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| --- | --- | --- | --- |
| **Target gene** | **Primer sequence (5’ - 3’)** | **GenBank accession number** | **Efficiency (%)** |
| *act1*  (ACLA\_095800) | F TTCCATTGTCGGTCGTCCC | XM\_001270072.1 | 102.5 |
| R GTGCCTCATCACCGACATAG |
| *aclasin* (ACLA\_006820) | F CAGGCTTGCCAATATACGAGC | XM\_001273349.1 | 102 |
| R GGACATGGGTGTGGAGGATG |
| *cat2*  (ACLA\_044200) | F GACTGGGAGCTGACCAAGAG | XM\_001275125.1 | 100.5 |
| R CGCGAGATTTTCTCGTAAGC |
| *hsp30*  (ACLA\_088240) | F CACCAAAGTTCGATGTGTGC | XM\_001272560.1 | 95.5 |
| R CTCGGAGTTGGTTTTGCTGT |

F: Forward, R: Reverse.