

logo

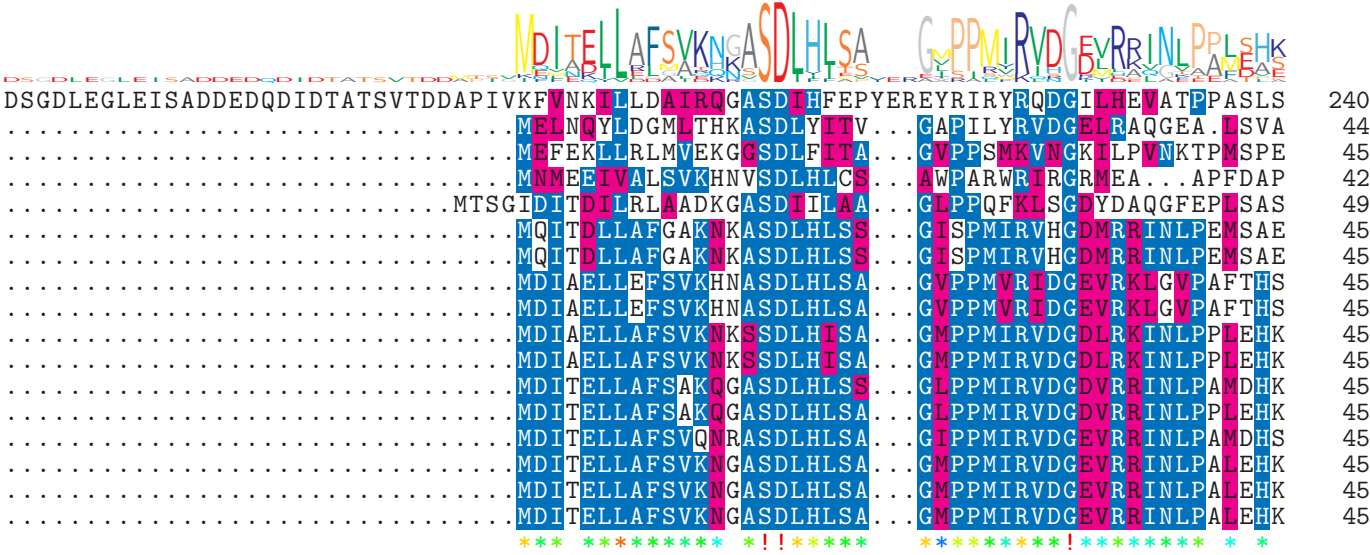
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NTDB id 1307 DR RS10055 WP 010888596.1		0
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NTDB id 1259 GC085 RS10305 WP 010947729.1		0
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1		0
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NTDB id 1040 HON27 RS13360 WP 000355489.1		0
consensus		

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NTDB id 1081 FDQ49 RS05335 WP 006581387.1		0
NTDB id 1060 ABD1 RS04380 WP 000355489.1		0
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consensus		

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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
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NTDB id	1202	PSJM300	01600	AFN	76400.1
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
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NTDB id	1177	A1552VC	RS01165	WP	000422572.1
NTDB id	1203	PSJM300	01605	AFN	76401.1
NTDB id	333735	C7V17	RS05890	WP	001324408.1
NTDB id	1307	DR	RS10055	WP	010888596.1
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NTDB id	1060	ABD1	RS04380	WP	000355489.1
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.....PDGKGRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ	GRRAVIEVLINTPLAADLIRKGEVHELKGLMKRSTDLGMQ
.....KQEGCVALFELLINTPAVGNLIREGKTHQLPHVITQTGQQVGMT	GCVALFELLINTPAVGNLIREGKTHQLPHVITQTGQQVGMT
.....ADGQGRVLAYEILLANPAVRALIREGKTFQITSVMTQTGAAREGMV	GRVLAYEILLANPAVRALIREGKTFQITSVMTQTGAAREGMV
.....HDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQ	GRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQ
.....HDGNGRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQ	GRVASHEILIANPAVRNLIRENKITQINSVLQTGQASGMQ
.....VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ	GGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ
.....VGGGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ	GGRVACHEIMLATPAIRNLIREDKVAQMYSIIQTGAAHGMQ
.....NKGGRVAAL EIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ	GGRVAAL EIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ
.....NKGGRVAAL EIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ	GGRVAAL EIMMCTGAIRNLIREDKIAQMYSIIQTGQAKGMQ
.....VGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLSGMQ	GGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLSGMQ
.....IGGGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLSGMQ	GGRVAAHEIMIGTPAIRNLIREDKVAQMYSIIQTGGSLSGMQ
.....EGGGRVAAHEIMIGIPAIRNLIREGKVAQMYSIIQTGATHGMT	GGRVAAHEIMIGIPAIRNLIREGKVAQMYSIIQTGATHGMT
.....NGGGRVAAHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGMT	GGRVAAHEIMIGIPAIRNLIRENKVAQMYSIIQTGANHGMT
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Sequence	Position
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TFQSFQQRVGEGR	326
TMDAYLAGLFRHLLITYDKGLERAIDPKEFARLANDASAGLGGASMTPAAPPVPAYNSASSSSTGRGGDFGRGAAAPATP	399
TMDQSLQSLVRQGLIAPEVARRRAQNSESMF	347
TMDQSLQSLVRQGLIAPEAARRRAQNSESMF	347
TMEQNAKQLIARGVVDAQEVQSKIETDLKAF	345
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TLDDHLTELVNKNIISKHTAHEAALNK.SLF	344
TLDDHLTELVNKNIISKHTAHEAALNK.SLF	344
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TLDMCLKGLVAKGLISRAREKAKIP.ENF	344
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TLDDSLKGLVSRGVISPQTARTAAKQP.ESFL	345
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NTDB id 1081 FDQ49 RS05335 WP 006581387.1		345
NTDB id 1060 ABD1 RS04380 WP 000355489.1		345
NTDB id 1040 HON27 RS13360 WP 000355489.1		345
consensus		

-  non conserved
-  similar
-  ≥ 50% conserved