

Use the NEBuilder[®] Protocol Calculator to calculate the optimal amounts of input DNA sequences given the length and concentration of each input fragment.

ENTER FRAGMENTS FOR ASSEMBLY

Vector ?

☐

Name

ID/Name

Length ?

basepairs

Concentration ?

ng/ μ l

Add Fragment

☒

backbone

2682

100

Delete

☐

insert

905

10

Delete

Clear All Fragments

SUGGESTED PROTOCOL

1. Set up the reaction on ice (see table).

Component	Volume	Amount
backbone	0.8 μ l	0.050 pmoles
insert	5.6 μ l	0.100 pmoles
Deionized H ₂ O	3.6 μ l	---
NEBuilder HiFi DNA Assembly Master Mix	10.0 μ l	---
Total	20.0 μ l	0.150 pmoles

☐ Maximize ?

2. Incubate samples in a thermocycler at 50°C for 15 minutes. Following incubation, store samples on ice or at –20°C for subsequent transformation.
Note: Extended incubation up to 60 minutes may help to improve assembly efficiency in some cases.
3. Transform NEB[®] 5-alpha, NEB Stable or NEB 10-beta Competent *E. coli* cells (provided in the cloning kit, bundle or purchased separately from NEB) with 2.0 μ l (10%) of the chilled assembled product, following the transformation protocol.

Related Resources



[NEBuilder[®] HiFi DNA Assembly Reaction Protocol](#)
[NEBuilder[®] HiFi DNA Assembly](#)

Related Products



[NEBuilder[®] HiFi DNA Assembly Master Mix](#)
[NEBuilder[®] HiFi DNA Assembly Cloning Kit](#)
[NEBuilder[®] HiFi DNA Assembly Bundle for Large Fragments](#)

Related Tools



[NEBuilder[®] Assembly Tool](#)