**Table S1.** Primer sequences used in the study.

|  |  |  |
| --- | --- | --- |
| **Gene** | **Sequence** | **Use (Product length in bp)** |
| *CrWOXB* | F-cacc\* atggtattccatctcgctttcg | Insert for RNAi (302) |
| R-gtgttggccgttccatggc |
| F-cagcggtgcttgcacgc | Expression analysis RT-PCR, RT-qPCR (283) |
| R-agccattcgtaggagacgaaga |
| F-tggtattccatctcgctttcg | Anti-sense In-situ probe synthesis (302) |
| R-taatacgactcactatagggctgcc\*gaactgattcagaca |
| F-taatacgactcactatagggtcggc\*agcattgtagaag | Sense In-situ probe synthesis (200) |
| R-ttataaagcagaacggcatagtag |
|  |  |  |
| *CrUBQ* | F-gatggccgtactcttgcagac | Expression analysis RT-PCR, RT-qPCR (348) |
| R-ggagacgaagcacgagatga |

Sequences left of \* are added for directional cloning or T7 promoter sequence. All sequences are 5’ to 3’