```
      hTRE2 KYDKGHRAGLPEDKGPEPVGINSSIDRFG--ILHETELPPVTAREAKKIR 68
    hRN-tre KYDRGREGAEIEPWEDADYLVYKVTDRFG--FLHEEELPDHNVAVERQKH 68
     hEPI64 ---LSSLGSDSEANG----FAERRIDKFG--FIVGSQ-GAEGALEEVPLE 77
DdDisgorgin -SINGNSGNTP---------ISNRRNVFGQSLTLNEKEQKKKDKNRKERE 422
  DdDrainin -VLKHQREYEE---------IQRKAKKT---LEREAKEKEKLDAIRKEKE 96
  hEvi5like --EEQNRLLEAD---------SKSMRSMNGSRRNSGSSLVSSSSASSNLS 80

      hTRE2 REMTRTSK-W-MEMLGEWETYKHS--------- 90
    hRN-tre LEIERTTK-W-LKMLKGWEKYKNTEKFHRR--- 96
     hEPI64 VLRQRESK-W-LDMLNNWDKWMAKKHKKIRLR- 107
DdDisgorgin RRMEESLKIWNEDMIPNWDKRKGTKKCREM--- 452
  DdDrainin RSLIDARKVWEEDIIPVWEKNSKKRDIKKIKDL 129
  hEvi5like HLEEDTWILW-GRIANEWEEWRRRKEKLLKEL- 111
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Du et al., Fig. S3