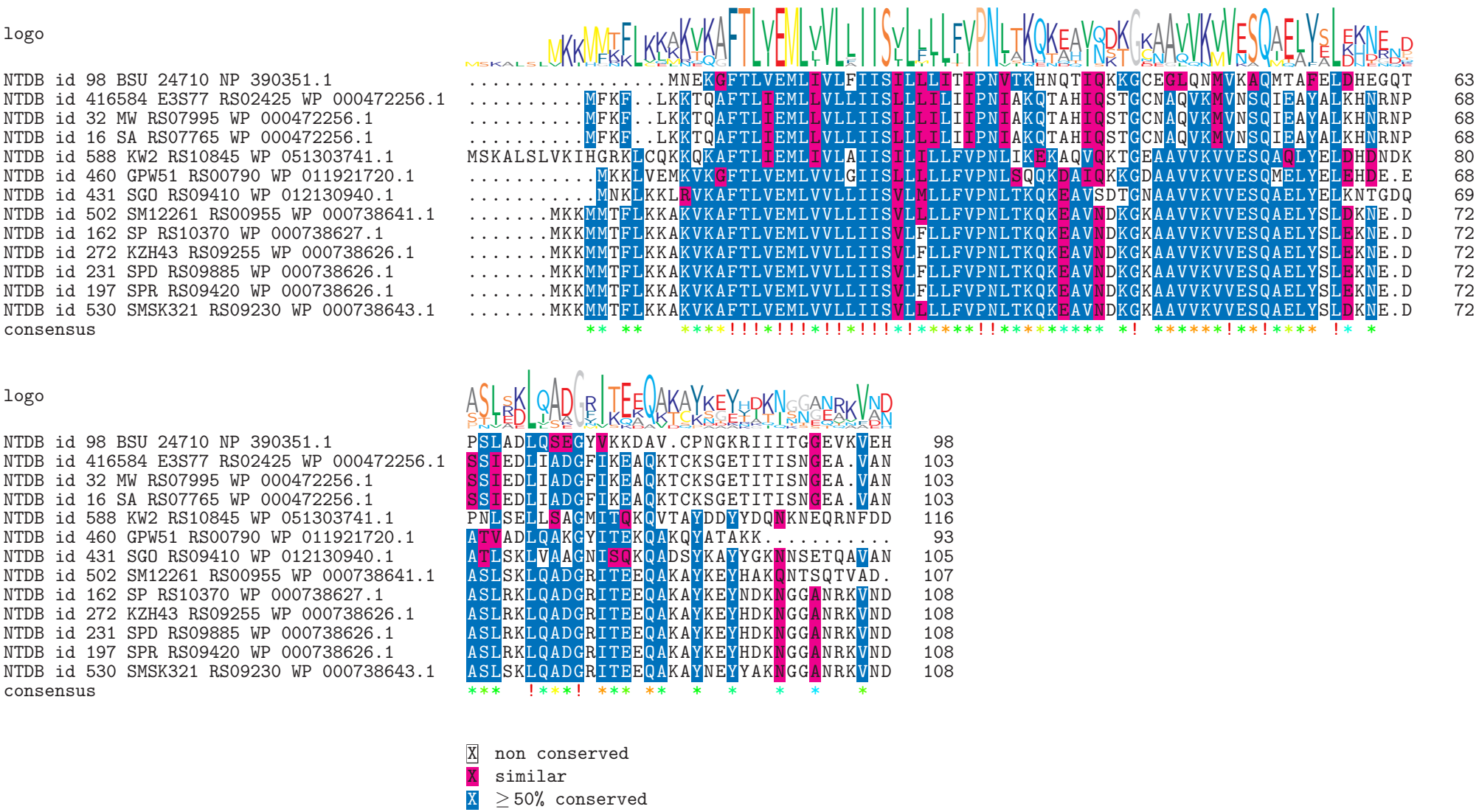
logo
NTDB id 98 BSU 24710 NP 390351.1 .....MNEKGFTLVEMLIVLFIISILLLITIPNVTKHNQTIQKKGCEGLQNMVKAQMTAFELDHEGQT 63
NTDB id 416584 E3S77 RS02425 WP 000472256.1 .....MFKF..LKKTQAFTLIEMLLVLLIISLLLILIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP 68
NTDB id 32 MW RS07995 WP 000472256.1 .....MFKF..LKKTQAFTLIEMLLVLLIISLLLILIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP 68
NTDB id 16 SA RS07765 WP 000472256.1 .....MFKF..LKKTQAFTLIEMLLVLLIISLLLILIPNIAKQTAHIQSTGCNAQVKMVNSQIEAYALKHNRNP 68
NTDB id 588 KW2 RS10845 WP 051303741.1 MSKALSLVKIHGRKLCQKKQKAFTLIEMLIVLAIISILILLFVPNLIKEKAQVQKTGEAAVVKVVESQAQLYELDHDNDK 80
NTDB id 460 GPW51 RS00790 WP 011921720.1 .....MKKLVEMKVKGFTLVEMLVVLGIISLLLLFVPNLSQQKDAIQKKGDAAVVKVVESQMELYELEHDE.E 68
NTDB id 431 SGO RS09410 WP 012130940.1 .....MNKLKKLRVKAFTLVEMLVVLLIISVLMLLFVPNLTKQKEAVSDTGNAAVVKVVESQAELYELKNTGDQ 69
NTDB id 502 SM12261 RS00955 WP 000738641.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNE.D 72
NTDB id 162 SP RS10370 WP 000738627.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.D 72
NTDB id 272 KZH43 RS09255 WP 000738626.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.D 72
NTDB id 231 SPD RS09885 WP 000738626.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.D 72
NTDB id 197 SPR RS09420 WP 000738626.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLFLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLEKNE.D 72
NTDB id 530 SMSK321 RS09230 WP 000738643.1 .....MKKMMTFLKKAKVKAFTLVEMLVVLLIISVLLLLFVPNLTKQKEAVNDKGKAAVVKVVESQAELYSLDKNE.D 72
consensus
logo
NTDB id 98 BSU 24710 NP 390351.1 PSLADLQSEGYVKKDAV.CPNGKRIIITGGEVKVEH 98
NTDB id 416584 E3S77 RS02425 WP 000472256.1 SSIEDLIADGFIKEAQKTCKSGETITISNGEA.VAN 103
NTDB id 32 MW RS07995 WP 000472256.1 SSIEDLIADGFIKEAQKTCKSGETITISNGEA.VAN 103
NTDB id 16 SA RS07765 WP 000472256.1 SSIEDLIADGFIKEAQKTCKSGETITISNGEA.VAN 103
NTDB id 588 KW2 RS10845 WP 051303741.1 PNLSELLSACMITQKQVTAYDDYYDQNKNEQRNFDD 116
NTDB id 460 GPW51 RS00790 WP 011921720.1 ATVADLQAKGYITEKQAKQYATAKK..... 93
NTDB id 431 SGO RS09410 WP 012130940.1 ATLSKLVAAGNISQKQADSYKAYYGKNNSETQAVAN 105
NTDB id 502 SM12261 RS00955 WP 000738641.1 ASLSKLQADGRITEEQAKAYKEYHAKQNTSQTVAD. 107
NTDB id 162 SP RS10370 WP 000738627.1 ASLRKLQADGRITEEQAKAYKEYNDKNGGANRKVND 108
NTDB id 272 KZH43 RS09255 WP 000738626.1 ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 231 SPD RS09885 WP 000738626.1 ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 197 SPR RS09420 WP 000738626.1 ASLRKLQADGRITEEQAKAYKEYHDKNGGANRKVND 108
NTDB id 530 SMSK321 RS09230 WP 000738643.1 ASLSKLQADGRITEEQAKAYNEYYAKNGGANRKVND 108
consensus
X non conserved
X similar
X ≥ 50% conserved