTACCGCTGGCACATCAACTT	GGCTAAGACAGTAGCCTTAG
<i>Spliced-Xbp1</i>	CTGAGTCCGAATCAGGTGCAG	GTCCATGGGAAGATGTTCTGG
<i>Srebp1</i>	GGAGCCATGGATTGCACATT	GGCCCGGGAAGTCACTGT
<i>Tbp</i>	AGAACAATCCAGACTAGCAGC A	GGGAACTTCACATCACAGCTC
<i>Tbp (h)</i>	CCACTCACAGACTCTCACAAC	CTGCGGTACAATCCCAGAACT
<i>Unspliced Xbp1</i>	AGCAGCAAGTGGTGGATTTG	GAGTTTTCTCCCGTAAAAGCTG A
<i>Wfs1</i>	CGGGAAGAAACGGACAGAGC	CGTAGGTAGTGTTTGCCAC
<i>Xbp1</i>	GACAGAGAGTCAAACAACTAACGT GG	GTCCAGCAGGCAAGAAGGT
<i>Serca</i>	GAGAACGCTCACACAAAGACC	CAATTCGTTGGAGCCCCAT
<i>Stim1</i>	TGACAGGGACTGTACTGAAGATG	TATGCCGAGTCAAGAGAGGAG
<i>Ncx</i>	CGCTGGGGAAGATGACGATG	TGGGACGAAGGCAAACAGAAC
<i>Ip3r</i>	CGTTTTGAGTTTGAAGGCGTTT	CATCTTGCGCCAATTCCCG
<i>Orai</i>	GATCGGCCAGAGTTACTCCG	TGGGTAGTCATGGTCTGTGTC

(h) show that the primer sequence is suitable for detection of human gene.