Targeted Mutagenesis Using RNA-guided Endonucleases in Mosses

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[Abstract] RNA-guided endonucleases (RGENs) have been used for genome editing in various organisms. Here, we demonstrate a simple method for performing targeted mutagenesis and genotyping in a model moss species, *Physcomitrella patens*, using RGENs. We also performed targeted mutagenesis in a non-model moss, *Scopelophilla cataractae*, using a similar method (Nomura *et al.*, 2016), indicating that this experimental system could be applied to a wide range of mosses species.

Keywords: Genome editing, RNA-guided endonucleases, CRISPR/Cas system, Targeted mutagenesis, Mosses

[Background] Targeted mutagenesis using RNA-guided endonucleases (RGENs) derived from the adaptive immune system, using the bacterial CRISPR (clusters of regularly interspaced palindromic repeats)/Cas (CRISPR-associated) systems, has dramatically advanced in recent years. In this method, the Cas9 endonuclease, derived from *Streptococcus pyogenes*, and an artificially designed single-chain guide RNA (sgRNA) are used. The Cas9-sgRNA complex recognizes the protospacer-adjacent motif (5'-NGG-3') and cleaves 3 bp upstream of the target site (Jinek *et al.*, 2012). Subsequently, random insertion and/or deletion mutations occur during the repair process for double-strand breaks (DSBs) in the DNA. As targeted mutagenesis using these RGENs is efficient as well as cost- and time-effective, it has been used for genome editing in various organisms, including many plant species. Here, we established a protocol for targeted mutagenesis using RGENs in mosses, and demonstrated it in a model and a non-model species (Nomura *et al.*, 2016).

Materials and Reagents

1. Washed and autoclaved cellophane (FUTAMURA CHEMICAL, catalog number: PS-1)
2. 9 cm plastic Petri dish (SANSEI MEDICAL, catalog number: 01-013)
3. 1.5 ml plastic tubes (FUKAEKASEI and WATSON, catalog number: 131-415C)
4. 15 ml plastic tubes (FUKAEKASEI and WATSON, catalog number: 1332-015S)
5. PCR tubes (NIPPON Genetics, FastGene, catalog number: FG-028DC)
6. 10 µl plastic pipette tips (FUKAEKASEI and WATSON, catalog number: 123R-254CS)
7. 200 µl plastic pipette tips (FUKAEKASEI and WATSON, catalog number: 123R-755CS)
8. 1,000 µl plastic pipette tips (FUKAEKASEI and WATSON, catalog number: 122-804B)
9. Protonemata of *Physcomitrella patens* (or other mosses)

10. pSCgRNA, pSCOE1-fcoCas9 (Nomura *et al.*, 2016, Figure 1A)
11. DH5 α -competent cells (Home-made)
12. NEBuffer 2.1 (New England Biolabs, catalog number: R0539S)
13. *Bbs*I (New England Biolabs, catalog number: R0539S)
14. Sterile Milli-Q water
15. GEL/PCR Purification Mini Kit (Favorgen Biotech, catalog number: FAGCK 001-1)
16. Ligation high ver.2 (TOYOBO, catalog number: LGK-201)
17. ScU6p Ins. check F primer 5'-GAGGATCACGGTGTACATGTCC-3'
18. Quick Taq[®] HS DyeMix (TOYOBO, catalog number: DTM-101)
19. ScU6p Seq. check F primer 5'-ATGTCAAACATAACCTGG-3'
20. Ampicillin (NACALAI TESQUE, catalog number: 02739-74)
21. Plasmid DNA Extraction Mini Kit (Favorgen Biotech, catalog number: FAPDE 001-1)
22. Agarose S (NIPPON GENE, catalog number: 312-01193)
23. DNA ladder markers (SMOBIO Technology, catalog number: DM3100)
24. NucleoBond[®] Xtra Midi (MACHEREY-NAGEL, catalog number: 740410.50)
25. G418 Disulfate (NACALAI TESQUE, catalog number: 16512-81)
26. BCDAT medium (Nishiyama *et al.*, 2000)
27. Glucose (Wako Pure Chemical Industries, catalog number: 049-31165)
28. Tks Gflex DNA polymerase (Takara Bio, catalog number: R060A)
29. Zero Blunt PCR Cloning Kit (Thermo Fisher Scientific, Invitrogen, catalog number: K275020)
30. CloneJET PCR Cloning Kit (Thermo Fisher Scientific, Thermo Scientific[™], catalog number: K1231)
31. LB agar, Miller (BD, Difco[™], catalog number: 244520)
32. LB broth, Miller (BD, Difco[™], catalog number: 244620)
33. Tris (hydroxymethyl) aminomethane (NACALAI TESQUE, catalog number: 35434-21)
34. Potassium chloride (Wako Pure Chemical Industries, catalog number: 163-03545)
35. Ethylenediaminetetraacetic acid (EDTA) (DOJINDO, catalog number: 345-01865)
36. Acetic acid (Wako Pure Chemical Industries, catalog number: 017-00256)
37. Ethidium bromide solution (NACALAI TESQUE, catalog number: 14631-94)
38. LB agar plate with 50 μ g/ml ampicillin (see Recipes)
39. Lysis buffer (see Recipes)
40. 50 x TAE buffer (see Recipes)
41. 1x TAE buffer (see Recipes)

Equipment

1. Plant growth chamber (SANYO, model: MLR-350HT)
2. Micropipettes (Eppendorf, model: Reference[®] 2)
3. Sterile tweezers (Electron Microscopy Sciences, DUMONT, catalog number: 0108-5-PO)

4. Thermal cycler (PCR Thermal Cycler Dice® Gradient, Takara Bio, model: TP600)
5. High-speed centrifuge (TOMY DIGITAL BIOLOGY, model: MX-300)
6. Block incubator (ASTEC, model: BI-515A)
7. Spectrophotometer (GE Healthcare, model: NanoVue)
8. Autoclave (TOMY DIGITAL BIOLOGY, model: SX300)
9. Agarose gel electrophoresis equipment (Cosmo Bio, model: i-MyRun.N)
10. Ultraviolet transilluminator (ATTO, models: DTB-20MP, TYPE-CX)
11. Heated incubator (SANYO, model: MIR-262)

Procedure

A. Design and construction of sgRNA expression vector

1. Selection of target sequence

- a. Choose the 20 bp target sequence in the gene or genomic region of interest (Figure 1B).
 - i. Check the protospacer-adjacent motif (PAM) sequence (5'-NGG-3') at the 3'-end of the target sequence.
 - ii. To induce a dysfunctional mutation in the gene, the target sequence should be designed near the 5'-end or in the functional domain.
 - iii. Avoid poly-T sequences, which work as transcriptional terminators for sgRNA; ensure that the G + C content is 40-80%.
 - iv. If the target site includes the recognition site of a restriction enzyme, it will be possible to perform restriction fragment length polymorphism (RFLP) analysis for genotyping.

Note: It has been reported that truncated sgRNA could decrease off-target effects (Fu et al., 2014). We confirmed that truncated sgRNA (18 or 17 bp target sequences) works in our system; however, we have not yet evaluated their targeting efficiency.

- b. Add the following adapter sequence for cloning (Figure 1B): 5'-TCTG-3' for sense oligonucleotides and 5'-AAAC-3' for anti-sense oligonucleotides.

2. Digestion of pSCgRNA vector (Figure 1C)

- a. Prepare the digestion mixture, as given below:

Reagent	Volume
pSCgRNA	3 µg
10x NEBuffer 2.1	10 µl
<i>Bbs</i> I	3 µl
Sterile Milli-Q water	Up to 100 µl

- b. Incubate the mixture at 37 °C for 3 h.
- c. Purify the *Bbs*I-digested pSCgRNA vector using the DNA purification kit (Favorgen GEL/PCR Purification Mini Kit).

3. Annealing of oligonucleotides (Figure 1C)

- a. Prepare the annealing mixture, as given below:

Reagent	Volume
Sense oligonucleotides (100 μ M)	1 μ l
Anti-sense oligonucleotides (100 μ M)	1 μ l
Sterile Milli-Q water	48 μ l
Total	50 μ l

- b. Incubate at 95 °C for 5 min in a thermal cycler.
c. Remove the tube from the thermal cycler and allow the mixture to cool down on the bench at 20-25 °C for 30 min.

4. Cloning of annealed oligonucleotides into pSCgRNA vector (Figure 1C)

- a. Prepare the ligation mixture, as given below:

Reagent	Volume
<i>Bbs</i> I-digested pSCgRNA vector (10 ng/ μ l)	1 μ l
Annealed oligonucleotides	1 μ l
Ligation High ver.2	2 μ l
Total	4 μ l

- b. Incubate the mixture at 16 °C for 3 h or more.
c. Transform DH5 α -competent cells using 2 μ l ligation mixture and spread on an LB plate with 50 μ g/ml ampicillin.
d. Incubate the LB plate at 37 °C for 14-16 h.
e. Mix a PCR reaction mixture, using the ScU6p Ins. check F primer (as the forward primer) and the antisense target sequence (as the reverse primer).

Reagent	Volume
2x Quick Taq® HS DyeMix	6 μ l
ScU6p check F primer (10 μ M)	0.25 μ l
Anti-sense oligonucleotides (10 μ M)	0.25 μ l
Sterile Milli-Q water	Up to 12 μ l

- f. Verify the insertion by performing a colony PCR with the following cycle:

94 °C	2 min	}	1 cycle
94 °C	30 sec		28 cycles
(T _m - 5) °C	30 sec		
68 °C	30 sec		
4 °C	Hold		1 cycle

Note: If the cloning is successful, a 950 bp band will be amplified and detected (Figure 1D).

We recommend testing at least 8-16 colonies.

- g. Grow the positive colonies in liquid LB medium with 50 µg/ml ampicillin.
- h. Purify the plasmids using a plasmid DNA mini prep kit (Favorgen Plasmid DNA Extraction Mini Kit).
- i. Confirm the inserted target sequence in the pSCgRNA vector using ScU6p Seq. check F primer using Sanger sequencing.

- C. Regeneration of protonemal cells and transient screening using antibiotic G418 (Geneticin).
1. After inoculating the transformed protoplasts in PRM/T on cellophane and on a PRM/B plate, incubate the mixture at 25 °C for 7 d in a plant growth chamber (Figure 2B).
 2. For transient screening, transfer the regenerated protonemata from the protoplasts in PRM/T on cellophane to new BCDAT plates with 20 µg/ml G418 (Figure 2C).
 3. Culture the plates in a plant growth chamber for 5 d (Figures 2C and 3).

Figure 3. Example of protonemata after transient screening (*PpFtsZ2-1* targeted). A. Representative example of dead protonemata; B. Representative example of surviving protonemata. Strain also shows the phenotype of *PpFtsZ2-1* mutation. Scale bars = 50 µm.

4. Transfer the surviving protonemata in PRM/T on cellophane to new BCDAT plates and culture again until small colonies have formed (Figure 2D).
5. Using tweezers, choose the protonemata at the edge of the colonies and transfer them to new, numbered BCDAT or BCDAT + 0.5% glucose plates (Figure 2D).
6. For genotyping, culture the protonemata until the colonies become 5-10 mm in diameter.

Note: There is a possibility that the acquired strain is a mosaic plant, which is a mixture of strains with different mutations or wild type. If so, we can isolate the monogenic strain from the protoplast, single protonemal branch, or leaf of the gametophore.

D. Genotyping

1. Green PCR
 - a. Transfer a pinch of fresh protonemata to PCR tubes (Figure 4A) and grind using the tip of the 200 µl pipette chip into 50 µl lysis buffer (see Recipes) (Figure 4B).

Figure 4. Preparation of the template for Green PCR. A. A pinch of fresh protonemata in the lysis buffer; B. Protonemata crushed by the tip of the 200 μ l chip in the lysis buffer.

- b. Incubate at 95 °C for 10 min.
- c. Spin down the debris at 3,500 $\times g$ for 10 sec and use the supernatant as the PCR template.
- d. Design the primer set using free online tool Primer 3 (<http://primer3.ut.ee/>) for amplifying the genomic DNA, including the target site for sequencing or for other genotyping such as RFLP.
- e. Prepare PCR reaction mixture, as given below:

Reagent	Volume
2x Gflex PCR buffer	10 μ l
Forward primer (10 μ M)	0.5 μ l
Reverse primer (10 μ M)	0.5 μ l
Template (lysate of plant)	0.5 μ l
Tks Gflex DNA polymerase	0.4 μ l
Water	Up to 20 μ l

- f. Amplify the PCR product, including the target site of the genome, with the cycle given below:

94 °C	1 min	1 cycle
98 °C	10 sec	} 30-40 cycles
68 °C	1 min/kb	
4 °C	Hold	1 cycle

2. Restriction fragment length polymorphism (RFLP) analysis

- a. If the target site includes the recognition site of a restriction enzyme, it will be possible to observe the targeted mutagenesis by RFLP.
- b. After confirming the PCR products by agarose gel electrophoresis, digest the PCR products using the corresponding restriction enzymes.
- c. After sufficient digestion, analyze the enzyme-treated and untreated PCR products by agarose gel electrophoresis. It is possible that restriction enzyme-tolerant PCR products have mutations.

Note: If a mixture of PCR products derived from wild type and mutant strains is used, the targeted mutagenesis can be checked using the T7 endonuclease I or cel1 assay, which can recognize and cleave non-perfectly matched DNA. As high throughput and

comprehensive genotyping method, high resolution melting analysis or next-generation sequencing is also applicable.

3. Sequencing of target sites in the genome
 - a. After confirming the PCR product by agarose gel electrophoresis, clone the PCR products into the sequencing vector using the Zero Blunt PCR Cloning Kit or the CloneJET PCR Cloning Kit.
 - b. After colony PCR and mini prep, check the DNA sequence of the inserted PCR product of several clones by Sanger sequencing.

Note: If the PCR product is present as a single band in monogenic plant, it will be possible to check by direct sequencing.

Data analysis

According to our evaluation based on the phenotypic change caused by the mutation of a target gene, the target mutagenesis efficiency in *P. patens* was approximately 45% to 68% (Nomura *et al.*, 2016). Since efficiency is dependent on the target sequence, we recommend using the 'focas' website (<http://focas.ayanel.com/>; Doench *et al.*, 2014; Xiao *et al.*, 2014; Osakabe *et al.*, 2016) for designing and predicting the on- and off-target efficiency of the target genome. In this case, the target sequence with an on-target score of 0.6 or higher and a low possibility of off-target effect is desirable.

Notes

1. It is possible to introduce long deletion mutations (~3 kb) through this method by using two gRNAs designed for distant sites on the genome (Nomura *et al.*, 2016).
2. In this method, the vector DNA is usually not integrated into the moss genome (Nomura *et al.*, 2016).
3. There is the possibility of the formation of polyploidy during the PEG-mediated protoplast transformation. DNA content in acquired strains should be assessed using a flow cytometer.
4. If this protocol is applied to other species, it may be necessary to modify some experimental conditions such as the transformation (e.g., PEG treatment time) and screening method (e.g., antibiotic concentration and duration of selection).

Recipes

1. LB agar plate with 50 µg/ml ampicillin
 LB agar, Miller 40 g
 Fill up to 1 L with Milli-Q water and autoclave (121 °C, 15 min)
 Add ampicillin to a final concentration of 50 µg/ml

2. Lysis buffer
 - 100 mM Tris-HCl (pH 9.5)
 - 1 M KCl
 - 10 mM EDTA
3. 50 x TAE buffer
 - 2 M Tris
 - 1 M acetic acid
 - 0.5 M EDTA (pH = 8.0)
4. 1 x TAE buffer
 - 20 ml 50 x TAE buffer
 - 980 ml Milli-Q water

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