
logo
NTDB id 375 SMU RS06880 WP 002262929.1 .....MTNVFESSPLFLRILLAVLIILLFFYFIFLNYREYKNNNQVKQLNAKVRSLITGHYTDKLKVEDNSDLSELVNNVNDLSE 80
NTDB id 414 AAK55818.1 838..2187( ) MLDLLKQTIFTRDFIFILILLGFIL.....VVTLLLLENRRDNIQLKQINQKVKDLIAGDYSKVLDMQGGSEITNITNNLNDLSE 80
consensus
logo
NTDB id 375 SMU RS06880 WP 002262929.1 VFRLTHENLAQEKNRLTSILSYMTDGVLATDRSGKITVINDMAQKQLNVTREQALECNILDILD.DDSYTYNDLITKTPELVLTR 164
NTDB id 414 AAK55818.1 838..2187( ) VIRLTQENLEQESKRLNSILFYMTDGVLATNRRGQIIMINDTAKKQLGLVKEDVLNRSILELLKIEENYELRDLITQSPELLLDS 165
consensus
logo
NTDB id 375 SMU RS06880 WP 002262929.1 RDEYDEFITLRIRFALNRRESGFISGLIAVLHDATEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDDGALTESVAPSFIKV 249
NTDB id 414 AAK55818.1 838..2187( ) QDINGEYLNLRVRFALIRRESGFISGLVAVLHDTTEQEKEERERRLFVSNVSHELRTPLTSVKSYLEALDEGALCETVAPDFIKV 250
consensus
logo
NTDB id 375 SMU RS06880 WP 002262929.1 SLDETNRMMRMITDLLSLSRIDNQTSHLDVELTNFTAFMNYILDRFDQIQSQQSTNKVYEIIRDYPDKSVWIEIDTDKMTQVIDN 334
NTDB id 414 AAK55818.1 838..2187( ) SLDETNRMMRMVTDLLHLSCIDNATSHLDVELINFTAFITFILNRFDKMKGQE.KEKKYELVRDYPINSIWMEIDTDKMTQVVDN 334
consensus
logo
NTDB id 375 SMU RS06880 WP 002262929.1 ILNNAIKYSPDGGKVTITMQTTDTQLILSISDQGLGIPKKDLPLIFDRFYRVDKARSRAQGGTGLGLAIAKEIVKQHKGFIWANS 419
NTDB id 414 AAK55818.1 838..2187( ) ILNNAIKYSPDGGKITVRMKTTEDQMILSISDHGLGIPKQDLPRIFDRFYRVDRARSRAQGGTGLGLSIAKEIIKQHKGFIWAKS 419
consensus
logo
NTDB id 375 SMU RS06880 WP 002262929.1 EEGEGSTFTIVLPYENDNDAIDEWEEDEDES 450
NTDB id 414 AAK55818.1 838..2187( ) EYGKGSTFTIVLPYDKDAVKEEVWEDEVED 449
consensus
X non conserved
X similar
X ≥50% conserved