

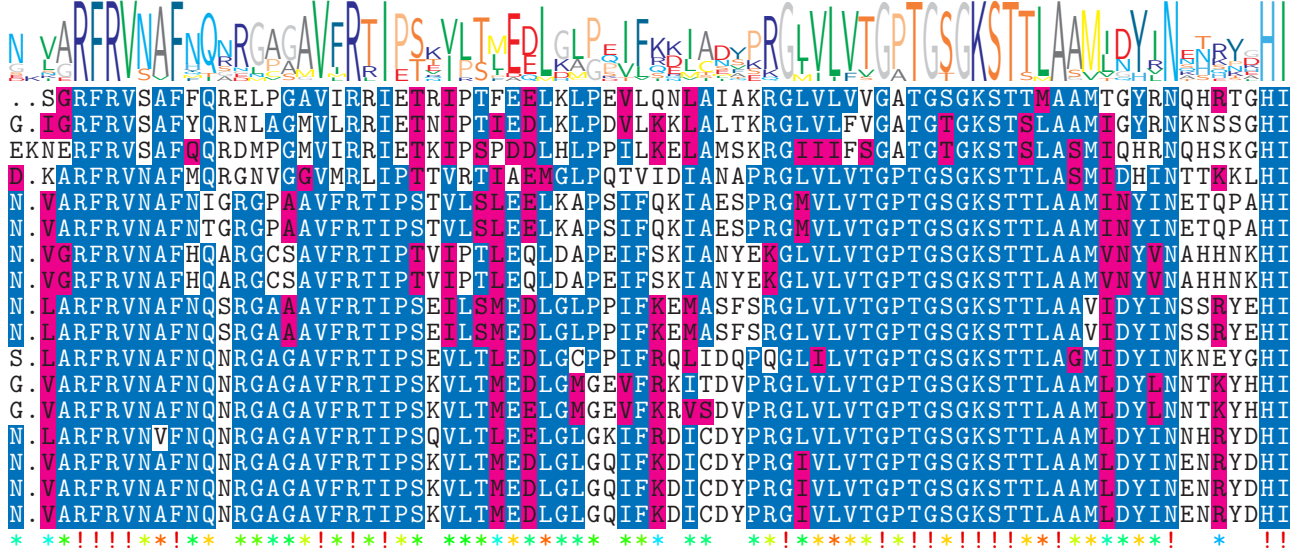
logo

NTDB	id	1177	A1552VC	RS01165	WP	000422572.1
NTDB	id	1203	PSJM300	01605	AFN76401.1	
NTDB	id	1019	ACIAD	RS04205	WP	004922051.1
NTDB	id	1307	DR	RS10055	WP	010888596.1
NTDB	id	1086	NMV	RS00245	WP	002218487.1
NTDB	id	1116	NGFG	RS10615	WP	003690200.1
NTDB	id	1185	GTF74	RS11940	WP	000350195.1
NTDB	id	1176	A1552VC	RS01160	WP	000350195.1
NTDB	id	1283	ACE17W	RS02630	WP	010947729.1
NTDB	id	1259	GCO85	RS10305	WP	010947729.1
NTDB	id	1112362	P3C55	RS07805	WP	003489919.1
NTDB	id	1202	PSJM300	01600	AFN76400.1	
NTDB	id	1194	PAKAF	RS01995	WP	003084552.1
NTDB	id	1020	ACIAD	RS04210	WP	011182164.1
NTDB	id	1081	FDQ49	RS05335	WP	006581387.1
NTDB	id	1060	ABD1	RS04380	WP	000355489.1
NTDB	id	1040	HON27	RS13360	WP	000355489.1
consensus						



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NTDB	id	1203	PSJM300	01605	AFN76401.1	
NTDB	id	1019	ACIAD	RS04205	WP	004922051.1
NTDB	id	1307	DR	RS10055	WP	010888596.1
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NTDB	id	1112362	P3C55	RS07805	WP	003489919.1
NTDB	id	1202	PSJM300	01600	AFN76400.1	
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NTDB	id	1060	ABD1	RS04380	WP	000355489.1
NTDB	id	1040	HON27	RS13360	WP	000355489.1
consensus						



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NTDB id 1177 A1552VC RS01165 WP 000422572.1
NTDB id 1203 PSJM300 01605 AFN76401.1
NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1307 DR RS10055 WP 01088596.1
NTDB id 1086 NMV RS00245 WP 002218487.1
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NTDB id 1176 A1552VC RS01160 WP 000350195.1
NTDB id 1283 ACE17W RS02630 WP 010947729.1
NTDB id 1259 GCO85 RS10305 WP 010947729.1
NTDB id 1112362 P3C55 RS07805 WP 003489919.1
NTDB id 1202 PSJM300 01600 AFN76400.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 1020 ACIAD RS04210 WP 011182164.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKTI	EQDALHSA	DSANDLRL	MLKTKRGDDY	...GSGSLQNVKI	DME		368
NTDB id 1203 PSJM300 01605 AFN76401.1	TFDQALYNLYVQCEIT	YEDALLHAD	SSNDLRL	MIKLGSETD	GEHLTSVS	.QGLTL	EMSDDDPGRSFR	
NTDB id 1019 ACIAD RS04205 WP 004922051.1	TFDQALFDLYDHQHI	SYADALKHAD	SPNNLRL	QIKLSEEANK	PLDNRRERMSEMER	KMTFDGQRRNLK		383
NTDB id 1307 DR RS10055 WP 010888596.1	TMDAYLAGLFR	HLITYDK	CLERAI	.DPKE	..FARLANDASAGL	.GGASMTPAAPPV	PAYNSASSSSTGRGGDFGRGAAA	395
NTDB id 1086 NMV RS00245 WP 002218487.1	TMDQSLQSLVRQGLI	IAPEVARRRA	QNSESM	..SF				347
NTDB id 1116 NGFG RS10615 WP 003690200.1	TMDQSLQSLVRQGLI	IAPEAARRRA	QNSESM	..SF				347
NTDB id 1185 GTF74 RS11940 WP 000350195.1	TMEQNAKQLI	ARGVVD	DAQEVQSKI	ELDLKA	..F			345
NTDB id 1176 A1552VC RS01160 WP 000350195.1	TMEQNAKQLI	ARGVVD	DAQEVQSKI	ELDLKA	..F			345
NTDB id 1283 ACE17W RS02630 WP 010947729.1	TLDQHLTEL	VNKNII	ISKHTA	HEAAL	.NKSL	..F		344
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NTDB id 1112362 P3C55 RS07805 WP 003489919.1	TLDQHLQDLVKRSL	ITRNQAREYAK	.DKRI	..FE				345
NTDB id 1202 PSJM300 01600 AFN76400.1	TLDQHLQDLVKRSL	ITRNQAREYAK	.DKRI	..FE				345
NTDB id 1194 PAKAF RS01995 WP 003084552.1	TLDQHLQDLVKRSL	ITRNQAREYAK	.DKRI	..FE				344
NTDB id 1020 ACIAD RS04210 WP 011182164.1	TLDQSLKGLI	AKGLIS	RENAREKAK	.IPEN	..F			344
NTDB id 1081 FDQ49 RS05335 WP 006581387.1	TLDQSLKGLVSRGVIS	PQTARTAAK	.QPES	..FL				345
NTDB id 1060 ABD1 RS04380 WP 000355489.1	TLDQSLKGLVSRGVIS	PQTARTAAK	.QPES	..FL				345
NTDB id 1040 HON27 RS13360 WP 000355489.1	TLDQSLKGLVSRGVIS	PQTARTAAK	.QPES	..FL				345
consensus	!*** *	!*	*****	*	*	*		

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NTDB id 1177 A1552VC RS01165 WP 000422572.1	PATPAAPAAGRSEPGRAGGFRR	368
NTDB id 1203 PSJM300 01605 AFN76401.1		381
NTDB id 1019 ACIAD RS04205 WP 004922051.1		383
NTDB id 1307 DR RS10055 WP 010888596.1	PATPAAPAAGRSEPGRAGGFRR	418
NTDB id 1086 NMV RS00245 WP 002218487.1		347
NTDB id 1116 NGFG RS10615 WP 003690200.1		347
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NTDB id 1202 PSJM300 01600 AFN76400.1		344
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NTDB id 1060 ABD1 RS04380 WP 000355489.1		345
NTDB id 1040 HON27 RS13360 WP 000355489.1		345
consensus		

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved