

Feb 01, 2024

# Molecular Cloning- Gibson and LR reactions

Nature Cell Biology

In 1 collection

DOI

dx.doi.org/10.17504/protocols.io.5jyl8p7n8g2w/v1

Melissa Hoyer<sup>1,2</sup>, Harper JW<sup>1,2</sup>

<sup>1</sup>ASAP; <sup>2</sup>Harvard Medical School



Melissa Hoyer Harvard Medical School

# Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN ACCESS



DOI: https://dx.doi.org/10.17504/protocols.io.5jyl8p7n8g2w/v1

External link: https://doi.org/10.1038/s41556-024-01356-4

Protocol Citation: Melissa Hoyer, Harper JW 2024. Molecular Cloning- Gibson and LR reactions. protocols.io

https://dx.doi.org/10.17504/protocols.io.5jyl8p7n8g2w/v1



#### Manuscript citation:

Melissa J. Hoyer, Cristina Capitanio, Ian R. Smith, Julia C. Paoli, Anna Bieber, Yizhi Jiang, Joao A. Paulo, Miguel A. Gonzalez-Lozano, Wolfgang Baumeister, Florian Wilfling, Brenda A. Schulman, J. Wade Harper (2024) Combinatorial selective ER-phagy remodels the ER during neurogenesis. Nature Cell Biology doi: 10.1038/s41556-024-01356-4

**License:** This is an open access protocol distributed under the terms of the **Creative Commons Attribution License**, which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited

Protocol status: Working

We use this protocol and it's working

Created: February 01, 2024

Last Modified: May 31, 2024

Protocol Integer ID: 94504

**Keywords:** ASAPCRN, er structures in axon, axonal er accumulation, endoplasmic reticulum, er remodeling by autophagy, Ir reactions the endoplasmic reticulum, er to degradative autophagy machinery, deficient neuron, phagy receptor fam134b, extensive er remodeling during differentiation, deficient neurons in vivodisplay, embedded receptor, neuron, er structure, degradative autophagy machinery, er proteome, axon, underlying er remodeling, vast proteomic landscape, autophagy, tractable induced neuron, monitoring extensive er remodeling, linking er, protein, selective er

#### **Funders Acknowledgements:**

**ASAP** 

Grant ID: ASAP-000282

#### Disclaimer

#### DISCLAIMER - FOR INFORMATIONAL PURPOSES ONLY; USE AT YOUR OWN RISK

The protocol content here is for informational purposes only and does not constitute legal, medical, clinical, or safety advice, or otherwise; content added to <u>protocols.io</u> is not peer reviewed and may not have undergone a formal approval of any kind. Information presented in this protocol should not substitute for independent professional judgment, advice, diagnosis, or treatment. Any action you take or refrain from taking using or relying upon the information presented here is strictly at your own risk. You agree that neither the Company nor any of the authors, contributors, administrators, or anyone else associated with <u>protocols.io</u>, can be held responsible for your use of the information contained in or linked to this protocol or any of our Sites/Apps and Services.



### Abstract

The endoplasmic reticulum (ER) has a vast proteomic landscape to preform many diverse functions including protein and lipid synthesis, calcium ion flux, and inter-organelle communication. The ER proteome is remodeled in part through membrane-embedded receptors linking ER to degradative autophagy machinery (selective ERphagy)1,2. A refined tubular ER network3,4 is formed in neurons within highly polarized dendrites and axons5,6. Autophagy-deficient neurons in vivodisplay axonal ER accumulation within synaptic ER boutons,7 and the ERphagy receptor FAM134B has been genetically linked with human sensory and autonomic neuropathy8,9. However, mechanisms and receptor selectivity underlying ER remodeling by autophagy in neurons is limited. Here, we combine a genetically tractable induced neuron (iNeuron) system for monitoring extensive ER remodeling during differentiation. With this system, we imaged fixed iNeuron cultures, imaged these via confocal fluorescence microscopy, and quantified ER structures in axons.

### **Materials**

Gateway TM LR Clonase TM II Enzyme mix (cat 11791020) TE Buffer Invitrogen (cat AM9858) Thermo Fisher One Shot™ OmniMAX™ 2 T1 Phage-Resistant Cells (Cat. no. C8540-03) Q5® Hot Start High-Fidelity 2X Master Mix (DNA Polymerase cat#M0494S) Gibson Assembly using NEBuilder® HiFi DNA Assembly Master Mix (cat# E2621S) dH2O S.O.C.E media; Luria-Bertani (LB) broth (cat) or agar QIAprep Spin Miniprep Kit (cat 27104) antibiotics

# **Troubleshooting**



### Gateway technology cloning

- LR reaction: Add pDONOR vectors and pDEST vectors to a Gateway<sup>TM</sup> LR Clonase<sup>TM</sup> II Enzyme mix supplemented with TE buffer. Gateway<sup>TM</sup> LR Clonase<sup>TM</sup> II Enzyme mix (cat 11791020) <a href="https://www.thermofisher.com/order/catalog/product/11791020">https://www.thermofisher.com/order/catalog/product/11791020</a>
  - 1–7  $\mu$ L entry clone (50–150 ng) 1  $\mu$ L destination vector (150 ng/ $\mu$ L) TE buffer pH 8.0, to 8  $\mu$ L
  - Incubate reaction overnight at room temperature. Add 1  $\mu$ L of the Proteinase K solution to each sample and incubate sample sample 37C for 10 minutes.
- Transformation: Take 1 μL of each LR reaction and add it to 50 μL of Thermo Fisher One Shot™ OmniMAX™ 2 T1 Phage-Resistant Cells (Cat. no. C8540-03). Incubate on ice for 30 minutes. Incubate at 42°C for 30 seconds to heat shock. Add 250 μL of S.O.C. Medium and incubate at 37°C for 1 hour with shaking for recovery. Plate 100 μL of each transformation onto selective LB agar plates.

### Gibson technology cloning

- 3 Design pDonor using insert and backbone fragments and the NEB builder website <a href="https://nebuilder.neb.com/">https://nebuilder.neb.com/</a>
- Perform any necessary PCR reactions using Q5® Hot Start High-Fidelity 2X Master Mix (DNA Polymerase cat#M0494S) and the protocol associated with that PCR enzyme. <a href="https://www.neb.com/en-us/products/m0494-q5-hot-start-high-fidelity-2x-master-mix#Product%20Information">https://www.neb.com/en-us/products/m0494-q5-hot-start-high-fidelity-2x-master-mix#Product%20Information</a>
- 5 Linearize backbone as designed in the NEB builder website (either PCR or restriction enzyme digest)
- Perform Gibson Assembly using NEBuilder ® HiFi DNA Assembly Master Mix (cat# E2621S) <a href="https://www.neb.com/en-us/products/e2621-nebuilder-hifi-dna-assembly-master-mix#Product%20Information">https://www.neb.com/en-us/products/e2621-nebuilder-hifi-dna-assembly-master-mix#Product%20Information</a>
  - 2-3 Fragment mix strategy:

Assembly: Recommended DNA Molar Ratio Vector:insert = 1:2

Total Amount of Fragments 0.03-0.2 pmol\*

NEBuilder HiFi DNA Assembly Master Mix 10  $\mu$ l

Deionized H2O Total Volume 20 µl\*\*\*\*

Incubate samples in a thermocycler at 50°C for 60 minutes

7 Transfrom Gibson reaction:



Take 2 μL of each Gibson reaction and add it to 50 μL of NEB 5-alpha(cat#C2987H). Incubate on ice for 30 minutes. Incubate at 42°C for 30 seconds to heat shock. Add 250 μL of S.O.C. Medium and incubate at 37°C for 1 hour with shaking for recovery. Plate 100 μL of each transformation onto selective LB agar plates.

### Prep and full plasmid sequencing of final products

8 At either stage (after Gibson or LR reaction). Take colonies and start overnights (LB+antibiotic E.coli cultures). Miniprep overnight cultures with QIAprep Spin Miniprep Kit (cat 27104). Test digest for good plasmid candidates that look like they have the correct insert. Full plasmid sequence plasmids to confirm proper plasmid construction.