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| **Table S3. Primer sequences used for quantitative-real time PCR (qPCR).** |
| **Gene**  | **Sequence** | **Accession number** | **Amplicon (bp)** |
| ***rps15*** | F TTCAGGAAATTCACCTACAGAGGR CCTCTGCCTGGCGCTATAC | NM\_001001819 | 93 |
| ***rps18*** | F AGATGGGAAATACAGCCAGGTR CCCCAGAAGTGACGGAGA | NM\_173234 | 111 |
| ***rpl11*** | F CTGTTCGAGGAGCCAAGGR TTGTTCTTTCTCAACTCGTACTCG | NM\_001002139 | 76 |
| ***bactin*** | F GATCTTCACTCCCCTTGTTCAR ATACCGGAGCCGTTGTCA | NM\_131031 | 91 |
| ***hif1aa*** | F GGACGGGTTTCTTTTGGTTCR CACTGTGACCAGTGAGTTCAATC | ENSDART00000044282 | 107 |
| ***hif1ab*** | F TGCAGTGCACCTGTCTCGR TGGAAGGCTAACATCTTTGGA | ENSDART00000018500 |  |
| ***nppa*** | F AGGACTGCTGCTCCTGGTTR GGCCTCCTCAAACTGCTG | NM\_198800 | 118 |
| ***nrg1*** | F ACCACAGCAAACCTATTCATCAR TCCTTCTCGCTCTCATTGC | NM\_001044911 | 98 |
| ***postnb*** | F GCGTTACCAAGGTCACTCGT R CCTCACCGAAATCTTCAATGACT  | ENSDART00000017098 | 149 |
| ***tgfb1a*** | F CAGAAGATGAAGAAACCTTGGAGR TGTCTAGCCCTGATATAGTGAAGAG | ENSDART00000060839 | 98 |
| Forward (F) and reverse (R) primer sequences, GeneBank (NM\_) or Ensmbl (ENSDART) accession numbers and amplicon length are shown. |