

Aug 04, 2021

Integration of a control brick

In 1 collection

Carolyn N Bayer¹, Maja Rennig¹, Anja Ehrmann¹, Morten Norholm¹

¹Technical University of Denmark



dx.doi.org/10.17504/protocols.io.bvman42e

SEGA

Carolyn Bayer Technical University of Denmark

ABSTRACT

SEGA, the Standardized Genome Engineering Architecture, is a comprehensive strain collection that enables genome engineering by combining only two reagents: a DNA fragment that can be ordered from a commercial vendor and a stock solution of bacterial cells followed by incubation on agar plates. Recombinant genomes are identified by visual inspection using green-white colony screening akin to classical blue-white screening for recombinant plasmids. The modular nature of SEGA allows precise multi-level control of transcriptional, translational, and post-translational regulation. The SEGA architecture simultaneously supports increased standardization of genetic designs and a broad application range by utilizing well-characterized parts optimized for robust performance in the context of the bacterial genome

This protocol describes to process of integrating a SEGA control brick. A control brick is integrated using *galK* counterselection.

DOI

dx.doi.org/10.17504/protocols.io.bvman42e

PROTOCOL CITATION

Carolyn N Bayer, Maja Rennig, Anja Ehrmann, Morten Norholm 2021. Integration of a control brick. **protocols.io**

https://dx.doi.org/10.17504/protocols.io.bvman42e

COLLECTIONS (i)

SEGA protocol collection

KEYWORDS

SEGA, genome engineering, recombineering, E. coli, synthetic biology

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

CREATED

Jun 08, 2021

LAST MODIFIED

Aug 04, 2021

BEFORE STARTING Transform a SEGA strain, harbouring the upstream gadget galk, with pSIM19 (Spectinomycin resistance). From now on cultures have to be kept at § 30 °C to retain the plasmid (temperature-sensitive or). preculture and DNA fragment- Day 1 1d 1d Prepare a PCR product of the control elements that need to be integrated and purify it from an agarose gel. Setup a preculture of the strain harbouring pSIM19 in LB medium supplemented with Spectinomycin [M10.05 mg/ml]. Incubate overnight at \$\triangle 250 \text{ rpm, 30°C} Recombineering-Day 2 1d Prepare: 3.1 Cold sterile water 3.2 Cold Glycerol [M]15 % volume 3.3 Pre-chilled centrifuge and tabletop centrifuge at § 4 °C 3.4 M63 agar plates supplemented with [M]0.2 Mass / % volume 2-deoxy-galactose, [M] 0.2 % volume glycerol and [M] 5 Milimolar (mM) L-rhamnose Inoculate ■50 mL LB-Medium supplemented with Spectinomycin ([M)0.05 mg/ml) with ■500 µl of the preculture from step 3 Incubate at **250 rpm, 30°C** until cultures reached an OD₆₀₀ of 0.5 Induce expression by transferring the culture to a shaking water bath at \$\to\$150 rpm, 42°C, 00:20:00

PROTOCOL INTEGER ID

PARENT PROTOCOLS

SEGA protocol collection

Part of collection

50562

mprotocols.io

Citation: Carolyn N Bayer, Maja Rennig, Anja Ehrmann, Morten Norholm (08/04/2021). Integration of a control brick. https://dx.doi.org/10.17504/protocols.io.bvman42e

08/04/2021

6	Transfer culture to prechilled □50 mL falcon tubes and put on ice for ⊙00:15:00		
7	Spin the culture down at 34000 x g, 4°C, 00:05:00 and discard the supernatant		
8	Add 1 mL of ice cold water, resuspend and transfer to a 1.5 ml tube		
9	30s Spin at (3)11000 x g, 4°C, 00:00:30 in a tabletop centrifuge		
10	Wash pellet twice with ☐1 mL ice cold water ⑤ go to step #9		
11	Resuspend the pellet in Good μl cold glycerol ([M]15 % volume)		
	11.1 Unused cells can be stored at 8-80 °C		
	This is not possibe for <i>E. coli</i> Nissle		
12	Electroporate $\Box 50~\mu I$ of cells with $\Box 200~ng$ of purified PCR product from step 2 or $\Box 2~\mu I$ of a [M] 100 Micromolar (μM) single-stranded oligonucleotide		
13	Transfer cells into ☐50 mL LB medium in a 250 ml baffled conical shake flask and recover overnight at ☐250 rpm, 30°C		
	Alternatively, cells can be recovered in ■10 mL medium for © 04:30:00		
Plating- Day 3 1d			
Wash The covered cells twice with 1X M9 salts. Centrifuge at 11000 rpm, 20°C, 00:00:30			
protoc	cols.io 3	08/04/2021	

 $\textbf{Citation:} \ \, \textbf{Carolyn N Bayer, Maja Rennig, Anja Ehrmann, Morten Norholm (08/04/2021).} \ \, \textbf{Integration of a control brick.} \\ \underline{\textbf{https://dx.doi.org/10.17504/protocols.io.bvman42e}}$

Make a dilution series and plate **100 μl** of the 1:100 - 1:1000 dilution on M63 agar supplemented with [M]**0.2 Mass / % volume** 2-deoxy-galactose, [M]**0.2 % volume** glycerol and [M]**5 Milimolar (mM)** L-rhamnose.

In case the cells were recovered for **© 04:30:00** dilution 1:10 and 1:100 need to be plated

16 incubate the plates at § 30 °C for (§ 48:00:00 to (§ 72:00:00

1d

5d

Cells can be incubated at § 37 °C if it is the last step of recombination and pSIM19 is no longer needed in the cells

Screening- Day 4-5

Screen for positive colonies by "green-white screening" on a blue-light table and perform colony PCR on the colorless colonies to identify the correct recombinants