

logo

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NTDB id 1384 A4U84 RS08525 WP 010787006.1
NTDB id 221535 BTV66 RS07740 WP 010906472.1
NTDB id 1366 NTHI RS01950 WP 011271972.1
NTDB id 1355 HI 0298 AAC21962.1
NTDB id 1168 A1552VC RS11070 WP 000957200.1
NTDB id 1293 VP RS12240 WP 005479695.1
NTDB id 1401 DSB67 RS12665 WP 010643259.1
NTDB id 1251 GC085 RS07725 WP 011946523.1
NTDB id 1058 ABD1 RS01615 WP 001274986.1
NTDB id 1018 ACIAD RS01685 WP 004920473.1
consensus

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[illegible]

logo

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consensus

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QRNQQQQRFLRLRYFALPLQEDEKQLWLGVDNSLANLSACDAFSFLTNGKHIEPVLLSTNELKRALLQSLPPKIQVEENIP.. 103
ERNQQQQSLLLRYFALPLKEENNRLWLGVDLSLNLSACETIAFITGKPVEPILLESSQLKELLQQLTPHQMVEEQVK.. 105
ERNQQQQSLLLRYFALPLKEENNRLWLGVDLSLNLSACETIAFITGKPVEPILLESSQLKELLQQLTPNQMQVEEQVK.. 105
CQQLGLRELITRDALPTAKQGNLLLLVADSDPTILLQAEEFFRFATGLQVELALADHRALQAATIRRLYGRSIGGAANQGK.. 151
CQQLGLRELITRHNAALPLNRTPSTLLLAVADPTNQAEEDFRFATGLQVELVLADFRELSTAIRRLYGRSLGHEKSGLK.. 151
CQTGLGLRDLITRHNAALPLNRSTSLLLAVADPTNLQAEEEDFRFATGLQVELVLADFRELTAATIRRLYGRSLGHEKSGLK.. 151
PVNLVNEKLIKRRHAMVPPLFSRGTNLYLATDDPSKQASLKEIQFHTGLNTHAIVVETDKLSALIDNLLTAKESQGLSEYVE.. 160
PKDLVDQKLLIKHRIPLLIQRGGIILYVATSNPSNIEAIDAIRFNSKLLVEPVIVEHHKLEKVLGHFAEESSFDND.. 158
LKDIDEKLIKQYRIMPLVVRHGHLVYATSNPTNIEAMDIAIRFNSKLLKVEPIIVEHDKLERLLSEHFVEETHFNFDTE.. 158

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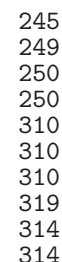
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[illegible]

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consensus

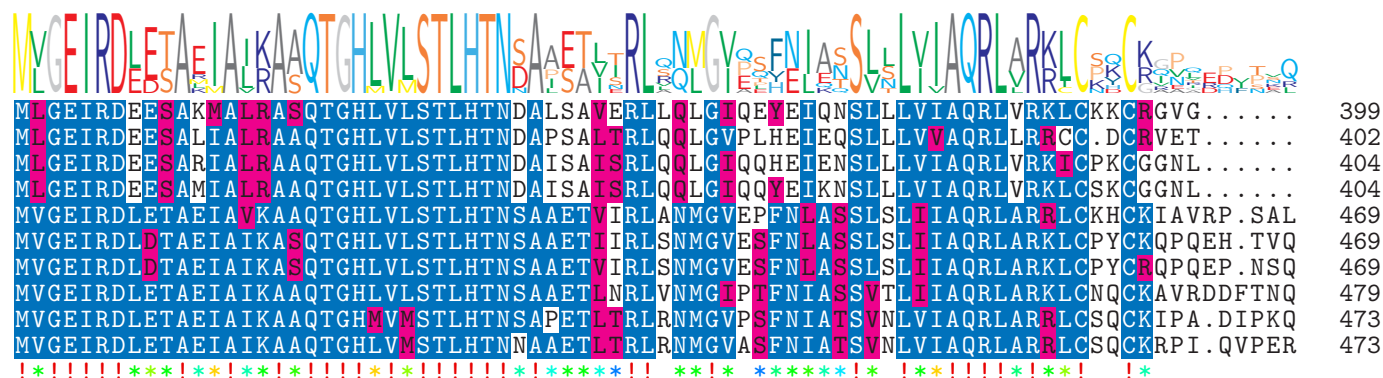
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consensus

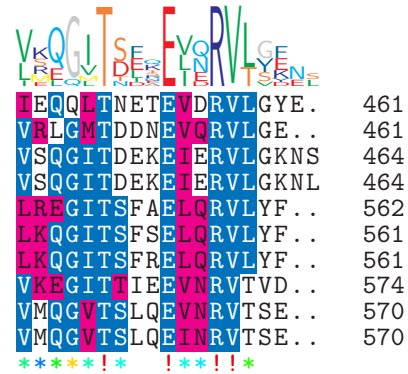
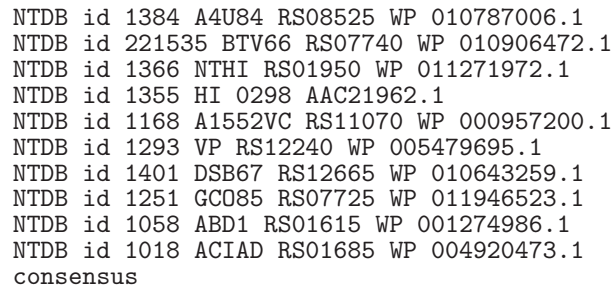
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consensus

```



- non conserved
- similar
- $\geq 50\%$ conserved