|  |
| --- |
| Nucleotide position |
|  | 16 | 42 | 58 | 72 | 123 | 157 | 158 | 229 | 234 | 246 | 252 | 338 | 346 | 393 | 423 | 486 | 633 | 656 | 717 | 786 | 796 | 827 |
| A1A2A3A4A5A6A7A8A9A10A11A12A13A14A15A16A17A18A19A20A21A22A23A24A25A26E1E2E3E4E5N1N2 | G.AA............................. | G.........................TTTTT.. | T....................G........... | G.........................AA..... | T.........................CCCCCCC | G...............................T | C.........T.....T....T........... | C.....................T.......... | A...............................G | A..............................GG | A.........................TTTTT.. | G.TT............................. | A..............G................. | G..................T...........A. | T.........................CCCCCAC | A.............................G.. | T..............................G. | C...TTTTT........................ | G.AA............................. | G..............................A. | A...............................G | T...AAAAA........................ |

Supplementary Table 2 Segregating sites between the concatenated sequences of the *nad2* gene of the haplotypes of *E. multilocularis* identified in this study (haplotypes A11 to A26) compared with the sequence of the already described haplotypes of *E. multilocularis* by Nakao et al, 2009 (haplotypes A1-A10 excluding O1). Nucleotide positions are numbered from the first nucleotide of the gene.