

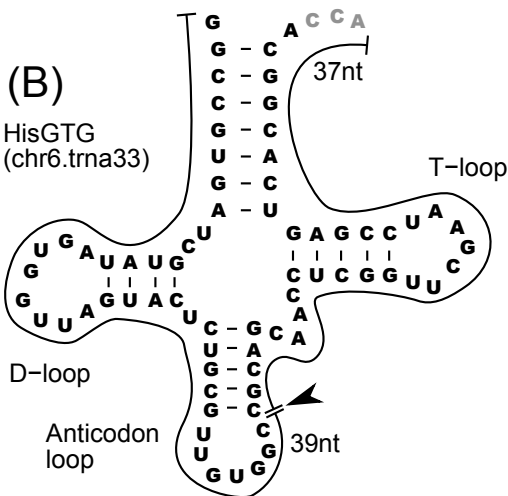
(A)

a) n=482 -----CGCAGCAACCTCGGTTCTGATCCGAGTCACGGCACCA
 b) n=374 GGCCGTGATCGTATAGTGGTTAGTACTCTGCGTTGTGGC
 c) n=222 -----CGCAGCAACCTCGGTTCTGAATCCGAGTCACGGCACCA
 d) n=180 -----CGCAGCAACCTCGGTTCTGATCCGAGTCACGGCA
 e) n=163 TGCCGTGATCGTATAGTGGTTAGTACTCTGCGTTGTGGC
 f) n=141 -----CGCAGCAACCTCGGT----ATCCGAGTCACGGCACCA
 HisGTG GCCGTGATCGTATAGTGGTTAGTACTCTGCGTTGTGGC CGCAGCAACCTCGGTTCTGAATCCGAGTCACGGCA



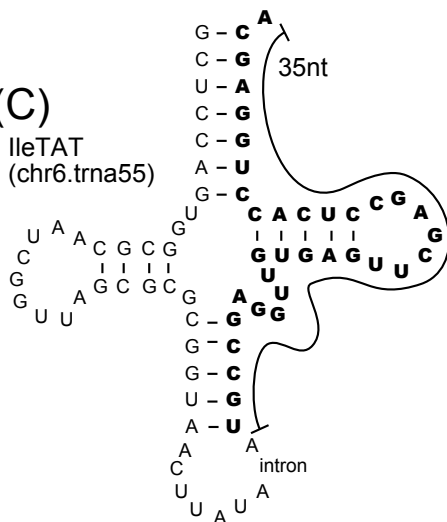
(B)

HisGTG
(chr6.trna33)



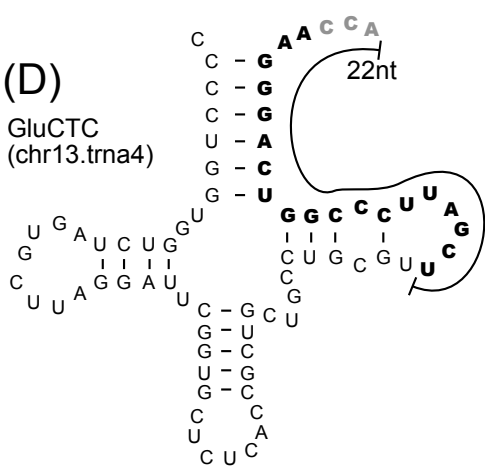
(C)

IleTAT
(chr6.trna55)



(D)

GluCTC
(chr13.trna4)



(E)

AspGTC
(chr1.trna63)

