|  |  |  |  |
| --- | --- | --- | --- |
| **Gene** | **S/AS** | **Sequence (5’-3’)** | **Amplicon size (bp)** |
| *Igf1* | S | GCTTGCTCACCTTTACCAGC | 301 |
| AS | AAATGTACTTCCTTCTGGGTCT |
| *Socs2* | S | ATATCCGTTAAGACGTCAGCTGG | 82 |
| AS | TATGATAGAAATCCAATCTGAATTTCCC |
| *Cis* | S | ACAGAAGATGCCGGAGGGTACATTC | 152 |
| AS | TTGACAAGCAGTTAGAGTCCAGCCG |
| *Ar* | S | TACCAGCTCACCAAGCTCCT | 174 |
| AS | GATGGGCTTGACTTTCCCAG |
| *Era* | S | ATGATTGGTCTCGTCTGGCGCT | 458 |
| AS | AGCAGGTCATAGAGGGGCACAACG |
| *Mstn* | S | ACCCATGAAAGACGGTACAAG | 288 |
| AS | TCATCACAGTCAAGCCCAAAG |
| *Stat5a* | S | GAAACCTCTGGAATCTGAAGCC | 159 |
| AS | CTGCTTTCGCAAGTACAGGAGT |
| *Stat5b* | S | GAGAATTTGCCAGGACGGAA | 51 |
| AS | CACGCCATCAAACCACTGC |