# On the Degree of Freedom in Multilevel Evolutionary Models

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Small Degree of Freedom

Conclusion

## **Evolution of Biotic Systems**

Biotic systems:

- Large degree of freedom
  - $\rightarrow$  Great variety in possible adaptations
- Constraint and structure
  - → RNA/protein folding is not random
  - → Random mutation produces statistical order
  - → Adaptations might be conceivable, not equally achievable

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# How to Model Evolution

Simplicity versus Complexity in Model

- Small degree of freedom + predefined structure
  - → Strengthening/weakening of predefined structure e.g.) Predator X <sup>axy</sup>/<sub>xy</sub> Prey Y
  - → Novel structure cannot emerge (structures are predefined)
- Large degree of freedom + biological constraint
  - $\rightarrow$  Novel structure can emerge

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# The Purpose of This Talk

- We compare the two modeling approaches  $\rightarrow$  small vs. large degree of freedom (N = 2!)
- Kinds of adaptations depend on the degree of freedom available to evolution

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Conclusion

# Model with Large Degree of Freedom

Evolution of Complexity in RNA-like Replicator Systems



 $C + T \xrightarrow{\alpha} C + 2T$ 

Takeuchi & Hogeweg (2008) Biol Direct 3:11

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Introduction

Large degree of freedom

Small Degree of Freedom

Conclusion

## Result



#### Arrows denote replication

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## Result



#### Arrows denote replication

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## Result





#### Arrows denote replication

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## Result



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#### Arrows denote replication

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Introduction

Large degree of freedom

Small Degree of Freedom

#### Result



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On the Degree of Freedom in Multilevel Evolutionary Models

Introduction

Large degree of freedom

Small Degree of Freedom

Conclusion

#### Mess of Data

51 :ls -lh sequence-prototype.seq
-rw-rr 1 nobuto binf 251M Sep 7 21:54 sequence-prototype.seq
nobuto@rna:~/tmp
52 :head -n 100 sequence-prototype.seq
t0 t0+ GCAUACGAUUUGCCUCUUUCGCGUUUCCUGGCGUUACAUUAUUUUCUUUC
to co- GAAAGAAAAUAAUGUAACGCCAGGAAACGCGAAAGAGGCAAAUCGUAUGC p104838 c2 (((((H)S)((H)S)M)S)R)
t500 t0+ GCAUACGAUUUGCCUCUUUCGCGUUUCCUGGCGUUACAUUAUUUUCUUUC
t500 c0- GAAAGAAAAUAAUGUAACGCCAGGAAACGCGAAAGAGGCAAAUCGUAUGC p34502 c2 (((((H)S)((H)S)M)S)R)
t500 c1+ GAAAGAAAAUAAUGUAACGCCAGGAAACGCGUAAGAGGCAAAUCGUAUGC p3382 c3 (((((H)S)((H)S)M)S)R)
t500 t1- GCAUACGAUUUGCCUCUUACGCGUUUCCUGGCGUUACAUUAUUUUCUUUC
t500 p0+ GCAUACGAUNUGCCUCUUUCGCGUUUCCUGGCGUUACAUUAUUUUCUGUC p884_c4 (((((((H)S)I)S)I)S)R)
C500 p0- GACAGAAAAUAAUGUAACGCCAGGAAAGAGGCAAAGGGCAAAUCGUAUGC p1477 c4 (((H)S)(((H)S)(H)S)M)S)R)
t500 p1+ GCAUACGAUUUGCCULUUUCGCGUUULAUGGCGUUACAUUAUUUUUUUC p586 c5 (((((((H)S)I)S)I)S)R)
L500 p1- GAAAGAAAAUAAUGUAAACGCCAAAGGGGAAAGAGGCAAAUCGUAUGC p1180 c5 (((((H)S)I)S)R)
t500 t2+ GCAAACGAOUUGCCUCUUUCGCGUUUCCUGGCGUUACUUUCCUUUC
L500 c2- GAAAGAAAAUAAUGUAALGCCAGGAAAGGCGAAAGAGGCAAAUCGUUUUGC p844 c6 (((((H)S)((H)S)M)S)R)
±500 ±3+ GA0ACGA000GCC0C000CGCG000CC0GGCG00AC0000C0000 p384 c7 ((((((A)S)1)S)1)S)1)S)R)
t500 c3- AAAAGAAAAUAAAGGAAAUAAGGAAAAGGAAAAGGAAAAGGAAAGGAGGAAGGAGGAGGAGGAAGGAGGAAGGAGA
±500 ±4+ GCH0ALGAOUUGCCUCUUULGCGUUULCUGGCGUUALCAUUAUUAUUAUUAUUAUUAUUAU
t500 c4 GAAAGAAUAAUGAALGULAGGAAAUGULGAAAAGAGULAAAUUGUAUGU p/55 c8 ((((H)S)((H)S)M)S)R)
t500 t5+ GLORICGUOUGLCUOUGLGUOUCLGGGGGOOLGOOLGOOLGOOL p3/9 c9 ((((((h)5/1)5/1)5/1)5/1)
t500 c5- GAAALAAAAUAAUGUAALGULAGGAAAUGUGAAAUGUGAAUGUAAUUGUAUGU p/30 c9 ((((H)S)((H)S)M)S)R)
1500 t6+ 6tH0H66H0006Ct000066C60000Ct066C6000H00H00H00H00H00H00H00H00H00H00H00H00
t500 cb- GAAAUAAAAUAAAUAUGAALGULAAGAAAUGGGAAAAGAGULAAAUUGUAUGU p/24 c10 (((((h)S)(h)S)R))
$t_{500}$ t/+ 6cH0HcGH0006cL0C000C6C60000CC066c600Hc00000L000c p345 cl1 ((((((H)S)(1)S)1)S)1)S)R)
CS00 67- GHAHGAHAHOUHAHGGUAHAGGCHAHAGGGAHAHGGGAHAHGGGAHAHGUGUAHAGUA p/37 611 ((((H/S)((H/S)M)S)R)
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Small Degree of Freedom

Conclusion

# Bioinformatic pattern detection



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# Finding Meaningful Observables





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#### Visualization with "designed" observables



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# Simplicity & Complexity of a Complex Model

#### Simplicity:

- \* General results are simple
  - $\rightarrow$  importance of the results

#### Complexity:

\* Results are unforeseeable

general ecological organization, let alone sequence & structure

\* Recognition of results is nontrivial

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# Model with Small Degree of Freedom

- System has predefined structure
- Only two parameters can evolve

$$\mathsf{R} \underbrace{\overset{k_L}{\longrightarrow}}_{\mathbb{I}^*} \mathsf{L}$$

- R : replicase
- L : parasites in template state
- L\*: parasites in folded state

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#### Survival Region in a Well-mixed System



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## **Multilevel Selection**

#### compartmentalization (vs. spatial self-organization)



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## Survival Region with Compartmentalization



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# Long-term Evolutionary Trend



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Conclusion

## Transition Happens in Evolutionary Trend



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Introduction	

Small Degree of Freedom



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# Simplicity & Complexity of a Simple Model

#### Simplicity:

∗ Recognition of results is trivial ← only 2 parameters change omplexity:

#### Complexity:

- \* Results (i.e. adaptations) are unforeseeable
- \* Results are intricate/subtle

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# Multilevel Evolutionary Models

- Adaptations (i.e. results) are unforeseeable.
- Depending on the degree of freedom available to evolution → Evolution generates different kinds of adaptation.

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# Complex Simplicity vs. Simple Complexity

- Small degree of freedom → Evolution "sensitively" detects possible adaptations (despite strong restriction)
  - $\rightarrow$  Trivial to recognize

(predefined structures & a few parameters)

- → Intricate/Subtle in adaptive effect
- Large degree of freedom → Evolution "inventively" generates best adaptations (also attainability & maintainability)
  - $\rightarrow$  Non-trivial to recognize

(lack of search images & predefined observables)

 $\rightarrow$  Obviously adaptive in hindsight and, thus, important

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