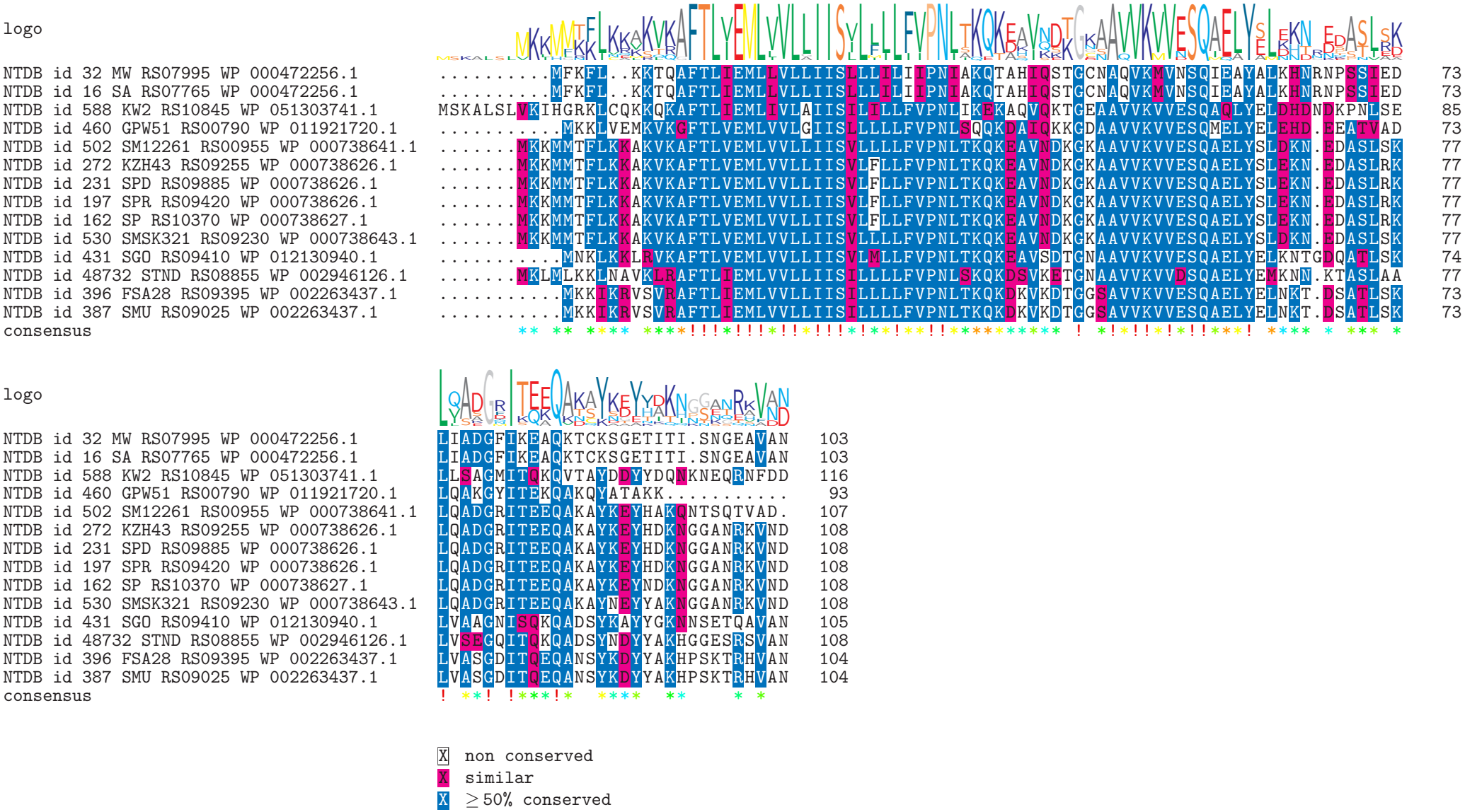
logo
NTDB id 32 MW RS07995 WP 000472256.1 ....MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED 73
NTDB id 16 SA RS07765 WP 000472256.1 ....MFKFL..KKTQAFTLIEMLLVLLIISLLLILIIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNPSSIED 73
NTDB id 588 KW2 RS10845 WP 051303741.1 MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILLLLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDKPNLSE 85
NTDB id 460 GPW51 RS00790 WP 011921720.1 ..........MKKLVEMKVKGFTLVEMLVVLGIISLLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHD.EEATVAD 73
NTDB id 502 SM12261 RS00955 WP 000738641.1 .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK 77
NTDB id 272 KZH43 RS09255 WP 000738626.1 .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK 77
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NTDB id 162 SP RS10370 WP 000738627.1 .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKN.EDASLRK 77
NTDB id 530 SMSK321 RS09230 WP 000738643.1 .......MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKN.EDASLSK 77
NTDB id 431 SGO RS09410 WP 012130940.1 ..........MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQATLSK 74
NTDB id 48732 STND RS08855 WP 002946126.1 .......MKLMLKKLNAVKLRAFTLIEMLVVLLIISILLLLFVPNLSKQKDSVKETGNAAVVKVVDSQAELYEMKNN.KTASLAA 77
NTDB id 396 FSA28 RS09395 WP 002263437.1 ..........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK 73
NTDB id 387 SMU RS09025 WP 002263437.1 ..........MKKIKRVSVRAFTLIEMLVVLLIISILLLLFVPNLTKQKDKVKDTGGSAVVKVVESQAELYELNKT.DSATLSK 73
consensus ** ** **** ****!!!*!!!*!!*!!!*!**!**!!*********** ! *!*!!*!*!!***! **** * *** *
logo
NTDB id 32 MW RS07995 WP 000472256.1 LIADGFIKEAQKTCKSGETITI.SNGEAVAN 103
NTDB id 16 SA RS07765 WP 000472256.1 LIADGFIKEAQKTCKSGETITI.SNGEAVAN 103
NTDB id 588 KW2 RS10845 WP 051303741.1 LLSAGMITQKQVTAYDDYYDQNKNEQRNFDD 116
NTDB id 460 GPW51 RS00790 WP 011921720.1 LQAKGYITEKQAKQYATAKK..... 93
NTDB id 502 SM12261 RS00955 WP 000738641.1 LQADGRITEEQAKAYKEYHAKQNTSQTVAD. 107
NTDB id 272 KZH43 RS09255 WP 000738626.1 LQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 231 SPD RS09885 WP 000738626.1 LQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 197 SPR RS09420 WP 000738626.1 LQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 162 SP RS10370 WP 000738627.1 LQADGRITEEQAKAYKEYNDKNGGANRKVND 108
NTDB id 530 SMSK321 RS09230 WP 000738643.1 LQADGRITEEQAKAYNEYYAKNGGANRKVND 108
NTDB id 431 SGO RS09410 WP 012130940.1 LVAAGNISQKQADSYKAYYGKNNSETQAVAN 105
NTDB id 48732 STND RS08855 WP 002946126.1 LVSEGQITQKQADSYNDYYAKHGGESRSVAN 108
NTDB id 396 FSA28 RS09395 WP 002263437.1 LVASGDITQEQANSYKDYYAKHPSKTRHVAN 104
NTDB id 387 SMU RS09025 WP 002263437.1 LVASGDITQEQANSYKDYYAKHPSKTRHVAN 104
consensus ! **! !***!* **** ** * *
X non conserved
X similar
X ≥ 50% conserved