```
         10        20        30        40        50        60
IIGGRECEKNSHPWQVAIYHYXSSFQCGGVLVNPKWVLTAAHCKNDNYEVWLXGRHNLFE
                           P              PP
    CC CCCC                C              CCCCC
    S   SSS                S              SS S S           S
                      TT TT               TT
    E   EEE                E              EE E

         70        80
NENTAQFFGVTADFPHPGFN

          C C
S     S
```

Rows :
1. Phylogeny (P)
2. COC (C)
3. SVM (S)
4. Threading (T)
5. Consensus (E)

Protein: 1HIA_A