

SUPPLEMENTARY TABLE S1. OLIGONUCLEOTIDES USED FOR CONVENTIONAL  
REVERSE TRANSCRIPTASE-POLYMERASE CHAIN REACTION

<i>Primer name</i>	<i>Target</i>	<i>Position</i>	<i>Annealing temperature</i>	<i>Product size</i>	<i>Sequence</i>
GAPDH sense	GI 7669491	555	55°C	352 bp	GGCCAAGGTC ATCCATGA
GAPDH ras		907			TCAGTGTAGC CCAGGATG
TBX5 sense	GI 31652233	1303	55°C	108 bp	AGCAGTGACTIONCCTACCAGAAC
TBX5 ras		1411			TGACATTCTGTGCAGCTCCAT
TBX20 sense	GI 31652229	486	50°C	470 bp	AGGTACCGCTACGCCTAC
TBX20 ras		893			GTCAGTGAGCCTGGAGGA
MLC2a sense	GI 20380769	4	60°C	446 bp	GAGGAGAATGGCCAGCAGGAA
MLC2a ras		451			GCGAACATCTGCTCCACCTCA
AFP sense	GI 20379786	1518	55°C	255 bp	ACTCCAGTAAACCCCTGGTGTTG
AFP ras		1772			GAAATCTGCAATGACAGCCTCA
TTR sense	GI 18089144	228	55°C	153 bp	AAAACCAGTGAGTCTGGAGAGC
TTR ras		380			GTTGGCTGTGAATACCACCTCT
WNT1 sense	GI 16936523	306	55°C	190 bp	TATTGTGAACGTAGCCTCCT
WNT1 ras		495			ATCCCGAACTGCCACTTG
FGF5 sense	GI 73486654	570	55°C	253 bp	ATCCACGAAGCCAATATGTTA
FGF5 ras		822			GTGAAAGAAAGTTCTGGCTGCT