```
            !                 !
masqhvpapsstasafpfslsgteafafpelttrTLLGSILTVVASILIFEQINYrnkkq  60
                                  <<<<transmembrane>>>>

hlpgptwtipiigkflnslhptmeVYKAQFAQGELSAFSvfhlflVMAVSNRVARKIFns 120
                         <<h-A>>><b1-1>>     <b1-2>><<h-B>>

pDYAEPclvasakqvllpenwvflhgkVHAEYRRALNVLFTRkalsiYLPIQEKIYRRYF 180
 <b1-5                      <<<<<<h-C>>>>>>     <<<<<<h-D>>>>

NEwmndpdtepkpYMMVMRDlnmetslrvfcgehmseeeqfaisdkyWLITCALQlvnfp 240
>>            <<b3-1>>                            <<h-F>>>

lalpgtkvynaikarKIAMTILEDVSAkskrsmaegneptclidaWMKEMIDAkagkges 300
               <<<<<h-G>>>>                     <<h-H>>>
     !
saeerralsrefsdheiaMVVLSFLFASQDAMTSALVNTFELLADHPEILAKVREeqyyv 360
                  <<<<<<<<<<<<<<h-i>>>>>>>>>>>>>><<h-J>>
                         *  *                    !
rhndvdaplsfnqmdemvYTRAMVKEALRLIPpvifvPYLTLkPFPILpdYTVPKgtMLI 420
                  <<<<<<h-K>>>>>     <b1-4 <b2-1  <b2-2  b1-
                                                 *
PSlwpsCHDETiypEPDSFVPERwlekggsaeanpknYMVWGAGAHKCIGFEYAYMHLAA 480
3>    <h-K'   <meander>              <<Cys pocket>><<<<<<h-L

CVGTASVTMDWHHKItpdtgKFQViatLFPKDglHLKFtprpkpsf 526
>>>>>>>>>>>b3-3     b4-1   b4-2   b3-2
```