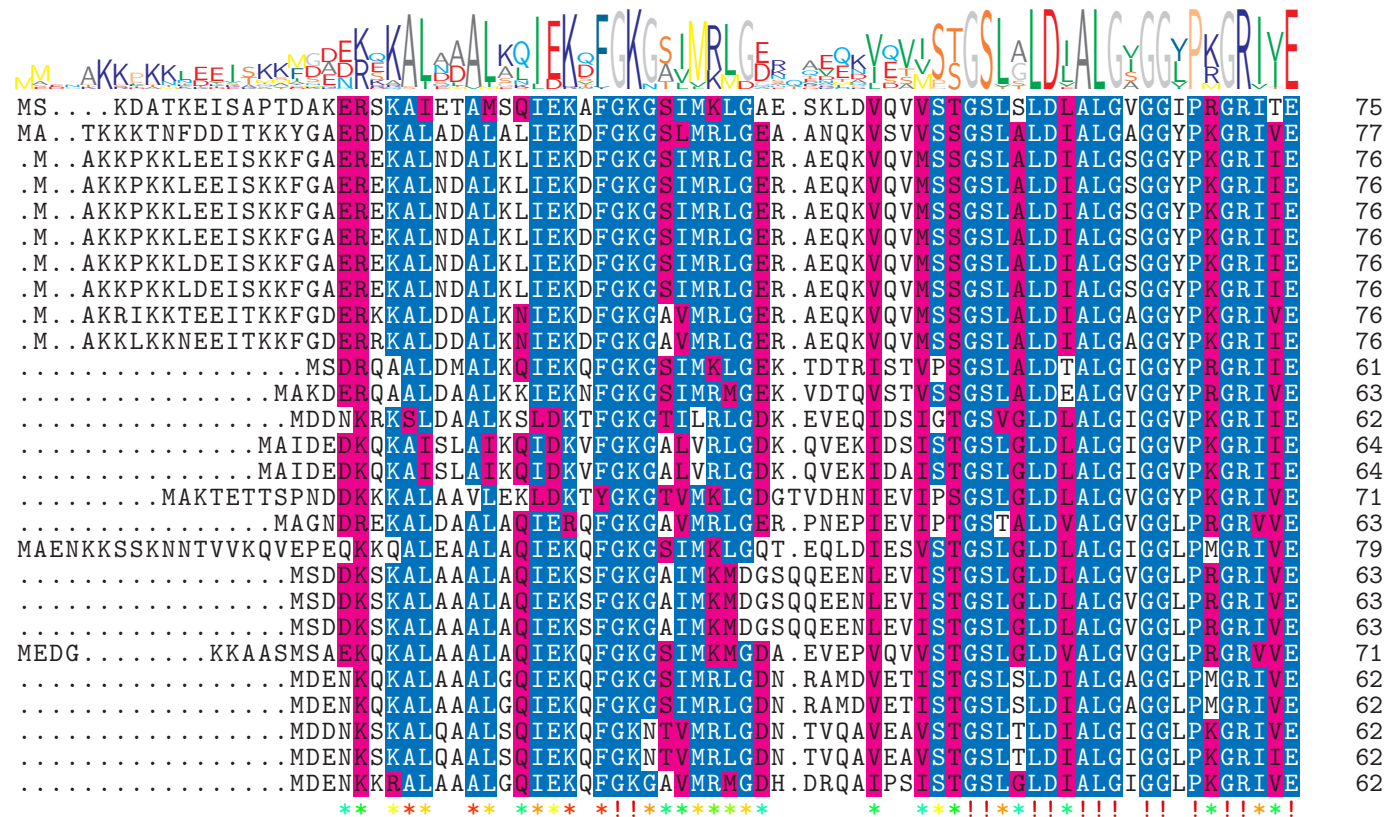


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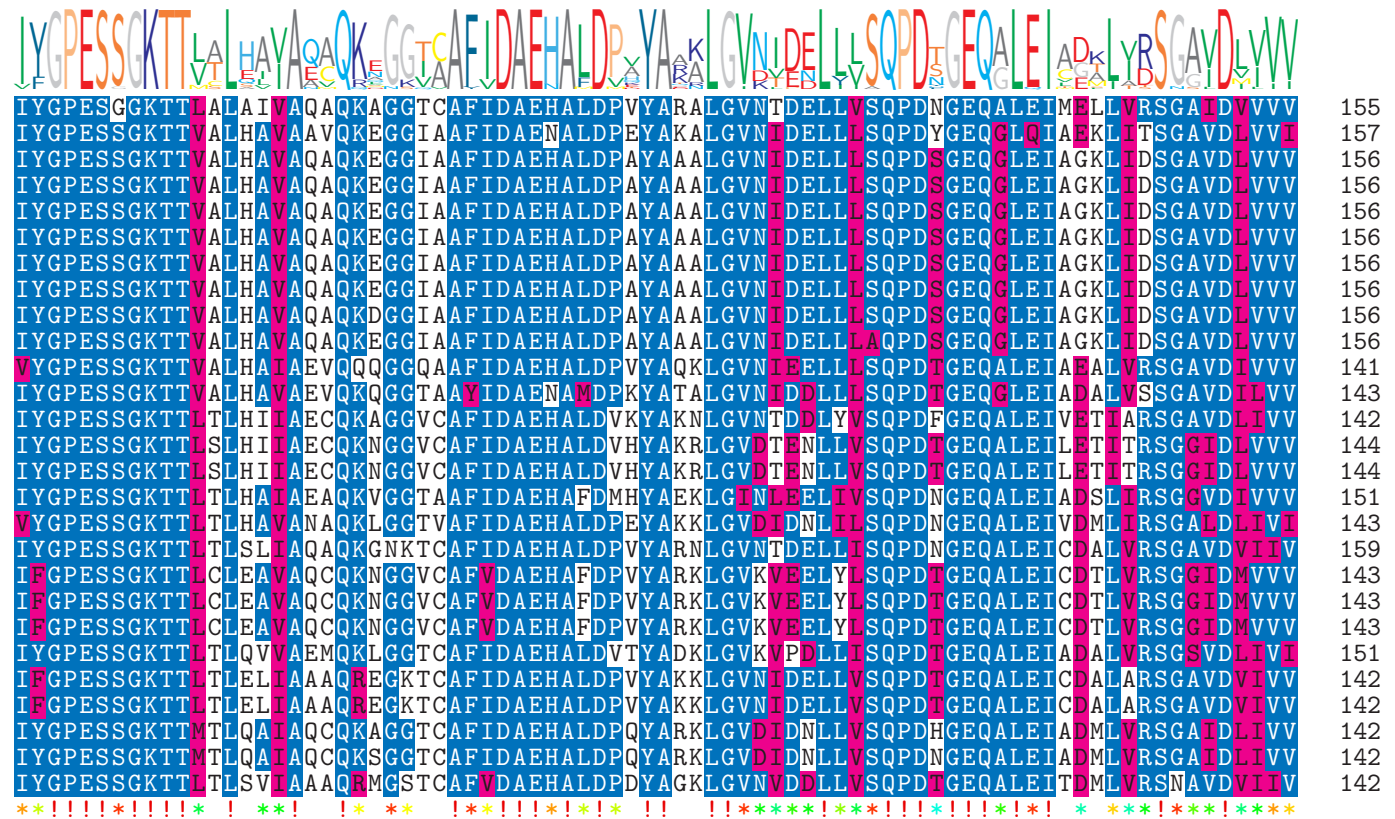
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consensus

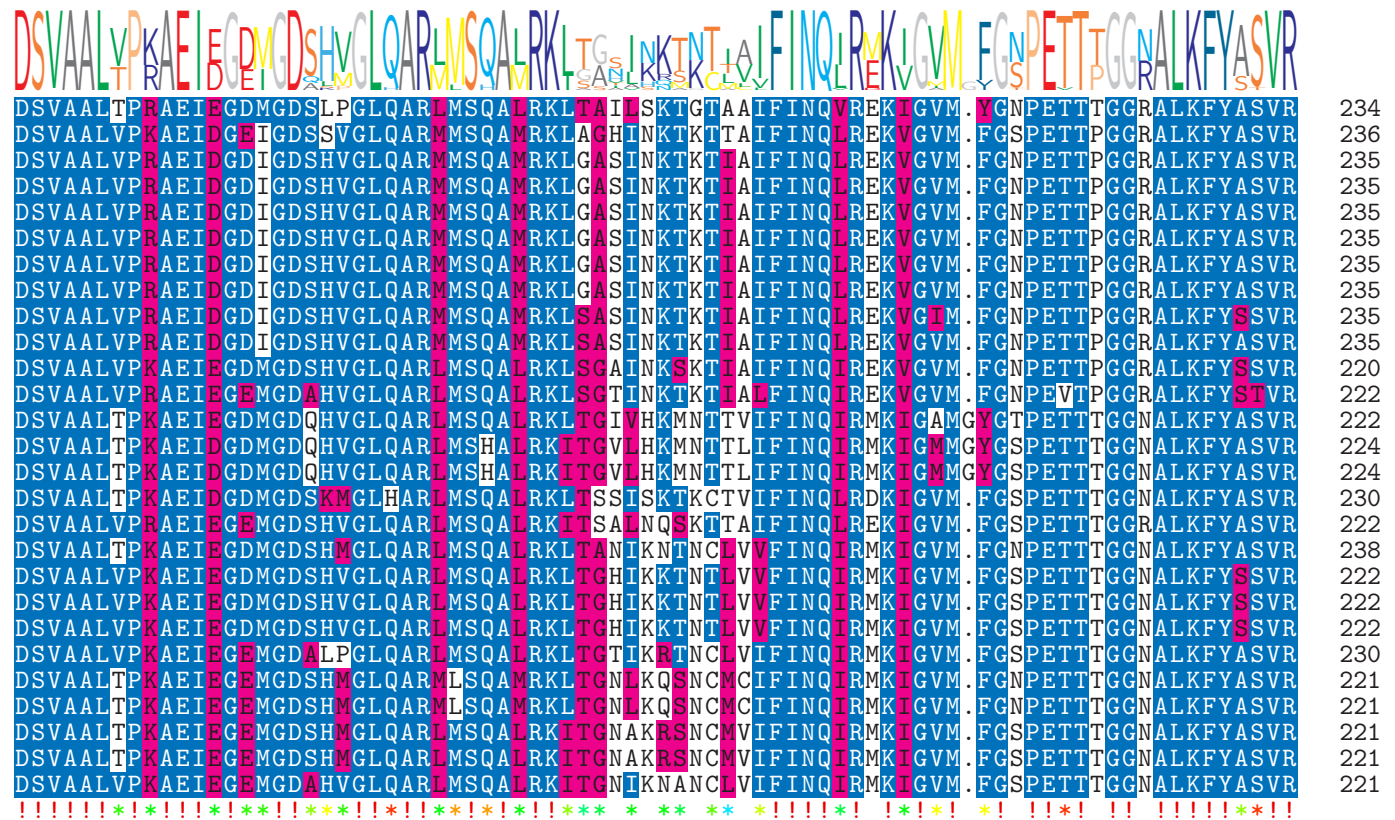
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consensus

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NTDB id 228 SPD RS09265 WP 001085462.1	LDVRGNTQIKGTG.DQKETNVGKETKIKVVKNKVAPPFKEAVVEIMYGEGISKTGELLKIASDLDIKKAGAWYSYKDEK	314
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NTDB id 115 BSU 16940 NP 389576.2	LEVRRRAETIKNGT.....DMIGNRARIKVVKNKVAPPFKVAEVDIMYGGGISRTGELVDMAVEKDIINKSGSWYSYGSER	297
NTDB id 625 LCA RS02525 WP 011374196.1	LDVRKVVAT....L.KQNEEPIGNRVKVVKNKVAPPFRQAEDVDMFGEGLSREGELIDYGVKLDIVDKSGAWFSYKDKK	297
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NTDB id 1313 DR RS12030 WP 010888966.1	IGQGKEKTIAYIAERPMEQEIRDRVMAAIRAGNAGEAPALAPAPAAPE.....AAEA.....	363
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NTDB id 1130 NGFG RS03960 WP 003688695.1	IGQGKDNVRVWLKENPEISDEIDAKIRALNGVEMHITEGTQDE.....TDGER.....PEE.....	348
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consensus			

-  non conserved
-  similar
-  ≥ 50% conserved