

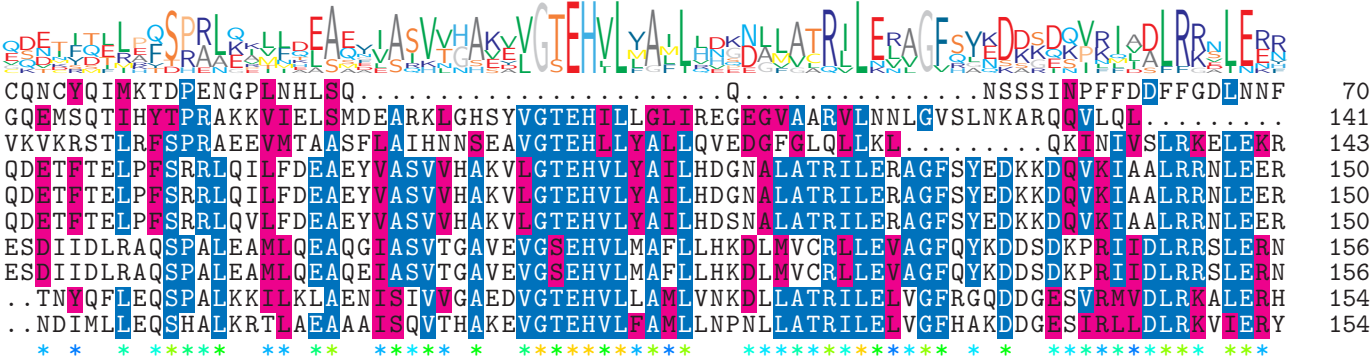
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NTDB id 85 BSU 00860 NP 387967.1
NTDB id 610 V4T04 RS10165 WP 012897346.1
NTDB id 290 KZH43 RS10025 WP 001109677.1
NTDB id 289 SPD RS10700 WP 001109677.1
NTDB id 287 SP RS11210 WP 001109712.1
NTDB id 329 STU RS10020 WP 011225298.1
NTDB id 297 STER RS00545 WP 011680614.1
NTDB id 154811 AMR84 RS09415 WP 000020286.1
NTDB id 377 SMU RS09275 WP 002262344.1
consensus



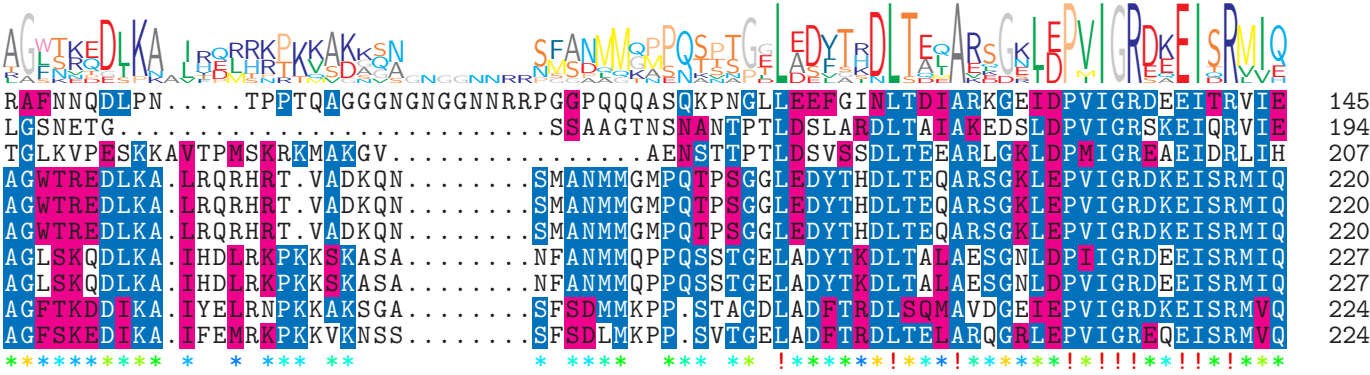
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ILFIDELHTIILICAGGMDSVNDAANILKPALARGDFQMGVATTYHIEYQKYIEKDEALERRLARINVDQPSVDEAIAILQG	367
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Sequence	Position
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LKATYEEKHHRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLIDEAATVQNKAKHVKAADSDS...PADKALMD	452
LKATYEEKHHRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLIDEAATVQNKAKHVKAADSDS...PADKALMD	452
LKATYEEKHHRVQITDEAVETAVKMAHRYLTSRHLPDSAIDLIDEAATVQNKAKHVKAADSDS...PADKALMD	452
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LKPAYEAFHNVTISDEAVMTAVKVAHRYLTSRKNLPDSAILDLDEASATVQVRIKK...NAPSLL	454
LKKSRYETYYHVIISDAAVMAVKGMAHRYLTSRKNLPDSAILDLDEASATVQGLVKK...KAPAYL	454

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consensus

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NTDB id 378 SMU RS02690 WP 002263569.1
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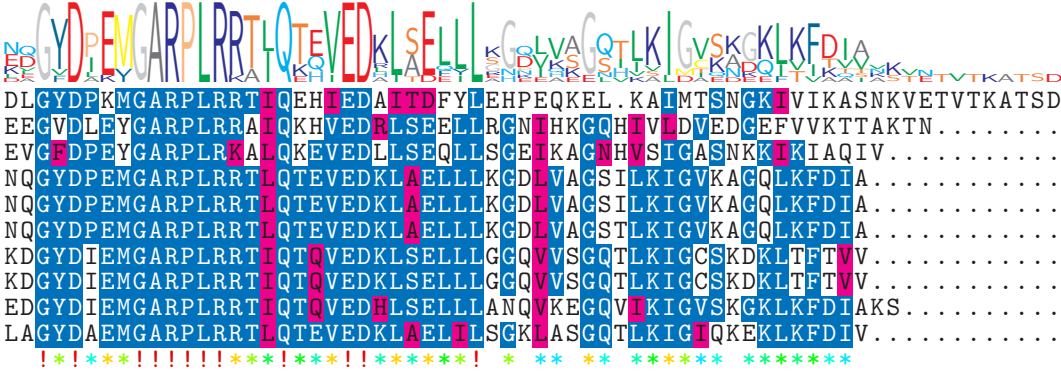
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688
752
761
756
756
756
762
762
759
759

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consensus



753
810
816
810
810
810
816
816
815
813

[X] non conserved
[X] similar
[X] ≥ 50% conserved