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NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 287 SP RS11210 WP 001109712.1
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
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NTDB id 245 SPD RS03860 WP 000882517.1
NTDB id 243 SPR RS03635 WP 000882517.1
NTDB id 241 SP RS04015 WP 000882523.1
consensus

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consensus



Sequence	Position
MKFENIKYTPPTLDRILEKAE EYAHQYQYGTIESAHLAAMATTSGSIAYSLLAGMNV DSSDLLIDL.....EDLSSH	72
....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSYVAGATLNDYPYEMDRLE....EVALELTETDYS	70
....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSYVAGATLNDYPYEMDRLE....EVALELTETDYS	70
....MNYSKALNECIESAYMVAGHFGARYLESWHLLIAMSNHYSYVAGATLNDYPYEMDRLE....EVALELTETDYS	70
...MTDYSLKMQEVFRLAQFEAARFESSYLESWHVLAMVEIDSSVAGLSFAEF EADVFEDYQAAAILAIGKKPKSSS	76
...MTIYSRKMQAIFHRAQLEAERFESPFL ETWHVLLAMVEVPGSVAYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
...MTIYSRKMQAIFHRAQLEAERFESPFL ETWHVLLAMVEVTGVSAYLTFTDFEDRIHSEEIETA AVLAMEKRPKDLS	76
..MMFGRFTERAQKVLALAQEEALRLGHNNIGTEHILLGLVREGEGIAAKALQALGLGSEKIQKEVESLI.....GR	70
.....MRCQHCQVNEATIRLNMQVN.....S.	21
.....MLCQNCNINEATIHLYTSVN.....G.	21
.....MLCQNCNINEATIHLYANVN.....G.	21
.....MLCQNCNINEATIHLYTNLN.....G.	21
.....MLCQNCNINEATIHLYTNLN.....G.	21
.....MLCQNCNINEATIHLYTNLN.....G.	21
.....MLCQNCNINEATIHLYTNLN.....G.	21

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consensus

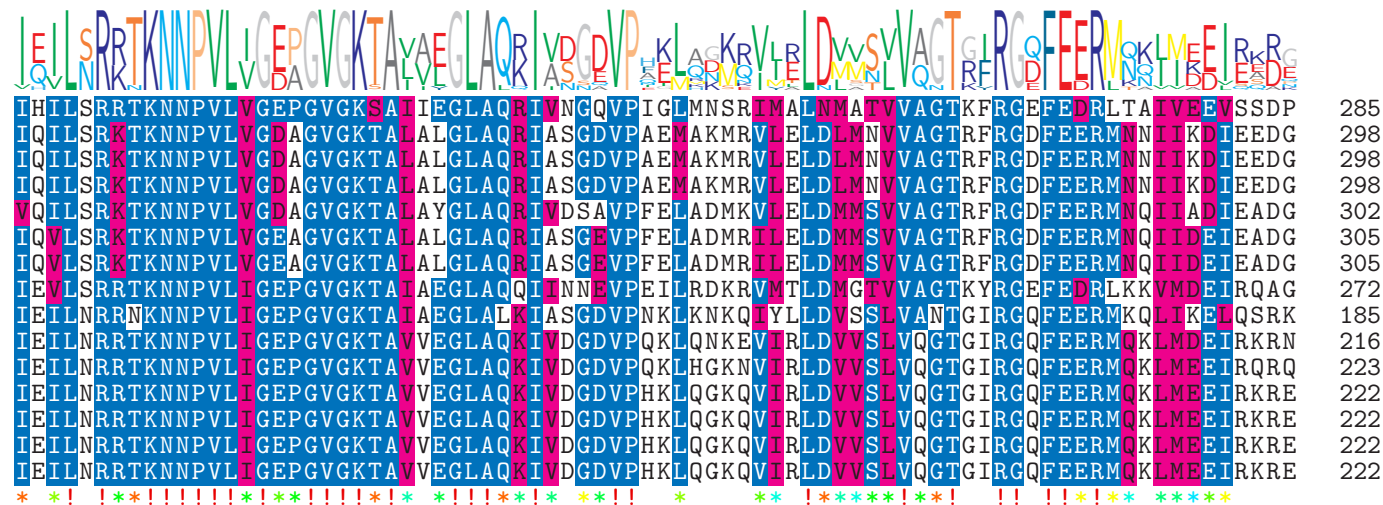
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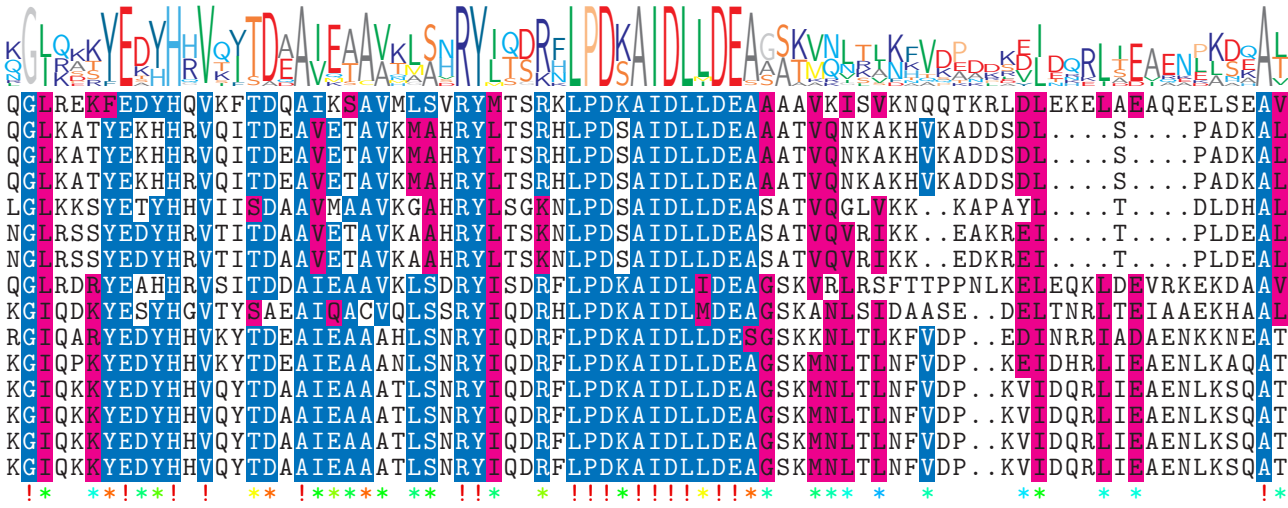
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consensus					



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NTDB id	610	V4T04	RS10165	WP	012897346.1
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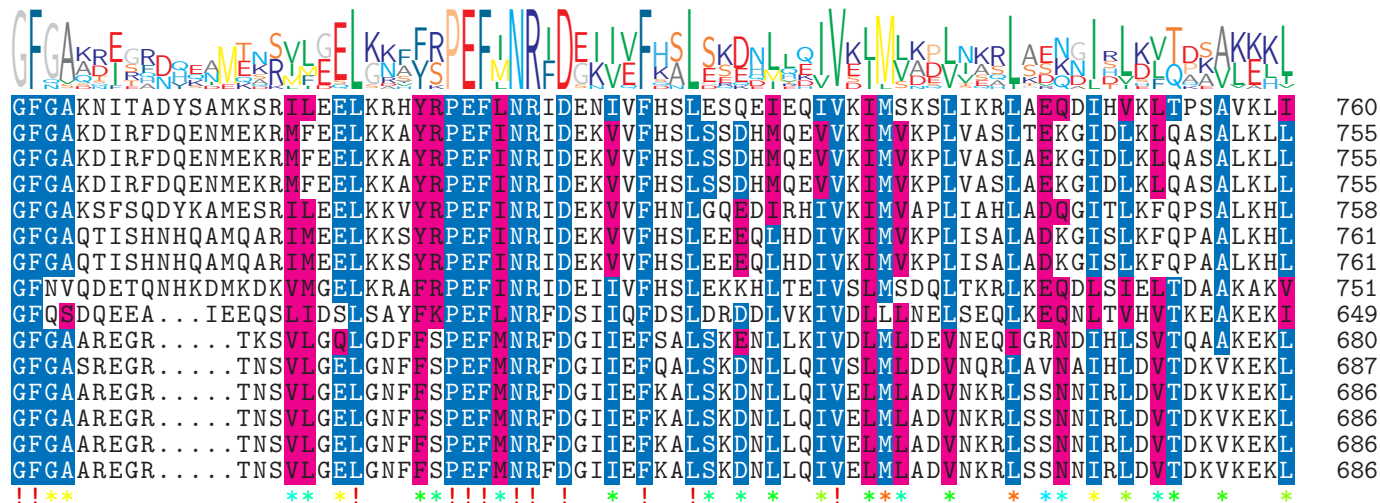


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