

A

NleH1: 1 MLSPYSVNLGCSWNSLTRNLTSPDNRVLSSVRDAAVHSDNGAQVKVGNRTYRVVATDNKF
 NleH2: 1 MLSPSSINLGCWNSLTRNLTSPDNRVLSSVRDAAVHSDSGTQVTVGNRTYRVVATDNKF
 ***** * ***** * ***** * ***** * ***** * *****

61 CVTRESHSGCFTNLLHRLGWPKEISRKIEVMLNASPVSAAMERGIVHSNRDLPVDYA
 61 CVTRESHSGCFTNLLHRLGWPKEISRKIEAMLNTPSPVSTTIERGSVHSNRDLPVDYA
 ***** * ***** * ***** * ***** * ***** * *****

121 PPELPSVDY-----NRLSVPGNVIGKGNVYVEDAEDATKVLKMF TTSQSNEEV
 121 QPELPPADY TQSELPRVSNKSPVPGNVIGKGNVYVEDMEDT TKVLKMF TISQSHEEV
 ***** ** * ***** * ***** * ***** * ***** * *****

171 TSEVRCFNQYYGAGSAEKIYGNNGDIIGIRMDKINGESLLN ISSLPAQAEHAIYDMFDRL
 181 TSEVRCFNQYYGSGSAEKIYNDNGNVIGIRMNKINGESLLDIPSLPAQAEQAIYDMFDRL
 ***** * ***** * ***** * ***** * ***** * *****

231 EQKGILFVDTTETNVLYDRAKNEFNPIDISSYNVSDRSWSSESQIMQSYHGGKQDLISVVLSKI
 241 EKKGILFVDTTETNVLYDRMRNEFNPIDISSYNVSDISWSEHQVMQSYHGGKLDLISVVLSKI
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