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NTDB id 1019 ACIAD RS04205 WP 004922051.1
NTDB id 1307 DR RS10055 WP 010888596.1
NTDB id 1116 NGFR RS10615 WP 003690200.1
NTDB id 1086 NMV RS00245 WP 002218487.1
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 GC085 RS10305 WP 010947729.1
NTDB id 1194 PAKAF RS01995 WP 003084552.1
NTDB id 232212 CCZ28 RS19935 WP 140220535.1
NTDB id 1202 PSJM300 01600 AFN76400.1
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

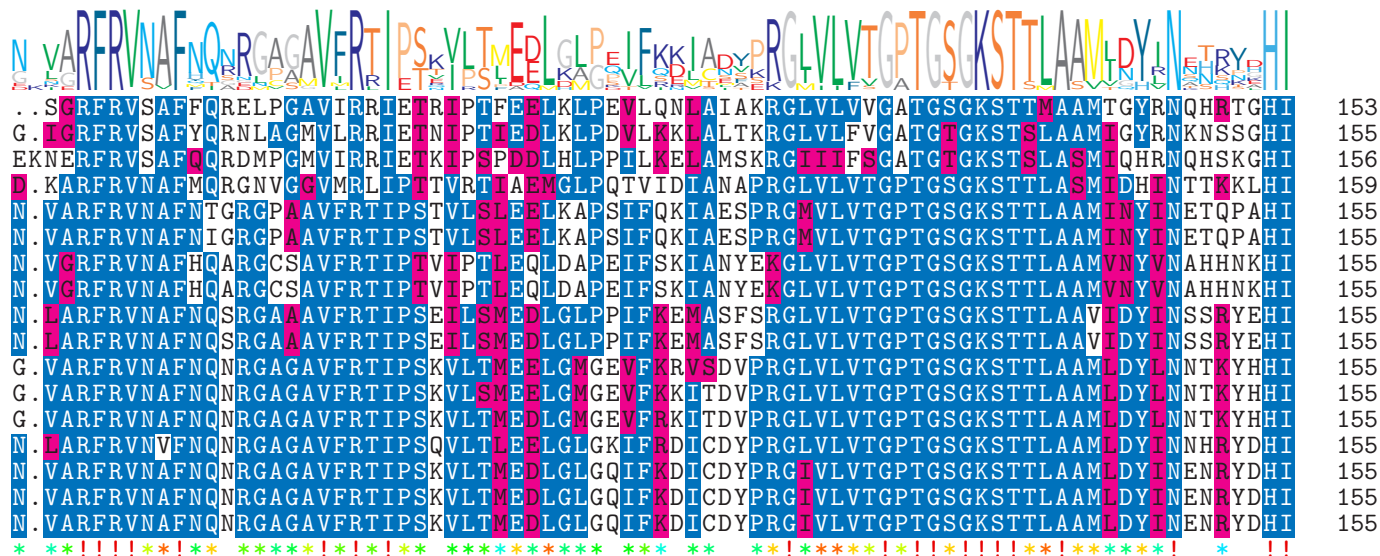
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NTDB id 1060 ABD1 RS04380 WP 000355489.1
NTDB id 1040 HON27 RS13360 WP 000355489.1
consensus

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Accession	Gene	Protein	Sequence	Score
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NTDB id 1283	ACE17W	RS02630 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAA	344
NTDB id 1259	GCO85	RS10305 WP 010947729.1	TLDQHLTELVNKNIISKHTAHEAA	344
NTDB id 1194	PAKAF	RS01995 WP 003084552.1	TLDMCLKGLVAKGLISRENAREKA	344
NTDB id 232212	CCZ28	RS19935 WP 140220535.1	TLDMCLKALLSKGLISRDAAREKA	344
NTDB id 1202	PSJM300	01600 AFN76400.1	TLDSCLKGLLAKGLISRDSAKEKA	344
NTDB id 1020	ACIAD	RS04210 WP 011182164.1	TLDQSLKQLLAKGLINLPTARAVA	344
NTDB id 1081	FDQ49	RS05335 WP 006581387.1	TLDQSLKGLVSRGVISPQTARTAA	345
NTDB id 1060	ABD1	RS04380 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAA	345
NTDB id 1040	HON27	RS13360 WP 000355489.1	TLDQSLKGLVARGVISPQTARTAA	345
consensus			! * * * * * ! * * * * * * * *	

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NTDB	id	1177	A1552VC	RS01165	WP	000422572.1		368
NTDB	id	1203	PSJM300	01605	AFN76401.1		381	
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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