Table S1. Primers used for amplification and sequencing of the complete mitochondrial genome of *Ilisha elongate.*

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| Forward | Sequences (5′ to 3′) | Reverse | Sequences (5′ to 3′) | Length |
| LY-F0 | GGATAAGGAGCAGGTATCAG | LY-R0 | GAAGTGTATTCCTTAGCCTCT | 1,488 bp |
| LY-F1 | TTCACCGAAGCCATAGATAA | LY-R1 | GAACTGGCACGATGTAGG | 1,711 bp |
| LY-F2 | GAGTTACACCTCTAAGTCTCA | LY-R2 | GGCATATCAGGCACAACC | 1,455 bp |
| LY-F3 | CCTCTCAGCACTCTTCCT | LY-R3 | TGAAGACGCAGATGAATAGT | 1,700 bp |
| LY-F4 | CTCCATCATCCTAGCAACAT | LY-R4 | GATTGAGGAGACGGTGTT | 1,814 bp |
| LY-F5 | TCTTAGCCAACTCCTCACTA | LY-R5 | CAGAGCATTGTCCGTAGAA | 1,229 bp |
| LY-F6 | AACAGACCACCGAATAGTAG | LY-R6 | GGCGATTAGTTGAATAAGGAG | 1,041 bp |
| LY-F7 | CGACTAACTGCTAATCTAACC | LY-R7 | GGAAGCCATCTGTGTTGT | 1,026 bp |
| LY-F8 | GCCGCTTGATACTGACAT | LY-R8 | TTAGTCGTTCTGCTTGGTT | 1,286 bp |
| LY-F9 | CCTAACCTGCTGACTACTAC | LY-R9 | AGACCAACGGATGACTGT | 1,347 bp |
| LY-F10 | CTATCATACACGAGAACACCTA | LY-R10 | TCCGAGGCATAAGCAGAT | 1,162 bp |
| LY-F11 | TTCCTCCTTATTCGCCTTC | LY-R11 | GTGTCGCTTGTTGTGATAG | 970 bp |
| LY-F12 | CTCCTTGTAGCCATAGAACTA | LY-R12 | GCGATGTCAGAGGTGTAG | 1,093 bp |
| LY-F13 | AACGACGCATTAGTAGACC | LY-R13 | GGAGTTAGAGGTGGAAGTTAA | 1,204 bp |
| LY-F14 | CCTTCATTATCATCGGACAA | LY-R14 | ACCATTAGTACACCACCATT | 939 bp |
| LY-F15 | TAAGAAACCACCAACCAGTT | LY-R15 | TTACGCCGAATGTATCAACT | 960 bp |