

logo

NTDB id 1028 TT RS08215 WP 011173991.1	MSVLTIGDKRLGAALLDAGLLTDEELQRALERHREVGGS LAEVLVDMGLLSERRIAQTIEDRFGIPLVELHRVEIPPKVK	80
NTDB id 1306 DR RS10060 WP 027479822.1	.MALSIGDRRLGAILLDQGYLGDNLDLQRALERHSEVGGRLADVLIDSGMVGEKRIARAIEEALGIPLVNLAVQPDPAAAL	79
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 95668 NFHSH190041 RS17605 WP 261923012.1	0
NTDB id 1018 ACIAD RS01685 WP 004920473.1	0
NTDB id 1058 ABD1 RS01615 WP 001274986.1	0
consensus		

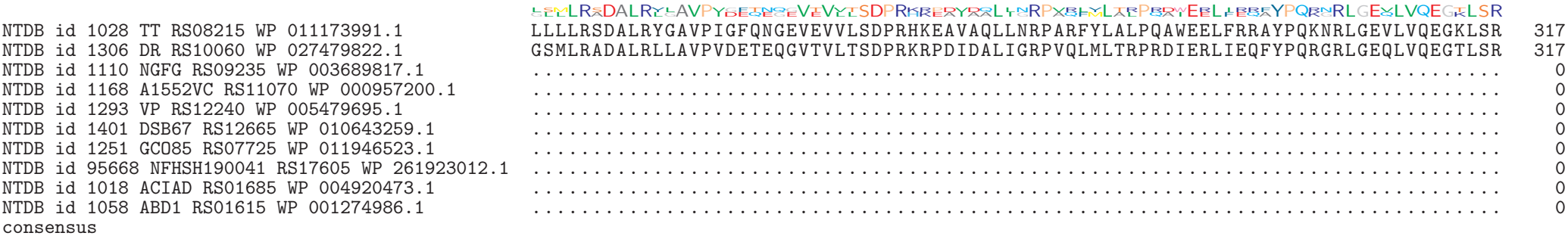
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NTDB id 1028 TT RS08215 WP 011173991.1	RSIRPQTALNLQAFPFALDEEAGVVRVAFNLPLDLSLEEVEDLTGLVVEPYQTTKSAFLYALAKHYPELGLPVPPP..P	158
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NTDB id 1168 A1552VC RS11070 WP 000957200.1	0
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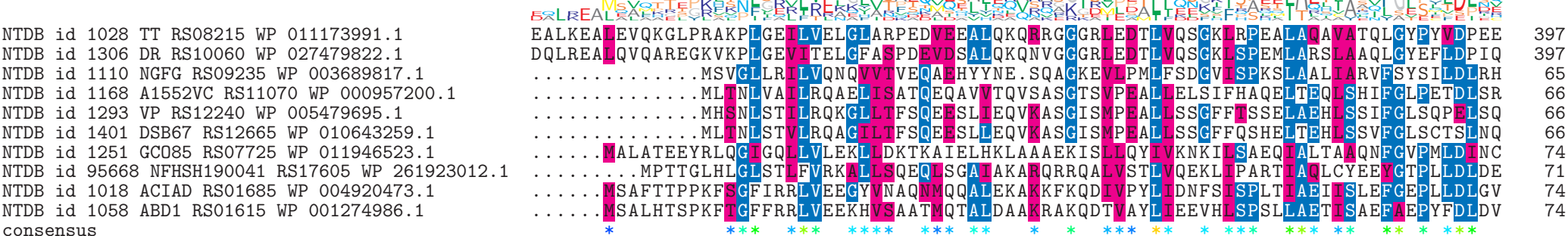
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NTDB id 1028 TT RS08215 WP 011173991.1	SGEGQKDLKLGELLQKGWISREALEEALVEQEKTGDLLGRILV.RKGLPEEALYRALAEQKGLEFLESTEGIVPDPSSAA	237
NTDB id 1306 DR RS10060 WP 027479822.1	SDAGRTGGKLGRLITHGYITDAQLQVALDAQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSPGFHPGEEVL	237
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NTDB id 1251 GC085 RS07725 WP 011946523.1	0
NTDB id 95668 NFHSH190041 RS17605 WP 261923012.1	0
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consensus		

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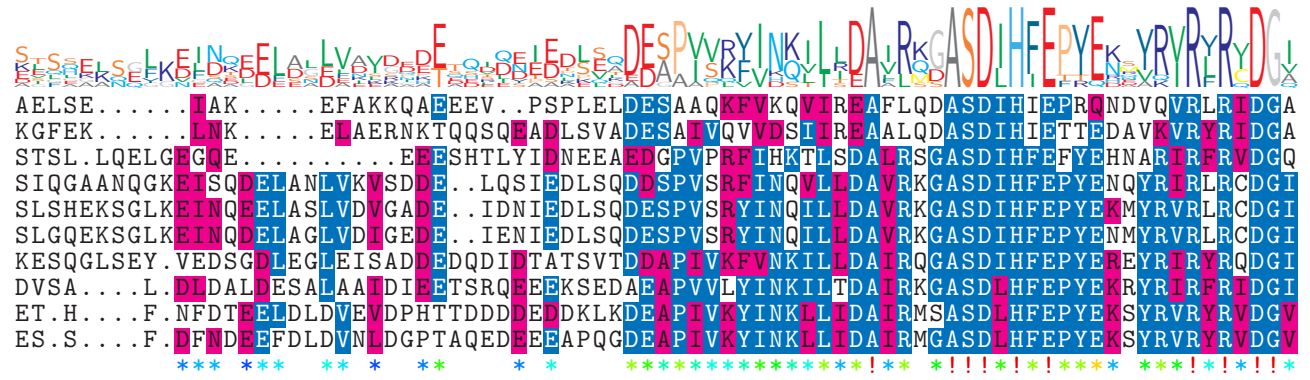


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 consensus



544
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209
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228
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 consensus



622
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289
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307
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 consensus



702
699
369
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387
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logo

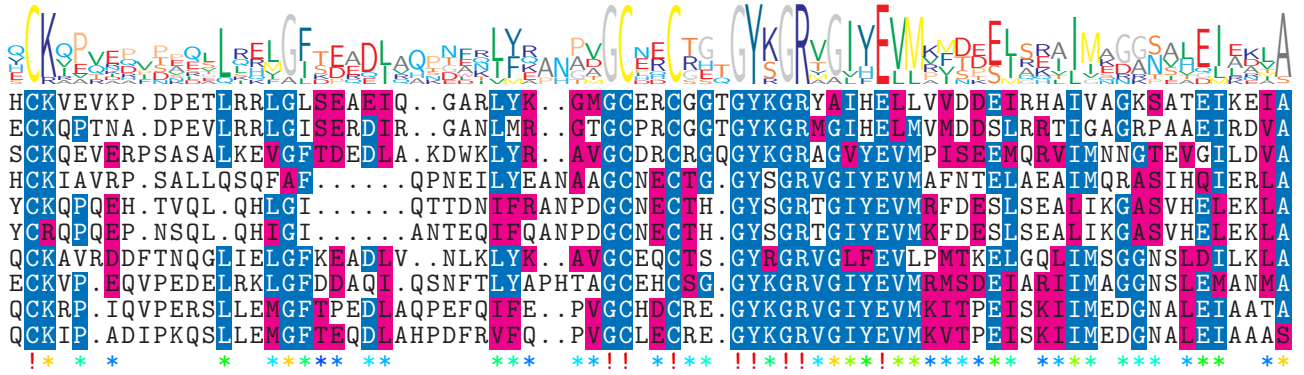
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 consensus



782
779
449
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467
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logo

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 consensus



857
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538

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 consensus



889
887
558
562
561
561
574
569
570
570

X non conserved
 X similar
 X ≥ 50% conserved