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 ... MEFEKLLRLMVEKGGSDLFITAGVPPSMKVNGKILPVNKT PMSPEMTRETIVHGVMNEQQRR EFTENHECNFAISAR 76
 ... MDFNDLNLMAKHKASD LFI TAGVEPSLKINGEITPVSKTKLDGMLAQLIDSITTTQKQRQEFIE THECNFAILNR 76
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 ... MDIAELLEFSVKHN ASDLHL SAGVPPMVRIDGEVRKLGVP AFTHSDVHRLIFEIMNDAQRSEYE EKLEVD FFSFELP 76
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[illegible]

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consensus

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consensus



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NTDB id 1177 A1552VC RS01165 WP 000422572.1	TFDQALYQLVVDKISEQDALHSADSANDLRLMLKTKR.GDDY...GSGSLQNVKI.....DME.....	368
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consensus	!*** * ** ***** * *	

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consensus		

- ☒ non conserved
- ☒ similar
- ☒ ≥ 50% conserved