

# Gibson Assembly® HiFi HC 1-Step Kit and Master Mix (4X)



## Quick Reference Manual

For additional product information, an online primer design tool, and detailed protocol visit [sgidna.com/hifi\\_kit](http://sgidna.com/hifi_kit).

**Catalog Numbers** GA1100-4XS, GA1100-4X10, GA1100-4X50, GA1100-4X10M, GA1100-4X50M, GA1100-4XB03, GA1100-4XB05, GA1100-4XB10, GA1100-4XB15, GA1100-4XB20, GA1100-4XB25

## Products and Storage Conditions

### Gibson Assembly® HiFi 1-Step Kit

Quantity	Component	Cat. GA1100-4XS (5 Reactions)	Cat. GA1100-4X10 (10 Reactions)	Cat. GA1100-4X50 (50 Reactions)	Storage Temperature
		Volume			
1 tube (each)	GA HiFi HC 1-Step Master Mix A (4X)	25 µL	50 µL	250 µL	Aliquot and store at –20°C
	GA Positive Control	10 µL (2 Control Rxns)	10 µL (2 Control Rxns)	25 µL (5 Control Rxns)	

### GA HiFi HC 1-Step Master Mix (4X)

**IMPORTANT:** Upon receipt, place GA HiFi HC 1-Step Master Mix (4X) on ice to thaw. Briefly vortex and centrifuge the thawed master mix. Then, aliquot the master mix to reduce the number of freeze-thaw cycles. Properly aliquoted GA HiFi HC 1-Step Master Mix is stable for up to 6 months when stored at –20°C.

Catalog Number	Volume	Number of Reactions
GA1100-4X10MM	50 µL	10
GA1100-4X50MM	250 µL	50
GA1100-4XB03	3 mL	1200
GA1100-4XB05	5 mL	2000
GA1100-4XB10	10 mL	4000
GA1100-4XB15	15 mL	6000
GA1100-4XB20	20 mL	8000
GA1100-4XB25	25 mL	10,000

### Guidelines for Assembly

- For a typical Gibson Assembly® HiFi HC 1-Step reaction, combine 25–50 ng of vector with approximately 10–300 ng of insert. For best results, we recommend balancing the molar ratio of the DNA fragments. For fragments >1 kb, use an equimolar ratio. For DNA fragments ≤1 kb, we recommend using a 5-fold molar excess of insert. To precisely determine the pmol or ng of DNA for a fragment of a given size, use the following formulas:

$$\text{pmol DNA} = [\text{ng DNA} / (660 \times \# \text{ of bases})] \times 1000$$

$$\text{ng of DNA} = [\text{pmol DNA} \times (660 \times \# \text{ of bases})] / 1000$$

Refer to the table on page 2 for approximate pmol of DNA for a given fragment size and amount.

- Keep GA HiFi HC 1-Step Master Mix (4X) on ice at all times.
- For the assembly of multiple fragments, create a master mix of fragments in the proper ratios to minimize pipetting error.

### Gibson Assembly® HiFi HC 1-Step Method

- Thaw GA HiFi HC 1-Step Master Mix (4X) on ice.
- Dilute your DNA fragments with nuclease-free water in PCR tubes to a total volume of 7.5 µL according to the “Guidelines for Assembly”.
- Vigorously vortex the master mix for 15 seconds immediately before use** after it is thawed.
- In a tube on ice, combine 7.5 µL of DNA fragments and 2.5 µL of GA HiFi HC 1-Step Master Mix (4X). Mix the reaction by pipetting up and down.
- (Optional) For the positive control, combine the following reagents in a tube on ice. Mix the reaction by pipetting up and down.

GA Positive Control	5 µL
Nuclease-free water	2.5 µL
GA HiFi HC 1-Step Master Mix (4X)	2.5 µL
- Vortex and spin down all reactions.
- Incubate the reactions at 50°C for 1 hour.
- After the incubation is complete, store the reactions at –20°C or dilute reactions for downstream applications such as PCR or *E. coli* transformation (see the protocols on the following pages).
- (Optional) Analyze the assembly reaction by performing gel electrophoresis with 5–10 µL of the reaction on an 0.8–2% agarose gel.

### Transformation Guidelines

We recommend transformation with *E. coli* 10G chemically competent cells (Lucigen Cat. No. 60107) or TransforMax™ EPI300™ Electrocompetent *E. coli*. (Lucigen Cat. No. EC300110). If you use competent cells other than the recommended cells, follow the transformation protocol provided with the competent cells. Use cells with a transformation efficiency  $\geq 1 \times 10^9$  CFU/µg pUC19.

Because some of the ingredients in the buffer mix can negatively impact the survival of some competent cells, we recommend diluting the assembly reaction before performing the transformation. Dilute HiFi HC assemblies up to 5-fold. You may need to empirically determine the optimal level of dilution, depending on the type of cells used.

## Transformation with *E. coli* 10G Chemically Competent Cells (Recommended)

1. Pre-chill 15 mL disposable polypropylene culture tubes (one tube for each transformation reaction).
2. Thaw cells on ice for 5–15 minutes.
3. Add 40  $\mu$ L of thawed, competent cells to each cold tube.
4. Add 2  $\mu$ L of the diluted assembly reaction to each cold tube of competent cells. Mix by briefly stirring (do not pipet up and down).
5. Incubate the cells and DNA on ice for 30 minutes. Do not mix.
6. Heat shock the mixture in a 42°C water bath for 45 seconds.
7. Return tubes to ice for 2 minutes.
8. Add 950  $\mu$ L room temperature recovery media to the cells in the culture tube.
9. Incubate the tubes with shaking at about 250 rpm for 90 minutes at 37°C to allow cells to recover.
10. Proceed to **Plating Procedure**.

## Transformation with TransforMax™ EPI300™ Electrocompetent *E. coli*

1. Add 1 mL SOC media to 1.5-mL microcentrifuge tubes (one tube per reaction). Place tubes on ice for 10 minutes.
2. Chill clean electroporation cuvettes on ice.
3. Pipet 30  $\mu$ L of EPI300™ cells directly between the slit of the cuvettes on ice (one cuvette per reaction).
4. Add 2  $\mu$ L of the diluted assembly reaction to the cells in the cuvette. Mix by pipetting up and down gently two times.
5. Incubate cuvette on ice for one minute.
6. Gently tap cuvette on a benchtop two times to make sure all contents are at the bottom of the cuvette in between the slit.
7. Insert the cuvette into a BioRad Electroporator or equivalent, and press PULSE. Pulse Settings for EPI300™ cells are 1200 V, 25  $\mu$ F, 200  $\Omega$ , 0.1 cm cuvette.
8. During the pulse ( $\approx$ 2 seconds), remove 800  $\mu$ L SOC from a pre-chilled 1.5 mL tube (step 1). Immediately add the SOC to the cuvette after the pulse.
9. Mix the cells and SOC by pipetting up and down. Add the mixture back into the tube containing the remaining SOC.
10. Incubate the cells for 1 hour at 37°C with shaking at 200 rpm.
11. Proceed to **Plating Procedure**.

Additional product information and resources are available at [sgidna.com/hifi\\_kit](http://sgidna.com/hifi_kit)

**Technical Services:** For technical assistance, please contact technical services at [techservices@sgidna.com](mailto:techservices@sgidna.com).

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Gibson Assembly® US Patent Nos. 7,776,532 and 8,435,736, and 8,968,999.

## Plating Procedure

1. Pre-warm LB plates in an incubator upside down for 10–15 minutes.
2. After the incubation, plate 1/10–1/100 of the transformation reaction (10–100  $\mu$ L out of 1 mL) onto LB agar plates with appropriate antibiotics. See “Recommended Plating Volume” below.
3. (Optional) For the positive control, plate 1/100 volume of the transformed reaction onto LB plates containing 100  $\mu$ g/mL ampicillin or carbenicillin with 40  $\mu$ g/mL X-Gal and 0.1 mM IPTG.
4. Incubate plates at 37°C upside down, overnight.
5. Pick colonies for screening.

## Recommended Plating Volume

Always plate two plates (one low and one high volume).

Number of fragments	Plating volume (fraction of the total transformation mixture)	For example, we typically plate*...
1–2	1/50	2 $\mu$ L and 20 $\mu$ L
3–5	1/10	10 $\mu$ L and 100 $\mu$ L

\*Based on a 1000  $\mu$ L transformation mixture

## Reference Material

### Amount of DNA to use in Gibson Assembly® reaction

Refer to the following table for approximate pmol of DNA for a given fragment size and amount.

Fragment size	ng of DNA	pmol of DNA
0.5 kb	20 ng	0.061
	40 ng	0.121
1 kb	10 ng	0.015
	25 ng	0.038
5 kb	10 ng	0.003
	25 ng	0.008
8 kb	25 ng	0.005
	50 ng	0.009
10 kb	25 ng	0.004
	50 ng	0.008
15 kb	50 ng	0.005
	100 ng	0.010
20 kb	50 ng	0.004
	100 ng	0.008
30 kb	50 ng	0.003
	100 ng	0.005