# Self-contained RNA inhibition with trans-acting ribozymes

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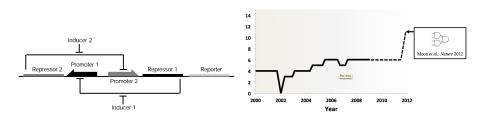
## Promise of Synthetic Biology

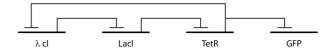
Complexity of eukaryotes $\not\approx \#$ of protein coding genes		
Oryza sativa (rice)	470 million	51,00
Gallus gallus (chicken)	1 billion	20,000-23,00
Canis familiaris (dog)	2.4 billion	19,00
Mus musculus (mouse)	2.5 billion	30,00
Homo sapiens	2.9 billion	20,000-25,000

The root of complexity is believed to be the regulation of these genes. However, the creation of novel protein regulatory elements is too difficult. Re-writers RNA-world may be the key to getting a handle on regulation.

#### Motivation

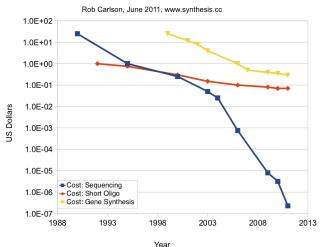
Toggle switch and repressilator in 2000 (images).





Exponential decrease in the cost of enabling technologies should result in exponential growth of circuit complexity.

#### Cost Per Base of DNA Sequencing and Synthesis



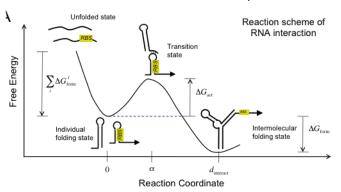
#### Pitfalls in current promoter-repressor pair design

- Orthogonal Limited number of repressors (until very recently)
- Predictable Gene circuit evolves away
- Safe shRNA toxicity in gene therapy
- Reliable 40 hour toggle switch breakdown
- Designable Protein structure prediction too difficult
- Cooperativity Unpredictable behavior when juxtaposed

## Types of riboregulation

Why shRNA sucks, because of cleaving mechanism. But this can be avoided.

Self-contained action. removes some dependencies.



## Choice of trans-acting ribozymes

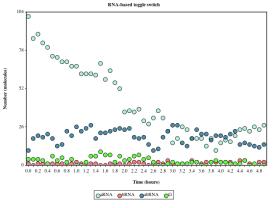
Proof of functional completeness Include reasoning for not having apatamer effected ribozymes. More difficult to design and predict than simple oligonucleotide effectors. Watson Crick base pairing dominates free energy minimization

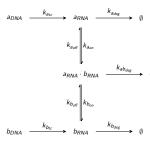
## General riboregulation model

Stochastic model - Gillespie algorithm The probability Rate of trans cleaving

## Toggle switch model riboregulation

No possible bistable point. First limitation.





## Parameter Comparison

## Implications and further work

If these

#### Souces

- Pray, L. A.; http://www.nature.com/scitable/topicpage/ eukaryotic-genome-complexity-437
- Arkin, A. and Weiss, Ron; Principles of Synthetic Biology Fall 2013;
   Lecture 3
- Carlson, Rob; DNA cost curves;
   http://www.synthesis.cc/2011/06/new-cost-curves.html
- Stanton, B.C. et. al.; Genomic Mining of prokaryotic repressors for orthogonal logic gates; http://www.nature.com/nchembio/ journal/vaop/ncurrent/full/nchembio.1411.html
- Martin, J.N., et. al.; Lethal toxicity caused by expression of shRNA in the mouse striatum: implications for therapeutic design; http: //www.nature.com/gt/journal/v18/n7/full/gt201110a.html
- Bongarets; GFP as a Marker for Conditional Gene Expression in Bacterial Cells http: //www.ifr.ac.uk/Safety/molmicro/pubs/bongaerts2002.pdf
- Anderson, J.C.; org.devicecourse Gillespie module
- © Gillespie, Daniel T. (1977). Exact Stochastic Simulation of Coupled Stack Field & Ryan Tsoi (UC Berkeley)

  RNA circuits

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## Blocks of Highlighted Text

#### Block 2

Pellentesque sed tellus purus. Class aptent taciti sociosqu ad litora torquent per conubia nostra, per inceptos himenaeos. Vestibulum quis magna at risus dictum tempor eu vitae velit.

#### Block 3

Suspendisse tincidunt sagittis gravida. Curabitur condimentum, enim sed venenatis rutrum, ipsum neque consectetur orci, sed blandit justo nisi ac lacus.

## Multiple Columns

#### Heading

- Statement
- 2 Explanation
- Example

Lorem ipsum dolor sit amet, consectetur adipiscing elit. Integer lectus nisl, ultricies in feugiat rutrum, porttitor sit amet augue. Aliquam ut tortor mauris. Sed volutpat ante purus, quis accumsan dolor.

## Table

Treatments	Response 1	Response 2
Treatment 1	0.0003262	0.562
Treatment 2	0.0015681	0.910
Treatment 3	0.0009271	0.296

Table: Table caption

#### **Theorem**

## Theorem (Mass-energy equivalence)

 $E = mc^2$ 

#### Verbatim

### Example (Theorem Slide Code)

```
\begin{frame}
\frametitle{Theorem}
\begin{theorem}[Mass--energy equivalence]
$E = mc^2$
\end{theorem}
\end{frame}
```

## **Figure**

Uncomment the code on this slide to include your own image from the same directory as the template .TeX file.

#### Citation

An example of the \cite command to cite within the presentation:

This statement requires citation [Smith, 2012].

#### References



John Smith (2012)

Title of the publication

Journal Name 12(3), 45 - 678.

## The End