

logo

NTDB id 173489 AZ909 RS06190 WP 002579657.1	MANIDA	EKLKATIESAMS	NIEKQFGKGSVMKLG	DH.NVTTMDAISTGCLD	LDIALGIGGV	PKGR	62					
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NTDB id 625 LCA RS02525 WP 011374196.1	MAKD	ERQAALDAALKKIEKN	FGKGSIMRMGEK	.VDTQVSTVSSGSLALD	EALGVGGY	PKGR	60					
NTDB id 1313 DR RS12030 WP 010888966.1	MSKDATKEISAPTD	AKERSKATETAMSQIEKA	FGKGSIMKLGA	E.SKLDVQVSTGSLSLD	LALGVGGI	PKGR	72					
NTDB id 1245 Cj1673c YP 002345041.1	MDDN	KRKSLDAALKSLDKT	FGKGTILRLGDK	.EVEQIDSIGTGSVGLD	LALGIGGV	PKGR	59					
NTDB id 1214 NCTC11637 00359 SQJ03206.1	MAIDE	DKQKATISLAIKQID	KVFGKGALVRLGDK	.QVEKIDAISTGSLGLD	LALGIGGV	PKGR	61					
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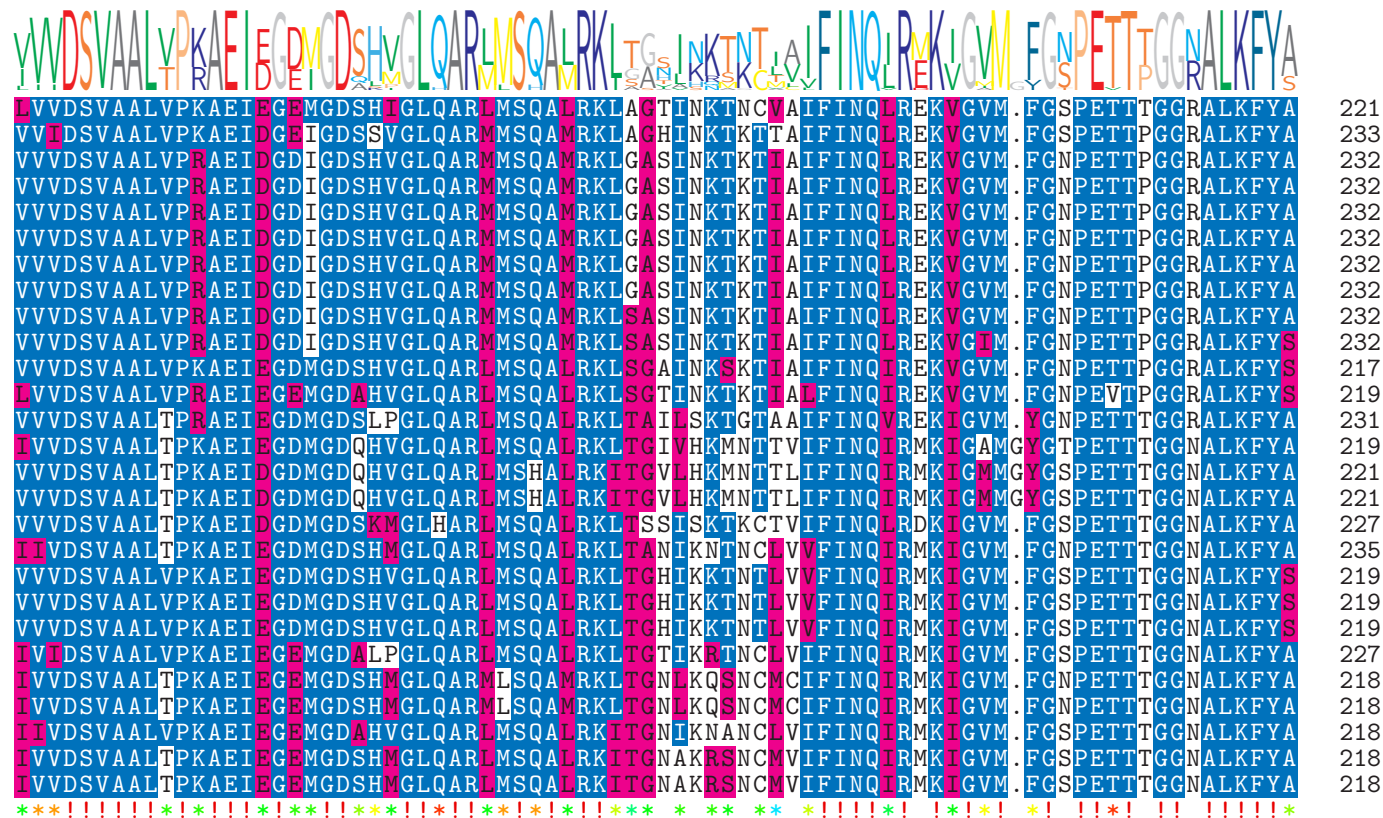
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NTDB id 194 SPR RS08825 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDA	153
NTDB id 159 SP RS09750 WP 001085462.1	IIEIYGPESSGKTTVALHAVAQAQKEGGIAAFIDA	153
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NTDB id 1214 NCTC11637 00359 SQJ03206.1	IIEIYGPESSGKTTVALHAVAQAQKAGGTC	141
NTDB id 1211 C694 RS00795 WP 000963128.1	IIEIYGPESSGKTTVALHAVAQAQKAGGTC	141
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consensus

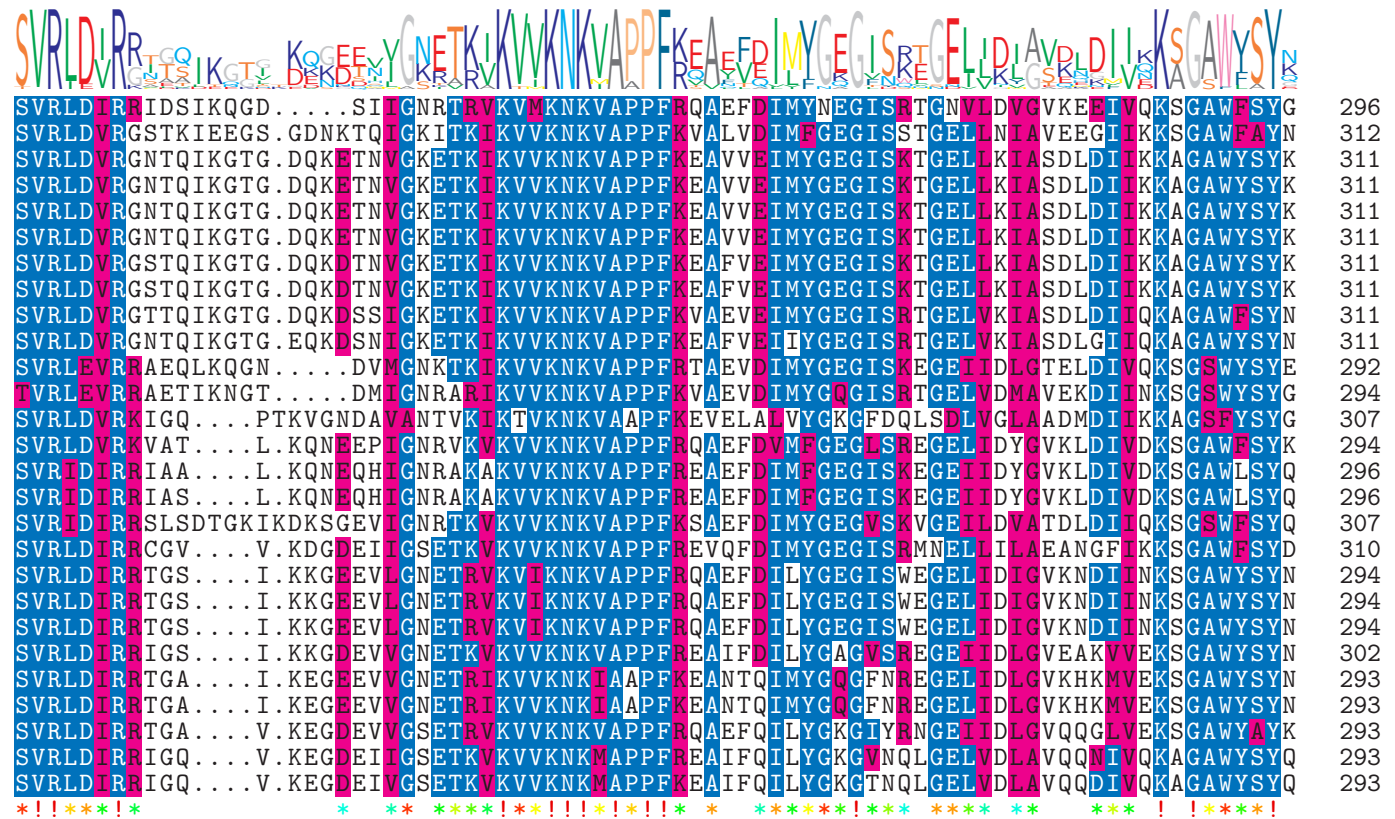
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consensus

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NTDB id 600 KW2 RS01765 WP 011675413.1	DEKIGQGAEKAKNYLKEHQDVFEIDHKKVRAAHGL	LDD	SEVAETEEETTAS	387
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NTDB id 228 SPD RS09265 WP 001085462.1	DEKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGL	IDG	EEVSEQDTENKK	384
NTDB id 194 SPR RS08825 WP 001085462.1	DEKIGQGSSENAKKYLAEHPEIFDEIDKQVRSKFGL	IDG	EEVSEQDTENKK	384
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NTDB id 527 SMSK321 RS11065 WP 001085454.1	DEKIGQGSSENAKKYLADNPEVFAEIDHQRVVRVQFGL	IDG	EEAAE...SKK	380
NTDB id 499 SM12261 RS08315 WP 078228268.1	DEKIGQGSSENAKKYLADNPEIFDEIDHQRVVRVQFGL	IDG	EEIAE...SKK	380
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NTDB id 1123 OK783 RS03845 WP 003688695.1	GAKIGQGGKDNVRVWLKENPEISDEIDAKIRALNGVEMHI	TEGT	...QDET	348
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NTDB id 1142 GTF74 RS11550 WP 000344154.1	GDKIGQGGKANACKYLLKENPEIAKTLDDKKLREMLLNPENM	QLI	AETSSAA...DD	354
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consensus	***!!!***** ** ***** ***** ** *		*	

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NTDB id 173489 AZ909 RS06190 WP 002579657.1		359
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved