Conformational design of self-organizing ribozymes

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RNA nanotechnology - self-assembled RNA



Peixuan Guo, Nature Nanotechnology (2010)

Ribozymes



The hairpin ribozyme



The modified hairpin ribozyme



Pieper et al. Biol. Chem. (2007)

The modified hairpin ribozyme



Pieper et al. Biol. Chem. (2007)

Summary

- small self-pocessing molecule
- minimal version \Rightarrow multimeric structures
- equilibrium dependent on sequence optimization

Goals

- predict the dynamics of the system
- design & optimize self-splicing ribozymes
- design & optimize multimeric structures

Challenges

- size of conformation space multimerization, catalytic activity
- coarse graining influences dynamics
- intra-/inter-molecular dynamics
- rule-based (network expansion/shrinking steps)

Network expansion



Stochastic simulation



Network shrinking, network expansion



Theoretical Approach

- rule-based landscape modeling
 - hypergraph
 - \Rightarrow vertices: RNA secondary structures
 - $\Rightarrow\,$ edges: transition rates & cleavage/ligation reactions
 - rule set
 - \Rightarrow sampling of RNA structures
 - \Rightarrow estimate transition rates
 - \Rightarrow stochastic simulation
 - \Rightarrow check for catalytic activity
- sequence/landscape optimization

Current Work

- store predefined set of helices (e.g.: len(h) > n)
- generate suboptimal structures
- allow overlappings (zippering-, elongating- helices)
 ⇒ nearly done :-)
- stochastic folding kinetics on helix-lists

Future Implementations

- generate a landscape with stochastic simulation (expansion, shrinking)
- intermolecular interactions
- catalytic motif finder

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Backup Slide: modified hairpin ribozyme



Pieper et al. Biol. Chem. (2007)



sequence length











