

Table S2. Primers used for assessment of the RNAi-mediated gene silencing by RT-qPCR.

Target gene	Primer sequence (5'→3')	Primers efficiency, %	Amplicon size (from cDNA), bp
<i>asp</i>	AAGTCGATTGGATCGTCTTTC AATTTAGGATGATCCGGCTG	105.1	155
<i>dgt6</i>	AACAGCTTACTCGCACCTGC GCATGGGATCGTTGATCTTG	101.3	118
<i>Eb1</i>	TCTGCACAGGTGCAGCTTACTGTCA CTTCTTGAAGCCCGCCTGCA	102.9	140
<i>Klp10A</i>	GCTGAGCGAACACGAGATGT CAGTGTGGCATTAAACGGTGC	100.2	115
<i>mars</i>	CAGACGACGGTTAAAGAAGACA GCGAGACACTAACAAACGGA	105.9	121
<i>mast</i>	AGACGCTGAGAAATAAACTAGATGC GCAGCTTTGGTGCATGTGTA	106.6	147
<i>mei-38</i>	TTTGGCCAGTCGGACTTCGA CTGTTCCCGGCCGATCTTCT	101.7	165, 168
<i>Patronin</i>	TTTTCAAATACAACCTCAGGAGGCA ATTGTGAAGGCGTCGATGGT	99.0	82
<i>RpL32</i> (as reference)	CTAAGCTGTCGCACAAATGG AGGAACTTCTTGAATCCGGTG	100.0	148