Prediction of RNA and DNA binding sites: preliminary presentation

Quirin Heiss^{1,2} Pandu Raharja ^{1,2} Julian Schmidt ^{1,2}

¹Technische Universität München

²Ludwig-Maximilians-Universität München

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Outline

- CAFA.
- ▶ Background: RNA and RNA binding proteins.
- ► Background: datasets.
- Project plan.

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.

Methods: abstraction level

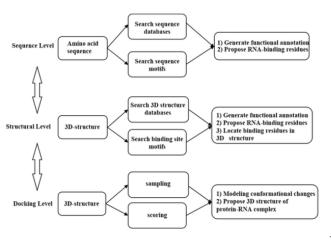


Figure 1. Strategies for RNA-binding site and RBP prediction.

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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 percnent (%).

Chemical and physical features:

- Hydrophobicity.
- Electrostatic patches.
- Cleft Size.