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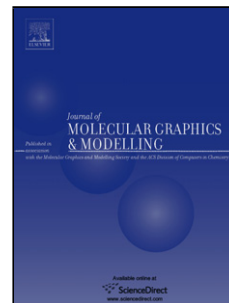
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# Protein Secondary Structure Prediction: A Survey of the state of the art

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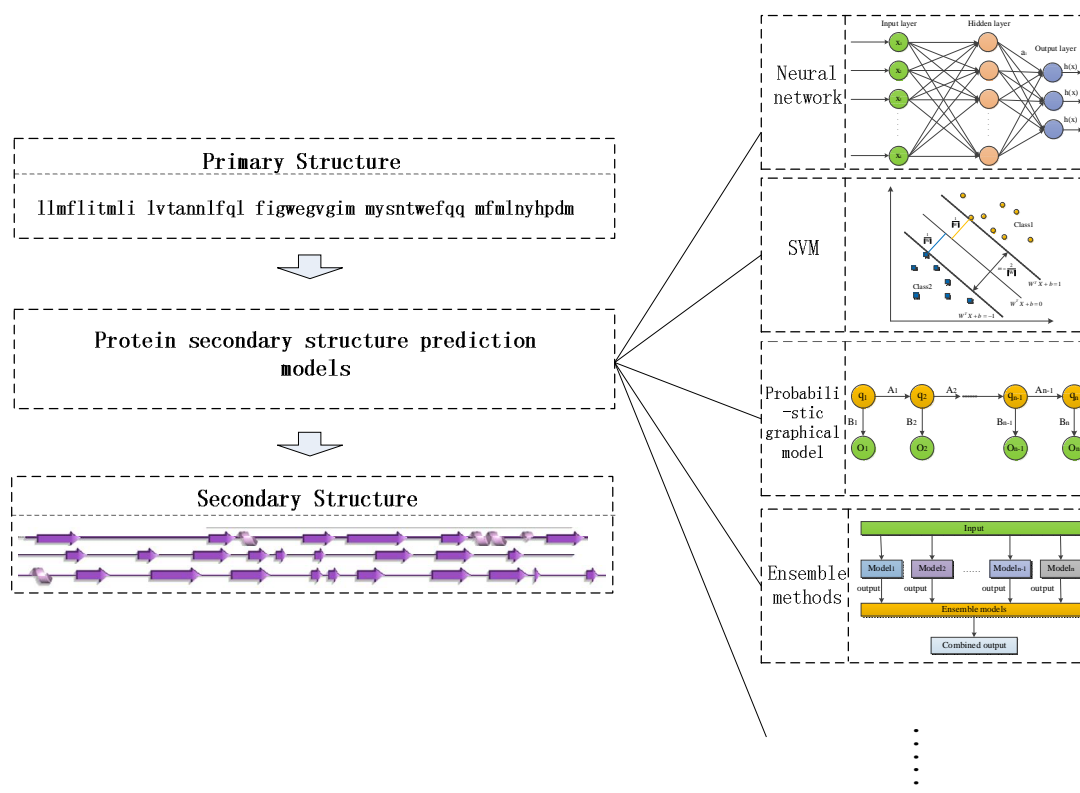
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Graphical abstract



## Highlights:

1. The background and related knowledge of protein secondary structure prediction is introduced.
2. The often-used prediction accuracy assessments methods are described.
3. The recent algorithmic developments of protein secondary structure prediction are reported in detail.
4. The corresponding tendencies and challenges are summarizes.
5. We conclude there are still further improvements and extensions in this field.

**Abstract:** Protein secondary structure prediction (PSSP) is a fundamental task in protein science and computational biology, and it can be used to understand protein 3-dimensional (3-D) structures, further, to learn their biological functions. In the past decade, a large number of methods have been proposed

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