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NTDB id 1086 NMV RS00245 WP 002218487.1
NTDB id 1116 NGFG RS10615 WP 003690200.1
NTDB id 1185 GTF74 RS11940 WP 000350195.1
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NTDB id 1259 GCO85 RS10305 WP 010947729.1
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consensus

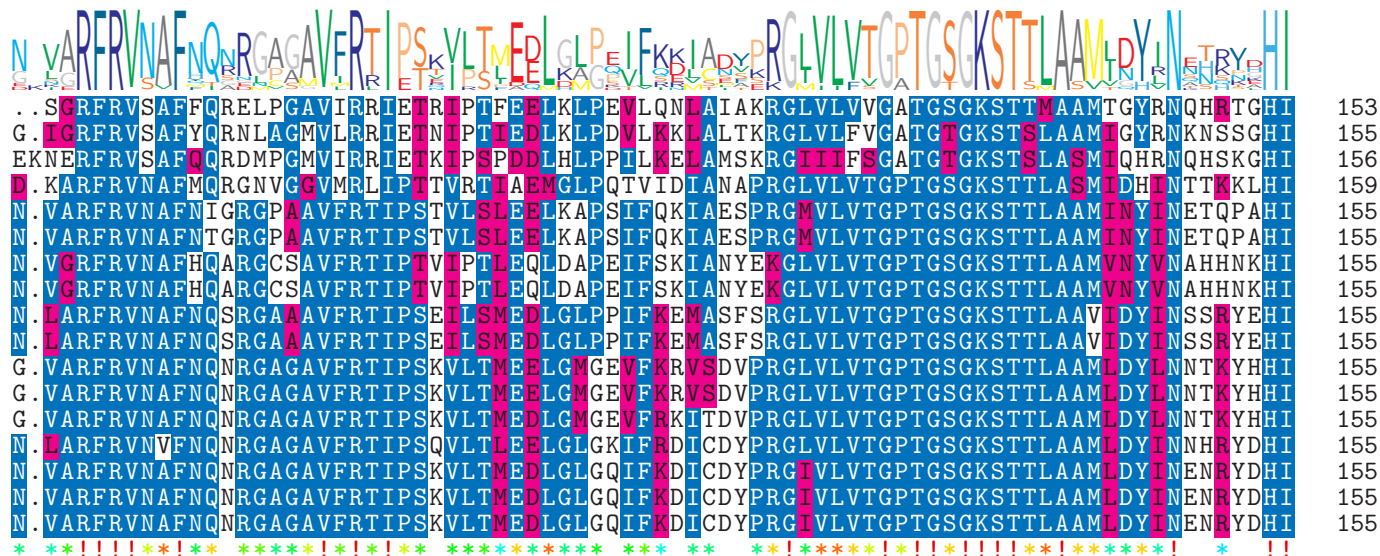
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consensus

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NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

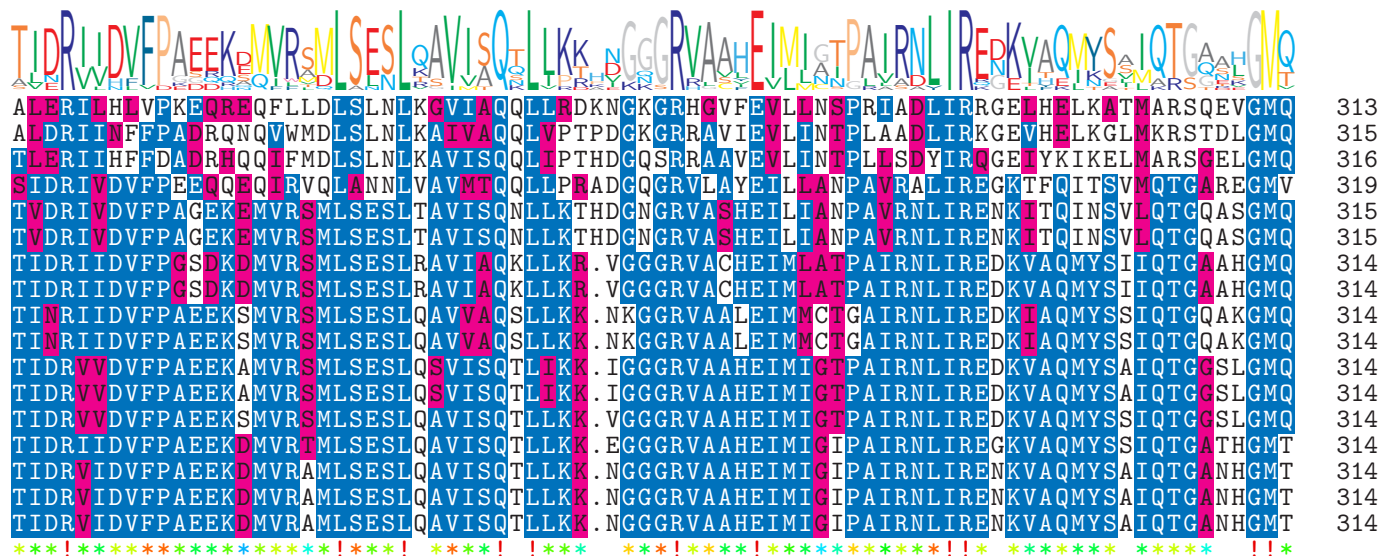
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NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
NTDB id 1081 FDQ49 RS05335 WP 006581387.1
consensus

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Sequence	Position
TFDQALYQLVDDKISEQDALHSADSANDLRLMLKTKRGDDY... GSGSLQNVKI..... DME.....	368
TFDQALYNLYVQGETIYEDALLHADSSNDLRLMIKLGSETDGEHLTSVS. QGLTL..... EMSDDDPGRSFR.....	381
TFDQALFDLYDHQHISYADALKHADSPNNLRLQIKLSEEANKPLDNRERMSEMER..... KMTFDGQRRNLK.....	383
TMDAYLAGLFRHLLITYDKLERAID.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	1203	PSJM300	01605	AFN76401.1		381	
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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
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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