

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	MTCQNCMINEAT.....	12
NTDB id 281 KZH43 RS03550 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 245 SPD RS03860 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 243 SPR RS03635 WP 000882517.1	MTCQNCKINDST.....	12
NTDB id 241 SP RS04015 WP 000882523.1	MTCQNCKINDST.....	12
NTDB id 1139572 DQM87 RS03300 WP 002983882.1	MTCQNCNINEST.....	12
NTDB id 378 SMU RS02690 WP 002263569.1	MTCQNCKINEAT.....	12
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGTAAKALQALGLGSEKIQKEVESLIG.....R	70
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYTPTLDRILEKAEYYAHQYQYGTIESAHLLEAMATTSGSLAYSLLAGMNVDSDDLIDL.....EDLSSH	72
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYS	70
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAAILAIGKKPKSSS	76
NTDB id 297 STER RS00545 WP 011680614.1	...MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVTGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLLETWHVLLAMVEVPGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
consensus	**.....*	

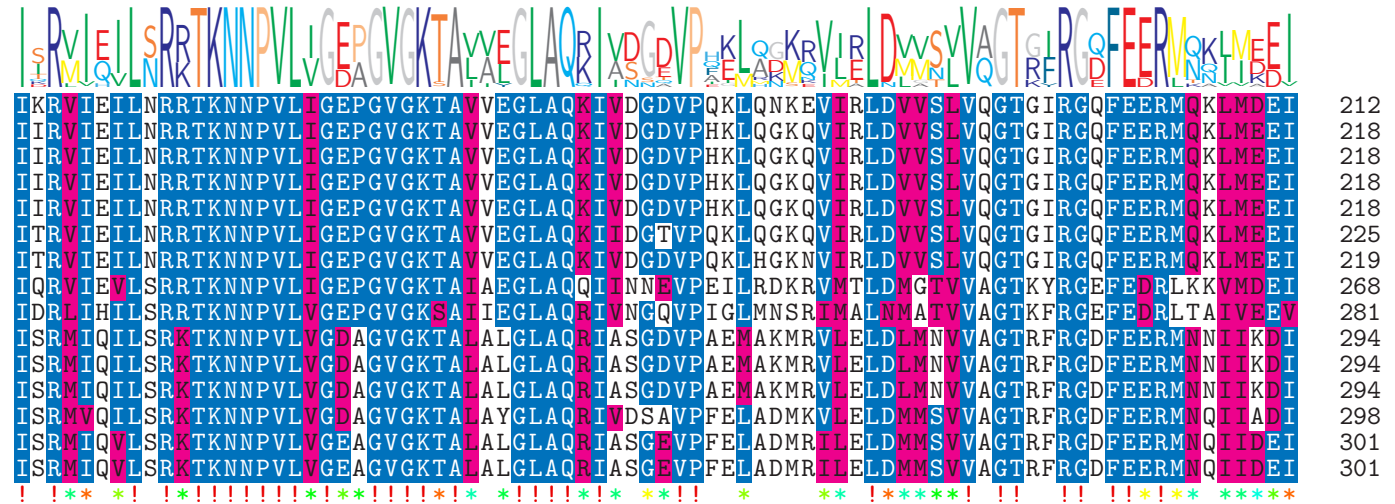
logo

NTDB id 601 KW2 RS02615 WP 021036805.1	IHL YTSVNGQKQKIDLCQNCYQI...MKSQGTEALFGAGNASMGNS.....DEPFN.PFNDIFSATQQGQ	72
NTDB id 281 KZH43 RS03550 WP 000882517.1	IHL YTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLN.....NRDF.DPFGDFFNDLNNF	69
NTDB id 245 SPD RS03860 WP 000882517.1	IHL YTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLN.....NRDF.DPFGDFFNDLNNF	69
NTDB id 243 SPR RS03635 WP 000882517.1	IHL YTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLN.....NRDF.DPFGDFFNDLNNF	69
NTDB id 241 SP RS04015 WP 000882523.1	IHL YTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDLN.....NRDF.DPFGDFFNDLNNF	69
NTDB id 1139572 DQM87 RS03300 WP 002983882.1	IHL YTSVNGKQKQVLDLCQNCYQI...MKSDPANSTLNLGTPGYRAQ.....DRSTSPFFDDFFGD LNNF	73
NTDB id 378 SMU RS02690 WP 002263569.1	IHL YANVNGQKQKIDLCQNCYQI...MKTDPENGPLNHLSSQQ...N.....SSSINPFFDDFFGD LNNF	70
NTDB id 85 BSU 00860 NP 387967.1	GQEMSQTIHYTPRAKKVIELSMDEARKLGHSYVCTEHLILGLIREGEGVAARVLNNLGVSLNKARQQVLQL.....	141
NTDB id 610 V4T04 RS10165 WP 012897346.1	VKVKRSTLRFSPRAEVEVMTAASF LAIHNNSEAVCTEHLIYALLQVEDGFG LQLLKL.....QKINIVSLRKETEK R	143
NTDB id 290 KZH43 RS10025 WP 001109677.1	QDETFTELPFSSRRLQILFDEAEYVASV VHAKVLCTEHLVYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEER	150
NTDB id 289 SPD RS10700 WP 001109677.1	QDETFTELPFSSRRLQILFDEAEYVASV VHAKVLCTEHLVYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEER	150
NTDB id 287 SP RS11210 WP 001109712.1	QDETFTELPFSSRRLQVLFDEAEYVASV VHAKVLCTEHLVYAILHDSNALATRILERAGFSYEDKKDQVKIAALRRNLEER	150
NTDB id 377 SMU RS09275 WP 002262344.1	N..DITMLEQSHALKHTLA EAAAI SQVTHAKEVCTEHLVLFAMLLNPNLLATRILELVGFHAKDDGESIRLLDLRKVTER Y	154
NTDB id 297 STER RS00545 WP 011680614.1	ESDIIDLRAQSPALEAMLQEAQEIASVTGAVEVGESEHVLMAFILHKLDMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERN	156
NTDB id 329 STU RS10020 WP 011225298.1	ESDIIDLRAQSPALEAMLQEAQGIASVTGAVEVGESEHVLMAFILHKLDMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERN	156
consensus	* * * * * * * * * * * * * *	

```

NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
NTDB id 241 SP RS04015 WP 000882523.1
NTDB id 1139572 DQM87 RS03300 WP 002983882.1
NTDB id 378 SMU RS02690 WP 002263569.1
NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

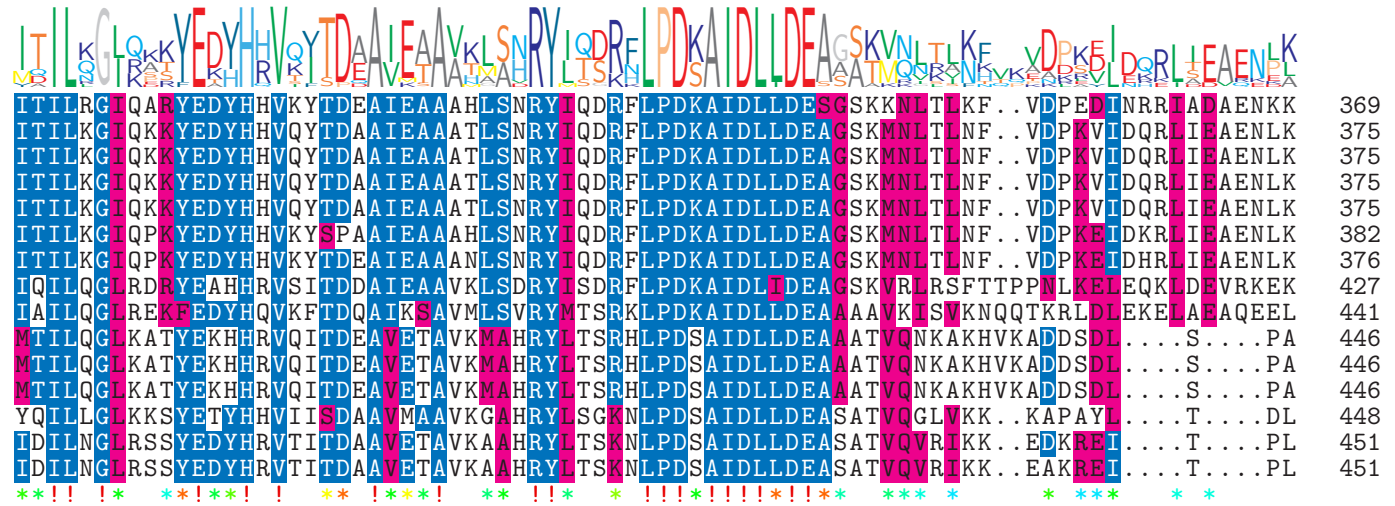
```



```

NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
NTDB id 241 SP RS04015 WP 000882523.1
NTDB id 1139572 DQM87 RS03300 WP 002983882.1
NTDB id 378 SMU RS02690 WP 002263569.1
NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

```



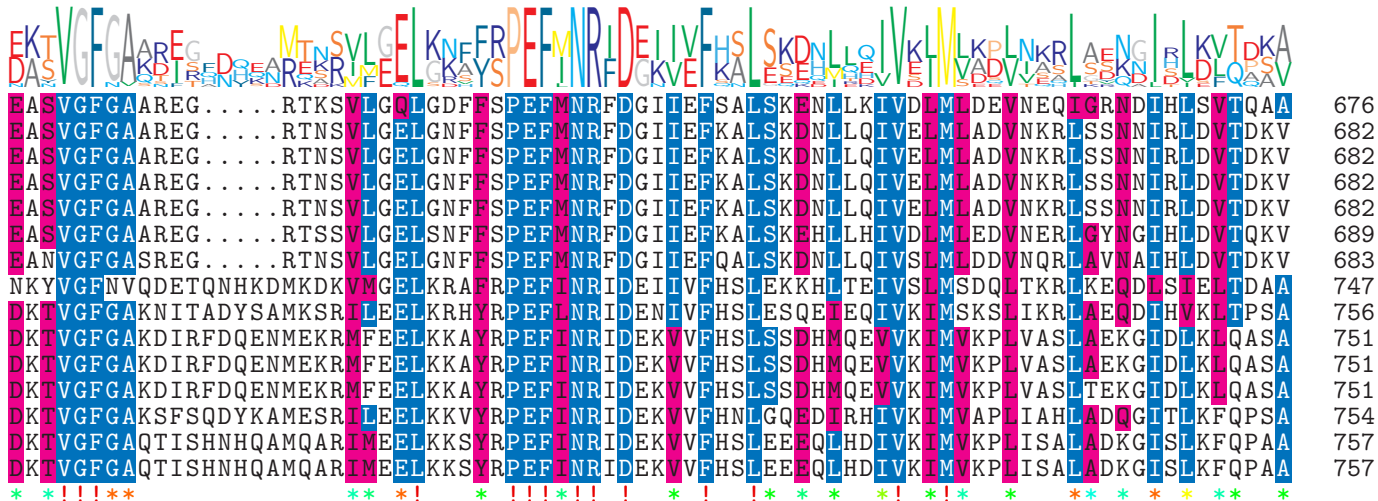
NTDB id	601 KW2 RS02615 WP 021036805.1	NEATKAEDFEKAAHFRDQISKLRELQK.....QEVTDEDMPVITEKDTEQIVEEQTKTIPVGDLKEKEQTQLINLADLLK	443
NTDB id	281 KZH43 RS03550 WP 000882517.1	SQATREEDFEKAAAYFRDQIAKYKEMQK.....KKITDQDTPIISEKTIEHIIIEQKTNIPVGDLKEKEEQQSLIHLAEDLK	449
NTDB id	245 SPD RS03860 WP 000882517.1	SQATREEDFEKAAAYFRDQIAKYKEMQK.....KKITDQDTPIISEKTIEHIIIEQKTNIPVGDLKEKEEQQSLIHLAEDLK	449
NTDB id	243 SPR RS03635 WP 000882517.1	SQATREEDFEKAAAYFRDQIAKYKEMQK.....KKITDQDTPIISEKTIEHIIIEQKTNIPVGDLKEKEEQQSLIHLAEDLK	449
NTDB id	241 SP RS04015 WP 000882523.1	SQATREEDFEKAAAYFRDQIAKYKEMQK.....KKITDQDTPIISEKTIEHIIIEQKTNIPVGDLKEKEEQQSLIHLAEDLK	449
NTDB id	1139572 DQM87 RS03300 WP 002983882.1	AQATRDEDYERAAAYFRDQITYKEMQA.....QKVDEQDIPITITEKTIEAIVEEQKTNIPVGDLKEKEEQQSLVNLANDLK	456
NTDB id	378 SMU RS02690 WP 002263569.1	AQATRDEDYEKAAAYFRDQIAKYKEMQS.....AKLDKENTPVITEKNIEVIVEEQKTNIPVGELKEKEEQQSLIHLADDLK	450
NTDB id	85 BSU 00860 NP 387967.1	DAAVQSSEFEKAAASLRDTEQILREQVEDTKKSWKEKQGENSEVTVDIDAMVVSSWTGVVPVSKIAQTETDKLINMENILH	507
NTDB id	610 V4T04 RS10165 WP 012897346.1	SEAVIKLDIKASRTKEKAVEKAIADKIYKFS....VKEDKRQEVTDAVAIVASTLTGVPITQMTKSESDDLINLEKELH	516
NTDB id	290 KZH43 RS10025 WP 001109677.1	DKALMDGKWQKAAQLIAKEE.....E....VPV.YKDLVTESDILTTL SRLSGIPVQKLTQTD AKKYLNLEAEHL	511
NTDB id	289 SPD RS10700 WP 001109677.1	DKALMDGKWQKAAQLIAKEE.....E....VPV.YKDLVTESDILTTL SRLSGIPVQKLTQTD AKKYLNLEAEHL	511
NTDB id	287 SP RS11210 WP 001109712.1	DKALMDGKWQKAAQLIAKEE.....E....VPV.YKDLVTESDILTTL SRLSGIPVQKLTQTD AKKYLNLEAEHL	511
NTDB id	377 SMU RS09275 WP 002262344.1	DHALITNDYQLAKRLLKKEK.....K....PLLKKRAEVKEEDILATLSKLSGIPVKKLTQADSKKYLNLERELH	514
NTDB id	297 STER RS00545 WP 011680614.1	DEALISGDIGA AVKQYKANQ.....K....AKFPKPALVDADQIMQTL SRLSGIPVEKMTQTD SKRYLNLESELH	517
NTDB id	329 STU RS10020 WP 011225298.1	DEALISGDIGA AVKQYKANQ.....K....AKFPKPALVDADQIMQTL SRLSGIPVEKMTQTD SKRYLNLESELH	517
consensus		! * * * * *	!

[illegible]

```

NTDB id 601 KW2 RS02615 WP 021036805.1
NTDB id 281 KZH43 RS03550 WP 000882517.1
NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
NTDB id 241 SP RS04015 WP 000882523.1
NTDB id 1139572 DQM87 RS03300 WP 002983882.1
NTDB id 378 SMU RS02690 WP 002263569.1
NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 377 SMU RS09275 WP 002262344.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
consensus

```



logo

NTDB id 601 KW2 RS02615 WP 021036805.1	KEKLVLDLGYNPAMGARPLRRTIQENTEDSIAIFYEHPEYKQLVADLIDDKIVISNQTQETAETTDDEEVP	748
NTDB id 281 KZH43 RS03550 WP 000882517.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDTTIDYYLENPSEKDLKAVMTSKCNIIQIKSAKKAIEVKSSEKEK	752
NTDB id 245 SPD RS03860 WP 000882517.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDTTIDYYLENPSEKDLKAVMTSKCNIIQIKSAKKAIEVKSSEKEK	752
NTDB id 243 SPR RS03635 WP 000882517.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDTTIDYYLENPSEKDLKAVMTSKCNIIQIKSAKKAIEVKSSEKEK	752
NTDB id 241 SP RS04015 WP 000882523.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDTTIDYYLENPSEKDLKAVMTSKCNIIQIKSAKKAIEVKSSEKEK	752
NTDB id 1139572 DQM87 RS03300 WP 002983882.1	KEKLVLDLGYDPKMGARPLRRTIQDYIEDAITDYYLEHPTEKQLRALMTNSENITIKAVKEGDSFLNEESL	760
NTDB id 378 SMU RS02690 WP 002263569.1	KEKLVLDLGYDPKMGARPLRRTIQEHTIEDAITDFYLEHPEQKELKAIMTSNCKIVIKASNKVETVTKATSD	753
NTDB id 85 BSU 00860 NP 387967.1	KAKVAEEGVLDLEYGARPLRRRIQKHVEDRLSEELLRGNIHKGQHVLVDVEDGEFVVVKTAKTN	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	VKLIAEVCFDPEYGARPLRKALQKEVEDLLSEQLLSGEIKAGNHVSTIGASNKKIKIAQIV	816
NTDB id 290 KZH43 RS10025 WP 001109677.1	LKLLANQGYDPEMGARPLRRTIQTEVEDKLAELLKKGDLVAGSILKIGVKACQLKFDIA	810
NTDB id 289 SPD RS10700 WP 001109677.1	LKLLANQGYDPEMGARPLRRTIQTEVEDKLAELLKKGDLVAGSILKIGVKACQLKFDIA	810
NTDB id 287 SP RS11210 WP 001109712.1	LKLLANQGYDPEMGARPLRRTIQTEVEDKLAELLKKGDLVAGSTLKIGVKACQLKFDIA	810
NTDB id 377 SMU RS09275 WP 002262344.1	LKHLALAGYDAEMGARPLRRTIQTEVEDKLAELLSGKLASGQTIKIGIQKEKIKFDIV	813
NTDB id 297 STER RS00545 WP 011680614.1	LKHLAKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTIKIGCSKDKITFTVV	816
NTDB id 329 STU RS10020 WP 011225298.1	LKHLAKDGYDIEMGARPLRRTIQTQVEDKLSELLLGGQVVSGQTIKIGCSKDKITFTVV	816
consensus	* * !*** *!!!!!!***! *!! *** *	

- non conserved
- similar
- ≥ 50% conserved