

Fabrication of DNA constructs by Gibson Assembly and Golden Gate reactions

Tamara Matute, Isaac Nuñez, Peter Von Dassow, Fernan Federici

Abstract

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Protocol

Introduction and rationale

Step 1.

Golden Gate assembly creates vectors by combining vectors containing level 0 parts (e.g. promoters, CDSes, terminators) and an acceptor vector. The first step involves the creation of libraries of level 0 parts by 'domesticating' DNA sequences of interest (e.g a promoter or a new fluorescent protein).

Domestication of DNA sequences into level 0 part vectors

Step 2.

Level 0 parts for Golden Gate reactions were generated by Gibson Assembly. Parts can be generated by PCRing natural sequences (e.g. genomic DNA) or existing vectors; or by fully synthesising (e.g. gblocks from IDT) DNA fragments of interest.

For PCRed parts, reactions were performed with Phusion® High-Fidelity DNA Polymerase following this conditions:

Thermocycling

98°C x 30 sec

 $(98^{\circ}\text{C} \times 10 \text{ sec} + 60^{\circ}\text{C} \times 30 \text{ sec} + 72^{\circ}\text{C} \times 2 \text{ min}) \times 35 \text{ cycles}$

72°C x 10 min

4°C

PCR mix:

1ul primer forward Gibson

1ul primer reverse Gibson

1,33ul template (concentration: 1ng/ul)

8 ul Buffer HF 5X

0,8ul dNTPs (10uM)

0,4ul Phusion® High-Fidelity DNA Polymerase (2U/ul)

27,46 ul H20

Commercially synthetisised DNA fragments were ordered containing the 20-30bp overhangs used for Gibson assembly into the level 0 part vector.

Gibson assembly step for the creation of level 0 part vector

Step 3.

Desired PCR products were identified by gel electrophoresis and extracted with the purification kit Wizard SV Gel & PCR Clean-Up System (Promega). DNA concentration were corroborated with a Take3[™] Micro-Volume Synergy[™] HTX before performing Gibson reactions.

The volumes used for each part were calculated by eq 1, 2 and 3.

Eq.1

$$X [pMol/\mu l] = \frac{Concentration [ng/\mu l] * 1000}{650 [\frac{gr/mol}{bp}] * Lenght [bp]}$$

Eq.2

$$Vol_{ratio\ i} = \frac{X}{pmol_{reg}}$$

Volume ratio [ul] = pmol/required pmol

where required pmol is:

0,01 for parts >2500 bp

0,03 for parts >200 bp and <2500 bp

Volume to be added was calculated as:

Eq.3

$$Vol_{piece\ i} = rac{Vol_{total}*Vol_{ratio\ i}}{\sum\limits_{i}^{n}Vol_{ratio\ i}}$$

1,5 μ l of parts were combined with 4,5 μ l of Gibson Master Mix in ice. This reaction was incubated at 50 $^{\circ}$ C for 1 hour.

Reagents:

Gibson Mix (1.33X):

100 ul 5X Isothermal Buffer

2 ul T5 Exonuclease (1 U/ul)

6,25 ul Phusion® High-Fidelity DNA Polymerase (2U/ul)

50 ul Taq DNA ligase (40U/ul)

216,75 ul H20

Store at -20°C.

5X Isothermal Buffer:

25% PEG-8000

500 mM Tris-HCl pH 7,5

50 mM MgCl2

50 mM DTT

1 mM dATP

1 mM dTTP

1 mM dCTP

1 mM dGTP

5 mM NAD

H20

Store at -80°C

Transformation of Level 0 donor vectors

Step 4.

These reactions were transformed into chemically competent TOP10 cells prepared by the OOW protocol (http://www.openwetware.org/wiki/TOP10_chemically_competent_cells), minipreped and sequenced. These level 0 parts were stored as vectors for the following Golden Gate reactions.

Golden Gate assembly of promoters and fluorescent proteins combinations **Step 5.**

Final combinatorial assemblies of promoters, CDSes and terminators here described were produced by Golden Gate. For this, a general aceptor vector was created that accepts level 0 parts for the assembly of transcriptional units (e.g. promoter-CDS-terminator of interest).

Eq.4

$$Vol_{required}[ul] = \frac{X * Length *650}{[concentration] *10^6}$$

The assembly was performed as follows:

 $(37^{\circ}C \times 3 \text{ min} + 16^{\circ}C \times 5 \text{ min}) * 40 \text{ cycles}.$

50°C x 5 min.

4°C

Transformation of Golden Gate reactions

Step 6.

 $3\mu l$ of these reactions were used to transform TOP10 Escherichia coli cells following the step described above.

Positive colonies were identified by colony PCR and sequenced.