Table S2. Primers used for amplification and sequencing of the complete mitochondrial genome of *Setipinna tenuifilis*

|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| Forward | Sequences (5′ to 3′) | Reverse | Sequences (5′ to 3′) | Length |
| HJ-F0 | TGGTCCTAGCCTTAATATCAG | HJ-R0 | TTGGTGTATGACTGTTGTGA | 1,466 bp |
| HJ-F1 | TCCGAGACAGCCTATACA | HJ-R1 | CAGCAGGTAGATGGAAGG | 1,340 bp |
| HJ-F2 | GTTACAACTCTAAGCCACAA | HJ-R2 | AAGTGCTAGTGTTAGTGGTAG | 1,384 bp |
| HJ-F3 | GTATTCACAGGAGGCTTCA | HJ-R3 | GCTTAGTAGTGCGGTTAGT | 1,572 bp |
| HJ-F4 | AACACTCACAGCACTAACA | HJ-R4 | GTAACGACAGGAGTAAGAGTA | 1,315 bp |
| HJ-F5 | GACCTCACCATCTTCTCAC | HJ-R5 | ACGGCTTCTACTACGATTG | 1,912 bp |
| HJ-F6 | CCTACGAATCCTCTACCTTAT | HJ-R6 | TATTGCTACGGCGACTTC | 1,310 bp |
| HJ-F7 | CAGCAGCATTCGTGTTAC | HJ-R7 | ATTAGTCCGAGTGTGAGAAG | 1,321 bp |
| HJ-F8 | CAACTGTCCTATCACTCGTA | HJ-R8 | CCTGCGTTAAGTCGTTCT | 1,116 bp |
| HJ-F9 | CAGAGGTAGCGTGAACAA | HJ-R9 | TGATGGTGAGGAGGATAATG | 1,214 bp |
| HJ-F10 | TCCACTAACAACAACCTGAT | HJ-R10 | ATGTCTTGTTCGTCGTTGA | 1,613 bp |
| HJ-F11 | TAGCCTTCCTTCACATCTG | HJ-R11 | GAGTCATTGGTCTTGGTTAG | 1,404 bp |
| HJ-F12 | AGCCGAATAAGCGAATACA | HJ-R12 | CGAGGATGGAGAATAACAGT | 1,183 bp |
| HJ-F13 | CCTCCACTTATTATTCCTCCA | HJ-R13 | CATTAGGAAGGACTCGTCAA | 1,533 bp |
| HJ-F14 | TCAGCATTACTCGGTGTTC | HJ-R14 | AGAAGTGGTGAGGTTGAAC | 1,292 bp |