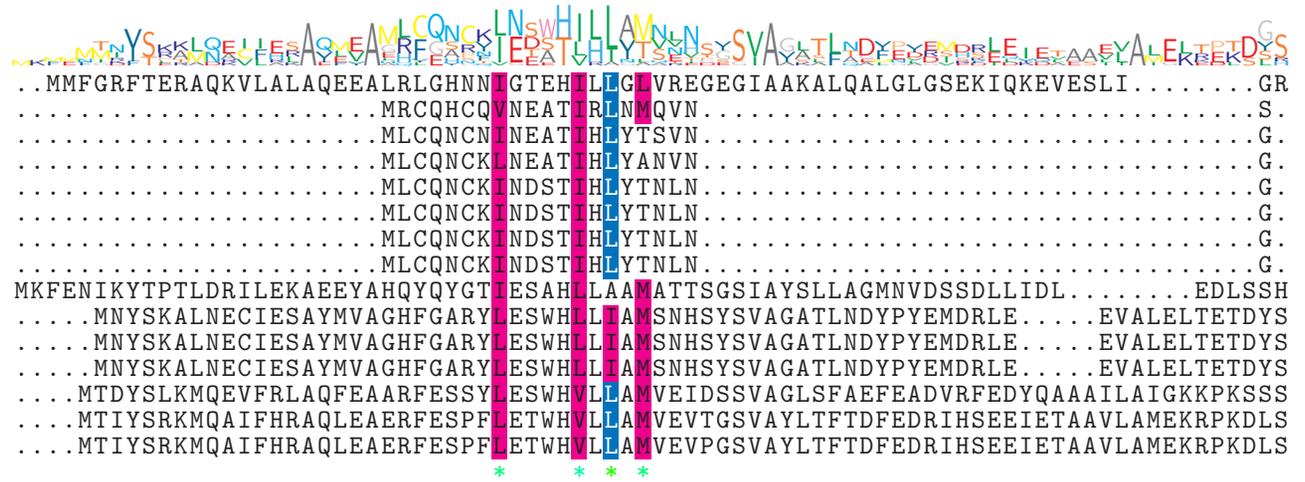


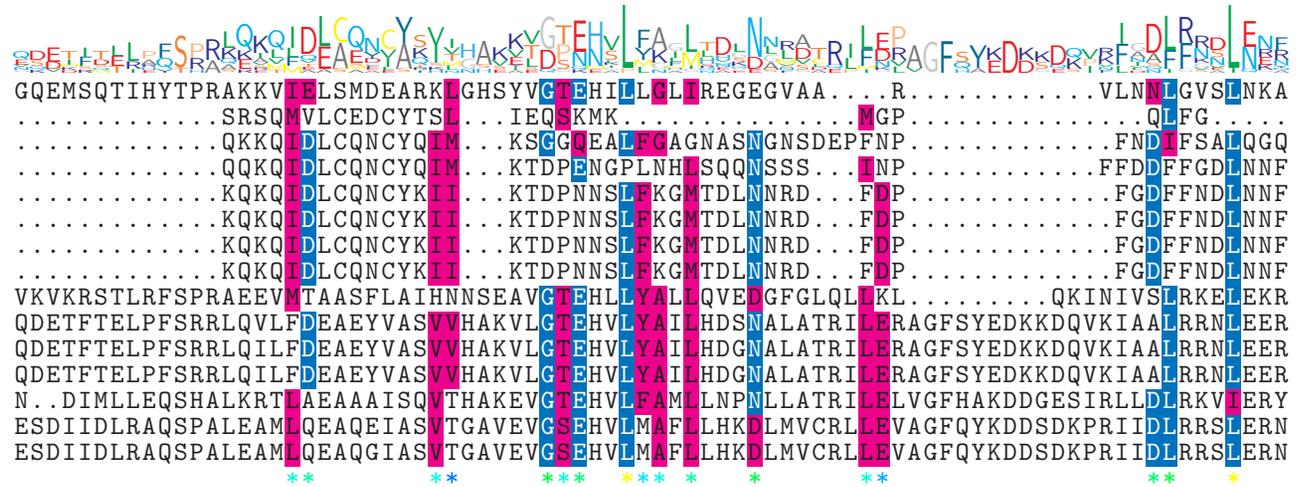
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 consensus



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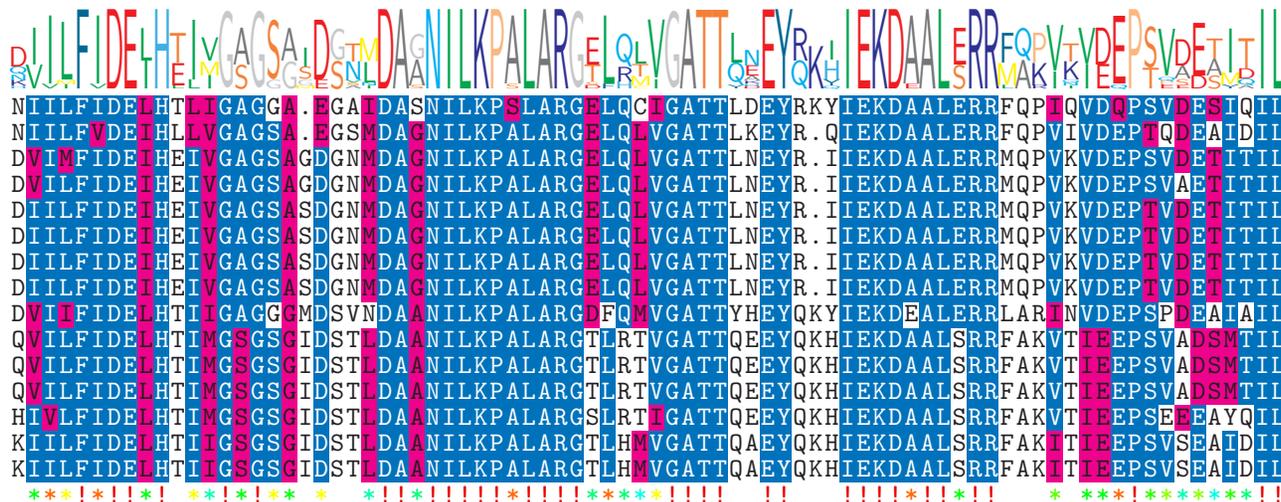
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 consensus



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consensus

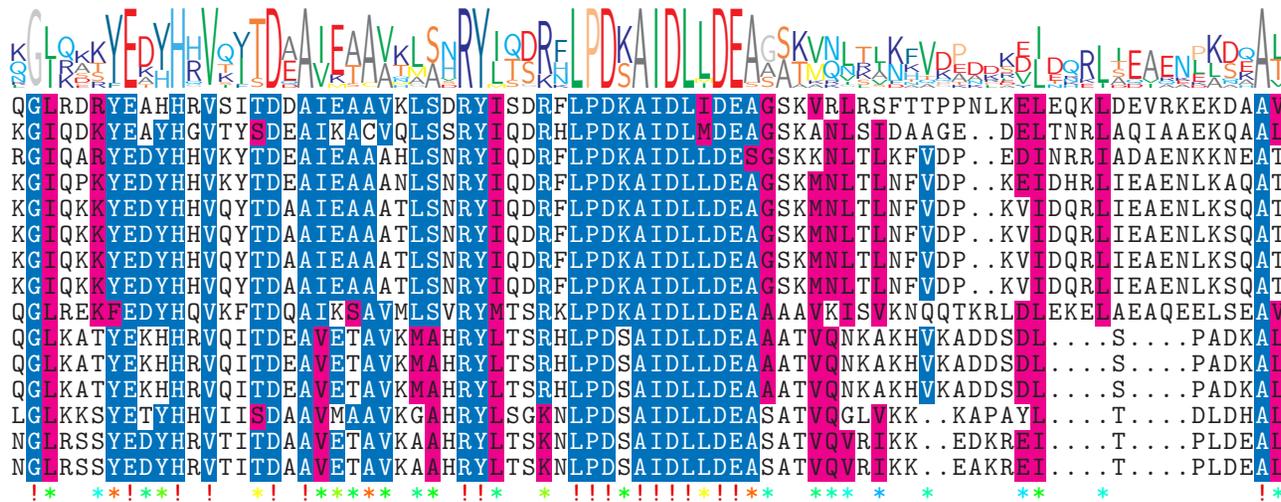


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consensus



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logo



NTDB id 85 BSU 00860 NP 387967.1	AEEGV	DLEYGARPLRRA	I	QKHVEDR	L	SEEL	LRGNIHK	Q	QHTIVL	D	VEDG	GEFVVKTTAKTN.....	810
NTDB id 985320 MHI08 RS06455 WP 342500091.1	AELGYHP	AFGARPLRRTIQ	E	HEVEDQ	M	TEILL	LEEEQLS	G	FTVDVE	.	HDEIVV	VKKG.....	700
NTDB id 601 KW2 RS02615 WP 021036805.1	VDLGYN	PAMGARPLRRTIQ	E	NTEDS	I	ADFYI	EHPEYKQLVADLI	.	DDKIVIS	N	QTQETAETTDEEVPAE		748
NTDB id 378 SMU RS02690 WP 002263569.1	VDLGYDPK	MGARPLRRTIQ	E	HEIEDA	I	TDYFL	EHPEQKELKAIMT	.	SNGKI	V	IKASNK	.VETVTKATSD.	753
NTDB id 281 KZH43 RS03550 WP 000882517.1	VDLGYDPK	MGARPLRRTIQ	D	YIEDT	I	TDYYL	ENPSEKDLKAVMT	.	SKGNI	Q	IKSACK	.AEVKSSEKEK.	752
NTDB id 245 SPD RS03860 WP 000882517.1	VDLGYDPK	MGARPLRRTIQ	D	YIEDT	I	TDYYL	ENPSEKDLKAVMT	.	SKGNI	Q	IKSACK	.AEVKSSEKEK.	752
NTDB id 243 SPR RS03635 WP 000882517.1	VDLGYDPK	MGARPLRRTIQ	D	YIEDT	I	TDYYL	ENPSEKDLKAVMT	.	SKGNI	Q	IKSACK	.AEVKSSEKEK.	752
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NTDB id 610 V4T04 RS10165 WP 012897346.1	AEVGF	DPEYGARPLR	K	ALQKEVEDL	L	SEQLL	SGEIKAGNHVS	I	GASNKK	I	KIAQIV.....		816
NTDB id 287 SP RS11210 WP 001109712.1	ANQGYD	PEMGARPLRRT	L	QTEVEDK	L	AELLL	KGDLVAGSTL	K	IGVKAGQL	K	FDIA.....		810
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NTDB id 377 SMU RS09275 WP 002262344.1	ALAGYD	AEMGARPLRRT	L	QTEVEDK	L	AELLL	SGKLASGQT	L	KIGIQKEK	L	KFDIV.....		813
NTDB id 297 STER RS00545 WP 011680614.1	AKDGYD	IEMGARPLRRT	I	QTVEDK	L	SELLL	GGQVVSQT	L	KIGCSKDK	L	TFTVV.....		816
NTDB id 329 STU RS10020 WP 011225298.1	AKDGYD	IEMGARPLRRT	I	QTVEDK	L	SELLL	GGQVVSQT	L	KIGCSKDK	L	TFTVV.....		816
consensus	**	!***	*!!!!!!***!	*!!	***	**		*	*	*	*	*	

- non conserved
- similar
- ≥ 50% conserved