

# **Gibson assembly**

# **Ruihong Wang**

### **Abstract**

Gibson is a PCR-based multi-fragment assembly technique. So whether PCR works has large impacts on final product.

Citation: Ruihong Wang Gibson assembly. protocols.io

dx.doi.org/10.17504/protocols.io.mzpc75n

Published: 15 Feb 2018

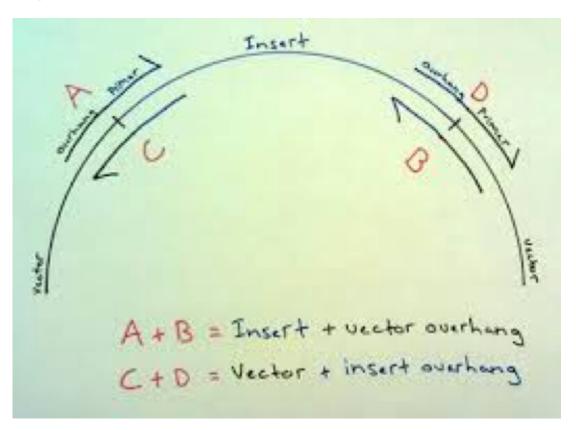
#### **Materials**

► Gibson Assembly Master Mix - 10 rxns <u>E2611S</u> by <u>New England Biolabs</u>

## **Protocol**

## Primer design

# Step 1.



Step 2.

Construct a complete map of insert in the desired destination vectors in Snapgene or Ape.

## Step 3.

Identify the connection region between inserts and vector.

## Step 4.

Design primer for amplifying insert and vectors. Target melting temperature Tm set to 60 or higher.

#### Step 5.

Order primers A, B, C and D.

Notes: A has two parts. Starting from 5', the overhang is the reverse complement of Rev primer of backbone, and the primer region is a standard primer to amplify insert (Fwd primer of insert). B is same and has reverse complement of backbone Fwd primer followed by standard Rev primer of insert.

So primer A= Backbone Rev primer (rev complement)+ Insert Fwd primer

Primer B= Backbone Fwd primer (rev complement)+ Insert Rev primer

Primer C is reverse complement of A.

Primer D is reverse complement of B.

#### Step 6.

For more fragment to assemble, increase number of primers correspondingly.

## PCR

#### Step 7.

Amplify insert with A and B primers. Amplify backbone with C and D primers. Use 25 or 50 uL reaction volume. Use DMSO or nested/ touchdown PCR if necessary.

#### Step 8.

Add 0.25 uL DpnI and incubate at 37 C for 1hr (or longer depending on your paranoid) to remove template.

#### Step 9.

Run a gel to verify the size of PCR product.

#### Step 10.

Use PCR cleanup or Gel extraction to get clean product.

#### **Step 11.**

Quantify the concentration by Nanodrop

# Gibson Assembly

#### Step 12.

#### **P** NOTES

## Ruihong Wang 02 Feb 2018

For 2 or 3 fragments assembly, use 0.02-0.5 pmols of DNA. For more fragments, use 0.2-1.0 pmols.

pmols=(weight in ng) x 1000 / (base pairs x 650 daltons)

## **Step 13.**

Put insert and vectors into reaction and add DI water to 10 uL.

## **Step 14.**

Add 10 uL 2X Gibson Master mix.

## **Step 15.**

Incubate at 50 C for 15-60 minutes depending on the fragments number. 2-3 fragments use 15 mins. 4-6 use 60 mins.

### **Step 16.**

Store in -20 C.

#### **Transformation**

### **Step 17.**

Mix 1 uL of reaction and 10 uL of competent cells, do not vortex. Keep on ice for 30 mins.

### **Step 18.**

Heat shock in 42 C water bath for 45 sec.

## Step 19.

Keep on ice for 5 mins.

## Step 20.

Add 300 uL SOC (pre-warm to Room temperature). Put in 250 rpm shaker in 37 C for 1hr.

#### Step 21.

Use 50-100 uL of recovered solution to plate out depending on copy number and plate size.

#### Step 22.

Harvest colony next day and pick single colony for liquid culture in corresponding antibiotics.

#### Step 23.

Sequencing and digestion to verify the assembly.