

logo

NTDB id 601 KW2 RS02615 WP 021036805.1	MLCQNCNINEAT.....I	13
NTDB id 281 KZH43 RS03550 WP 000882517.1	MLCQNCNKINDST.....I	13
NTDB id 245 SPD RS03860 WP 000882517.1	MLCQNCNKINDST.....I	13
NTDB id 243 SPR RS03635 WP 000882517.1	MLCQNCNKINDST.....I	13
NTDB id 241 SP RS04015 WP 000882523.1	MLCQNCNKINDST.....I	13
NTDB id 51540 H7678 RS07595 WP 003060519.1	MLCQNCNINEST.....I	13
NTDB id 378 SMU RS02690 WP 002263569.1	MLCQNCNKINEAT.....I	13
NTDB id 85 BSU 00860 NP 387967.1	..MMFGRFTEAQQVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLIG.....RGQEMS	75
NTDB id 610 V4T04 RS10165 WP 012897346.1	MKFENIKYTPTLDRILEKAEEYAHQYQYGTIESAHLAAMATTSGSIAYSLLAGMNVDSDDLIDL.....EDLSSHVKVKR	77
NTDB id 287 SP RS11210 WP 001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 290 KZH43 RS10025 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 289 SPD RS10700 WP 001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNSHSYSVAGATLNDYPYEMDRLE.....EVALELTETDYSQDET	75
NTDB id 377 SMU RS09275 WP 002262344.1	MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLLAMVEIDSSVAGLSFAEFADRVRFEDYQAAAAILAIGKKPKSSSN..D	79
NTDB id 297 STER RS00545 WP 011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFLWTWHVLLAMVEVTGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDI	81
NTDB id 329 STU RS10020 WP 011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFLWTWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETAAVLAMEKRPKDLSESDI	81
consensus	**.....*	*

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NTDB id 245 SPD RS03860 WP 000882517.1	HLYTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDL...NNR.....DFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 243 SPR RS03635 WP 000882517.1	HLYTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDL...NNR.....DFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 241 SP RS04015 WP 000882523.1	HLYTNLNGKQKQIDLCQNCYKI...IKTDPNNSLFKGMTDL...NNR.....DFD.PFGDFFNDLNNFRPSSN..TPP	77
NTDB id 51540 H7678 RS07595 WP 003060519.1	HLYTNVNGKQKQVDLCQNCYQI...MKTDPTNPILGGLNGGQRAQER.....SSNPFFDDFFGDLLNNFRAFNG..LPN	81
NTDB id 378 SMU RS02690 WP 002263569.1	HLYANVNGCQKQIDLCQNCYQI...MKTDPENGPLNHL...SQQNSS.....SINPFFDDFFGDLLNNFRAFNNQDLPN	80
NTDB id 85 BSU 00860 NP 387967.1	QTIHYTPRAKKVIELSMDEARKLGHSYVGTTEHILLGLIREGEGVAARVLNNLGVSLNKRQVQLQL.....LGSNET....	147
NTDB id 610 V4T04 RS10165 WP 012897346.1	STLRFSPRAEEVMTAASFLAIHNNSEAVGTEHLLYALLQVEDGFGQLLLKL.....QKINIVSLRKELEKRTGLKVPESKK	153
NTDB id 287 SP RS11210 WP 001109712.1	TELPFSRRLQVLFDEAEYVASVVHAKVLGTEHVLVYAILHDSNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 290 KZH43 RS10025 WP 001109677.1	TELPFSRRLQILFDEAEYVASVVHAKVLGTEHVLVYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 289 SPD RS10700 WP 001109677.1	TELPFSRRLQILFDEAEYVASVVHAKVLGTEHVLVYAILHDGNALATRILERAGFSYEDKKDQVKIAALRRNLEERAGWTREDLKA	160
NTDB id 377 SMU RS09275 WP 002262344.1	MLLEQSHALKRTLAEEAAISQVTHAKEVGTTEHVLVYAILHDSNALATRILELVGFHAKDDGESIRLLDLRKVTIERYACFSKEDIKA	164
NTDB id 297 STER RS00545 WP 011680614.1	DLRAQSPALEAMLQEAQEIASVTGAVEVGEHVLMAFLLHKDLMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
NTDB id 329 STU RS10020 WP 011225298.1	DLRAQSPALEAMLQEAQGIASVTGAVEVGEHVLMAFLLHKDLMVCRLLEVAGFQYKDDSDKPRIIDLRRSLERNAGLSKQDLKA	166
consensus	* * * * *	* * *

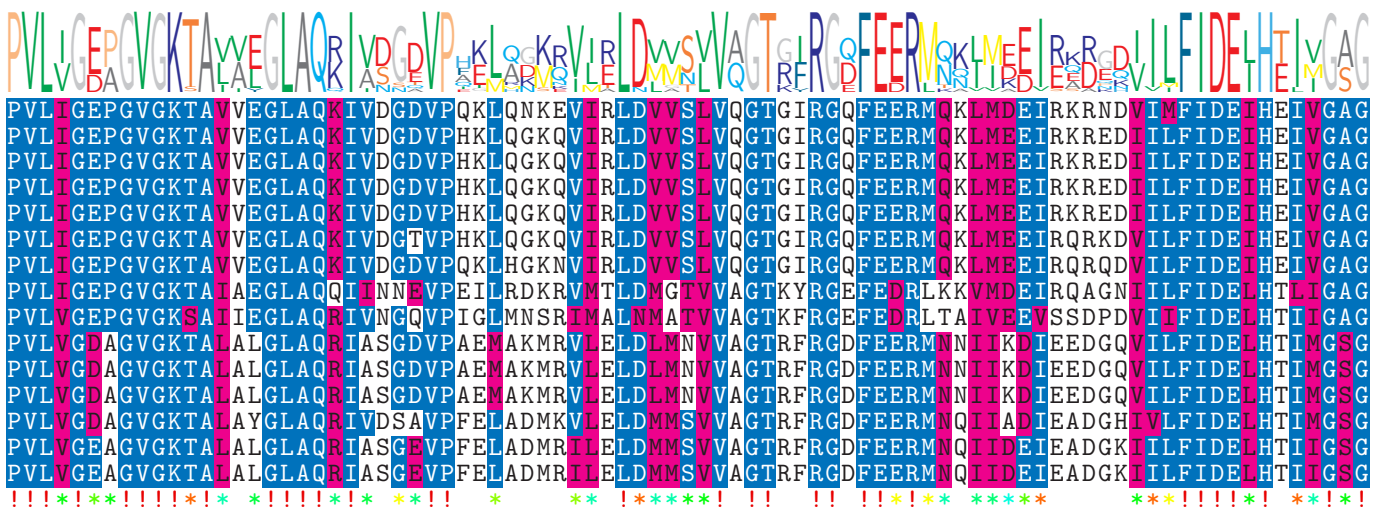
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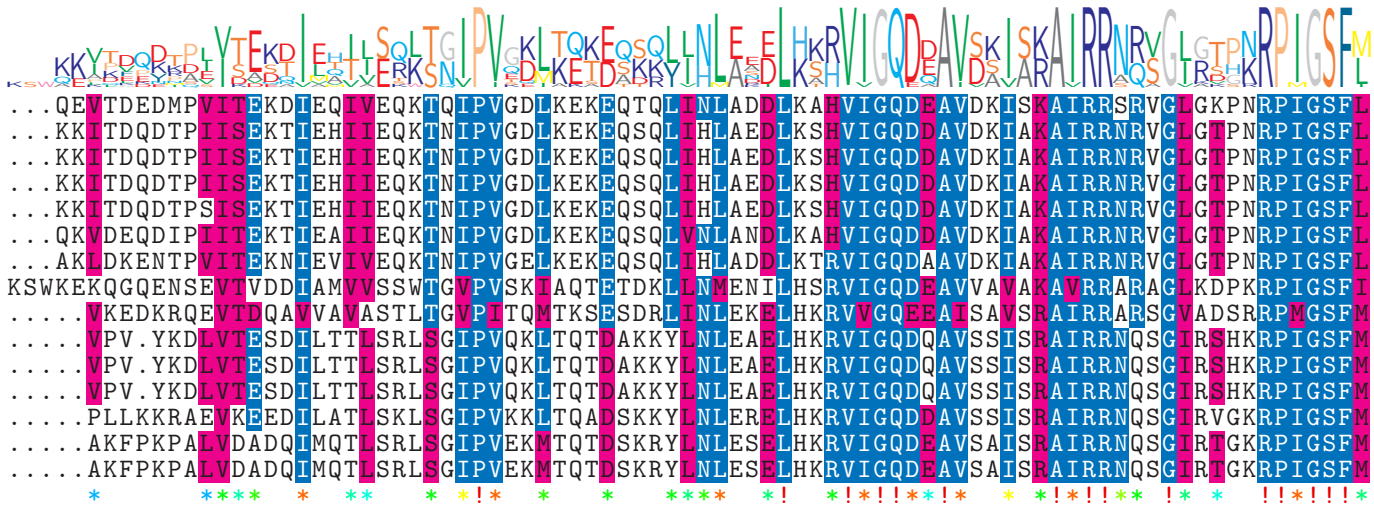


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NTDB	id	601	KW2	RS02615	WP	021036805.1	AAAHLSNRYIQDRFLPDKAIDLLDEAGSKMNLTLKF..VDPEDINRRITADAENKKNEATKAEDFEKAAHFRDQISKLRELQK...	396
NTDB	id	281	KZH43	RS03550	WP	000882517.1	AAATLSNRYIQDRFLPDKAIDLLDEAGSKMNLTLNF..VDPKVIDQRLIEAENLKSQATREDFEKAAYFRDQIAKYKEMQK...	402
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NTDB	id	241	SP	RS04015	WP	000882523.1	AAATLSNRYIQDRFLPDKAIDLLDEAGSKMNLTLNF..VDPKVIDQRLIEAENLKSQATREDFEKAAYFRDQIAKYKEMQK...	402
NTDB	id	51540	H7678	RS07595	WP	003060519.1	AAAHLSNRYIQDRFLPDKAIDLLDEAGSKMNLTLNF..VDPKVIDQRLIEAENLKSQATREDFEKAAYFRDQIAKYKEMQK...	402
NTDB	id	378	SMU	RS02690	WP	002263569.1	AAANLSNRYIQDRFLPDKAIDLLDEAGSKMNLTLNF..VDPKVIDQRLIEAENLKSQATREDFEKAAYFRDQIAKYKEMQK...	402
NTDB	id	85	BSU	O0860	NP	387967.1	AAVKSLSRYISDRFLPDKAIDLIDEAGSKVRLRSFTTPPNLKELEQKLDVVRKEKDAAVQSQEEFEKAASLRDTEQLRREQVEDTK	457
NTDB	id	610	V4T04	RS10165	WP	012897346.1	SAVMLSVMRYMSTRKFLPDKAIDLLDEAGAAVKSLSVKNQQTKRDLLEKELAEAAQEELSEAVIKLDTKASRTKEKAVEKIADKIYKFS	471
NTDB	id	287	SP	RS11210	WP	001109712.1	TAVKMAHRYLTSRHLPDS AIDLLDEAG AATVQNKAKHVKA DSDL....S....PADKALMDGKWKQAAQLIAKEE.....E	467
NTDB	id	290	KZH43	RS10025	WP	001109677.1	TAVKMAHRYLTSRHLPDS AIDLLDEAG AATVQNKAKHVKA DSDL....S....PADKALMDGKWKQAAQLIAKEE.....E	467
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NTDB	id	377	SMU	RS09275	WP	002262344.1	AAVKGAAHRYLSGKNLPDS AIDLLDEAG SATVQGLVKK..KAPAYL....T....DLDHALITNDYQLAKRLKKKEK.....K	469
NTDB	id	329	STER	RS00545	WP	011680614.1	TAVKAAHRYLTSKNLPDS AIDLLDEAG SATVQVRIKK..EDKREI....T....PLDEALISGDIGAAVKQYKANQ.....K	472
NTDB	id	297	STU	RS10020	WP	011225298.1	TAVKAAHRYLTSKNLPDS AIDLLDEAG SATVQVRIKK..EAKREI....T....PLDEALISGDIGAAVKQYKANQ.....K	472
consensus							* ! * * ! ! * * ! ! ! * ! ! ! * ! ! ! * * * * * * * * * * * * * * * * ! * * * * * * * * * * * *	

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NTDB id 378 SMU RS02690 WP 002263569.1
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NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
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



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NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 329 STU RS10020 WP 011225298.1
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NTDB id	601	KW2	RS02615	WP	021036805.1	HMFLQIILEDGRLTDAQGRTVSVFKD	SLIIMTSNAGTGK..VEASVVGFGAAREG....RTKS	VLGQLGDFFSPEFMNRRFDGIIEFS	641
NTDB id	281	KZH43	RS03550	WP	000882517.1	HMFLQVLDDGRLTDGQGRTVSVFKDAIIIMTSNAGTGK..TEASVVGFGAAREG....RTNSVLGELGNFFSPEFMNRRFDGIIEFK			647
NTDB id	245	SPD	RS03860	WP	000882517.1	HMFLQVLDDGRLTDGQGRTVSVFKDAIIIMTSNAGTGK..TEASVVGFGAAREG....RTNSVLGELGNFFSPEFMNRRFDGIIEFK			647
NTDB id	243	SPR	RS03635	WP	000882517.1	HMFLQVLDDGRLTDGQGRTVSVFKDAIIIMTSNAGTGK..TEASVVGFGAAREG....RTNSVLGELGNFFSPEFMNRRFDGIIEFK			647
NTDB id	241	SP	RS04015	WP	000882523.1	HMFLQVLDDGRLTDGQGRTVSVFKDAIIIMTSNAGTGK..TEASVVGFGAAREG....RTNSVLGELGNFFSPEFMNRRFDGIIEFK			647
NTDB id	51540	H7678	RS07595	WP	003060519.1	HMFLQVLDDGRLTDGQGRTVSVFKDTIIIMTSNAGTGK..SEASVVGFGAAREG....RTSSVLGELSNFFSPEFMNRRFDGIIEFK			654
NTDB id	378	SMU	RS02690	WP	002263569.1	HMFLQVLDDGRLTDGQGRTVSVFKDTIIIMTSNAGTGK..SEANVVGFGASREG....RTNSVLGELGNFFSPEFMNRRFDGIIEFK			648
NTDB id	85	BSU	00860	NP	387967.1	NILLQVLEDGRLTDSKGRVDFRNTILIMTSNVLGASELKRNVYGFNVQDETQNHKMDKDKVMGELKRAFRPEFINRIDEIIVFH			712
NTDB id	610	V4T04	RS10165	WP	012897346.1	NIMLQILDDGFVTDTKGRKVDVFRNTIIIMTSNVLGATALRDDKTVGFGAKNITADYSAMKSRIIEELKRHYRPEFLNRIDENIVFH			721
NTDB id	287	SP	RS11210	WP	001109712.1	NVLLQVLDDGVLTDISKGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVVFH			716
NTDB id	290	KZH43	RS10025	WP	001109677.1	NVLLQVLDDGVLTDISKGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVVFH			716
NTDB id	289	SPD	RS10700	WP	001109677.1	NVLLQVLDDGVLTDISKGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAKDIRFDQENMEKRMFEELKKAYRPEFINRIDEKVVVFH			716
NTDB id	377	SMU	RS09275	WP	002262344.1	NILLQVLDDGVLTDISRGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAKSFSQDYKAMESRIIEELKKVYRPEFINRIDEKVVVFH			719
NTDB id	297	STER	RS00545	WP	011680614.1	NILLQVLDDGVLTDISRGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAQTISHNHQAMQARIMEELKKSYPPEFINRIDEKVVVFH			722
NTDB id	329	STU	RS10020	WP	011225298.1	NVLLQVLDDGVLTDISRGRKVDVFSNTIIIMTSNVLGATALRDDKTVGFGAQTISHNHQAMQARIMEELKKSYPPEFINRIDEKVVVFH			722
consensus						* !!!*!!! *!! !! ! !* ****!!! ! *			

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NTDB id	601	KW2	RS02615	WP	021036805.1	ALSKENLLKIVDLMIDEVNEQIGRNDIHLSVTQA	AAKEKLVDLGYNPAMGARPLRRTIQENIEDS	IADFYIEHPEYKQL.VADLID	725
NTDB id	281	KZH43	RS03550	WP	000882517.1	ALSKDNLLQIVELMLADVNRKLSSNNIRLDVTDKVK	KEKLVDLGYDPKMGARPLRRTIQDYIEDTITD	YYLENPSEKDL.KAVMTS	731
NTDB id	245	SPD	RS03860	WP	000882517.1	ALSKDNLLQIVELMLADVNRKLSSNNIRLDVTDKVK	KEKLVDLGYDPKMGARPLRRTIQDYIEDTITD	YYLENPSEKDL.KAVMTS	731
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NTDB id	241	SP	RS04015	WP	000882523.1	ALSKDNLLQIVELMLADVNRKLSSNNIRLDVTDKVK	KEKLVDLGYDPKMGARPLRRTIQDYIEDTITD	YYLENPSEKDL.KAVMTS	731
NTDB id	51540	H7678	RS07595	WP	003060519.1	ALSKDNLLHIVDLMIEDVNRKLSYNGIHLDTQKV	KEKLVDLGYDPKMGARPLRRTIQDYIEDAITD	YYLEHPTTEKDL.RAIMTS	738
NTDB id	378	SMU	RS02690	WP	002263569.1	ALSKDNLLQIVSLMLDDVNQRLAVNAIHLDTQKV	KEKLVDLGYDPKMGARPLRRTIQEHIEDIAITD	FYLEHPEQKEL.KAIMTS	732
NTDB id	85	BSU	00860	NP	387967.1	SLEKKHLTEIVSLMSDQLTKRLKEQDLSIELTDA	AAKVAEAGVLDLEYGARPLRRAIQKHVEDRLSE	ELLRGNHKGQHTVL	797
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NTDB id	377	SMU	RS09275	WP	002262344.1	NLGQEDIRHIVKIMVAPLIAHLADQGITLKFQPS	ALKHLALAGYDAEMGARPLRRTLQTEVEDKLA	ELISGKLASGQTL	804
NTDB id	297	STER	RS00545	WP	011680614.1	SLEEEQLHDIVKIMVKPLISALADKGISLKFQPA	AALKHLAKDGYDIEMGARPLRRTIQTQVEDK	LSELLGGQVVSGQTL	807
NTDB id	329	STU	RS10020	WP	011225298.1	SLEEEQLHDIVKIMVKPLISALADKGISLKFQPA	AALKHLAKDGYDIEMGARPLRRTIQTQVEDK	LSELLGGQVVSGQTL	807
consensus						!* *** *! *!* * *			

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NTDB id 601 KW2 RS02615 WP 021036805.1	DKI	V	I	S	N	Q	T	Q	E	.	T	A	E	T	T	D	E	E	V	P	A	E	.	748	
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NTDB id 245 SPD RS03860 WP 000882517.1	K	G	N	I	Q	I	K	S	A	K	K	A	E	V	K	S	S	E	K	E	K	.	.	.	752
NTDB id 243 SPR RS03635 WP 000882517.1	K	G	N	I	Q	I	K	S	A	K	K	A	E	V	K	S	S	E	K	E	K	.	.	.	752
NTDB id 241 SP RS04015 WP 000882523.1	K	G	N	I	Q	I	K	S	A	K	K	A	E	V	K	S	S	E	K	E	K	.	.	.	752
NTDB id 51540 H7678 RS07595 WP 003060519.1	NSN	I	M	I	K	A	A	K	K	D	E	T	V	T	A	K	Q	D	K	T	D	E	N		763
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NTDB id 85 BSU 00860 NP 387967.1	D	G	E	F	V	V	K	T	T	A	K	T	N	.	.	.	.	.	.	.	.	.	.	.	810
NTDB id 610 V4T04 RS10165 WP 012897346.1	N	K	K	I	K	I	A	Q	I	V	.	.	.	.	.	.	.	.	.	.	.	.	.	816	
NTDB id 287 SP RS11210 WP 001109712.1	A	G	Q	L	K	F	D	I	A	.	.	.	.	.	.	.	.	.	.	.	.	.	.	810	
NTDB id 290 KZH43 RS10025 WP 001109677.1	A	G	Q	L	K	F	D	I	A	.	.	.	.	.	.	.	.	.	.	.	.	.	.	810	
NTDB id 289 SPD RS10700 WP 001109677.1	A	G	Q	L	K	F	D	I	A	.	.	.	.	.	.	.	.	.	.	.	.	.	.	810	
NTDB id 377 SMU RS09275 WP 002262344.1	K	E	K	L	K	F	D	I	V	.	.	.	.	.	.	.	.	.	.	.	.	.	.	813	
NTDB id 297 STER RS00545 WP 011680614.1	K	D	K	L	T	F	T	V	V	.	.	.	.	.	.	.	.	.	.	.	.	.	.	816	
NTDB id 329 STU RS10020 WP 011225298.1	K	D	K	L	T	F	T	V	V	.	.	.	.	.	.	.	.	.	.	.	.	.	.	816	
consensus	*	*																							

- ⓧ non conserved
- ✖ similar
- ⓧ ≥ 50% conserved