

Protocol

RNAi-Mediated Loss of Function of *Xenopus* Immune Genes by Transgenesis

Eva-Stina Edholm¹ and Jacques Robert

Department of Microbiology and Immunology, University of Rochester Medical Center, Rochester, New York 14620

Generation of transgenic frogs through the stable integration of foreign DNA into the genome is well established in *Xenopus*. This protocol describes the combination of transgenesis with stable RNA interference as an efficient reverse genetic approach to study gene function in *Xenopus*. Initially developed in the fish medaka and later adapted to *Xenopus*, this transgenic method uses the I-SceI meganuclease, a “rare-cutter” endonuclease with an 18 bp recognition sequence. In this protocol, transgenic *X. laevis* with knocked down expression of a specific gene are generated using a double promoter expression cassette. This cassette, which is flanked by I-SceI recognition sites, contains the shRNA of choice under the control of the human U6 promoter and a green fluorescent protein (GFP) reporter gene under the control of the human EF-1 α promoter. Prior to microinjection the plasmid is linearized by digestion with I-SceI and the entire reaction is then microinjected into one-cell stage eggs. The highly stringent recognition sequence of I-SceI is thought to maintain the linearized plasmid in a nonconcatamered state, which promotes random integration of the plasmid transgene in the genome. The injected embryos are reared until larval stage 56 and then screened for GFP expression by fluorescence microscopy and assessed for effective knockdown by quantitative RT-PCR using a tail biopsy. Typically, the I-SceI meganuclease transgenesis technique results in 35%–50% transgenesis efficiency, a high survival rate (>35%) and bright nonmosaic GFP expression. A key advantage of this technique is that the high efficiency and nonmosaic transgene expression permit the direct use of F₀ animals.



MATERIALS

It is essential that you consult the appropriate Material Safety Data Sheets and your institution’s Environmental Health and Safety Office for proper handling of equipment and hazardous materials used in this protocol.

RECIPES: Please see the end of this protocol for recipes indicated by <R>. Additional recipes can be found online at <http://cshprotocols.cshlp.org/site/recipes>.

Reagents

Agarose gel DNA extraction kit
Annealing buffer for shRNA (10 \times) <R>
Competent bacteria
DNA sequencing reagents
DNeasy Blood and Tissue Kit (QIAGEN) (optional; Step 15)

¹Correspondence: Eva-Stina_Edholm@urmc.rochester.edu
From the *Xenopus* collection, edited by Hazel L. Sive.

© 2018 Cold Spring Harbor Laboratory Press
Cite this protocol as *Cold Spring Harb Protoc*; doi:10.1101/pdb.prot101519

Ethanol (optional; see Step 10)
Ficoll (lyophilized powder, Type 400-DL)
Gentamicin (50 mg/mL)
GFP-specific PCR primers (optional; Step 15)
I-SceI-GFP-huU6-I-SceI plasmid (available upon request from the *Xenopus laevis* Research Resource for Immunobiology, <https://www.urmc.rochester.edu/microbiology-immunology/xenopus-laevis.aspx>)
I-SceI meganuclease (New England Biolabs R06943)
Luria Bertani (LB) ampicillin agar plates
Modified Barth's saline (MBS) <R>
PCR reagents for standard and qRT-PCR and target gene-specific primers
Phenol:chloroform:isoamyl alcohol (25:24:1) (optional; see Step 10)
Plasmid DNA isolation kits (mini and midi or maxi scale)
Restriction endonucleases appropriate for experimental design
shRNA oligonucleotides (designed according to Steps 1–2)
T4 DNA ligase
U6 promoter sequencing primer
Water (DNase/RNase-free) (optional; see Step 10)

Equipment

Agarose gel electrophoresis apparatus
Beaker (3 L minimum)
DNA sequencing apparatus
Fluorescence microscope
Heat block
Hot plate
Incubator
Microcentrifuge tubes (1.5 mL)
Microinjector (PLI-100, Harvard Apparatus or equivalent)
PCR apparatus (for standard and quantitative PCR)
Petri dishes (60 × 15 mm and 35 × 12 mm)
Scalpel (optional; Step 15)

METHOD

shRNA Design and I-SceI-GFP-huU6-shRNA-I-SceI Plasmid Preparation

1. Design shRNA oligonucleotides according to experimental goals. Include restriction sites at the 5' ends of each oligonucleotide to facilitate shRNA cloning between the huU6 promoter and the I-SceI restriction site of I-SceI-GFP-huU6-I-SceI. (See Fig. 1 for potential restriction sites, which include BbsI, HindIII, ClaI, SalI, XhoI, and KpnI.) Use the following guidelines to design effective shRNAs.
 - Determine whether the gene of interest has one or multiple splice variants and decide whether to target all splice forms of the gene or specific variants. Select exons for targeting accordingly.
 - Design the shRNA so that the first nucleotide is a purine.
 - Choose a sequence with low (30%–50%) GC content.
 - Avoid the first 50–100 nt downstream from the start codon and the 100 nt upstream of the stop codon. Also avoid 5'- and 3'-UTRs. These regions contain binding sequences for regulatory proteins that may affect the accessibility of the RNA target sequence.

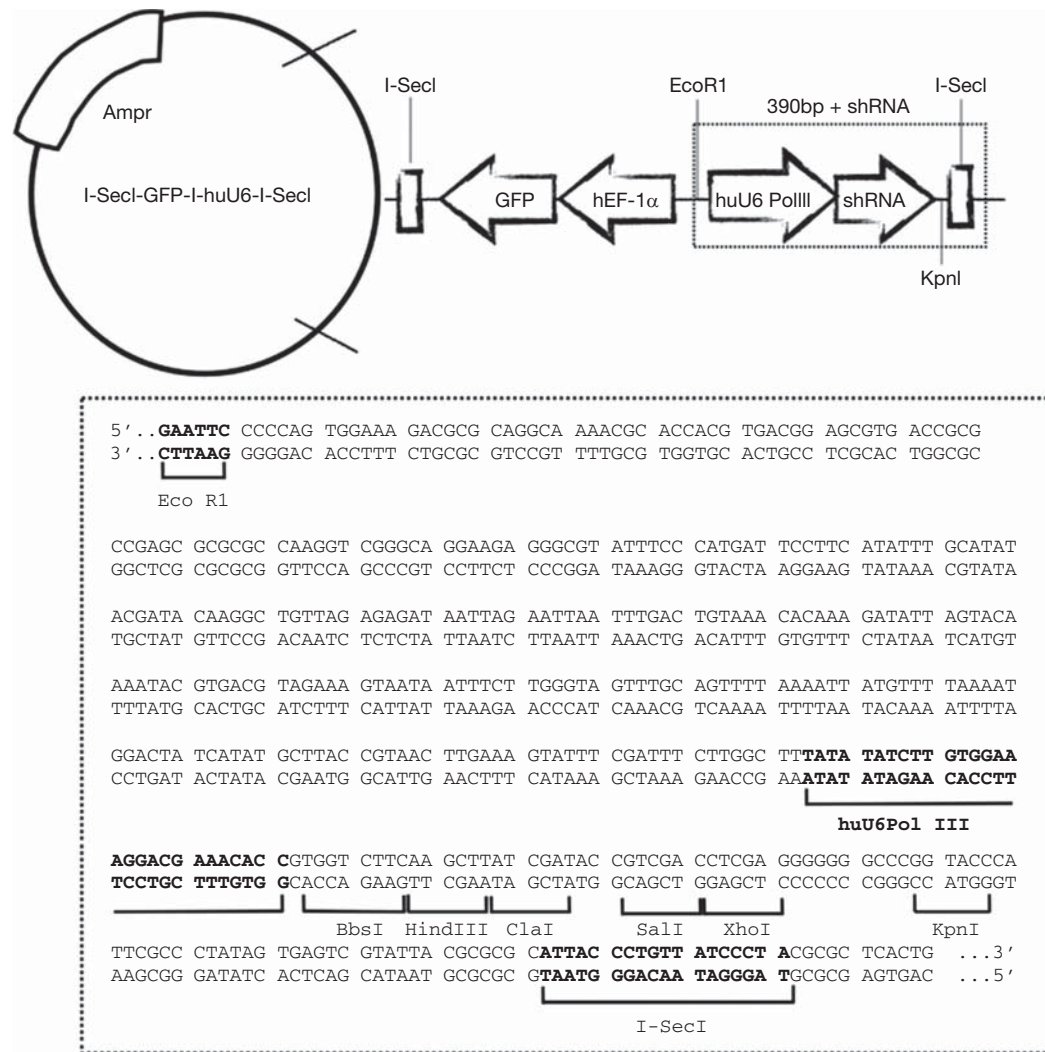


FIGURE 1. I-SceI-GFP-huU6-shRNA-I-SceI vector map and multiple-cloning site sequence. The sequence covering the shRNA cloning site includes the locations of potential restriction sites used to insert the shRNA into the vector.

- Design more than one shRNA per gene target.
- Ensure that the target sequence is specific for the gene of interest. Potential off targets can be identified by blast searching the shRNA sequence against the *X. laevis* genome available at <http://www.xenbase.org/entry/>.

There are several commercial and noncommercial websites available to assist in designing knockdown shRNAs and appropriate controls, including <http://www.invivogen.com/sirnazaward/>, <http://biodev.extra.cea.fr/DSIR/DSIR.html>, and <http://www.genelink.com/sirna/shRNAi.asp>.

shRNA oligonucleotide design is an empirical process and the gene silencing efficiency of each shRNA needs to be determined experimentally.

2. Design a control shRNA by randomly scrambling the target sequence.
It is important to check the control shRNA for potential off target effects by blast searching the shRNA sequence against the X. laevis genome.
3. Synthesize or commercially obtain 40 nmol of each shRNA oligonucleotide in the 5'-phosphorylated form.
4. Resuspend 4 nmol of each shRNA in 10 μ L 2 \times Annealing buffer and then combine the two oligonucleotides in a 1.5 mL microcentrifuge tube. Mix gently.

5. Boil 2 L of water in a beaker. When water reaches boiling point turn off the hot plate and add the 1.5 mL microcentrifuge tube containing the oligonucleotides. Leave for a minimum of 6 h (or overnight) to allow oligonucleotides to anneal.
6. Double digest 1 μ g of the I-SceI-GFP-huU6-I-SceI vector using restriction enzymes for the sites in the shRNA oligonucleotides. Digest for 1 h at 37°C. Heat inactivate digestion for 20 min at 65°C. Run the digest on a 1% agarose gel and perform gel purification of the digested vector using an agarose gel DNA extraction kit.
7. Ligate the annealed oligonucleotides from Step 5 into the digested I-SceI-GFP-huU6-I-SceI vector from Step 6 using T4 ligase. Combine 3.3 μ L annealed shRNA, 3.3 μ L digested I-SceI-GFP-huU6-I-SceI, 1 μ L T4 ligase and 2 μ L T4 10 \times buffer. Adjust volume to 20 μ L with molecular grade H₂O. Ligate overnight at 16°C.
8. Transform 2.5 μ L of the ligation reaction into competent bacteria using standard protocols. Select transformants on LB ampicillin agar plates.
9. Purify plasmid DNA from transformants using standard protocols and verify correct shRNA insertion by sequencing using U6 promoter primers.
10. Prepare plasmid DNA for digestion and microinjection using standard “midi- or maxi-prep” scale kits for isolation of pure plasmid DNA.

Plasmid DNA can be further purified by phenol:chloroform:isoamyl alcohol (25:24:1) extraction, ethanol precipitation, and resuspension in DNase/RNase-free water.

11. Digest 1 μ g I-SceI-GFP-huU6-shRNA-I-SceI with 10 units I-SceI meganuclease in a 25 μ L volume for 40 min at 37°C.

Store the I-SceI meganuclease at -80°C at all times (storage at -20°C leads to inefficient or no transgene incorporation into the genome). Freeze thawing of both the enzyme and the buffer significantly reduces transgenesis efficiency; therefore use single aliquots.

Microinjection of *Xenopus* Eggs

*The microinjection of fertilized *Xenopus* eggs is well documented and the precise methodology used will depend on the type of equipment available.*

12. Fertilize and dejelly eggs according to standard protocols and place eggs in a 35 \times 12 mm injection Petri dish containing 0.3 \times MBS solution with 4% Ficoll. Ensure the eggs are completely submerged in the solution such that they are not subject to surface tension. Inject each egg with 10 nL of the digest from Step 11 containing 80 pg DNA and 1 \times 10⁻³ U I-SceI.

Use the I-SceI-GFP-huU6-shRNA-I-SceI digest within 15–30 min. If more eggs are to be injected, set up fresh digests. It is crucial that injection is performed within 30 min after fertilization and within 30 min after plasmid digestion to increase the likelihood of early integration. To avoid mosaicism it is advised that eggs are not injected once the first signs of cell division are observed.

13. Transfer injected eggs to a 60 \times 15 mm Petri dish containing 0.3 \times MBS solution with 4% Ficoll and incubate for 4 h at 13°C.

This delays cell division to extend the time for transgene integration.

14. Transfer the embryos to Petri dishes containing 0.3 \times MBS with 50 μ g/mL gentamicin and rear the embryos at 18°C until hatching. Ensure that the embryos are completely submerged and not too crowded; between 50–100 embryos per dish is suitable.

To ensure optimal survival of the embryos it is important to change the medium daily and to remove any dead embryos, which might otherwise compromise embryo survival.

See Troubleshooting.

15. At developmental stage 58, screen tadpoles for GFP fluorescence using a fluorescence stereomicroscope.

Alternatively, amputate the distal third of the tail with a scalpel and extract genomic DNA. Perform polymerase chain reaction (PCR) for the GFP transgene with GFP specific primers.

E.-S. Edholm and J. Robert

16. Select tadpoles with successful insertion of the GFP-containing vector and perform quantitative PCR to determine expression of the target gene. Verify target gene knockdown by comparing expression levels in shRNA-injected tadpoles with scrambled shRNA-injected age-matched controls.

See Troubleshooting.

TROUBLESHOOTING

Problem (Steps 14–15): Knockdown of a specific gene results in embryonic lethality.

Solution: Clone the shRNA of choice in a vector containing an H1 RNA polymerase III promoter repressed by the Tet-repressor. The expression of the shRNA can then be induced by inhibiting the tetracycline-element-specific repressor (TetR) from binding and blocking transcription with the tetracyclin analogue, doxycycline.

Problem (Step 16): Low to no knockdown of the gene of interest.

Solution: Design multiple shRNAs targeting the same gene and assess efficiency.

DISCUSSION

Different strategies to generate transgenic *Xenopus* frogs have been developed with specific molecular and technical features. The meganuclease approach is straightforward and results in high transgenesis efficiency with uniform expression of the transgene as well as stable germline transmission (Ogino et al. 2006a,b). One key advantage of this technique is that, because substantial numbers of nonmosaic embryos with high levels of knockdown are produced, it allows F_0 animals to be used for experimentation within a month after microinjection (Edholm et al. 2013). Another advantage of this knockdown technique is that different levels of silencing can be obtained in different animals, which may result in variable phenotypes. In addition, I-SceI meganuclease-mediated transgenesis has also been optimized for use with isogenic clones (LG 15 and LG 6 [Nedelkovska and Robert 2012; Nedelkovska et al. 2013]).

RELATED INFORMATION

We recommend Jensen et al. (2012) and Bofill-De Ros and Gu (2016) for in-depth discussion of shRNA design. These articles highlight important points to consider and potential pitfalls. In addition, see Ishibashi et al. (2012) for a detailed protocol describing I-SceI meganuclease-mediated transgenesis in *X. tropicalis*.

RECIPES

Annealing Buffer for shRNA (10×)

Reagent	Concentration
Tris-HCl (1 M, pH 7.5)	5 mL
NaCl (5 M)	10 mL
EDTA (0.5 M)	1 mL
H ₂ O	34 mL

Sterilize by autoclaving, and store at 4°C.

Modified Barth's Saline (MBS)

CaCl₂ (0.1 M)
MBS salts (10×)

For a 1× solution of MBS, mix 100 mL of 10× MBS salts with 7 mL of 0.1 M CaCl₂. Adjust the volume up to 1 liter with H₂O. Store at room temperature.

MBS Salts (10×)

NaCl (880 mM)
KCl (10 mM)
MgSO₄ (10 mM)
HEPES (50 mM, pH 7.8)

Omit HEPES if MBS is to be used for oocyte maturation.

NaHCO₃ (25 mM)

Adjust pH to 7.8 with NaOH. Autoclave. Store at room temperature.

REFERENCES

- Bofill-De Ros X, Gu S. 2016. Guidelines for the optimal design of miRNA-based shRNAs. *Methods* 103: 157–166.
- Edholm ES, Albertorio Saez LM, Gill AL, Gill SR, Grayfer L, Haynes N, Myers JR, Robert J. 2013. Nonclassical MHC class I-dependent invariant T cells are evolutionarily conserved and prominent from early development in amphibians. *Proc Natl Acad Sci* 110: 14342–14347.
- Ishibashi S, Love NR, Amaya E. 2012. A simple method of transgenesis using I-SceI meganuclease in *Xenopus*. *Methods Mol Biol* 917: 205–218.
- Jensen SM, Schmitz A, Pedersen FS, Kjems J, Bramsen JB. 2012. Functional selection of shRNA loops from randomized retroviral libraries. *PLoS One* 7: e43095.
- Nedelkovska H, Robert J. 2012. Optimized transgenesis in *Xenopus laevis*/gilli isogenetic clones for immunological studies. *Genesis* 50: 300–306.
- Nedelkovska H, Edholm ES, Haynes N, Robert J. 2013. Effective RNAi-mediated β 2-microglobulin loss of function by transgenesis in *Xenopus laevis*. *Biol Open* 2: 335–342.
- Ogino H, McConnell WB, Grainger RM. 2006a. High-throughput transgenesis in *Xenopus* using I-SceI meganuclease. *Nat Protoc* 1: 1703–1710.
- Ogino H, McConnell WB, Grainger RM. 2006b. Highly efficient transgenesis in *Xenopus tropicalis* using I-SceI meganuclease. *Mech Dev* 123: 103–113.



Cold Spring Harbor Protocols

RNAi-Mediated Loss of Function of *Xenopus* Immune Genes by Transgenesis

Eva-Stina Edholm and Jacques Robert

Cold Spring Harb Protoc; doi: 10.1101/pdb.prot101519 originally published online January 30, 2018

Email Alerting Service

Receive free email alerts when new articles cite this article - [click here](#).

Subject Categories

Browse articles on similar topics from *Cold Spring Harbor Protocols*.

- [Analysis of Gene Expression](#) (180 articles)
- [Analysis of Gene Expression, general](#) (122 articles)
- [DNA Delivery/Gene Transfer](#) (305 articles)
- [DNA Delivery/Gene Transfer, general](#) (315 articles)
- [RNA Interference \(RNAi\)/siRNA](#) (121 articles)
- [Use of Reporter Genes](#) (115 articles)
- [Xenopus](#) (146 articles)
- [Xenopus Transgenics](#) (18 articles)

To subscribe to *Cold Spring Harbor Protocols* go to:
<http://cshprotocols.cshlp.org/subscriptions>