Coming up with a project

Teach the Teachers Workshop
May 16, 2009

Talk to people

Previous iGEM projects

igem.org

New organisms

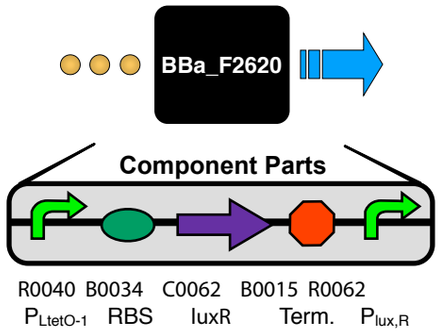
New parts and tools for future teams

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

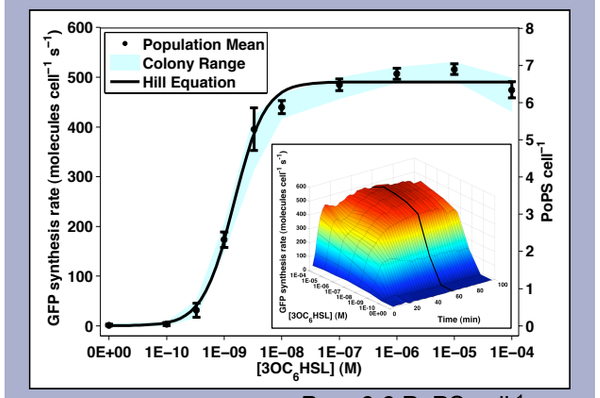
BBa_F2620

3OC$_6$HSL → 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 (P$_{LtetO-1}$). 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.



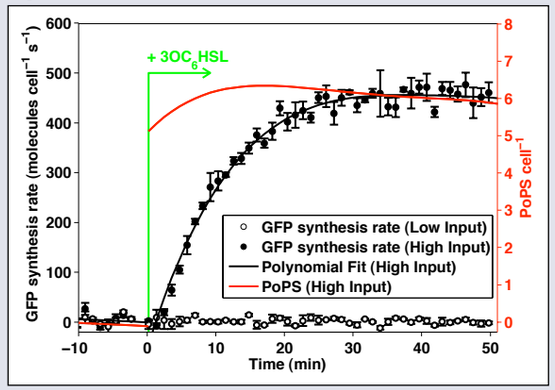
Static Performance*



$$P_{out} = \frac{P_{max}[3OC_6HSL]^n}{K^n + [3OC_6HSL]^n}$$

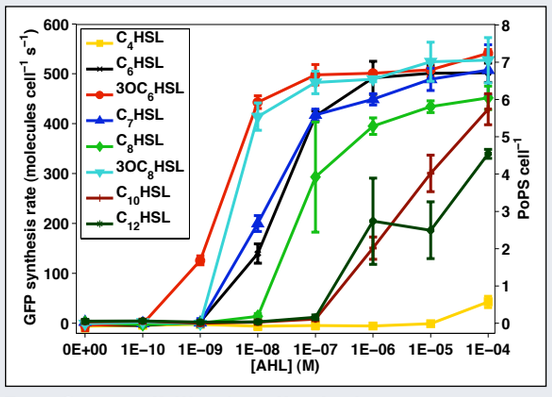
P_{max}: **6.6** PoPS cell^{-1}
K: **1.5E-09** M 3OC$_6$HSL
n: **1.6**

Dynamic Performance*

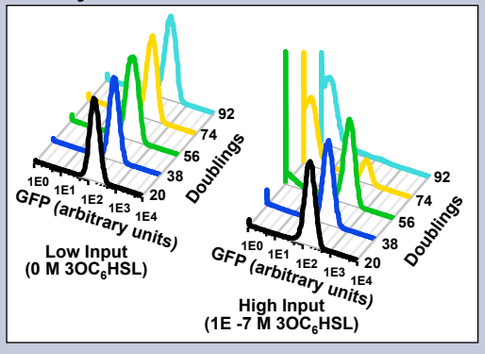


BBa_F2620 Response Time: ***<1** min*
BBa_T9002 Response Time: ***6±1** min*
Inputs: 0 M (Low), 1E-07 M (High) 3OC$_6$HSL

Input Compatibility*



Reliability**



Genetic: >**92**/>**56** culture doublings
Performance: >**92**/>**56** culture doublings
(low/high input during propagation)

Part Compatibility (qualitative)

Chassis: *MC4100*, *MG1655*, and *DH5α*
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, 37ºC
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

Reuse and existing parts

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

Teach the Teachers Workshop
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Standard assembly

Teach the Teachers Workshop
May 16, 2009

BioBrick standard parts

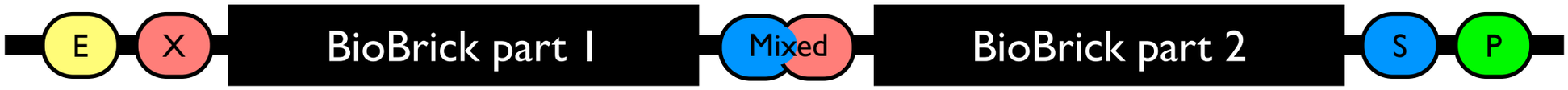
Knight, 2003

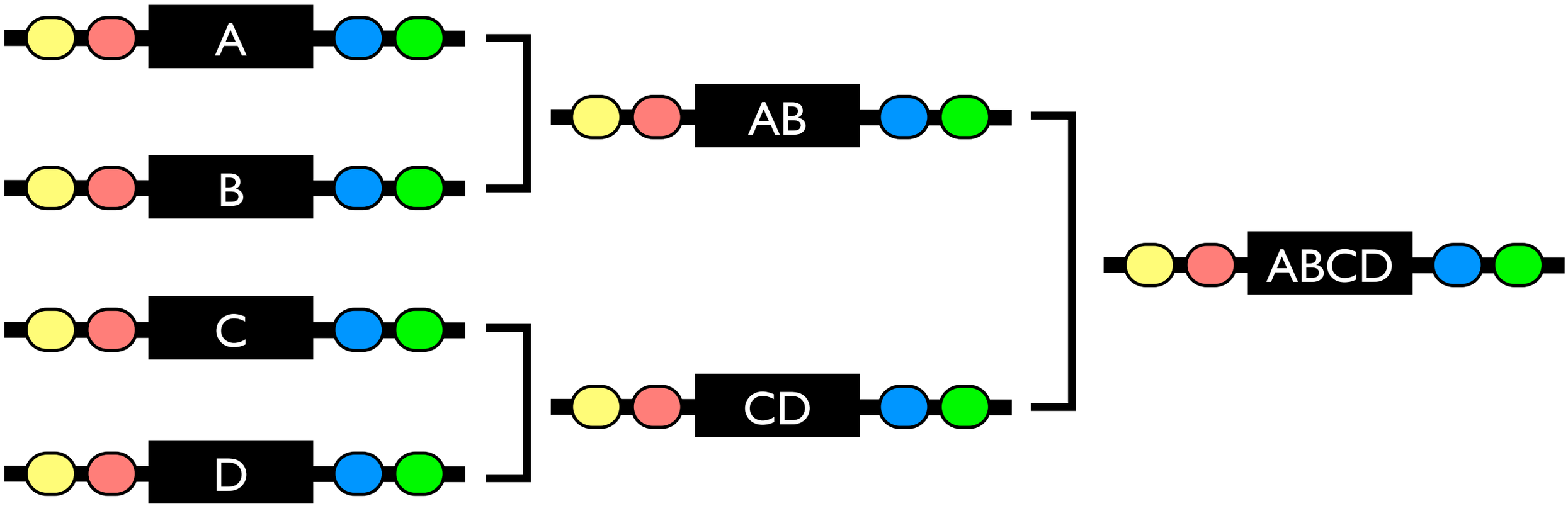
BioBrick standard assembly

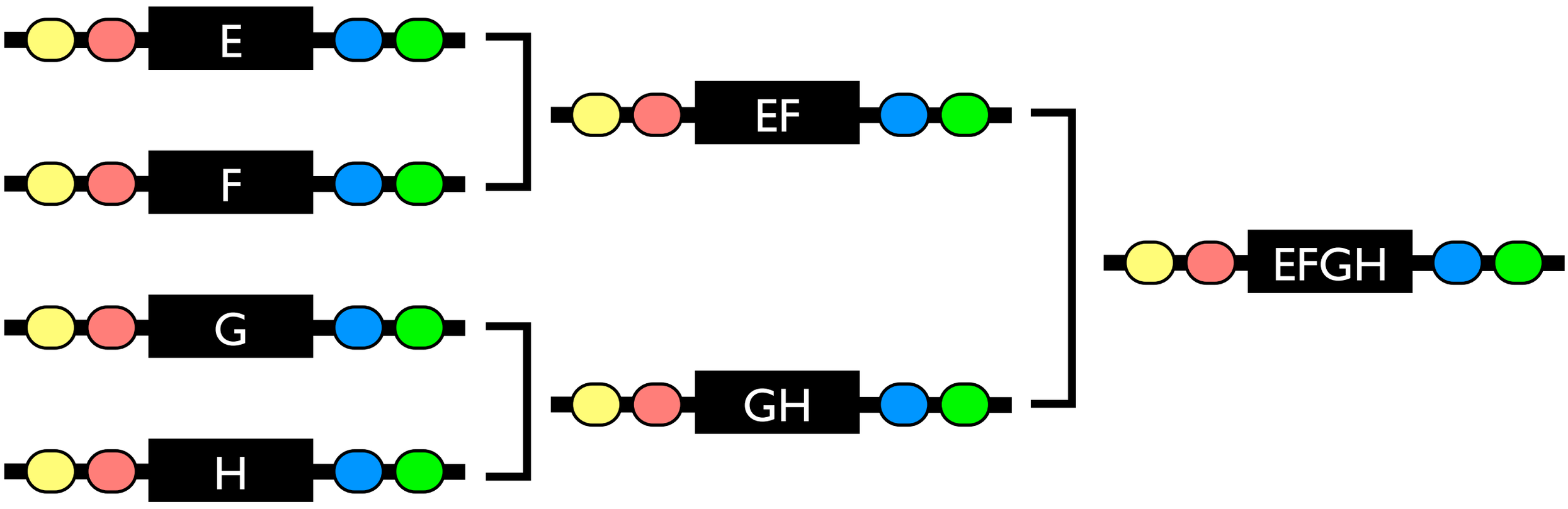
Digest with EcoRI and SpeI

Digest with XbaI and PstI

Ligate




A
B
C
D
AB
CD
ABCD


E
F
G
H
EF
GH
EFGH

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

ginkgobioworks

NEW ENGLAND BioLabs Inc.

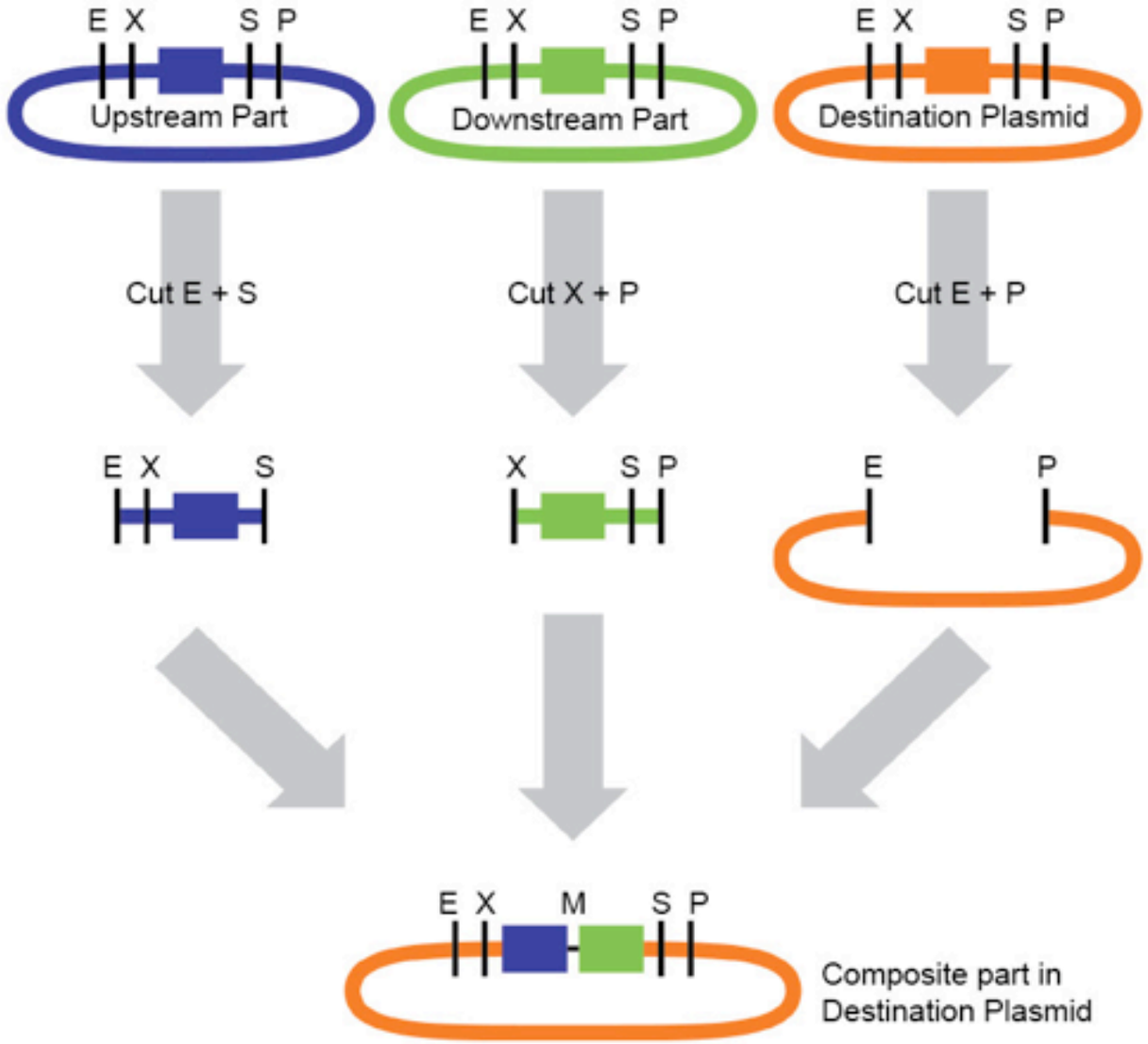
Get the enzymes cheaper

BioBrick™ Assembly Manual

ginkgobioworks

NEW ENGLAND BioLabs Inc.

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



E=EcoRI-HF™
X=XbaI
S=SpeI
P=PstI
M=Mixed site

BioBrick assembly overview

1 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.

2 Digest each of the parts with the appropriate restriction enzymes.

3 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

materials

consumables

A Restriction enzymes (EcoRI-HF, XbaI, SpeI, PstI), NEBuffer 2, BSA

B 10X T4 DNA Ligase Reaction Buffer, T4 DNA Ligase

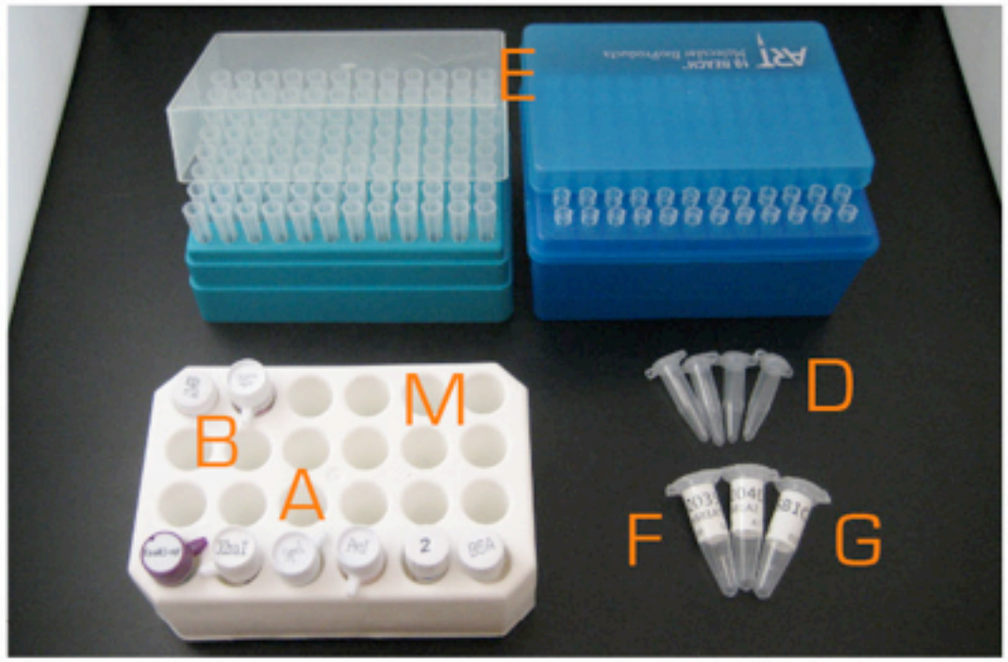
C H_2O (not shown)

D Small PCR tubes

E 2 µl, 200µl pipet tips

F Destination plasmid as purified DNA

G Upstream and downstream parts as purified DNA



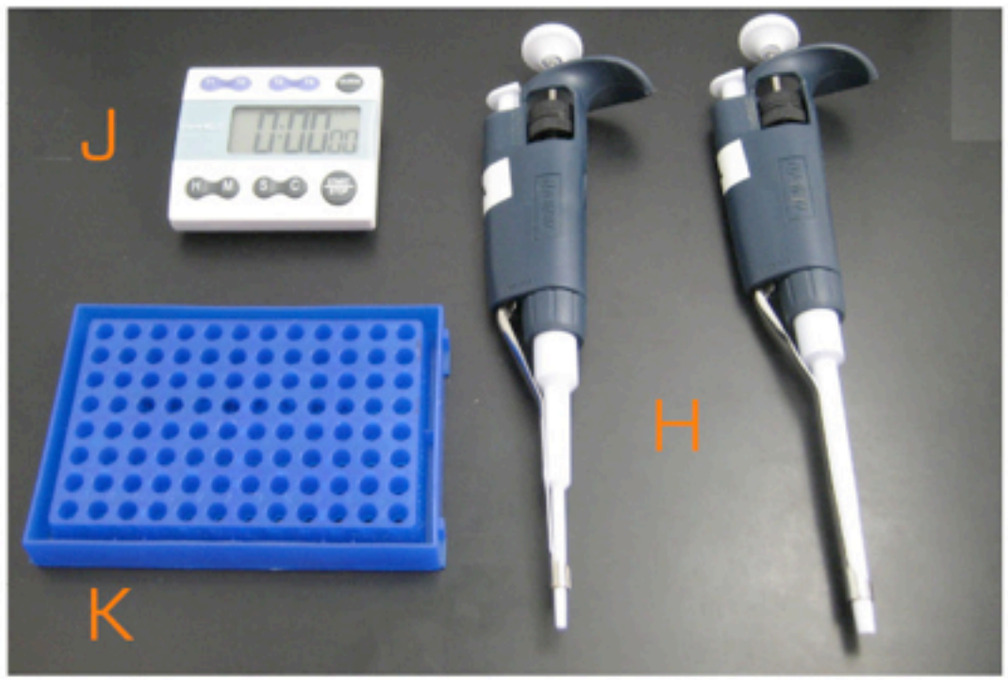
equipment

H 2 µl and 20 µl pipet

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

J Timer

K Rack for small PCR tubes



ginkgobioworks

BioBrick™ Assembly M

digest

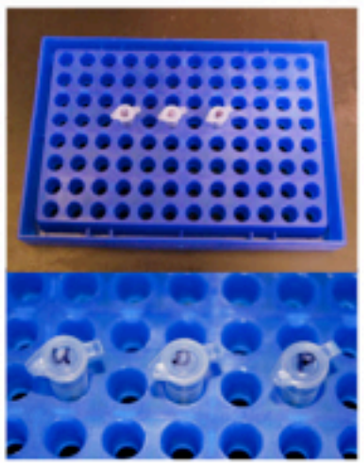
! This protocol assumes you have purified DNA for each of the BioBrick parts you want to assemble and also that you have purified DNA for the destination plasmid. The DNA could be produced from a DNA miniprep or a PCR amplification from a template. If the DNA was produced via a PCR amplification, the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

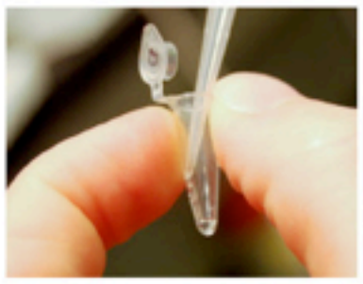
prepare reaction mix

1 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.

2 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).



3 To each tube, add H_2O 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.



4 Add 5 µl of NEBuffer 2 to each tube.

5 Add 0.5 µl of BSA to each tube.

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

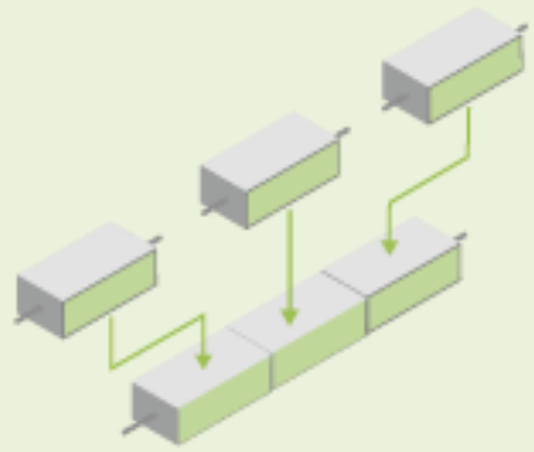
7 Add 1 µl of the second appropriate* restriction enzyme to each tube**.

* 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").

brick layer

A fast, reliable, BioBrick assembly service from the Ginkgo bots



The details

- All assemblies use BioBrick™ assembly standard 10.
- Orders can be submitted via email.
- Ginkgo engineers will consult on the selection of parts.
- Assemblies are sent in two weeks. Faster assembly is available for standard registry parts.

Coming Summer 2009

Keep me informed...

Email address

Submit

Plasmid backbones

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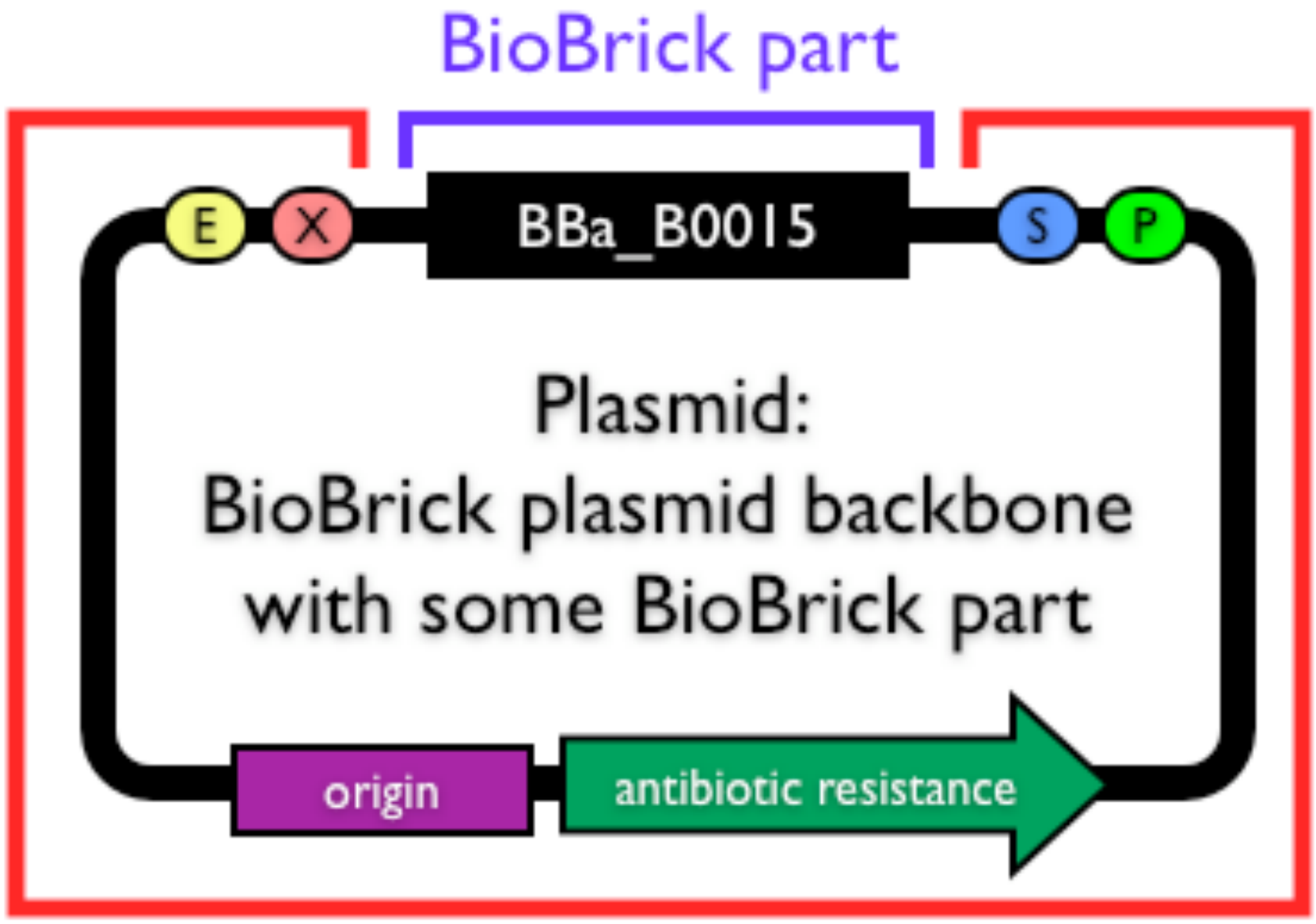

BioBrick plasmid backbone