

Supplementary TABLE 4. Primers for qRT-PCR analysis.

Gene_id	Gene name	Primer sequence			
	<i>Actin</i>	F	ACAGTGTCTGGATTGGAGGGTC	R	CATTTGGAGAACTCAGAAGCACT
LOC103950162	<i>PbMET1</i>	F	CTGCCTGTTCTTGCCGCCCT	R	GCTCGACCCAAGTGCAGAACCT
LOC103965404	<i>PbCMT3</i>	F	CGCGCTCTCTCTCTGTCCACC	R	AGTCACTGGGTTCTCGGCGG
LOC103926943	<i>PbDRM2</i>	F	GCGGCGAGGTTCTTCTGCGG	R	CGATTTCCAAGCTCCGCGGG
LOC103944304	<i>PbROS1</i>	F	TGGGGAGAACGGAATTGAGCGT	R	TCCGGCTGCAATTTGGGCTCCA
LOC103961774	<i>PbPin-likes 3</i>	F	ATCAGTCCACGAAAGGTCC	R	CACAGCAGTGTATGGCAGTGA
LOC103959162	<i>PbPin-likes 5</i>	F	TGCCTGCAATCGGTATTCTCAT	R	GCGGCAGGGTAAACTGAATC
LOC103960175	<i>PbPin-likes 6</i>	F	ATGAAGAAGAGACAGCACGCA	R	GGTGGTTGAAGAACTTGCTGAG
LOC103942180	<i>PbGH3.1-1</i>	F	GACCAGTGCTGCCTAACGA	R	CATCAACTCCTCAAACGTGCC
LOC103957346	<i>PbGH3.1-2</i>	F	AACTGGCAACAACGTGCCCT	R	TGCTAGCTTGGGATCCGGT
LOC103965914	<i>PbGH3.6</i>	F	GCAAACCCAGTGAAGTGTCCCT	R	TGAGTGATTTGCGGACAGAGAC
LOC103951583	<i>PbAPRR5</i>	F	GGTCCGCGGTACCTTTGG	R	TAGTTCAGCTGCCACCAG
LOC103950970	<i>PbARR6</i>	F	TCGGATCGGAGAATCAAGCG	R	CGCCTTCCATTTTACCGTCG
LOC103955055	<i>PbAHK2</i>	F	TCCGAGTTGCTTCAAACCCA	R	GTCACCTGTGTGGAGAGTGG
LOC103958138	<i>PbAHK4</i>	F	ATGCCAATCACCTTACTTGTTAGT	R	TGGGTTTGAGGCAACTCGC
LOC103927480	<i>PbCKX7</i>	F	GCTCAAGTAATCAGTCCACGAC	R	AGGGCCAGCGACGGT
LOC103952784	<i>PbCKX7-like</i>	F	TGCCCGTTGATCGAGTGG	R	TCTCGCCGCTAGCCTTC