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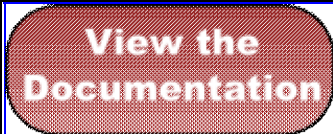
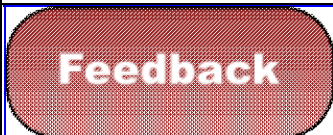
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StarORF Home Page

StarORF facilitates the identification of the protein(s) encoded within a DNA sequence. Using StarORF, the DNA sequence is first transcribed into RNA and then translated into all the potential ORFs (Open Reading Frame) encoded within each of the six translation frames (3 in the forward direction and 3 in the reverse direction). This allows students to identify the translation frame that results in the longest protein coding sequence.

Using StarORF

StarORF is accessible via the web. Press the "START" button to get started.

	<u>Start StarORF application</u> Start using the StarORF application.
	<u>Read StarORF documentation</u> StarORF user manual.
	<u>Send Us Feedback</u> Send us bug reports, comments, or questions. You can also send feedback via email to #include <emailstar.inc.html>

StarORF Screenshot

