

# An Introduction to Synthetic Biology

*“Hacking DNA for Fun and Profit”*



1. what?
2. applications
3. the tech
4. free and safe
5. go hack

1. what?

2. applications

3. the tech

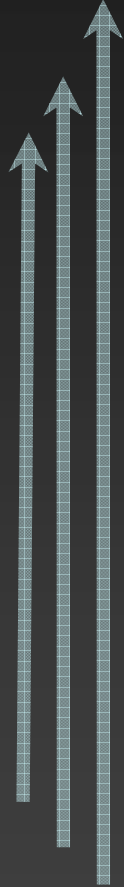
4. free and safe

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# Engineering vs Science

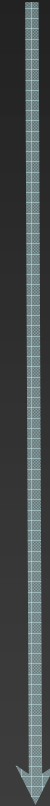
Understanding,  
Models

Science



Artifacts  
(cells)

Engineering



# Synthetic Biology

Synthetic Biology

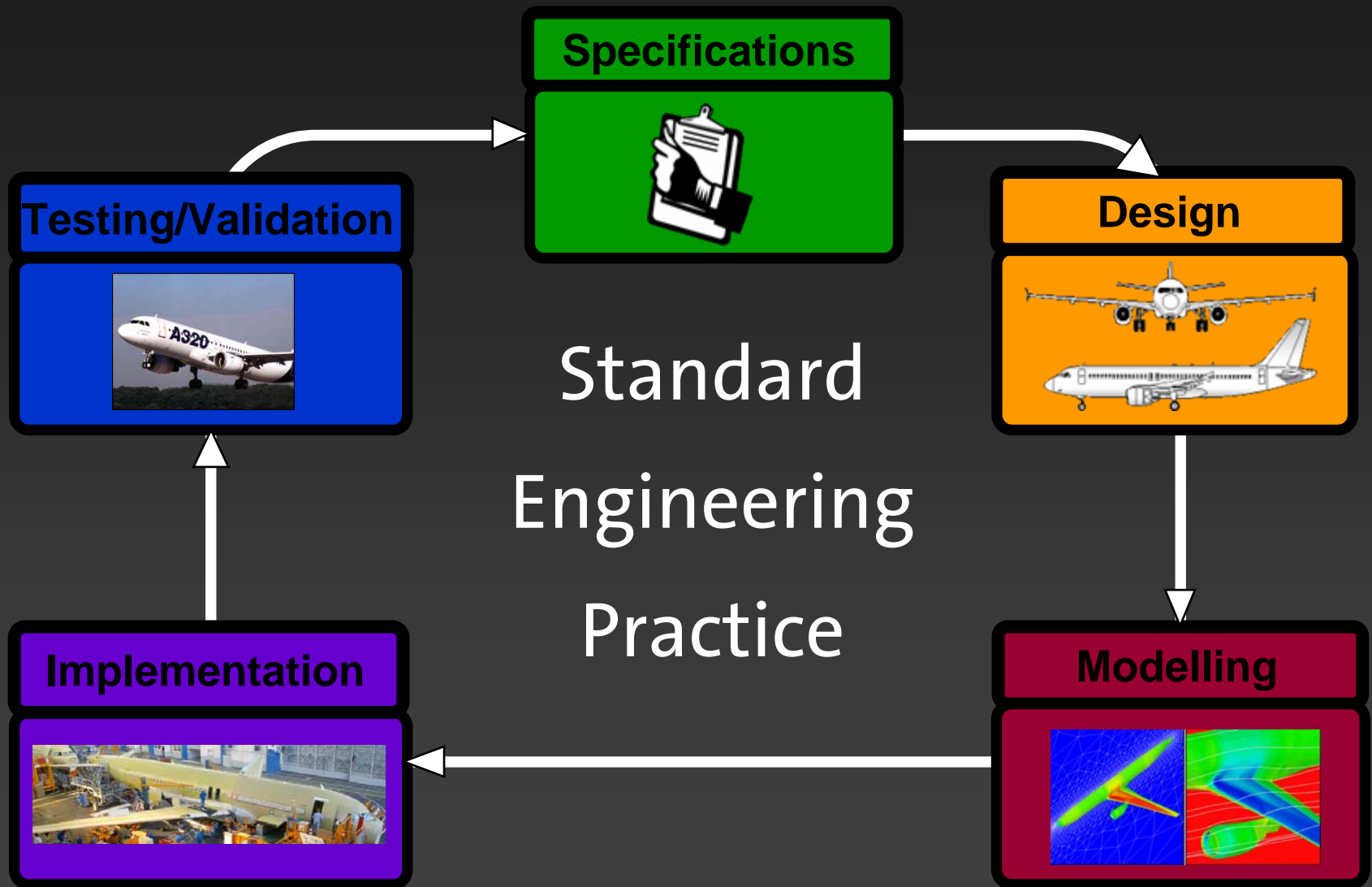
=

GeneticEngineering++



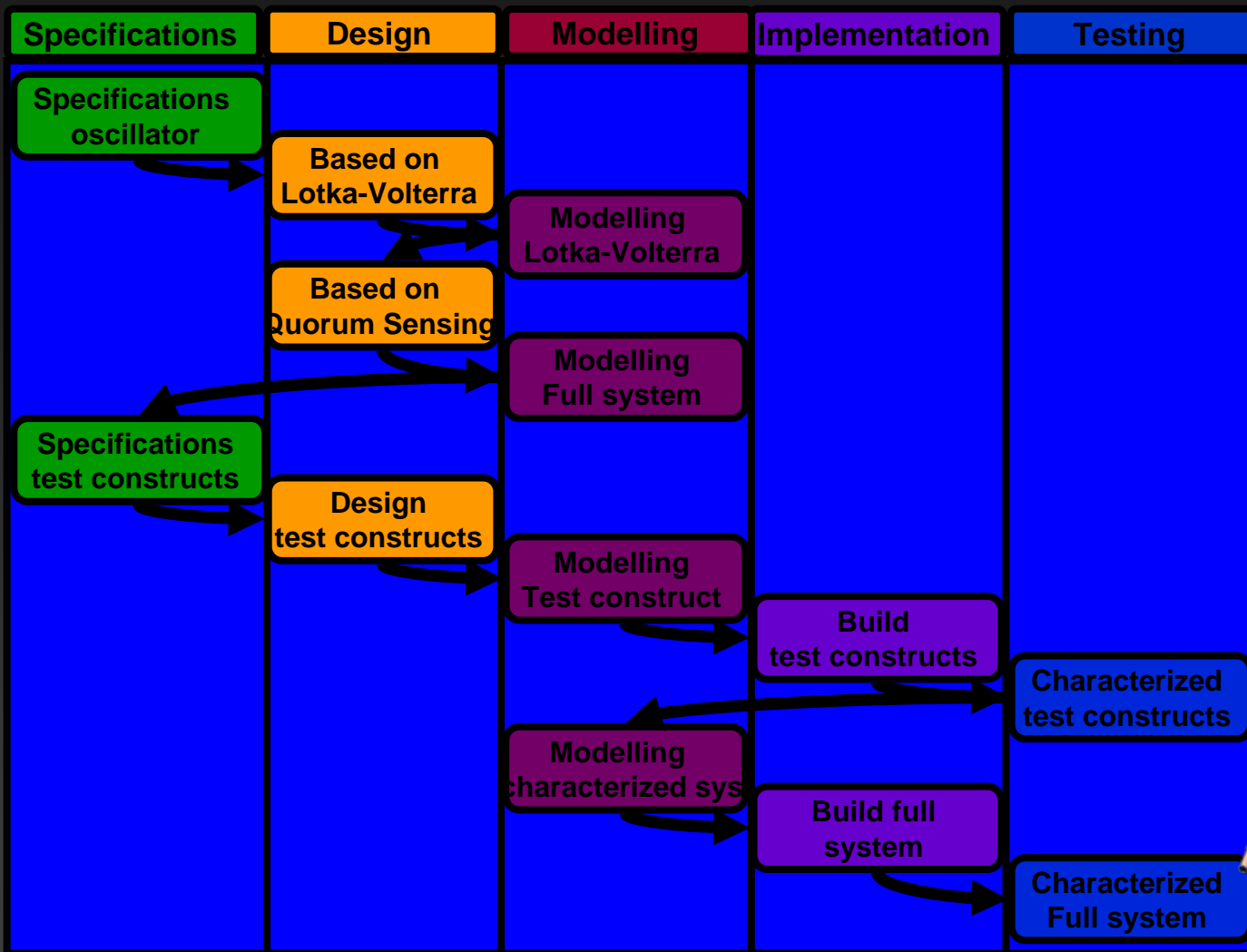
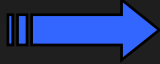
1. Recombinant DNA
2. PCR
3. Automated sequencing

1. Recombinant DNA
2. PCR
3. Automated sequencing
4. Automated construction
5. Standardization
6. Abstraction



The Imperial College of London, iGEM 2006

Start!



Our Goal!

1. what?

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# Biofuels

# Bioremediation

# Medicine



# Space ISRU

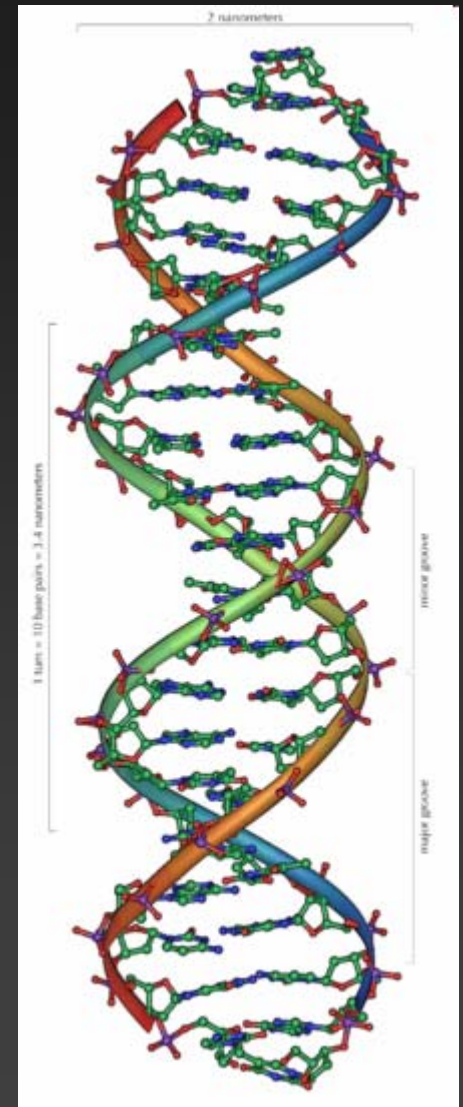
1. what?

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\$1/bp, 4wk

e. coli: 4k

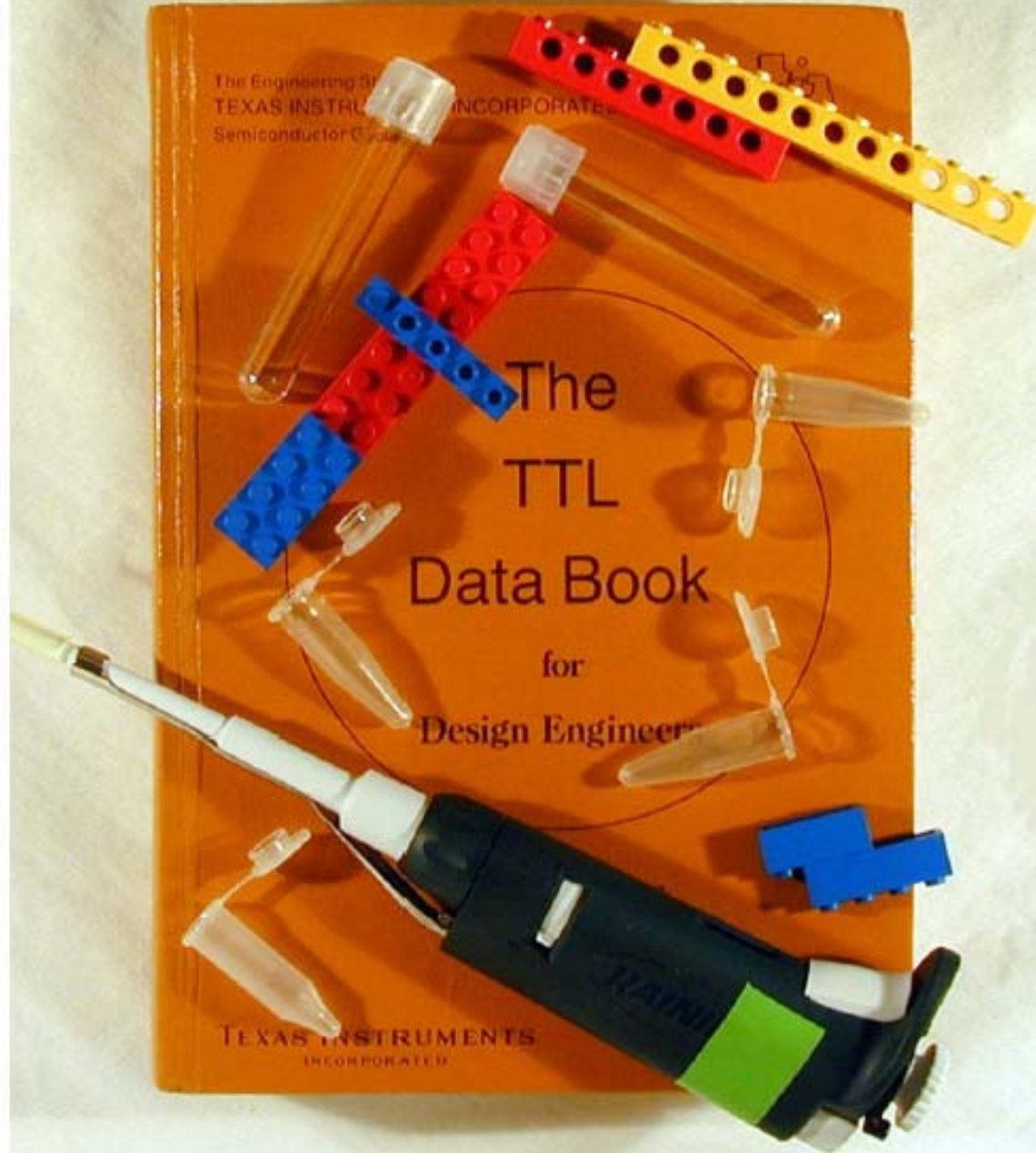
you and me: 3gbp

# Standardized Assembly

The Engineering Staff  
TEXAS INSTRUMENTS INCORPORATED  
Semiconductor Company

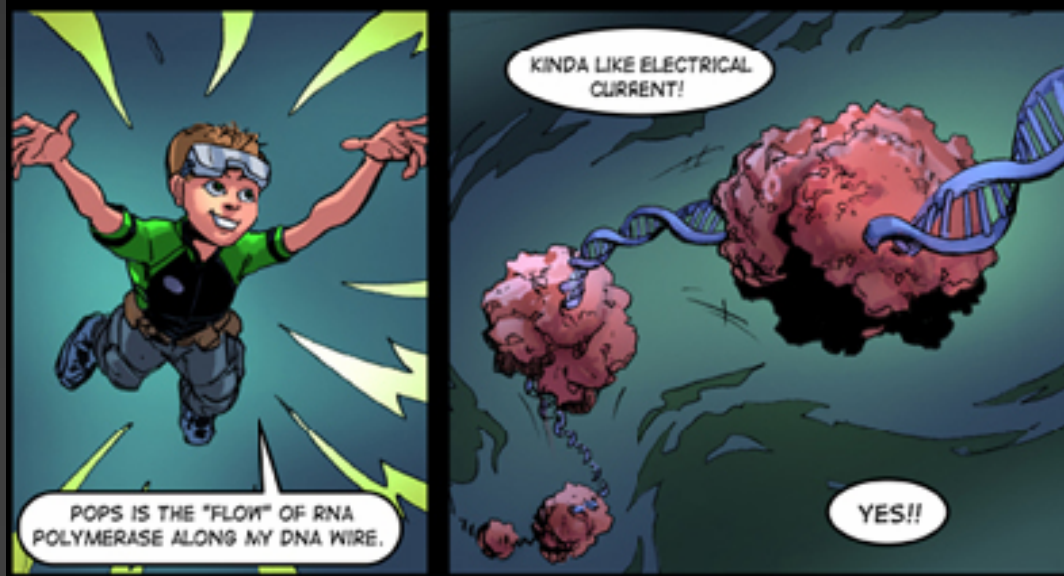
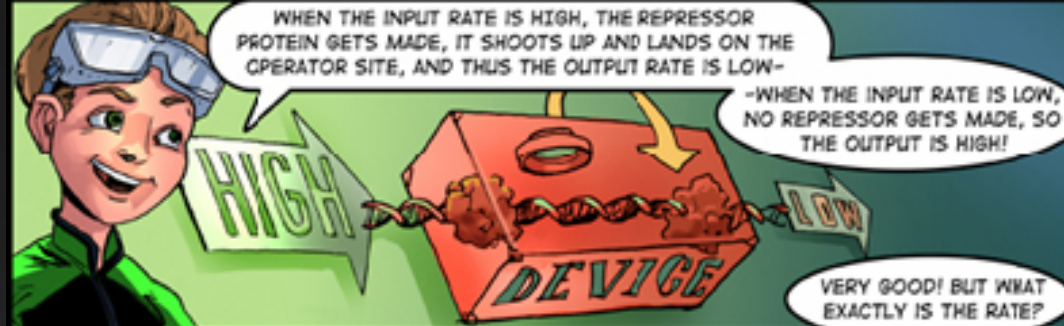
# The TTL Data Book for Design Engineers

TEXAS INSTRUMENTS  
INCORPORATED



# Common Signal Carrier





Polymerase  
Operations  
per Second



# Characterization

# BBa\_F2620

3OC<sub>6</sub>HSL → PoPS Receiver

[http://parts.mit.edu/registry/index.php/Part:BBa\\_F2620](http://parts.mit.edu/registry/index.php/Part:BBa_F2620)



F2620



Authors:  
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Last Update: 15 January 2007

## Description

A transcription factor (LuxR, BBa\_C0062) that is active in the presence of cell-cell signaling molecule 3OC<sub>6</sub>HSL is controlled by a TetR-regulated operator (BBa\_R0040). Device input is 3OC<sub>6</sub>HSL. Device output is PoPS from a LuxR-regulated operator. If used in a cell containing TetR then a second input signal such as aTc can be used to produce a Boolean AND function.

## Characteristics

**Input Swing:** 1E-9 to 1E-6 M 3OC<sub>6</sub>HSL, exogenous

**Output Swing:** 0±1 to 503±1 GFP molecules cfu<sup>-1</sup> s<sup>-1</sup>

**Switch Point:** 7±1 nM 3OC<sub>6</sub>HSL, exogenous

**LH Response:** 9 min (t<sub>50%</sub>), 27 min (t<sub>90%</sub>)

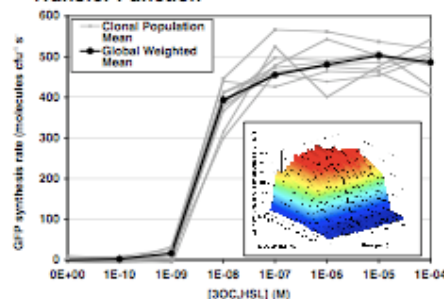
## Key Parts

BBa\_R0040: TetR-regulated operator

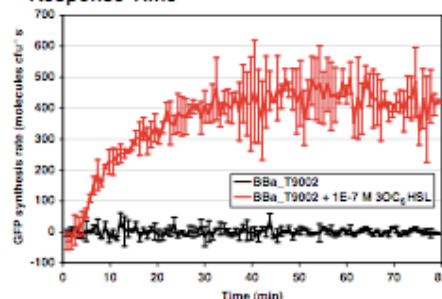
BBa\_C0062: luxR ORF

BBa\_R0062: LuxR-regulated operator

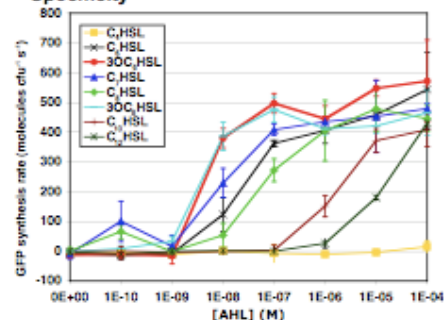
## Transfer Function\*



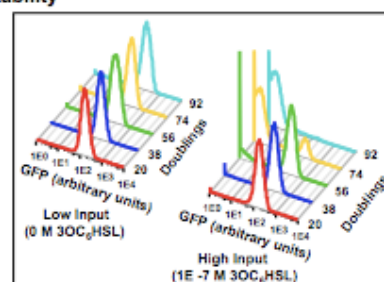
## Response Time\*



## Specificity\*



## Stability\*\*



## Demand (low/high input)

**Translational:**  
256/8048 ribosomes cfu<sup>-1</sup>  
3.8E3/1.2E5 charged tRNA cfu<sup>-1</sup> s<sup>-1</sup>

## Compatibility

**Chassis:** Compatible with MC4100, MG1655, and DH5α

**Plasmids:** Compatible with pSB3K3 and pSB1A2

**Devices:** Compatible with E0240, E0430 and E0434

Crosstalk with systems containing TetR (C0040)

**Signaling:** Crosstalk with input molecules similar to 3OC<sub>6</sub>HSL

## Stability (low/high input)

**Genetic:** >92/74 replication events\*\*

**Performance:** >92/74 replication events\*\*

## Conditions (abridged)

**Output:** Indirect via BBa\_E0240

**Vector:** pSB3K3

**Chassis:** MG1655

**Culture:** Supplemented M9, 37°C

**\*Equipment:** PE Victor3 plate reader

**\*\*Equipment:** BD FACScan cytometer

Signaling Devices

# Composition & Abstraction

## Part:BBa R0011

Designed by Neelaksh Varshney, Grace Kenney, Daniel Shen, Samantha Sutton

Entered: Antiquity

**Promoter (lacI regulated, lambda pL hybrid)**

Inverting regulatory region controlled by LacI (BBa\_C0010 , BBa\_C0012 , etc.) The PLlac O-1 promoter is a hybrid regulatory region consisting of the promoter P(L) of phage lambda with the cl binding sites replaced with lacO1. The hybrid design allows for strong promotion that can nevertheless be:

- repressed by LacI, the Lac inhibitor (i.e. repressor) ([BBa\\_C0012](#)) ([LUTZ97]).
- induced by [IPTG](#) in E.Coli DH5-alpha-Z1 (same paper reference) over a >600-fold range

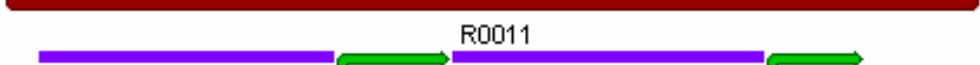
### Usage and Biology

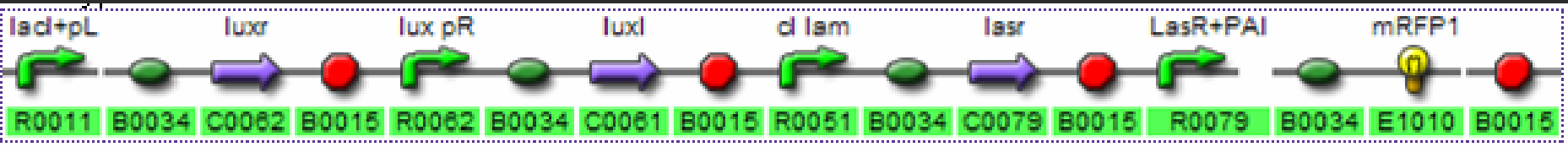
[\[edit\]](#)

Strong promoter. [jib, 5/24/04]

R0011 will be on in strains without lacI, off in strains that are lacIq (ie. [Part:BBa V1003](#)) and medium in strains that are lacI

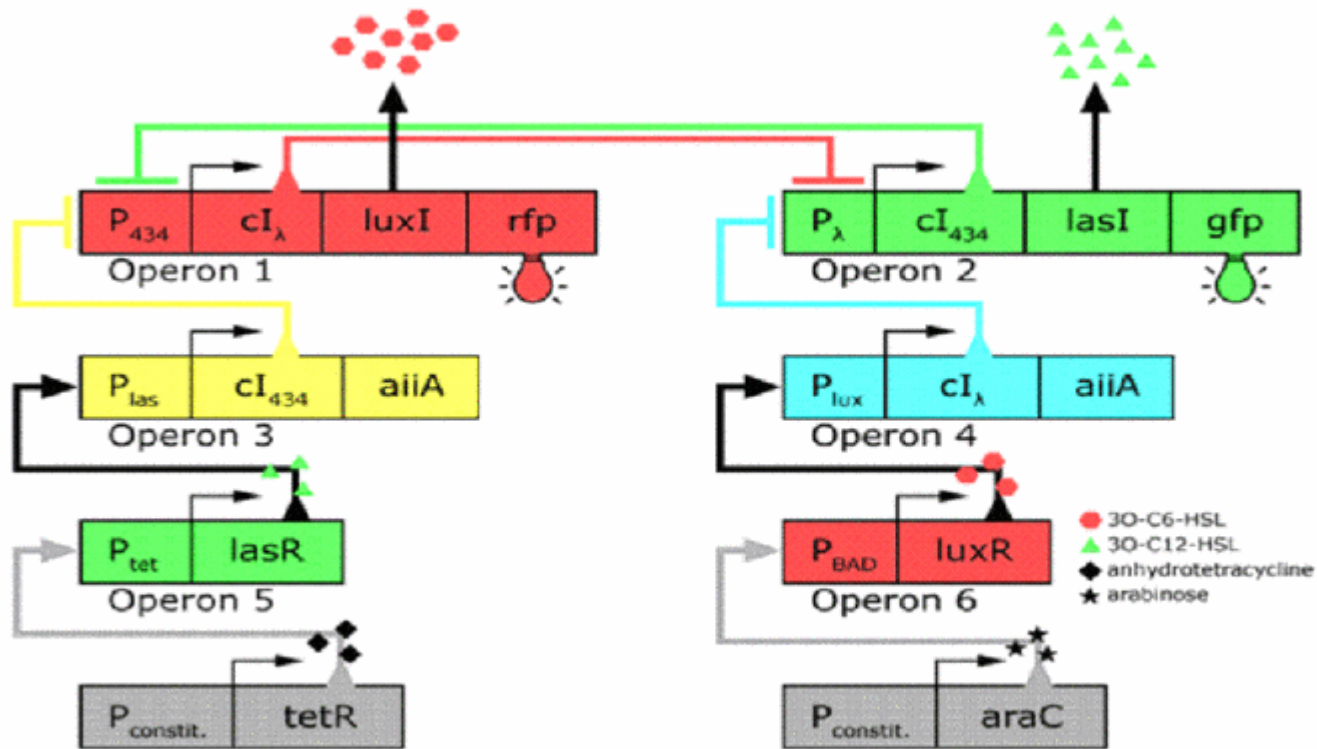
## Sequence and Features

Format:	Subparts	<a href="#">Ruler</a>	<a href="#">SS</a>	DS	Search:	Length: 55 bp	Context: Part only	<a href="#">Get selected sequence</a>		
	1	11	21	31	41	51	61	71	81	91
1	aattgtgagc ggataacaat tgacattgtg agcggataac aagataactga gcaca ttaacactcg cctattgtta actgtaacac tcgcctattg ttctatgact cgtgt									
										



<http://www.ccbi.cam.ac.uk/iGEM2006/index.php/Description>





**Figure 5.1.** Diagram of genetic circuitry of the proposed bi-stable switch system

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# Legal Frameworks

# Human Factors

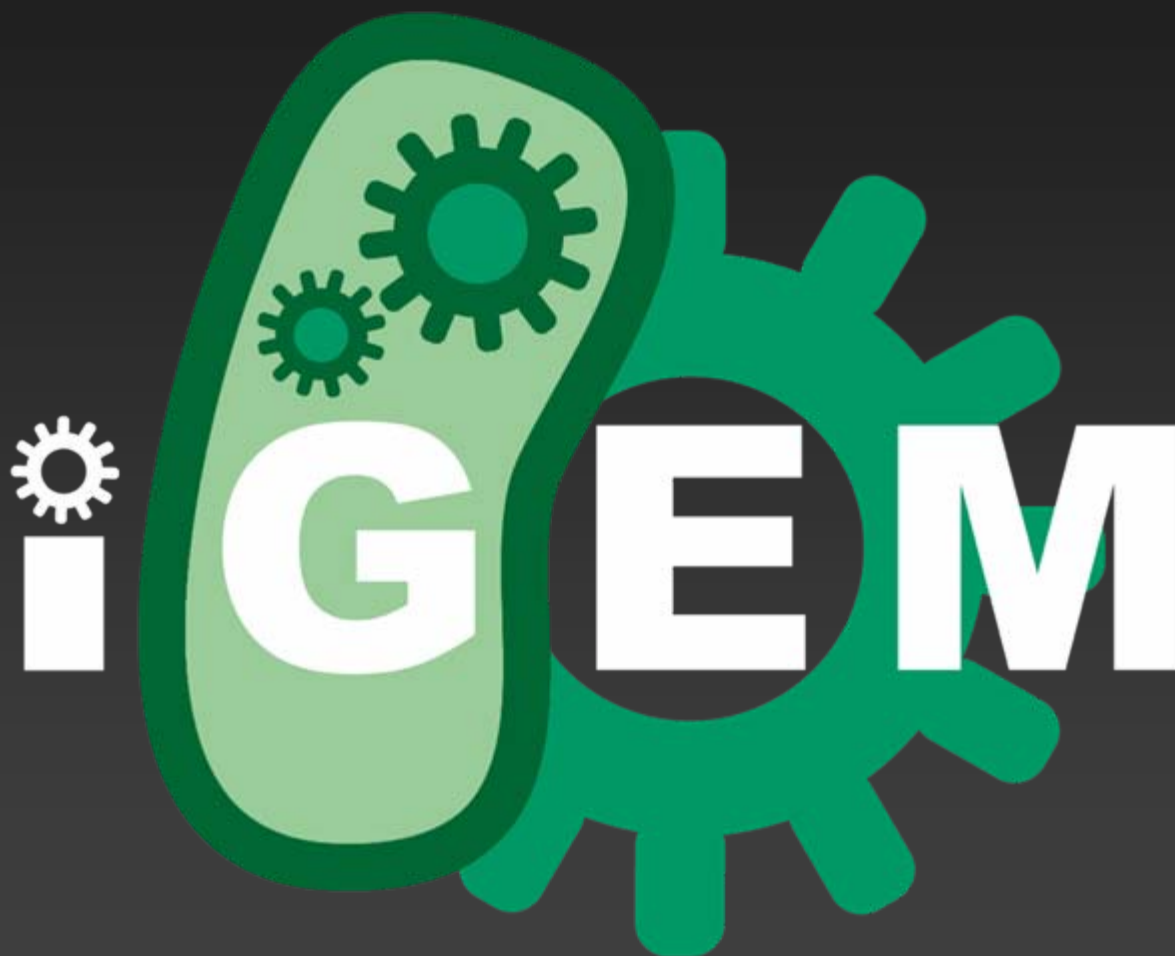
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## iGEM 2007 Wiki

International Genetically Engineered Machine Competition

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### iGEM Boogie

A mass of anxious iGEMMERS join a spontaneous dance on stage while waiting for the final Grand Prize decision from the Judges.

### Publish Your Project



IBE's [Journal of Biological Engineering](#)  
abstracts due Dec. 21!!!

Find more info on the [Publish](#) page

## iGEM 2007 has officially concluded



And it was a great success! 54 teams from around the world spent their summer engineering novel biological machines using and creating BioBrick standard biological parts, then gathered in November at the 2007 Championship

### iGEM?

Hundreds of undergraduates all over the world spend their summer making Synthetic Biology a reality by participating in the annual International Genetically Engineered Machine competition.

iGEM through the years

- [2008](#)
- [2007](#)
- [2006](#)

[Learn More](#)

### iGEM in the News

Here are some recent publications about iGEM teams:

- **Slate:** [Students from around the world synthesize new forms of life at the iGEM Jamboree](#)
- **NY Times:** [English, Algebra, Phys Ed ... and Biotech](#)
- **San Francisco Chronicle:** [High](#)



# http://parts.mit.edu

Main Page - Registry

http://parts.mit.edu/

GCaL EG@OWW MIT WM@MIT Parts iGEM PubMed Weather iK

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## Registry of Standard Biological Parts

jump to part  
BBa\_

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- Recent part changes

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- Add a Part
- Part Searches
- DNA Repositories
- Sequence Analysis
- Assembly Tool
- Help

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Browse Parts by Type

Browse Parts by School

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Help & Documentation

Users & Groups

### Registry Toolbox

- Add a part
- Search Parts
- DNA Repositories
- Sequence Analysis

### Latest News

- [8/01/06] We have contact information for the creators of parts. You can access this information when you access "Hard Information" of a part.
- [8/01/06] A table made for [yeast parts](#) is now available on the [Part Types](#) page

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## Main Page



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## OWW News

[OpenWetWare Announces ROD: ?Research on Demand?](#)

2008-04-01 13:30:17 EDT[[OpenWetWare](#)]

OWW is pleased to announce a new addition to our system: ROD: "Research on Demand". ROD enables the creation of research results that meet the demands of your publication and graduation schedules. By design, ROD is never 100% correct, and includes errors as subtle or as blatant as you would find in actual research results. [ 1 ]

# Questions!

[diybio.org](http://diybio.org)

[igem.org](http://igem.org)

[parts.mit.edu](http://parts.mit.edu)

[openwetware.org](http://openwetware.org)

[biohack.sf.net](http://biohack.sf.net)