

Prediction of RNA and DNA binding sites: preliminary presentation

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Outline

- ▶ CAFA.
- ▶ Background: RNA and RNA binding proteins.
- ▶ Background: datasets.

Definition

- ▶ **RNA/DNA Binding Protein prediction:** given a protein, determine whether a protein is RNA/DNA binding.
- ▶ **RNA/DNA binding site prediction:** given a protein sequence, determine side chains that bind with a DNA/RNA.

CAFA

1. Determine whether protein is RNA or DNA binding.
2. Determine binding site.

Methods: abstraction level

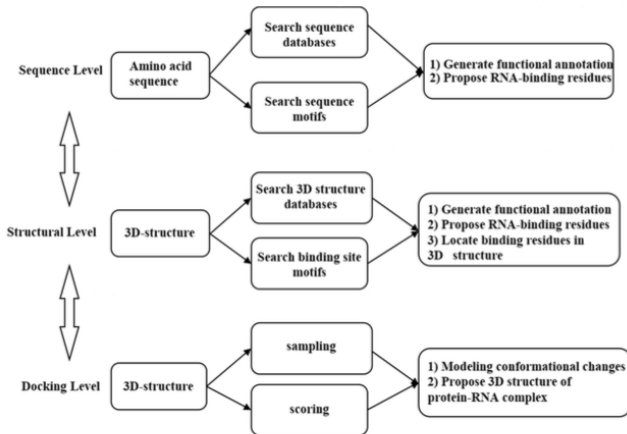


Figure 1. Strategies for RNA-binding site and RBP prediction.

Possible Features

Sequence-based features:

- ▶ **Amino acid composition.**
- ▶ **Sequence similarity**, such as MSA.
- ▶ **Evolutionary invormation**, such as PSSM.
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Structure-based features:

- ▶ **Secondary structure:** experimental (assigned using DSSPcont) or predicted.
- ▶ **Accessible surface area**, in percent (%).

Chemical and physical features:

- ▶ **Hydrophobicity.**
- ▶ **Electrostatic patches.**
- ▶ **Cleft Size.**

Methods: previously used algorithm

- ▶ **Naive Bayes (NB)** classifier.
- ▶ **Support Vector Machine (SVM)**.
- ▶ **Random Forest**.
- ▶ **Neural Network (NN)**.

Statistics

Name	Num
SwissProt (HUMAN)	20120
GO:0003676 (HUMAN)	1248
SwissProt (filtered out GO:0003676)	20005

Statistics (cont.)

Results of redundancy reduction:

Before	After
706	567