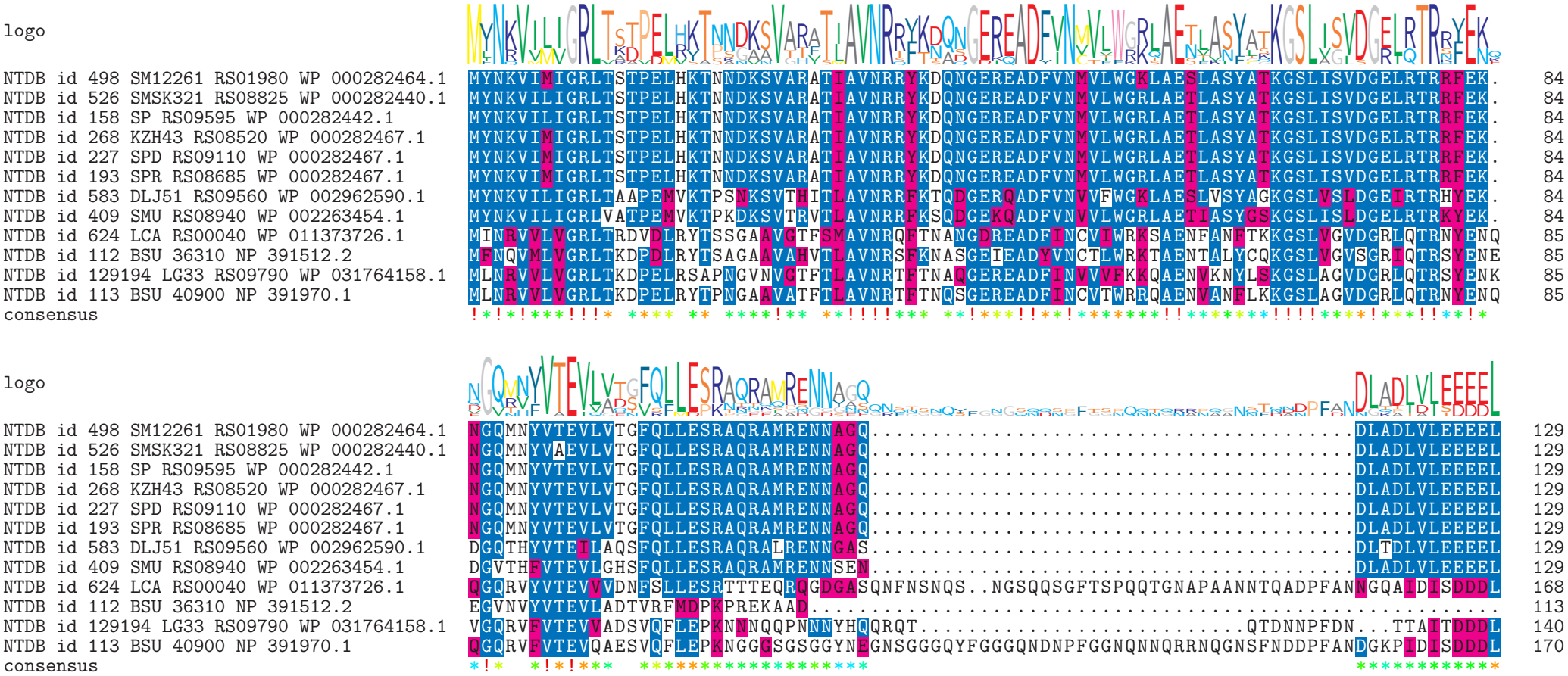

logo
NTDB id 498 SM12261 RS01980 WP 000282464.1 MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGKLAESLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 526 SMSK321 RS08825 WP 000282440.1 MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 158 SP RS09595 WP 000282442.1 MYNKVILIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 268 KZH43 RS08520 WP 000282467.1 MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 227 SPD RS09110 WP 000282467.1 MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 193 SPR RS08685 WP 000282467.1 MYNKVIMIGRLTSTPELHKTNNDKSVARATIAVNRRYKDQNGEREADFVNMVLWGRLAETLASYATKGSLISVDGELRTRRFEK. 84
NTDB id 583 DLJ51 RS09560 WP 002962590.1 MYNKVILIGRLTAAPEMVKTPSNKSVTHITLAVNRRFKTQDGERQADFVNVVFWGKLAESLVSYAGKGSLVSLDGEIRTRHYEK. 84
NTDB id 409 SMU RS08940 WP 002263454.1 MYNKVILIGRLVATPEMVKTPKDKSVTRVTLAVNRRFKSQDGEKQADFVNVVLWGRLAETIASYGSKGSLISLDGELRTRKYEK. 84
NTDB id 624 LCA RS00040 WP 011373726.1 MINRVVLVGRLTRDVDLRYTSSGAAVCTFSMAVNRQFTNANGDREADFINCVIWRKSAENFANFTKKGSLVGVDGRLQTRNYENQ 85
NTDB id 112 BSU 36310 NP 391512.2 MFNQVMLVGRLTKDPDLRYTSAGAAVAHVTLAVNRSFKNASGEIEADYVNCTLWRKTAENTALYCQKGSLVGVSGRIQTRSYENE 85
NTDB id 129194 LG33 RS09790 WP 031764158.1 MLNRVVLVGRLTKDPELRSAPNGVNVGTFTLAVNRTFTNAQGEREADFINVVVFKKQAENVKNYLSKGSLAGVDGRLQTRSYENK 85
NTDB id 113 BSU 40900 NP 391970.1 MLNRVVLVGRLTKDPELRYTPNGAAVATFTLAVNRTFTNQSGEREADFINCVTWRRQAENVANFLKKGSLAGVDGRLQTRNYENQ 85
consensus
logo
NTDB id 498 SM12261 RS01980 WP 000282464.1 NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 526 SMSK321 RS08825 WP 000282440.1 NGQMNYVAEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 158 SP RS09595 WP 000282442.1 NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 268 KZH43 RS08520 WP 000282467.1 NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 227 SPD RS09110 WP 000282467.1 NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 193 SPR RS08685 WP 000282467.1 NGQMNYVTEVLVTGFQLLESRAQRAMRENNAGQ.....DLADLVLEEEEL 129
NTDB id 583 DLJ51 RS09560 WP 002962590.1 DGQTHYVTEILAQSFQLLESRAQRALRENNGAS.....DLTDLVLEEEEL 129
NTDB id 409 SMU RS08940 WP 002263454.1 DGVTHFVTEVLGHSFQLLESRAQRAMRENNSEN.....DLADLVLEEEEL 129
NTDB id 624 LCA RS00040 WP 011373726.1 QGQRVYVTEVVVDNFSLLESRTTTEQRQGDGASQNFNSNQS..NGSQQSGFTSPQQTGNAPAANNTQADPFANNGQAIDISDDDL 168
NTDB id 112 BSU 36310 NP 391512.2 EGVNVYVTEVLADTVRFMDPKPREKAAD..... 113
NTDB id 129194 LG33 RS09790 WP 031764158.1 VGQRVFVTEVVADSVQFLEPKNNNQQPNNNYHQQRQT.....QTDNNPFDN...TTAITDDDL 140
NTDB id 113 BSU 40900 NP 391970.1 QGQRVFVTEVQAESVQFLEPKNGGGSGSGGYNFGNSGGGQYFGGGQNDNPFGGNQNNQRRNQGNSFNDDPFANDGKPIDISDDDL 170
consensus
logo
NTDB id 498 SM12261 RS01980 WP 000282464.1 PF 131
NTDB id 526 SMSK321 RS08825 WP 000282440.1 PF 131
NTDB id 158 SP RS09595 WP 000282442.1 PF 131
NTDB id 268 KZH43 RS08520 WP 000282467.1 PF 131
NTDB id 227 SPD RS09110 WP 000282467.1 PF 131
NTDB id 193 SPR RS08685 WP 000282467.1 PF 131
NTDB id 583 DLJ51 RS09560 WP 002962590.1 PF 131
NTDB id 409 SMU RS08940 WP 002263454.1 PF 131
NTDB id 624 LCA RS00040 WP 011373726.1 PF 170
NTDB id 112 BSU 36310 NP 391512.2 .. 113
NTDB id 129194 LG33 RS09790 WP 031764158.1 PF 142
NTDB id 113 BSU 40900 NP 391970.1 PF 172
consensus
X non conserved
X similar
X ≥50% conserved