

Golden Gate Assembly

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10 μ L reaction is transformed to 50 μ L chemically competent cell. Then stalled cell in sawed ice for 30 min, and allowed to recover for an hour in SOC medium.

1 Introduction

Pending...

2 Protocol

2.1 Reaction System

As an example, here, we use BsaI (type IIs restriction enzyme) in this reaction system. Below is an typical reaction system which is 20 μ L.

| Components | Volume |
|------------------------------|---|
| Fragments | 40 fmol or 100 ng ea. ⁽¹⁾ |
| BsaI | 10 U (2 μ L) |
| T4 ligase | 5 U (0.5 μ L) ⁽²⁾ |
| 10 \times T4 Ligase Buffer | 2 μ L |
| ddH ₂ O | Up to 20 μ L |

Table 1: Golden Gate reaction system

Note:

- (1) When do 1 ~ 2 insertions reaction, it is convenient adding 100 ng of each fragment into reaction. If performing multiple (> 3 insertions) ligation, we recommend adding 40 fmol of each fragment.
- (2) If assembling more than 3 fragments, high concentration of T4 ligase is necessary. I recommend 20 U (for NEB CEU, 1 U = 200 CEU).
- (3) For convenience, we often prepare 10 μ L reactions and the volume of each enzyme is 1 μ L.

2.2 Ligation Procedure

If performing one fragment insertion ligation, incubating reaction system 30 min at 37 °C. When insertions are more than 2, we recommend using thermocycler: 37 °C for 2 min, the 16 °C for 3 min, repeat for 10 - 100 cycles, depending on the number of insertions (10 cycles per insertion). Following, the reaction is heated to 50 °C for 5 min, which is for digesting non-specific ligation events, completely. Last, the reaction is incubated 5 min at 80 °C for inactivating the reaction enzyme. The

3 Design

3.1 Primer Design

The flank sequences of primer for introducing enzymatic cutting site into double sites of insertions are collected in Table 2.

| Enzyme Name | Sequence |
|-------------|--------------------------------|
| BsaI | GCATTAGGTCTCCNNNN |
| BbsI | AAGTGCGAAGACCANNNN |
| BfuAI | GGCAATACCTGCGTGANN NN |
| BtgZI | CGAATGGCGATGTTG TACTGCCNNNN |
| SapI | GGAATCGCTCTTCCNNN |
| BsmBI | ATAGCGCGTCTCCNNNN |

Table 2: Flank Sequence of Primer, the underlined nucleotides are denoted as the enzyme recognition sequence, and the sequence before recognition sequence is protection sequence, and the base N is denote as the sticky end.