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NTDB id 1436	BW25113 RS00530 WP 001025145.1		0
NTDB id 1168	A1552VC RS11070 WP 000957200.1		0
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consensus			

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consensus			

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NTDB	id 1306	DR RS10060 WP 027479822.1	SDAGRTGGKLGGERLITHGYITDAQLQVALDAQQQTGEALGATLISQRAITEDQLYEVLAEQEGTTFLPNPSGFHPGEEVL	237	
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consensus					

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consensus

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 MSQYT ICEQQ TERIF 15
 MQYS VCDMN SQQVF 14
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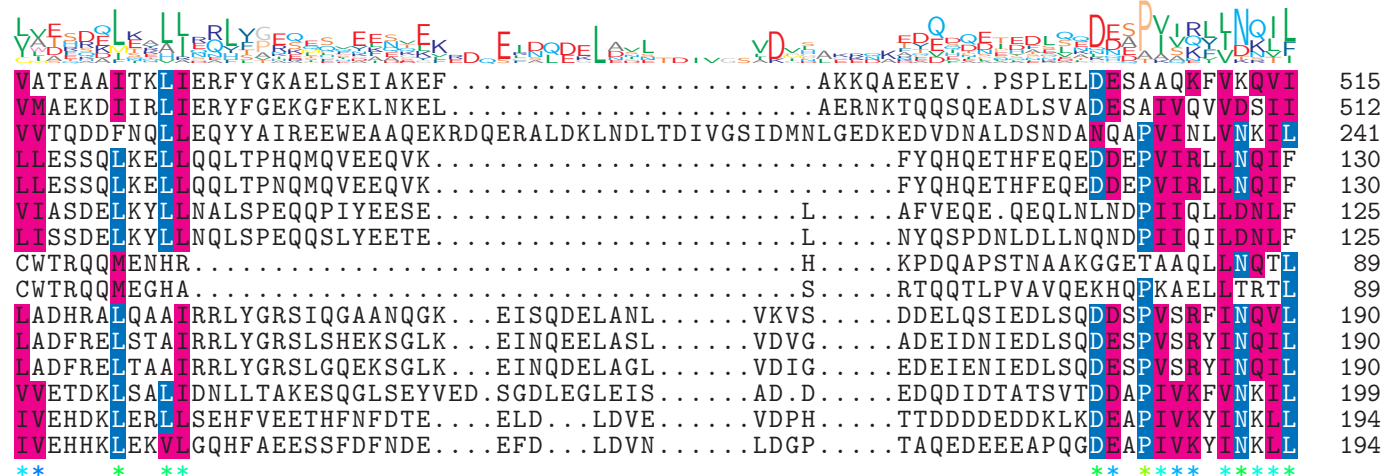
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GYEFLDPIQN PPDPKVALMPEATARRYVVPVLRQG NSLVAMKDPNRVFAFLDKLITG KEILPA	1455	1455
GVDSVDPEVAPIATQQMAELIGRFFPLDTCRRFKFLPLAQEGEDPPSVLVAWVDPDNLAQDEINRILRVKG FEIRRL	161	161
. TI SPDLWERNQQQSSLLRLRYFALPLKEEN NRLWLGVDSLSNLSACETIAFI TG KPVEPI	79	79
. TI SPDLWERNQQQSSLLRLRYFALPLKEEN NRLWLGVDSLSNLSACETIAFI TG KPVEPI	79	79
. DI SPSRWQKNCEEKQLILRYLAVPVQETE HKLWLAVDDENNITACEIFAFM TH KQIEPV	74	74
. DI SESLWQKNCEDEKHLILRYLAIPQIEND DTLWLAIIDMNNLSACEIFAFI AH KTIEPV	73	73
. MKDAQLNTLCQRHQAVLINSAS NSITVAVVDAPSHALLDALHFA TQ KQIDIV	52	52
. MNIPQLTALCLRYHGVLIDASE EVVHVAVDAPSHELLDALHFA TT KRIEIT	52	52
GLPETDLTRY DYANLCQQLGLRELITRYDAIPAKQG NLLLLAVSDPTLQAEEFRFA TG LQVELA	124	124
GLSQPELSQY EYASLCQQLGLRELITRHNAIPLHRT STLLLAVADPTNQQAEDDFRFA TG LQVELV	124	124
GLSCTSLNQY EYASLCQTLGLRDLITRHNAIPLNRTS STLILAVADPTNQAEDDFRFA TG LQVELV	124	124
GVPMLDINCI DVGITIPVNLVNEKLIKRIHAMVPLFSRG TNLVLYLATDDPSKQASLKEITQFH TG LNTHAI	132	132
GEPLLDLGVF DPALFLKDKIDEKLIQKYRIMPLVHRG HVLYVATSNPTNIEAMDATIRFN SK LKVEPI	132	132
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consensus

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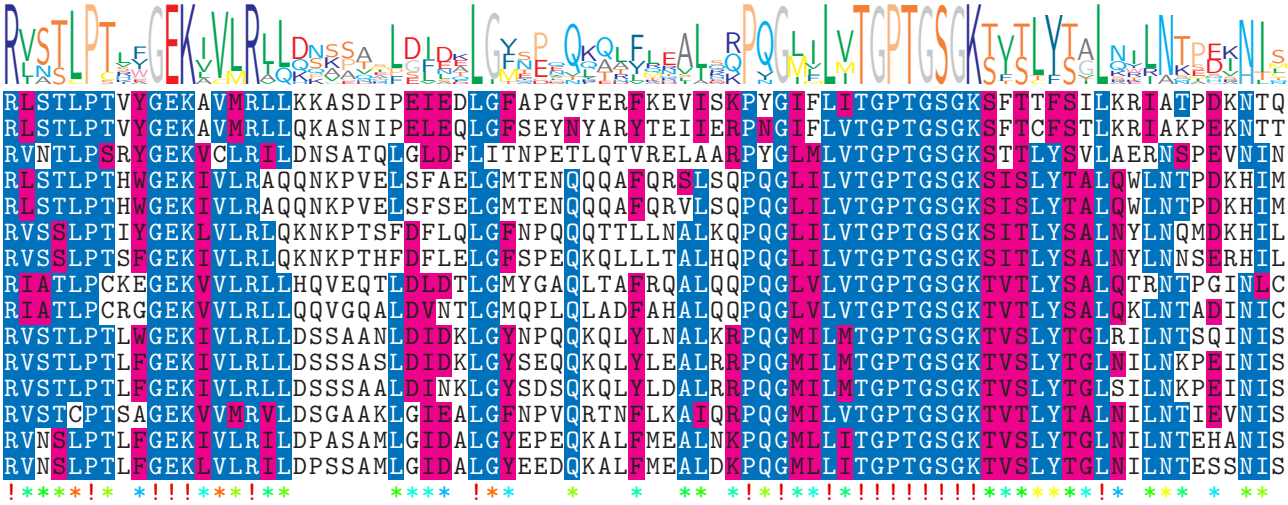
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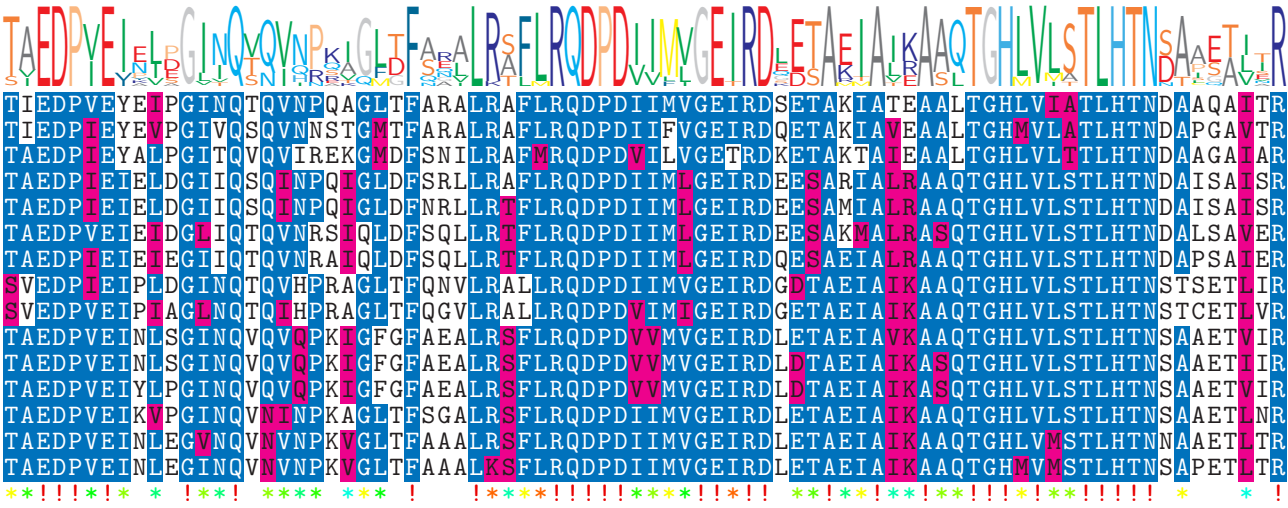
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consensus					



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consensus					



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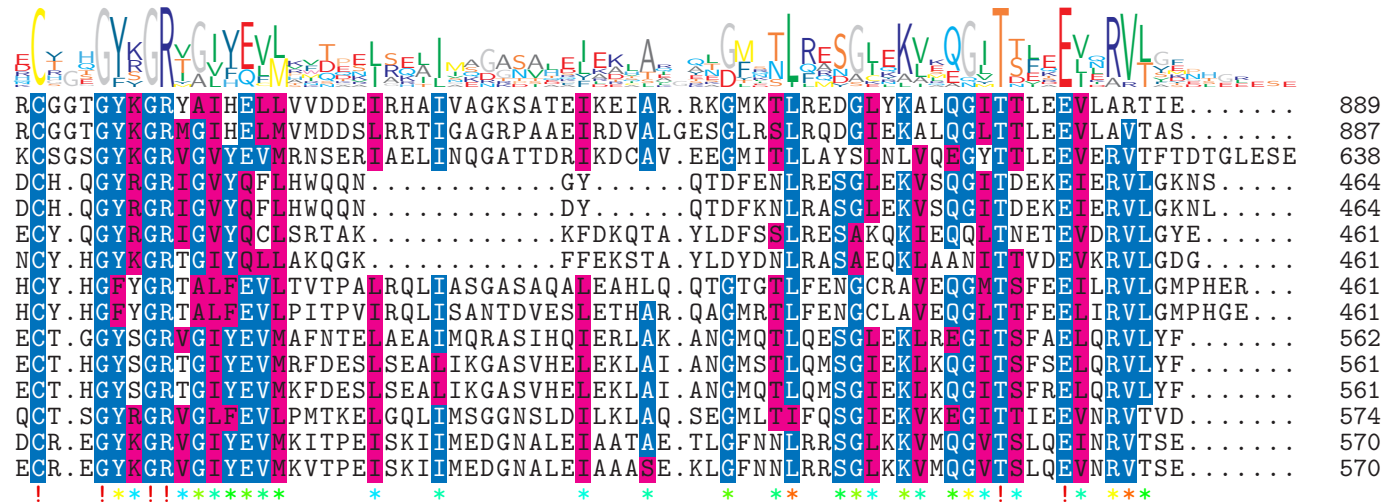
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consensus		

-  non conserved
-  similar
-  ≥ 50% conserved