logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 MDANTKKWLKRGGIVLGGALGLFVLAIVVDKS...TEKSKEEILELEESTYPLSDYFFVNGKKKDQLQSLPQIETHQHNTKKDYN 82
NTDB id 1218 CAA10657.1 2413..3543( ) ....MNKWLKGVIVFIGGFATITTISLVYHQKPKAPLNNQPSLLNDDEVKYPLQDYTFTQNPQPTNTESSKD..... 68
consensus
logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 TINKELEAQNKALEKALESAKQEAKKLPQVVVLKNTETPKSIGSVATPQPVAPIVPPRTRRLIKEQLELLNSRMSPFEPLKDAPK 167
NTDB id 1218 CAA10657.1 2413..3543( ) ATIKALQEQLKAALKALNS.....KEMNYSKEETFKSPPIDPNTTPPKKDFSLKQLDLLAARITPFKQSPKNYE 137
consensus
logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 A.....EIDYGVDSFENLKAQDKGTNENKLLRTITADKMIPAFLITPISSQIAGKVTAQVETDIFANMGRAVLIPKGSKVIGYY 246
NTDB id 1218 CAA10657.1 2413..3543( ) ENLIFPVDNPKGIDGFTNLKEKDIATNENKLLRTITADKMIPAFLITPISSQIAGKVIAQVESDIFAHMGKAVLIPKGSKVIGYY 222
consensus
logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 NNNNKIGEYRLNIAWTRIITPQGINIMLTNARGVDVKGYNGLIGKVIARNFSKYGMPLLTSTLSNGLLIALTSALANRQNKAGVG 331
NTDB id 1218 CAA10657.1 2413..3543( ) SNNNKMGEYRLDIVWSRIITPHGINIMLTNAKGADIKGYNGLVGELIERNFQRYGVPLLLSTLTNGLLIGITSALNNRGNKEGA. 306
consensus
logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 NPFFGDYLLMQLTRNTGMSLNQIVGQILADKARINPIIIIREGSRVFISPNLDIFIPKPRHGEVLAQFFKEKKPVVKSQEESYEE 416
NTDB id 1218 CAA10657.1 2413..3543( ) TNFFGDYLLMQLMRQSGMGINQVVNQILRDKSKIAPIVVIREGSRVFISPNTDIFFPIPRENEVIAEFLK 376
consensus
logo
NTDB id 96882 OO773 RS01610 WP 040499125.1 NEF 419
NTDB id 1218 CAA10657.1 2413..3543( ) ... 376
consensus
X non conserved
X similar
X ≥50% conserved