Coming up with a project

Talk to people



New organisms

New parts and tools for future teams

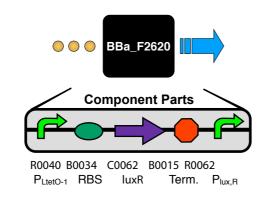
Most commonly used parts:
B0015 - a terminator
F2620 - an inducible promoter
B0034 - a RBS
R0011 - lac promoter
Plasmid backbones

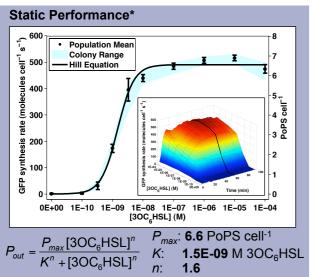
BBa_F2620

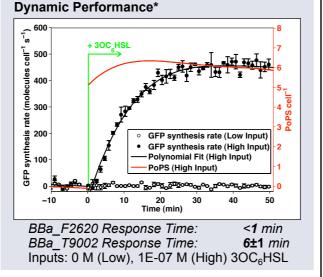
 $3OC_6HSL \rightarrow PoPS$ Receiver

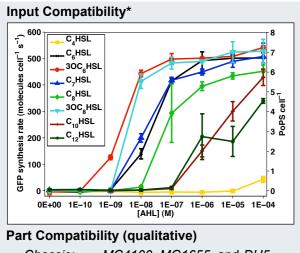
Mechanism & Function

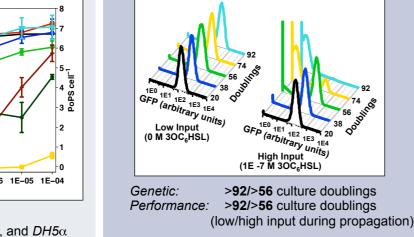
A transcription factor (LuxR) that is active in the presence of a cell-cell signaling molecule (3OC $_6$ HSL) is controlled by a regulated operator (PLtetO-1). Device input is 3OC $_6$ HSL. Device output is PoPS from a LuxR-regulated operator. If used in a cell containing TetR then a second input such as aTc can be used to produce a Boolean AND function.











Reliability**

Chassis: MC4100, MG1655, and DH5 α Plasmids: pSB3K3 and pSB1A2 Devices: E0240, E0430 and E0434

Transcriptional Output Demand (low/high input) *Nucleotides*: **0** / **6xNt** nucleotides cell⁻¹ s⁻¹

Polymerases: 0 / 1.5E-1xNt RNAP cell-1

(Nt = downstream transcript length)

Conditions (abridged)

Output: PoPS measured via BBa_E0240

Culture: Supplemented M9, 37°C

Plasmid: pSB3K3 Chassis: MG1655

*Equipment: PE Victor3 multi-well fluorimeter

**Equipment: BD FACScan cytometer

Authors: Barry Canton Ania Labno Updated: March 2008

Registry of Standard Biological Parts
making life better, one part at a time

License: Public

http://parts.mit.edu/registry/index.php/Part:BBa_F2620

Signaling Devices

Reuse and existing parts

Sunday, May 23, 2010

Let the students choose

Help them make smart choices

- Figure out what's practical: How many assembly stages could the team possibly do over the course of the summer? That sets an upper limit to the size of the system.
- Design the project so that different modules can be done in parallel.
- It doesn't have to be a brand new idea.

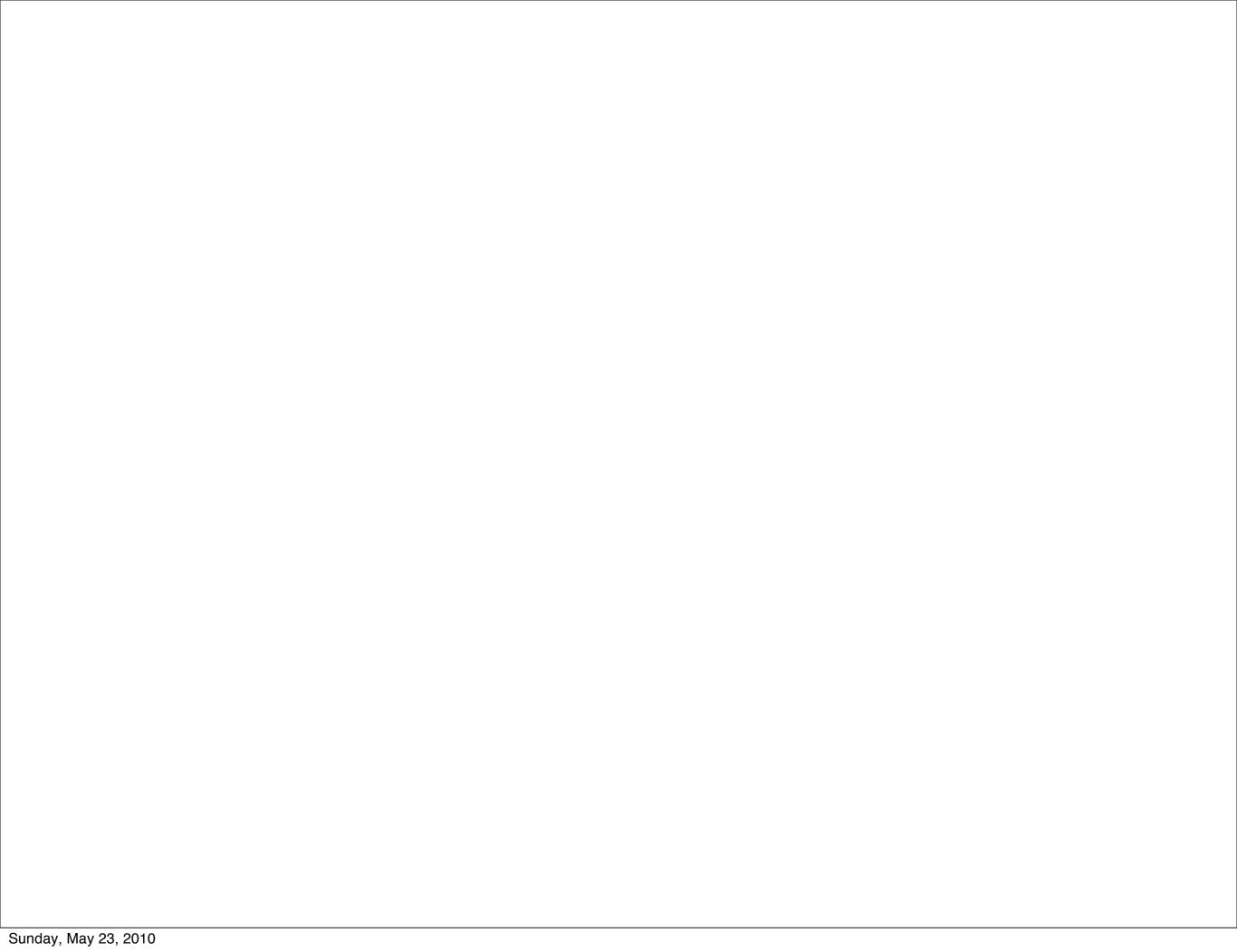
Describe your project on your team wiki



Sunday, May 23, 2010

Measurement

- Only some parts in the Registry have characterization data
- It can be hard to <u>compare the</u> <u>measurements</u> we do have
- Registry tour: <u>BBa_F2620</u> and <u>Anderson</u> families
- Controls
- Where to put the data: on the Registry

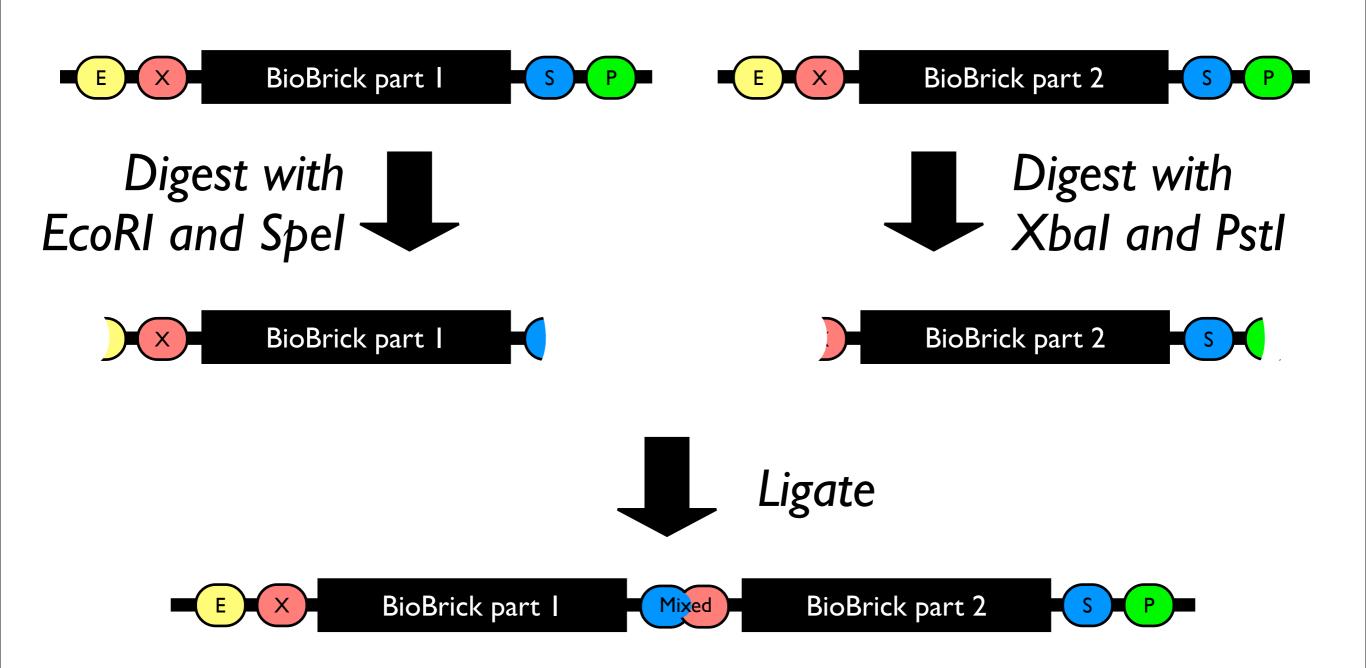


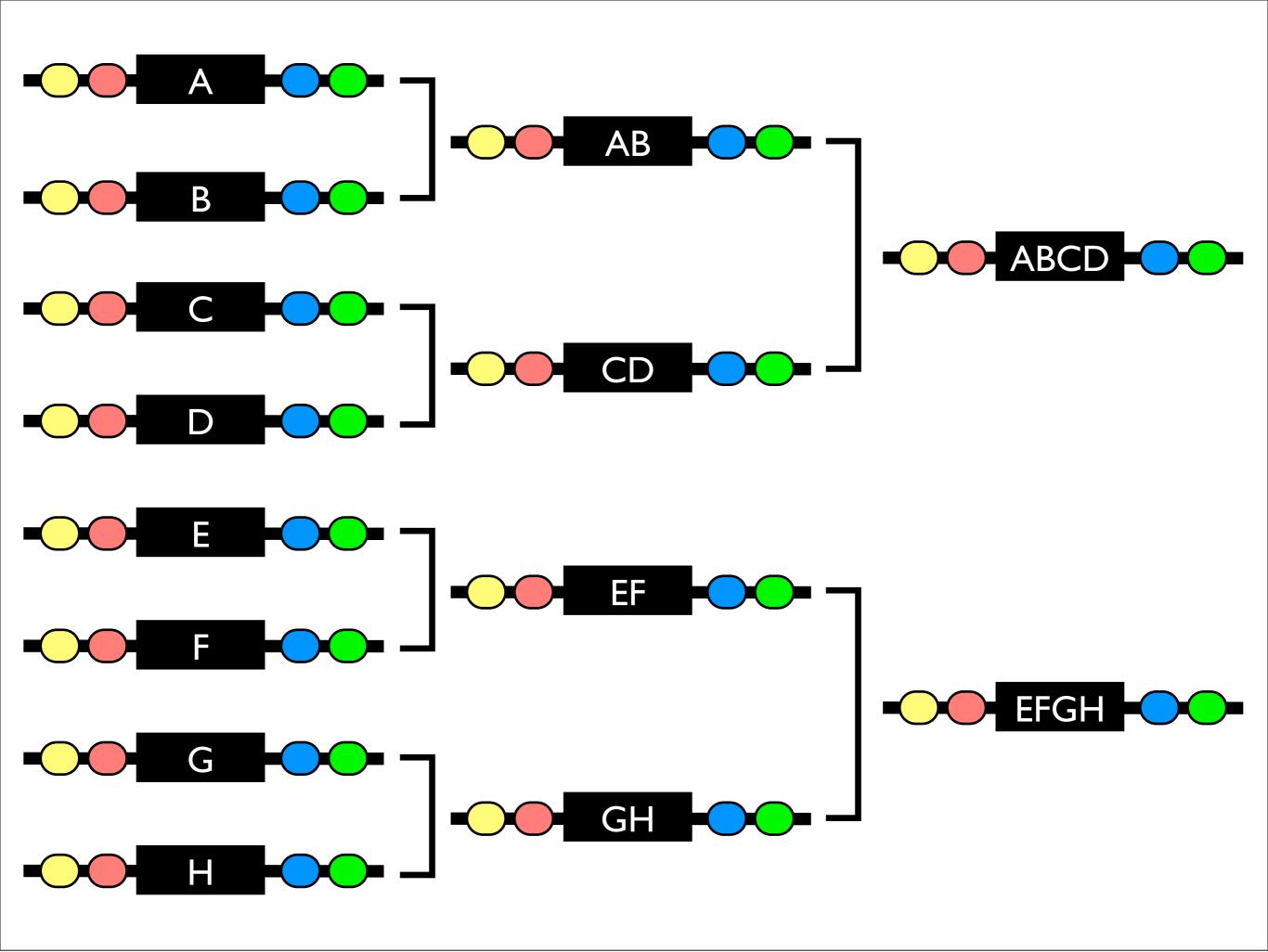
Standard assembly

BioBrick standard parts



BioBrick standard assembly





Why use the BioBrick standard?

- It is faster to build multi-part systems
- Assembling every two parts is the same
- You can reuse parts from the Registry
- Other people can reuse your parts
- It is required to win a prize at iGEM!

BioBrick™ Assembly Kit







Get the enzymes cheaper

BioBrick[™] Assembly Manual

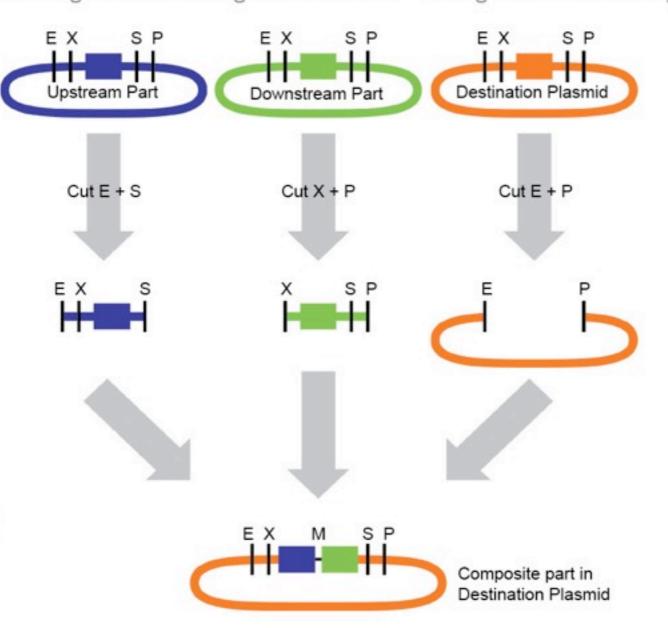




This manual describes the major steps of BioBrick form a circularized plasmid containing the composite assembly using BioBrick Assembly Standard 10. The part. The product of the ligation reaction can be used input to the protocol is DNA for the two parts to be to transform competent cells with the composite part. assembled and a destination plasmid. The manual in- To read more about the BioBrick system and browse cludes protocols for the digestion of the three input the BioBrick collection, visit the Registry of Standard DNA molecules and the ligation of the digested DNA to Biological Parts at http://partsregistry.org.

E=EcoRI-HF™ X=Xbal S=Spel P=Pstl M=Mixed site

BioBrick assembly overview



Start with two BioBrick parts and a BioBrick destination plasmid. The destination plasmid contains a toxic gene, ccdB, in the BioBrick cloning site and a different antibiotic resistance marker to the upstream and downstream parts.

Digest each of the parts with the appropriate restriction enzymes.

Mix the digests together and perform a ligation step. One of the ligation products formed will be the correctly assembled composite part in the destination plasmid. You can use the ligation mix to transform competent cells with the new composite part.

The BioBrick™ Assembly Kit from NEB and Ginkgo BioWorks has been designed for use with this manual. Download this manual from http://ginkgobioworks.com/support

Version 1.0

materials

consumables

▲ Restriction enzymes (EcoRI-HF, Xbal, Spel, Pstl), NEBuffer 2, BSA

10X T4 DNA Ligase Reaction Buffer, T4 DNA Ligase

H₂O (not shown)

Small PCR tubes

2 μl, 200μl pipet tips

Destination plasmid as purified DNA

Upstream and downstream parts as purified DNA

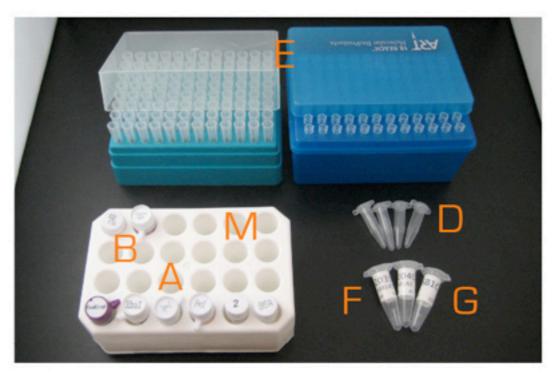
equipment

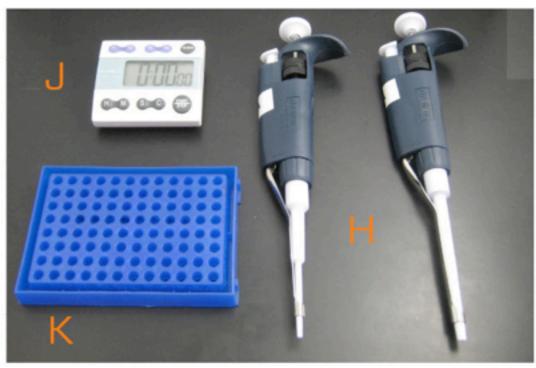
2 μl and 20 μl pipet

Incubator/water bath/thermocycler capable of holding 37°C and 80°C (not shown)

Timer

Rack for small PCR tubes





digest

This protocol assumes you have purified DNA for each of the BioBrick parts you want to assemble and also that you have purified DNA for the destination plasmid. The DNA could be produced from a DNA miniprep or a PCR amplification from a template. If the DNA was produced via a PCR amplification, the protocol assumes the DNA has been purified from the PCR enzymes that can reduce ligation efficiency.

The destination plasmid must have a different antibiotic resistance than the plasmids carrying the parts to be assembled, otherwise, many of the colonies obtained after transformation of competent cells will contain the input BioBrick parts, and not the composite BioBrick part. The toxic gene in the BioBrick cloning site of the destination plasmid ensures that cells transformed with undigested destination plasmid will not grow.

To each tube, add H₂O and 500 ng of the part or plasmid to be digested. Adjust the amount of water you add such that the total volume in each tube is 42.5 µl.

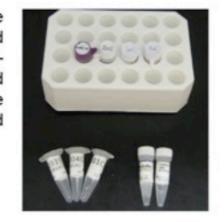


Add 5 µl of NEBuffer 2 to each tube.



prepare reaction mix

Remove the DNA for the upstream part, the downstream part and the destination plasmid along with NEBuffer 2 and BSA from the freezer to thaw. Thawing is fast if the tubes are immersed in room temperature water. You can also remove the enzymes from the freezer but leave them in a cold box so they remain close to -20°C.



5 Add 0.5 μl of BSA to each tube.



Add 1 µl of the first appropriate* restriction enzyme to each tube**.



You will need three PCR tubes, one for the digest of the upstream part, one for the downstream part, and one for the destination plasmid. You should label each tube (for example, U, D, P, for upstream part, downstream part, and destination plasmid respectively).



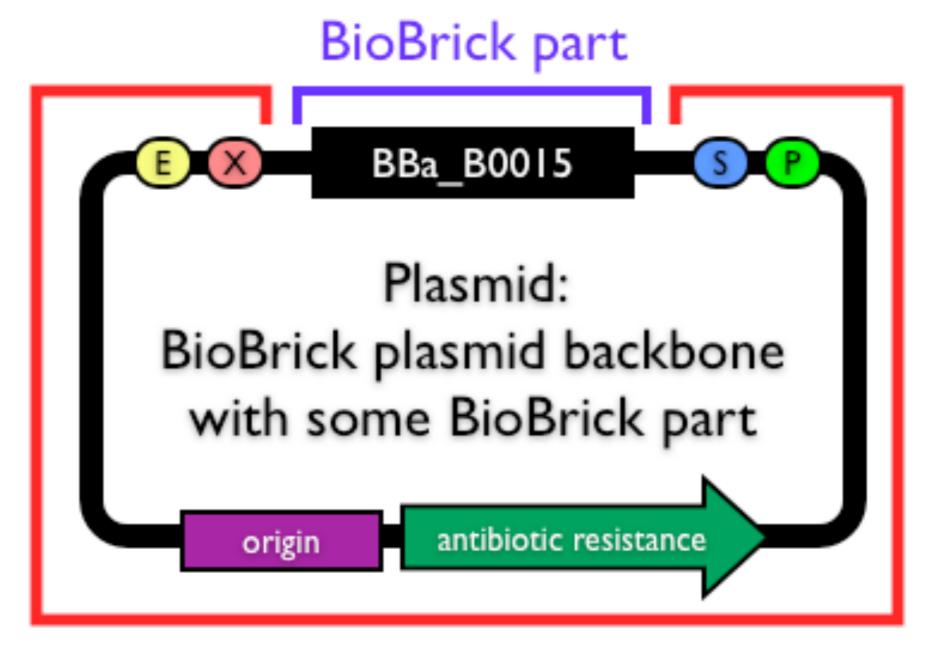
7 Add 1 µl of the second appropriate* restriction enzyme to each tube**.

Version 1.0

^{*} See the overview diagram on Page 1 for the appropriate restriction enzymes for each part and the destination plasmid.

^{**} When pipeting restriction enzyme, only touch the very end of the pipet tip into the restriction enzyme. Restriction enzymes are stored in a high percentage glycerol solution that sticks to the outside of the pipet tip. If you dip the tip deeply into the restriction digest you will add much more restriction digest than needed as well as increase the glycerol concentration of the digest mix. A high glycerol concentration (>5%) can result in non-specific cutting of the DNA (referred to as "star activity").

Plasmid backbones



BioBrick plasmid backbone