

logo

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|----------------|--------------------------------|--|----|
| | | MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGSLAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK | 80 |
| NTDB id 1028 | TT RS08215 WP 011173991.1 | .MALSIGDRRLGAILLDQGYLGDNDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL | 79 |
| NTDB id 1306 | DR RS10060 WP 027479822.1 | | 0 |
| NTDB id 1110 | NGFG RS09235 WP 003689817.1 | | 0 |
| NTDB id 1168 | A1552VC RS11070 WP 000957200.1 | | 0 |
| NTDB id 1293 | VP RS12240 WP 005479695.1 | | 0 |
| NTDB id 1401 | DSB67 RS12665 WP 010643259.1 | | 0 |
| NTDB id 1251 | GC085 RS07725 WP 011946523.1 | | 0 |
| NTDB id 221583 | B5U27 RS04575 WP 005745482.1 | | 0 |
| NTDB id 1058 | ABD1 RS01615 WP 001274986.1 | | 0 |
| NTDB id 1018 | ACIAD RS01685 WP 004920473.1 | | 0 |
| consensus | | | |

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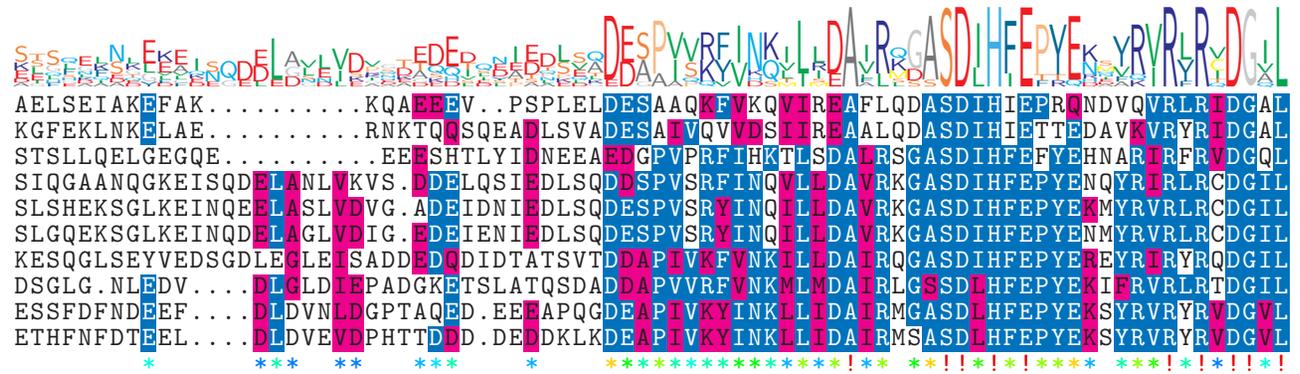
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|----------------|--------------------------------|--|-----|
| | | ALLPAEKAKELKAIPFALDEEAGVVRVAFLNPLDPLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLLEIYVPSGA | 158 |
| NTDB id 1028 | TT RS08215 WP 011173991.1 | RSIRPQTALNLQAFPPFALE..GDRLRVALVDPLSSFSIETLEDDSGFDIEPYQALREEVMWAIATHYPELGLLEIIVPSGA | 157 |
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| NTDB id 1058 | ABD1 RS01615 WP 001274986.1 | | 0 |
| NTDB id 1018 | ACIAD RS01685 WP 004920473.1 | | 0 |
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| | | SGEGQKDLKLGELLQKGWISREALEEALVEQEKTDGDLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVDPDPSAA | 237 |
| NTDB id 1028 | TT RS08215 WP 011173991.1 | SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL | 237 |
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| NTDB id 1168 | A1552VC RS11070 WP 000957200.1 | | 0 |
| NTDB id 1293 | VP RS12240 WP 005479695.1 | | 0 |
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| NTDB id 221583 | B5U27 RS04575 WP 005745482.1 | | 0 |
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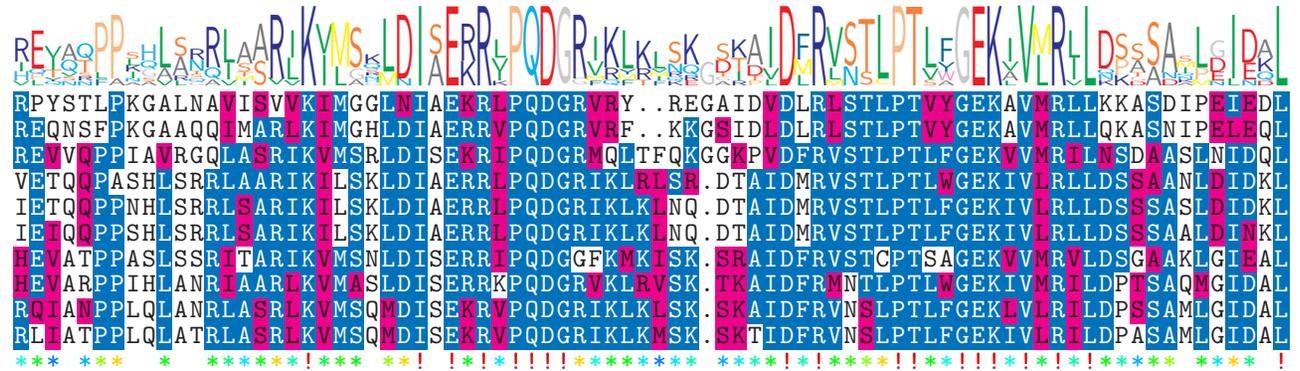
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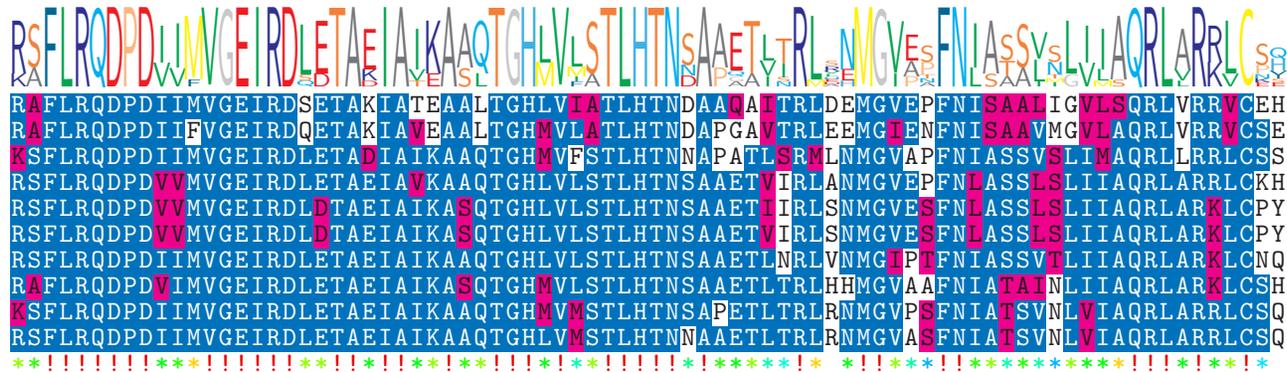
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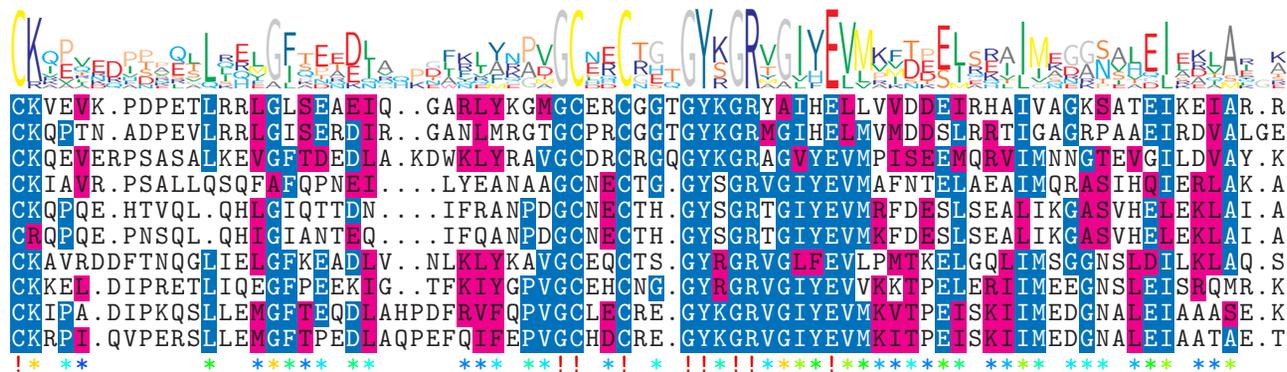
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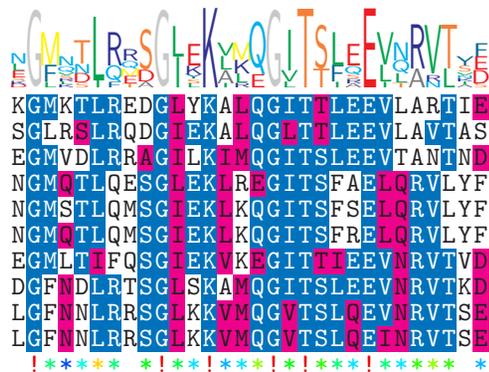
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non conserved
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 ≥ 50% conserved