

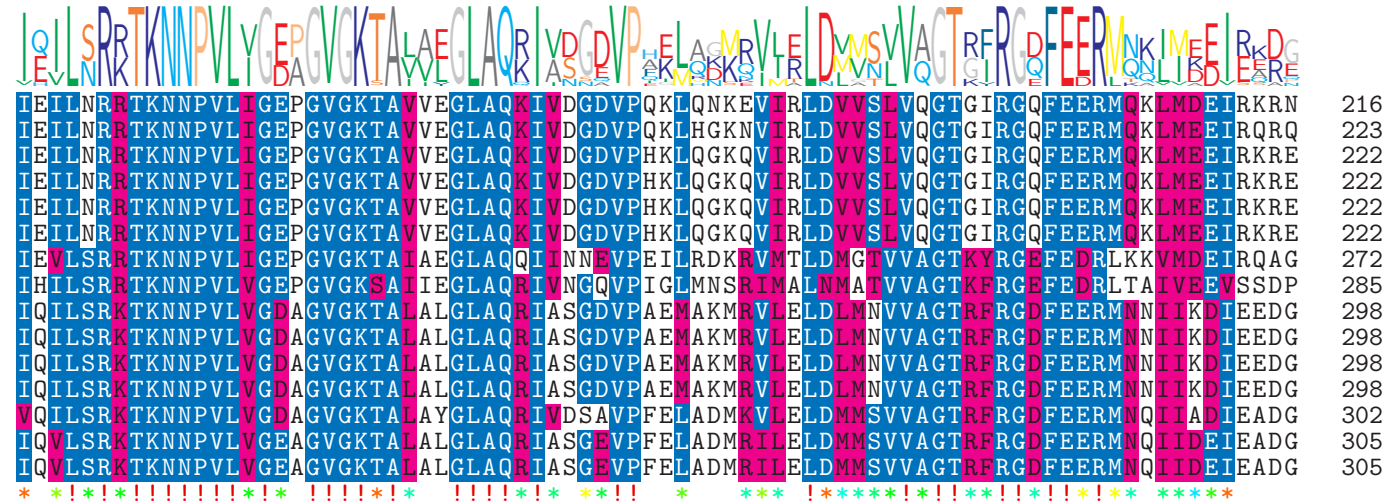
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NTDB id	378	SMU	RS02690	WP	002263569.1	MLCQNCKINEA.....	11
NTDB id	281	KZH43	RS03550	WP	000882517.1	MLCQNCKINDS.....	11
NTDB id	245	SPD	RS03860	WP	000882517.1	MLCQNCKINDS.....	11
NTDB id	243	SPR	RS03635	WP	000882517.1	MLCQNCKINDS.....	11
NTDB id	241	SP	RS04015	WP	000882523.1	MLCQNCKINDS.....	11
NTDB id	85	BSU	O0860	NP	387967.1	..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHLLCLVREGEGIAAKALQALGLGSEKIQKEVESLI.....GR	70
NTDB id	610	V4T04	RS10165	WP	012897346.1	MKFENIKYPTLDRILEKAEEYAHQYQYGTTESAHLLAAMATTSGSIAYSLLAGMNVDSDDLII.....DLEDLSSH	72
NTDB id	287	SP	RS11210	WP	001109712.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
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NTDB id	290	KZH43	RS10025	WP	001109677.1	MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHSYSVAGATLNDYPYEMDRLE....EVALELTETDYS	70
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NTDB id	377	SMU	RS09275	WP	002262344.1	MTDISLKMQEVRFLAQFEAAERFESSYLESWHLLAMVEIDSSVAGLSFAEFEADVRFEDYQAAAAILAIGKKPKSSS	76
NTDB id	297	STER	RS00545	WP	011680614.1	MTIYSRKMQAIFHRAQLEAERFESPFIETWHVLLAMEVETGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
NTDB id	329	STU	RS10020	WP	011225298.1	MTIYSRKMQAIFHRAQLEAERFESPFIETWHVLLAMEVEVPGSVAYLTFDTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
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NTDB	id	297	STER	RS00545	WP	011680614.1	ESDIIDLRAQSPALEAMQEAQEIASVTGAVEV	GSEHVLMAFLHLKDLMVCRILEVAGFQYKDDSD	KPRIIDRRSLERN	156
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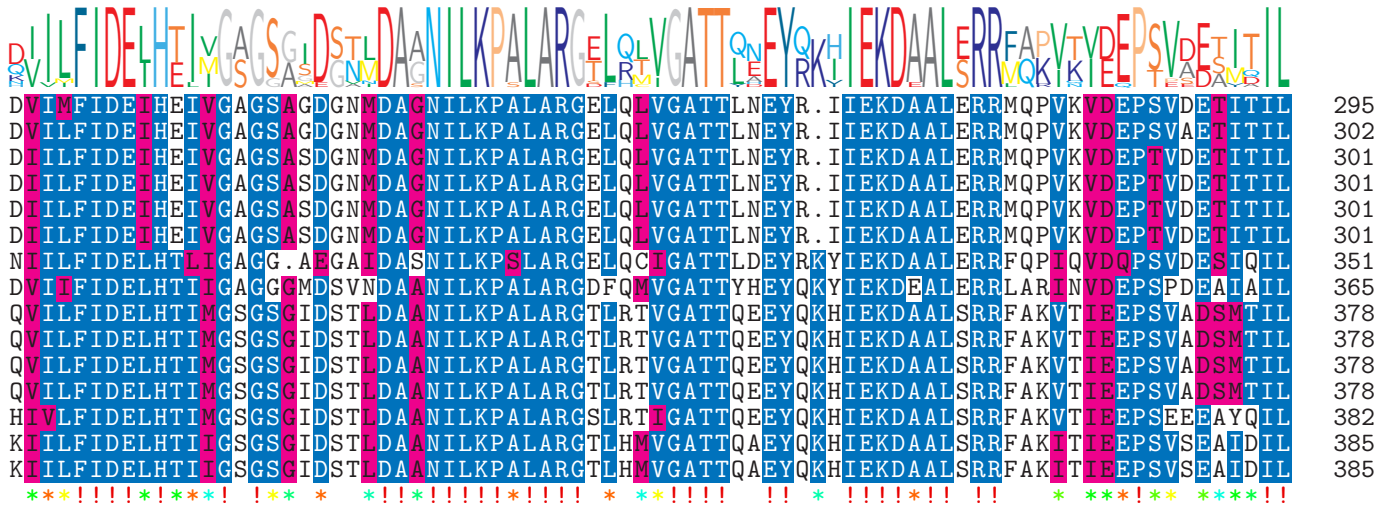
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NTDB id 1152078 MMH06 RS02950 WP 001109705.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
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NTDB id 329 STU RS10020 WP 011225298.1
consensus

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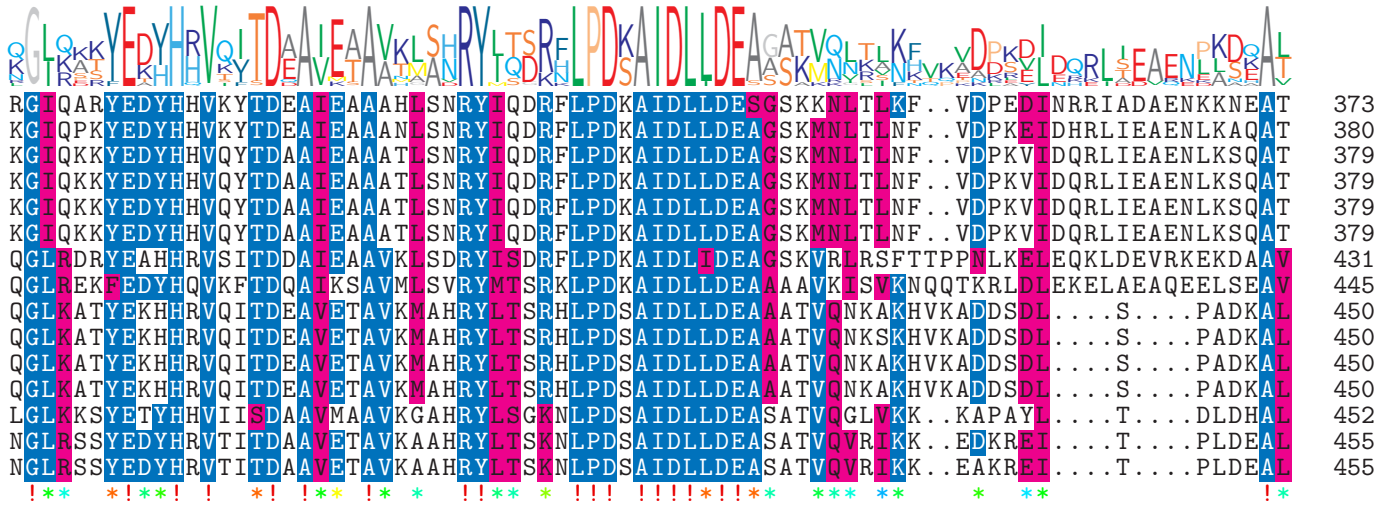
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