

Figure S1. Melting curves, melting peaks and amplification curves of the five selected candidate reference genes.

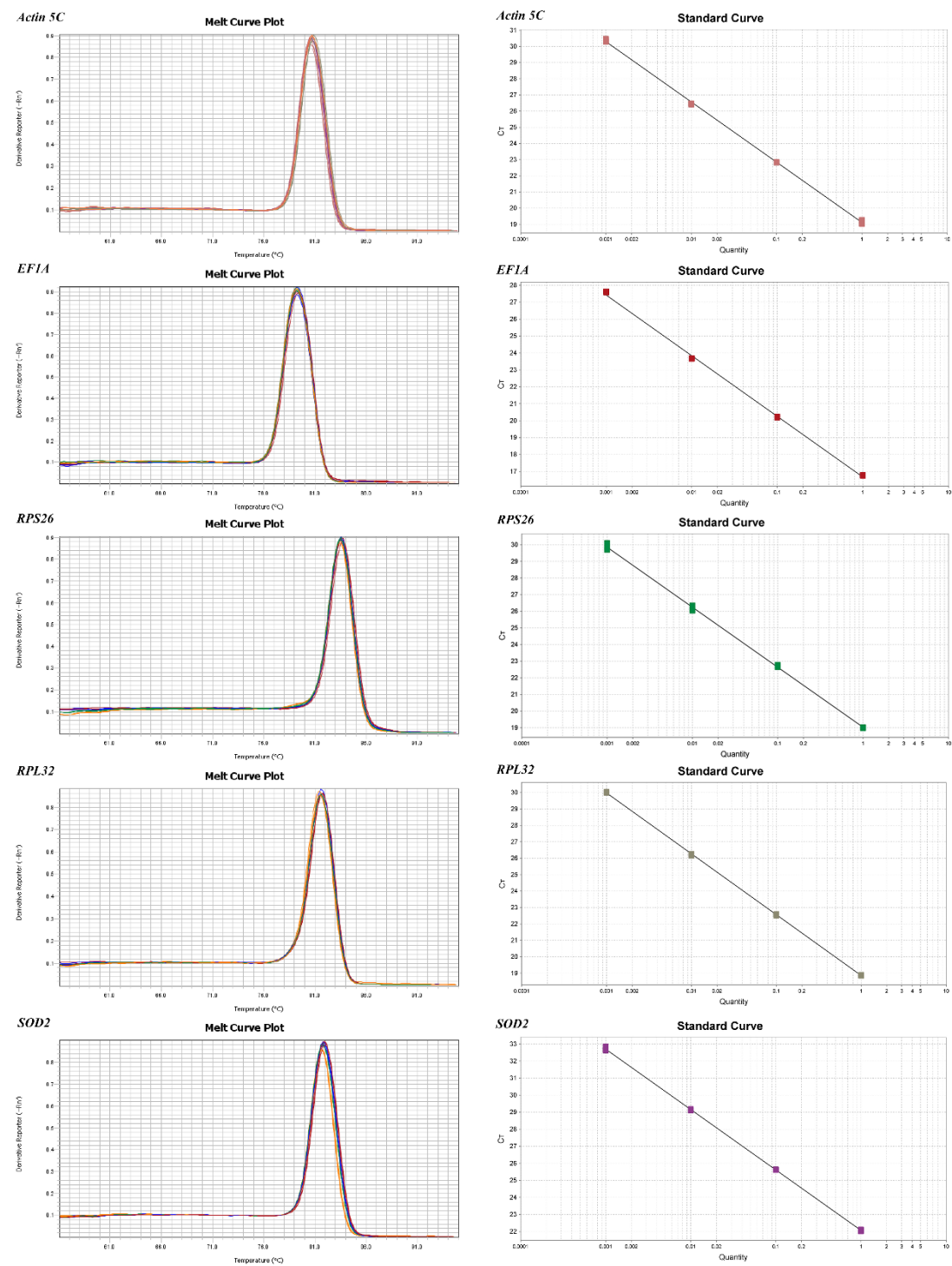


Table S1 Primers of the candidate reference genes used in this study.

Gene Name	Gene Symbol	Primer Sequences (F and R)	Amplicon Length (bp)	Amplification efficiency (%)	R ²
Actin 5C	<i>Actin5C</i>	GTATGCTTCTGGACGTACT GCAGAGCATAACCTTCGTAA	90	85.554	0.999
Elongation factor-1 alpha	<i>EF1A</i>	ACTGAACCCACCTTACAGTGA AACAGCAGCTGGATTGTAAC	93	90.204	0.999
Ribosome protein S26	<i>RPS26</i>	ATGACTTGCAAGCGAAGAAA CAGTTTGTACACCTCACTGG	77	89.314	0.999
Ribosome protein L32	<i>RPL32</i>	CAGTTTCTGATGCCCTCAAT ACATTGTGAACAAGCACCTT	92	86.062	1
Superoxide dismutase 2	<i>SOD2</i>	GGCTTGGCTTGGTTACAATA CAGTTGATGGTTCCAAAGGA	83	91.562	1

Table S2 Primers used for qRT-PCR analysis of *Myo* gene expression and dsRNA synthesis.

Gene Name	Primer Sequences (F and R)	Amplicon Length (bp)	Application
<i>Myo</i>	T7 + GACAGATGGGAGGAAACATCG T7 + TTGTGGGCTTGCTTGTTGAA	232	RT-PCR
<i>Muslta</i>	T7 + CACCCTCTCCACGAATTG T7 + TAGAAGATGCTGCTGTTTCA	193	RT-PCR
<i>Myo</i>	CCCGAGAAAAATGTCGTCTAT CCGATCCACTACCATTTCCT	91	qRT-PCR

T7 RNA polymerase promoter sequence: TAATACGACTCACTATAGGG