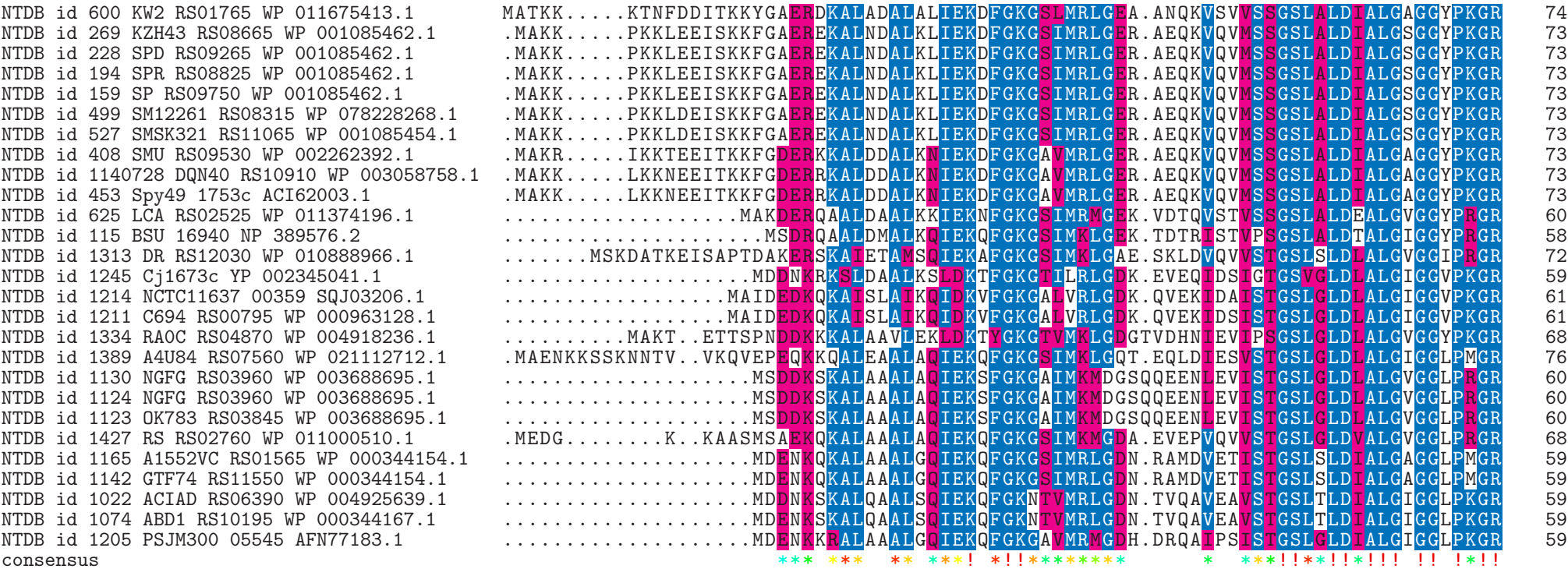


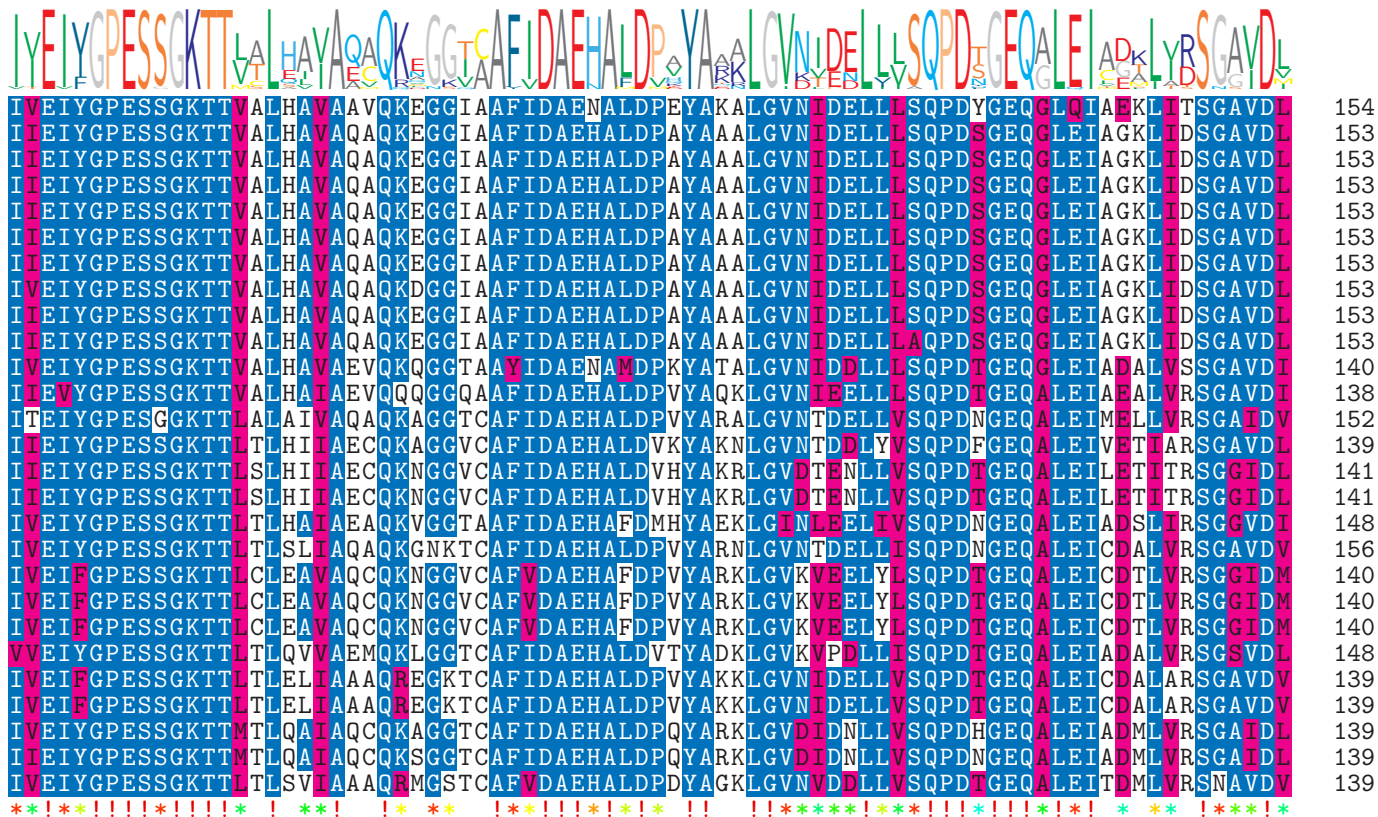
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NTDB id 1142 GTF74 RS11550 WP 000344154.1
NTDB id 1022 ACIAD RS06390 WP 004925639.1
NTDB id 1074 ABD1 RS10195 WP 000344167.1
NTDB id 1205 PSJM300 05545 AFN77183.1
consensus

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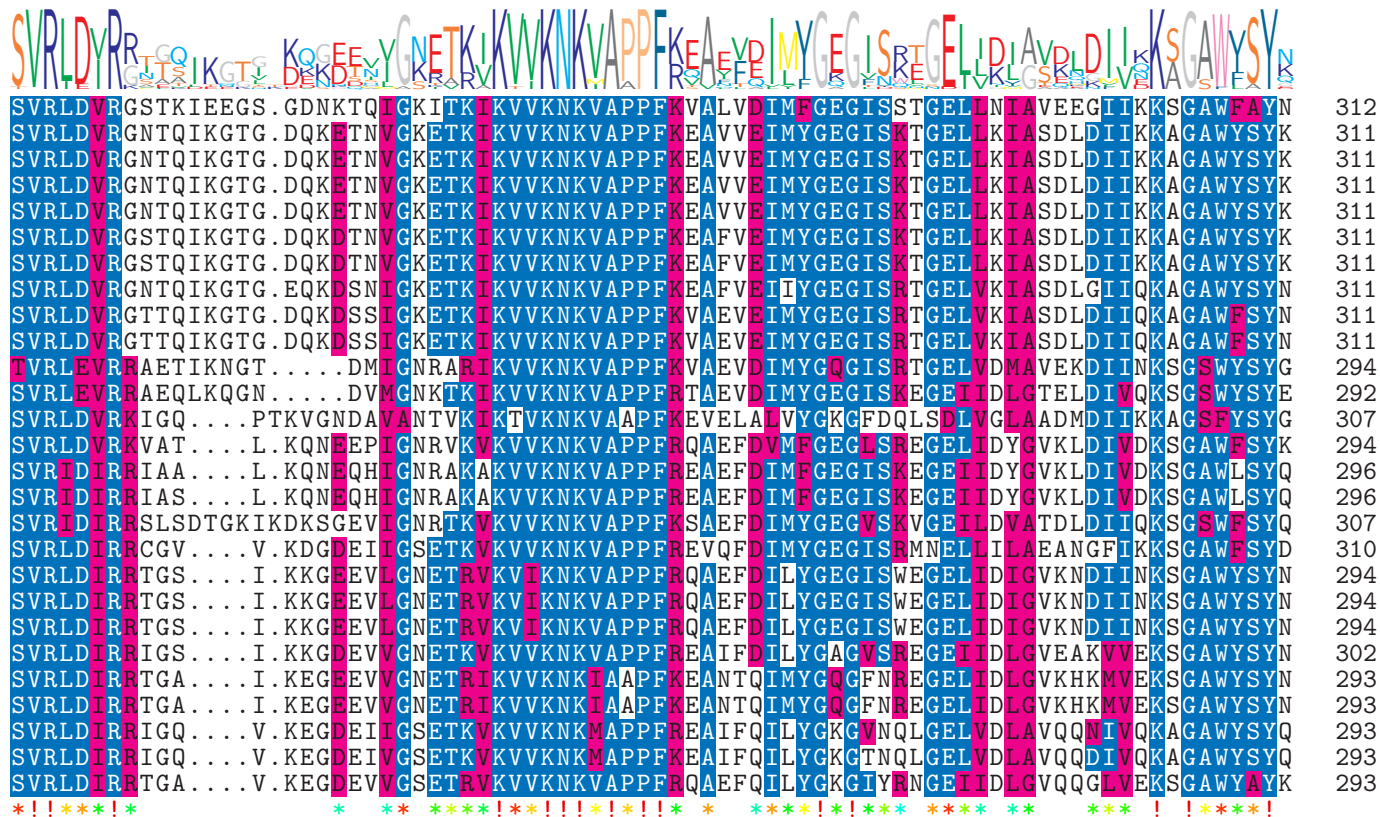
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NTDB id 269 KZH43 RS08665 WP 001085462.1	VVVDS	VAAALVPRAEIDGDI	GDSHVGLQARMSQAMRKL	GASINKTKTTAIFINQLREKVGVM	.FGNPETTPGGRALKFYA 232
NTDB id 228 SPD RS09265 WP 001085462.1	VVVDS	VAAALVPRAEIDGDI	GDSHVGLQARMSQAMRKL	GASINKTKTTAIFINQLREKVGVM	.FGNPETTPGGRALKFYA 232
NTDB id 194 SPR RS08825 WP 001085462.1	VVVDS	VAAALVPRAEIDGDI	GDSHVGLQARMSQAMRKL	GASINKTKTTAIFINQLREKVGVM	.FGNPETTPGGRALKFYA 232
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NTDB id 1245 Cj1673c YP 002345041.1	IVVDS	VAAALTPKAEIEGDMGDQHVGLQARLMSQALRKLTG	IVHKMNTTVIFINQIRMKIGAMGY	YGPETTTGGNALKFYA	219
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NTDB id 1211 C694 RS00795 WP 000963128.1	VVVDS	VAAALTPKAEIDGDMGDQHVGLQARLMSHALRKITGV	LHKMNTTLIFINQIRMKIGMNGY	CSPETTTGGNALKFYA	221
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NTDB id 1205 PSJM300 05545 AFN77183.1	IIVDS	VAAALVPKAEIEGEMGDAHVGLQARLMSQALRKITGN	IKNANCLVIFINQIRMKIGVM	.FGSPETTTGGNALKFYA	218
consensus		***!!!!!!*!*!!!!*!***!!!!*!!!!*!!!!*!***	* ** *		

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NTDB id 600 KW2 RS01765 WP 011675413.1
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NTDB id 194 SPR RS08825 WP 001085462.1
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consensus

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NTDB id 1245 Cj1673c YP 002345041.1	DKKIGQGS	ENAKSKA	ELKENPEI	ADEIT	KATQNSMC	TEGMI		SGSEDD..EGEE	343	
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NTDB id 1165 A1552VC RS01565 WP 000344154.1	GDKIGQGS	ANACKY	LKENPEI	AKTLD	KKLREML	LNPNEM		QLIAETSSAAD	353	
NTDB id 1142 GTF74 RS11550 WP 000344154.1	GDKIGQGS	ANACKY	LKENPEI	AKTLD	KKLREML	LNPNEM		QLIAETSSAAD	353	
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NTDB id 1074 ABD1 RS10195 WP 000344167.1	GNKIGQGS	NNVIRY	FEENN	QIA	FEIERN	IREQLLTTGTN		GAVQ.I.EDEE	349	
NTDB id 1205 PSJM300 05545 AFN77183.1	GSKIGQGS	ANAAKY	LKENPEI	GQIE	QQIRAKLL	TSAN		TKATPVAEDLAD	347	
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NTDB id 1205 PSJM300 05545 AFN77183.1			347
consensus			

- non conserved
- similar
- ≥ 50% conserved