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NTDB id 1040 HON27 RS13360 WP 000355489.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

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NTDB id 1040 HON27 RS13360 WP 000355489.1
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consensus

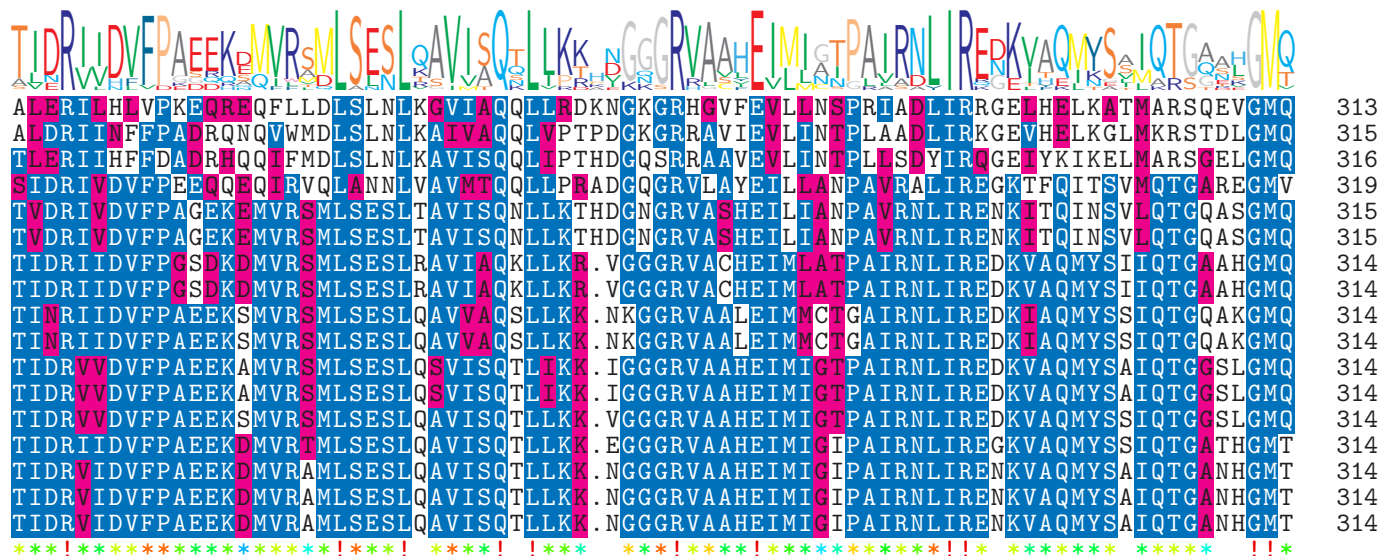
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NTDB id 1259 GCO85 RS10305 WP 010947729.1
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NTDB id 1040 HON27 RS13360 WP 000355489.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

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Protein	Sequence	Score
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TDFQALYNLYVQGETIYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS. QGLTL..... EMSDDDPGRSFR.....	381	
TDFQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEMER..... KMTFDGQRRNLK.....	383	
TMDAYLAGLFRHLLITYDKLERAIID.PKEFARLA.. NDASAGL.GGASMPAAPPVPAYNSASSSSSTGRGGDFGRGAAA	395	
TMDQSLQSLVRQGLIAPEVARRRRAQNSSEMSF.....	347	
TMDQSLQSLVRQGLIAPEAARRRAQNSSEMSF.....	347	
TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345	
TMEQNAKQLIARGVVDAQEVQSKIELDLKAF.....	345	
TLDQHLTELVNKNIISKHTAHEAALN.KSLF.....	344	
TLDQHLTELVNKNIISKHTAHEAALN.KSLF.....	344	
TLDMCLKGLVAKGLISRENAREKAKI.PENF.....	344	
TLDMCLKGLVAKGLISRENAREKAKI.PENF.....	344	
TLDSCCLKGLLAKGLISRDSAKEKAKQ.PENF.....	344	
TLDQSLKQLLAKGLINLPTARAVAKS.PESF.....	344	
TLDQSLKGLVARGVISPQTARTAAKQ.PESFL.....	345	
TLDQSLKGLVARGVISPQTARTAAKQ.PESFL.....	345	
TLDQSLKGLVSRGVISPQTARTAAKQ.PESFL.....	345	

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NTDB	id	1019	ACIAD	RS04205	WP	004922051.1		383
NTDB	id	1307	DR	RS10055	WP	010888596.1	PATPAAPAAGRSEPGRAGGFGR	418
NTDB	id	1086	NMV	RS00245	WP	002218487.1		347
NTDB	id	1116	NGFG	RS10615	WP	003690200.1		347
NTDB	id	1185	GTF74	RS11940	WP	000350195.1		345
NTDB	id	1176	A1552VC	RS01160	WP	000350195.1		345
NTDB	id	1283	ACE17W	RS02630	WP	010947729.1		344
NTDB	id	1259	GCO85	RS10305	WP	010947729.1		344
NTDB	id	366469	FF962	RS20575	WP	003084552.1		344
NTDB	id	1194	PAKAF	RS01995	WP	003084552.1		344
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
NTDB	id	1081	FDQ49	RS05335	WP	006581387.1		345
consensus								

- X non conserved
- X similar
- X $\geq 50\%$ conserved