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# Muscle

## Multiple Sequence Alignment (MSA)

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Survey link: [https://www.surveymonkey.com/r/QGFMBH8?channel=\[webpage\]](https://www.surveymonkey.com/r/QGFMBH8?channel=[webpage])

### Results for Job ID

muscle-I20250718-071908-0115-39479710-p1m

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# Nightingale

## COLOR SCHEME

clustal2



## LEGEND

A R N D C Q E G H I L K M F P S T W Y V B X Z

Color-coded legend for amino acids: A (blue), R (red), N (green), D (purple), C (pink), Q (orange), E (light green), G (light purple), H (teal), I (light blue), L (light blue), K (red), M (blue), F (blue), P (yellow), S (green), T (blue), W (teal), Y (blue), V (blue), B (blue), X (blue), Z (blue).

10 sequences



5JYQ\_2|CHAIN

3I40\_2|CHAIN

4E7U\_2|CHAINS

1ZNI\_2|CHAINS

4INS\_2|CHAINS

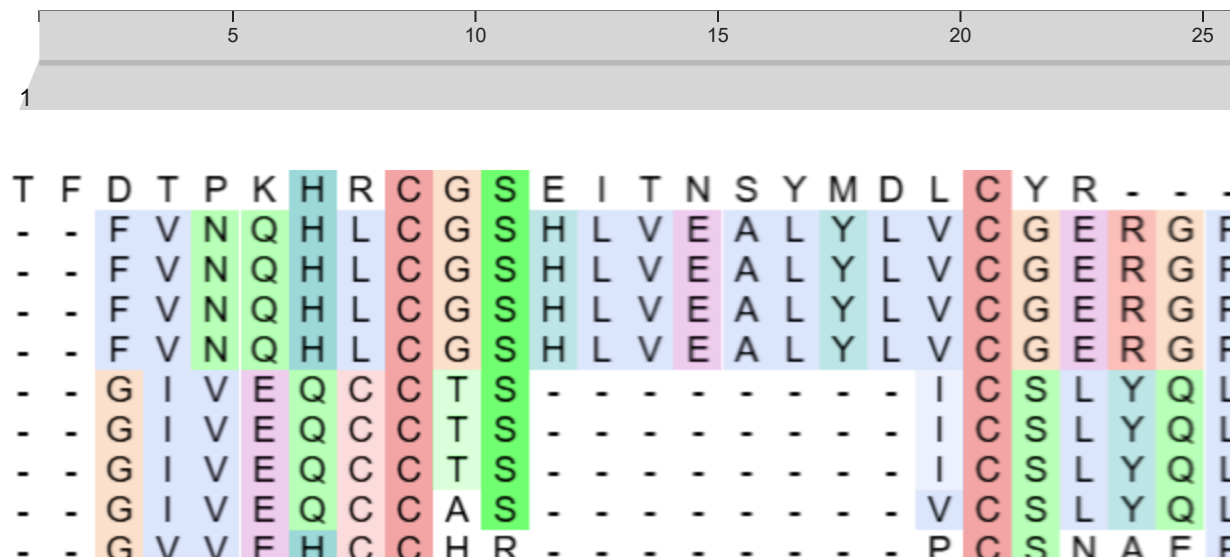
3I40\_1|CHAIN

1ZNI\_1|CHAINS

4INS\_1|CHAINS

4E7U\_1|CHAINS

5JYQ\_1|CHAIN



If you use this service, please consider citing the following publication: [The EMBL-EBI Job Dispatcher sequence analysis tools framework in 2024](#). More information about this bioinformatics application can be found in its [bio.tools](#) record.

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