

Table S1. Primers used for tissue specific expression analysis, qRT-PCR, gene silencing quantification, RACE sequencing and RNAi experiments.

Sequence Name		Primer sequence (5' to 3')	Tm (°C)	Product size (bp)
<i>RferOBPu1</i>	Forward	TCAAGGAACACTTGGTGTGT	54.3	185
	Reverse	TGTCATTTCGGTAAGCCAGTT	54.1	
<i>RferOBPu2</i>	Forward	GGCATTTCGCTTTCTGTCTCT	55.2	179
	Reverse	AAGTGTGGAAGGCTGTTTCT	54.4	
<i>RferOBPu3</i>	Forward	CTTCAGTGACGATCCCAAGTT	54.7	188
	Reverse	TTTCTTCGGGAGAGCCTTTG	55	
<i>RferOBP2374</i>	Forward	GCTTCAGCGACGATCCAAA	55.7	195
	Reverse	ATGCCGTTTCTTCAGGAGTG	55.2	
<i>RferOBP257</i>	Forward	TACAAACACTGCGTGGAAGA	54.2	204
	Reverse	CTGGTCCTCGTTTCCTTTGA	54.6	
<i>RferOBP29</i>	Forward	TCGCCAACACCATTGCTAAG	55.8	196
	Reverse	CCGTCCGTTGTCATCCATTT	55.5	
<i>RferOBP3199</i>	Forward	ATGTGCTCAACAGGAGAAGG	54.8	174
	Reverse	TTGGCTCTTTCGATCCAGTC	54.7	
<i>RferOBP3213</i>	Forward	CGTAAAGGCCATGTCTGATGA	54.8	213
	Reverse	ATGGTTGCCTCAACGTCTATT	54.3	
<i>RferOBP7073</i>	Forward	GGCCTTTGCATTCTGCATATC	54.6	205
	Reverse	AGGGTTGGCTTTGTAGTAGC	54.8	
<i>RferOBP8586</i>	Forward	ACATAGGTTGTCAGGGCATC	54.6	191
	Reverse	GACATGACGGATGTGTGTCT	54.6	
<i>RferOBP9136</i>	Forward	AACCGGCATAGATGGAAACA	54.3	224
	Reverse	GAGTTCCCTCCTTTCCAAGAC	55.1	
<i>RferOBP981</i>	Forward	CTTCAGCGACGATCCCAAAT	55.3	186
	Reverse	TCTTCGGGAGTGTCTTCTT	55.4	
<i>RferOBP9915</i>	Forward	CGCTGATGAAACGAACGTAAA	53.7	213
	Reverse	TCGCAATCGACAGCTACTT	54	
<i>RferOBP77</i>	Forward	GAAACTGAAACGCGGTGAAA	54	183
	Reverse	GAGCAAGCACTGACGATTTG	54.5	
<i>RferOBP382</i>	Forward	TGCTCTGTACGTCGATGATG	54.4	206
	Reverse	CGGTAACCTGTTTCGCTTTG	54.4	
<i>RferOBP446</i>	Forward	ACTTCGTGGACGACAATTCC	55.1	196
	Reverse	AGGTAAACGGTATCGCAAGC	55.2	
<i>RferOBP1768</i>	Forward	CCTCGCTTGCGGACTAATA	54.4	175
	Reverse	ACCAACACAAACCAGATGAT	52.1	
<i>RferOBP3937</i>	Forward	CTGCTAAACCTTTGACAGATG	51.4	199
	Reverse	GGTTGAAAGACGTTGTTGG	51.9	
<i>RferOBP3997</i>	Forward	GTCGAACCTACCACCAACTT	54.6	194
	Reverse	ACGGCAGCATCGTCTTTAT	54.4	
<i>RferOBP4661</i>	Forward	AAGAAGTCATCAACACTTGCC	53.3	172

	Reverse	GCCTCCAATTTGCGATATACA	53	
<i>RferOBP12010</i>	Forward	GCAGCCACTCGCATAGATAAA	55.1	214
	Reverse	CGCACTCCTGCAACAAAGTA	55.6	
<i>RferOBP14025</i>	Forward	CCCGAATGCTCTACGACAAT	54.8	184
	Reverse	TTCTCATCCTCGCCGTTTAC	54.7	
<i>RferOBP23691</i>	Forward	TATCACAACCTGCGGATCTGG	54.5	204
	Reverse	GCTTCCTTGTTTCAGTGTGTTC	54.2	
<i>RferOBP28119</i>	Forward	CCTGTGTTGCCGCTTTATTG	54.8	198
	Reverse	TGGAGATGCAGAGCGAATG	55	
<i>RferOBP29381</i>	Forward	GCACATGAACAATGCCAGTC	54.8	184
	Reverse	TGACCTGTCCCATCGTATCT	55	
<i>RferOBP33721</i>	Forward	ATGCATGGTTGATTGCCTTT	53.4	187
	Reverse	CACATTGCAACCCTCTGATTC	54.2	
<i>RferOBP10788</i>	Forward	CCTTTATGGGCCTGAGTTTCT	54.7	229
	Reverse	ACGTCAGCTTGTACATCACC	54.9	
<i>RferOBP12481</i>	Forward	TGTGAAGAAAGCTGCTGGTAT	54.3	172
	Reverse	GCATTTGGTCATTTGGTAGGC	54.8	
<i>RferOBP12511</i>	Forward	GTGAGGAATGCTTTCAAGGG	53.8	217
	Reverse	ATGCTGTCTCTTCAGGACTC	53.9	
<i>RferOBP14511</i>	Forward	GAAGCCGAGAATGCTACACA	54.9	192
	Reverse	CACGTCATCCAAACCCTTCT	55	
<i>RferOBP16551</i>	Forward	AAACTCAACTTTTCGGAGAGAA	51.7	182
	Reverse	AAGCATATCCTTAACGTGACT	51.4	
<i>RferOBP1689</i>	Forward	TTCAAGGCGAGTGCATGAT	54.7	210
	Reverse	CATCGACTTTTCGTTTCATCGC	54.5	
<i>RferOBP17793</i>	Forward	TGCGATATCCGCTCTTACAC	54.5	210
	Reverse	TCGTCGGTTTGTAGGTTTCC	54.8	
<i>RferOBP19755</i>	Forward	TGACGGATTACCACTGTTC	54.9	191
	Reverse	AGCGGTTTCCTGTTGAGTAT	53.9	
<i>RferOBP23</i>	Forward	CAATGCACGAGCTGGTACATTCG	58.6	168
	Reverse	TTTCTGATTACTGGCGGTAGGGT	57.9	
<i>RferOBP107</i>	Forward	TCGCAGGGCATATCGACATAGA	57.7	155
	Reverse	TTGCAAAGGCTCGGTCACAGT	59.6	
<i>RferTubulin</i>	Forward	GCTACCTTCATCGGCAACTC	55.8	196
	Reverse	CGGTGGCTTCTTGGTATTGT	55.3	
<i>Rferbeta-actin</i>	Forward	AAAGGTTCCGTTGCCCTGAA	57.3	129
	Reverse	TGGCGTACAAGTCCTTCCTG	57	

Sequence Name		Primer sequences (5' to 3')	T _m (°C)
<i>RferOBP23</i>	Forward	ATGTTTAAAACGTTACCGATAG	48.7
	Reverse	TAATACGACTCACTATAGGGTTAAATTAAGAAATAATG	55.4
<i>RferOBP107</i>	Forward	ATG TTCAGTTTAACCGA ACTAG	50.4
	Reverse	TAATACGACTCACTATAGGGTCAAATACATAGTATATATC	56.8
<i>RferOBP1768</i>	Forward	ATGTGTCGTTTTACTGCTATTTTA	50.9
	Reverse	TAATACGACTCACTATAGGGTTAAGCTCCAAAATGTTCTTTAAAC	61.2
<i>RferOBP3213</i>	Forward	ATGAGACAGCATTTCGATTTTTTTGGGT	57.7
	Reverse	TAATACGACTCACTATAGGGTTAAATTAACATGTATTCCGCAAAT	61.1
<i>RferOBPu1</i>	Forward	ATGAATAATTTGGTGGTTTTATTACTT	50.2
	Reverse	TAATACGACTCACTATAGGGTTACAGACCGAAATGTTTCATTATACAA	62.1