

Designing and Constructing CRISPR/Cas9

Soybean Precision Genomics Workshop 2014

Jean-Michel Michno

Mich0391@umn.edu

Designing a CRISPR/Cas9 for your *Glycine max* gene of choice
(Steps 1-3 credit Justin Anderson ande9112@umn.edu)

Note: All of our CRISPR/Cas9 information can be found at:

<http://stuparlab.cfans.umn.edu/protocols/crispcas9-glycine-max/>

- 1) Find your gene
 - a) [Go to www.arabidopsis.org/](http://www.arabidopsis.org/)
 - b) Search your gene of interest in the top right corner i.e. RNA Polymerase
 - c) Click on a locus ([AT#G#####](#))
 - d) Scroll down to the “Protein Data” section and click on the name ([AT#G#####](#))
 - e) Scroll down to the “Sequence” section and copy the entire amino acid sequence of your gene (including the numbers is fine)
- 2) Use BLAST to find your gene’s Soybean version
 - a) [Go to www.phytozome.net/](http://www.phytozome.net/)
 - b) Click on the large box with the list of species
 - c) Click on the little box next to *Glycine max*
 - d) Under “2. Choose a tool:” expand “BLAST search”
 - e) Select “proteome” instead of “genome: masked”
 - The bar under proteome should say BLASTP- protein query to protein db
 - f) Paste your amino acid sequence from part one into the Query sequence box (the numbers can be included and will not affect the BLAST)
 - g) Click “submit” to begin the BLASTp search
- 3) Pick a target gene from the BLAST results
 - a) Browse the list of BLAST results for a corresponding gene
 - b) Click the box “Gene Page” to open the details about this gene
 - c) Confirm the description, domain, or annotations fit your original interests
 - d) Open the “Sequences” tab near the top
 - e) Expand “Genomic sequence” to view the genomic context
 - f) The color key is in the upper right, Blue indicates CDS
 - g) Copy the entire genomic sequence
- 4) Use your Glyma to find CRISPR/Cas9 target sites
 - a) Go to <http://cfans-pmorrell.oit.umn.edu/CRISPR/>
 - b) Enter your Glyma number (v1.1 or v2.1a of the genome assembly) or sequence into the text box and click submit
 - c) Once submitted scroll down to find your designed CRISPR target region (this site is designed to target GN19NGG recognition sites only)
 - d) Pick a target site with a unique five to six base pair restriction enzyme that cuts between 15-19 bp for forward CRISPRs and 4-8bp for reverse complement CRISPR sites

e) If you are going to use Stupar Lab destination vectors order the displayed target oligos from your favorite site

5) Order your plasmids from Addgene.org (skip if you already order them)

- a) Go to <http://www.addgene.org/CRISPR/>
- b) In the right side of the page there will be an orange box with the title "CRISPR Depositing Labs" click the Robert Stupar link
- c) You will find a list of plasmids available, you MUST select the pBlu gRNA and one or more of the destination vectors

| Name | Backbone | Promoter | Non-bacterial selection | Bacterial Selection | DoubleNLS | Digestion Enzyme |
|-----------------|-------------------|----------|-------------------------|---------------------|-----------|------------------|
| pBlu gRNA | pBluescript KS(+) | U6 | NA | Ampicillin | NA | BbsI |
| Cas9 MDC32 | PMDC32 | 2x35S | Hygromycin | Kanamycin | no | EcoRI |
| Cas9 MDC123 | PMDC123 | 2x35S | Bar cassette | Kanamycin | no | EcoRI |
| G10 Cas9 MDC123 | PMDC123 | G10 | Bar cassette | Kanamycin | yes | EcoRI |
| gXL Cas9 MDC123 | PMDC123 | GmUbi XL | Bar cassette | Kanamycin | no | EcoRI |

d) Order selected plasmids and proceed to assembly

6) Assembly Round 1

a) Dilute your target oligos and anneal at 50C for 6 hours

| Annealing target oligos | Concentration | amount |
|---------------------------|---------------|--------|
| Forward target oligo | 100ng/ul | 10ul |
| Reverse target oligo | 100ng/ul | 10ul |
| PCR buffer | 10X | 2.2ul |
| Total | | 22.2ul |
| Anneal at 50C for 6 hours | | |

b) Digest pBlu gRNA with BbsI (New England Biolabs # R0539S)

| Digestion with BbsI | amount |
|---|--------|
| water | 7ul |
| NEB cutsmart buffer | 2ul |
| vector (1 microgram) | 10ul |
| BbsI (5,000 units/ml) | 1ul |
| Total | 20ul |
| Digest at 37C for 5 hrs then heat inactivate for 15 minutes | |

c) Run the product from step 6b on a gel and extract 3500 base pair band

d) Ligation of target oligos and BbsI cut pBlu/gRNA (NEB #M0202S)

| Ligation of target oligos and BbsI cut pBlu gRNA | |
|---|------------|
| Vector | 2ul |
| Target oligos | 2ul |
| T4 ligase buffer | 2ul |
| T4 ligase | 1ul |
| H2O | up to 20ul |
| Total | 20ul |
| Ligate overnight at 16C | |

e) Clone into Dh5 alpha (Life technologies #18258-012)

- i) Add 5ul of ligation to comp cells on ice
- ii) Heat shock at 42C for 30 seconds
- iii) Place cells immediately back on ice
- iv) Add 250 ul LB broth
- v) Grow cells for 30min-1hr at 37C
- vi) Plate 40-80 ul on LB agar with Ampicilli 100mg/L resistance grow overnight
- vii) Pick three colonies into 15ml tubes and grow in 3ml LB Amp 100mg/L overnight

f) Miniprep and Sequence

- i) Extract DNA from ecoli using a miniprep kit (Qiagen # 27104)
- ii) Sequence DNA using T3 primer: aattaaccctcactaaaggg

7) Assembly Round 2

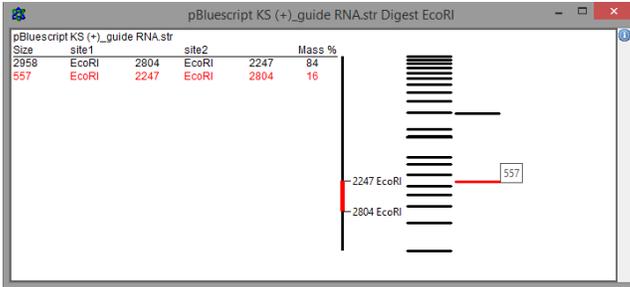
a) Digest sequenced pBlu/gRNA/insert and destination vector of choice using EcoRI (NEB # R0101S)

| Digestion with EcoRI | pBlu/gRNA/insert | Destination Vector |
|---|-------------------------|---------------------------|
| Water | 7ul | 7ul |
| NEB cutsmart buffer | 2ul | 2ul |
| Vector/insert | 10ul | 10ul |
| EcoRI | 1ul | 1ul |
| Total | 20ul | 20ul |
| Digest at 37C for 5 hrs then heat inactivate for 15 minutes | | |

b) CIP treat destination vector (NEB # M0289S)

- i) Add 2.5 ul of Antarctic Phosphatase Reaction Buffer directly to the ligation
- ii) Add 1 ul of Antarctic Phosphatase
- iii) Incubate at 37C for 1 hour
- iv) Incubate at 70C for 5 minutes

c) Run EcoRI cut pBlu/gRNA/insert on a gel



- d) Cut out 557 bp band using gel extraction kit (Qiagen # 28704)
 e) Ligation of gRNA insert and CIP treated destination vector

| Ligation of gRNA and Destination vector | |
|--|------------|
| Destination vector | 1ul |
| gRNA fragment (EcoRI cut) | 4ul |
| T4 ligase buffer | 2ul |
| T4 ligase | 1ul |
| H2O | up to 20ul |
| Total | 20ul |
| Ligate overnight at 16C | |

- f) Clone ligation into Dh5 alpha with Colony PCR
 i) Add 5ul of ligation to competent cells on ice
 ii) Heat shock at 42C for 30 seconds
 iii) Place cells immediately back on ice
 iv) Add 250 ul LB broth
 v) Grow cells for 30min-1hr at 37C
 vi) Plate entire solution on LB agar with Kanamycin 100mg/L resistance grow overnight
 vii) Pick three colonies (MAKING SURE TO REPEAT PIPETTE INTO YOUR COLONY PCR BEFORE PUTTING YOUR PIPETTE TIP INTO YOUR 15ml TUBE) and grow in 3ml LB Kanamycin 100mg/L overnight

| Colony PCR | |
|---|----------------|
| Qiagen HS+ MM | 10ul |
| primers F+R (10um) | 3ul |
| water | 7ul |
| one colony | repeat pipette |
| Step1: 95C for 5 min | |
| Step 2: 95C for 45 seconds | |
| Step 3: 52C for 45 Seconds | |
| Step 4: 72C for 1 minute | |
| Step 5: go back to step 2 for 30 cycles | |
| Step 6: 72C for 5 min | |

- g) Run colony PCR product on a gel to screen for present absent 550bp gRNA band
 h) Miniprep gRNA positive 15 ml tubes
 You have now completed CRISPR/Cas9 the next step is to either transform the vector yourself or send it the facility of your choice.