```
   1 TACCTAAAAA ATCCCAAACA TATAACTGAA CTCCTCACAC CCAATTGGAC(8F)
  51 CAATCTATCA CCCTATAGAA GAACTAATGT TAGTATAAGT AACATGAAAA
 101 CATTCTCCTC CGCATAAGCC TGCGTCAGAT CAAAACACTG AACTGACAAT
 151 TAACAGCCCA ATATCTACAA TCAACCAACA AGTCATTATT ACCCTCACTG(7F)
 201 TCAACCCAAC ACAGGCATGC TCATAAGGAA AGGTTAAAAA AAGTAAAAGG
 251 AACTCGGCAA ACCTTACCCC GCCTGTTTAC CAAAAACATC ACCTCTAGCA(6F/12R)
 301 TCACCAGTAT TAGAGGCACC GCCTGCCCAG TGACACATGT TTAACGGCCG
 351 CGGTACCCTA ACCGTGCAAA GGTAGCATAA TCACTTGTTC CTTAAATAGG(2F)
 401 GACCTGTATG AATGGCTCCA CGAGGGTTCA GCTGTCTCTT ACTTTTAACC
 451 AGTGAAATTG ACCTGCCCGT GAAGAGGCGG GCATGACACA GCAAGACGAG
 501 AAGACCCTAT GGAGCTTTAA TTTATTAATG CAAACAGTAC CTAACaaacc(14R)
 551 ctgttcttgg gtgggtgtgg gtataatact aagttgagat gatatcattt(4R)
 601 acgggggaag gcgctttgtg aagtaggcct tatttctctt gtcctttcgt
 651 acagggagga atttgaaggt agatagaaac cgacctggat tactccggtc
 701 tgaactcaga tcacgtagga ctttaatcgt tgaacaaacg aacctttaat(17R)
 751 agcggctgca ccatcgggat gtcctgatcc aacatcgagg tcgtaaaccc
 801 tattgttgat atggactcta gaataggatt gcgctgttat ccctagggta
 851 acttgttccg ttggtcaagt tattggatca attgagtata gtagttcgct
 901 ttgactggtg aagtcttagc atgtactgct cggaggttgg gttctgctcc
 951 gaggtcgccc caaccgaaat ttttaatgca ggtttggtag tttaggacct
1001 gtgggtttgt taggtactgt ttgcattaat aaattaaagc tccatagggt
1051 cttctcgtct tgctgtgtca tgcccgcctc ttcacgggca ggtcaatttc
1101 actggttaaa agtaagagac agctgaaccc tcgtggagcc attcatacag
1151 gtccctattt aaggaacaag tgattatgct acctttgcac ggttagggta
1201 ccgcggccgt taaacatgtg tcactgggca ggcggtgcct ctaatactgg
1251 tgatgctaga ggtgatgttt ttggtaaaca ggcggggtaa ggtttgccga
1301 gttcctttta ctttttttaa cctttcctta tgagcatgcc tgtgttgggt
1351 tgacagtgag ggtaataatg acttgttggt tgattgtaga tattgggctg
1401 ttaattgtca gttcagtgtt ttgatctgac gcaggcttat gcggaggaga
1451 atgttttcat gttacttata ctaacattag ttcttctata gggtgataga
1501 ttggtccaat tgggtgtgag gagttcagtt atatgtttgg gatttttag
1551 gtagtgggtg ttgagcttga acgctttctt aattggtggc tgcttttagg
1601 cctactatgg gtgttaaatt ttttactctc tctacaaggt tttttcctag
1651 tgtccaaaga gctgttcctc tttggactaa cagttaaatt tacaagggga
1701 tttagagggt tctgtgggca aatttaaagt tgaactaaga ttctatcttg
1751 gacaaccagc tatcaccagg ctcggtaggt ttgtcgcctc tacctataaa
1801 tcttcccact attttgctac atagacgggt gtgctctttt agctgttctt
1851 aggtagctcg tctggtttcg ggggtcttag ctttggctct ccttgcaaag(16F)
1901 ttatttctag ttaattcatt atgcagaagg tatagggggtt agtccttgct
1951 atattatgct tggttataat ttttcatctt tcccttgcgg tactatatct
2001 attgcgccag gtttcaattt ctatcgccta tactttattt gggtaaatgg
2051 tttggctaag gttgtctggt agtaaggtgg agtgggtttg gggctaggtt(10R)
2101 tagc
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Dataset S2. Sequence of the ASncmtRNA-2. The sequence of the human ASncmtRNA-2 revealed an IR of 545 nt (uppercase) linked to the 5' of the antisense 16S mitochondrial RNA (lowercase). The sequence of 44 nt at the 5' end (underlined) was determined by 5' RACE (see text). The sequences in bold correspond to the different primers utilized as shown in Fig. 2a. They are indicated at the right of the sequence. The sequence from position 552 to 1001 (italics) corresponds to the 450 nt of the loop (GenBank Access No EU863790).