Graveley Lab CRISPR Genome Editing followed by RNA-seq Biosample Preparation and Characterization Document

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Project: ENCORE Grant: U41HG009889 Sample Description: CRISPR genome editing of EIF4E in K562 cells Cell Line: K562 RNA ID: EIF4E-BGKcLV15-33 ENCODE BIOSAMPLE ACCESSION: ENCSR033XYV

This document contains the protocols used to clone target guide sequence, to generate lentiCRISPRv2-gRNA particles, transduction of K562 cells, harvesting of RNA, characterization of the RNA integrity and measurement of target knockdown efficiency by both qRT-PCR and Western blotting.

Target guide sequence cloning protocol - Adopted from ZhangLab's

In this portion of the protocol we will clone the target sequence into the lenti-CRISPRv2.

| Item | Info |
|---------------|----------------------|
| Target | EIF4E-human |
| gRNA ID | BGC0000390 |
| gRNA sequence | AAAGCTTACCTGTTCTGTAG |

In order to clone the target sequence into the lentiCRISPRv2 , synthesize two oligos of the following form. All plasmids have the same overhangs after BsmBI digestion.

| Oligo 1 5 | · _ | Target Sequence: 1 2 3 4 5 6 7 8 9 10 11 12 13 14 15 16 17 18 9 NGG PAM CACCGNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNNN | _ | 3′ | |
|-----------|-----|---|---|--------------|-----------|
| 3 | · - | CNNNNNNNNNNNNNNNNNNNNNNCAAA | _ | 5 ′ ← | — Oligo 2 |

Lentiviral vector digestion, oligo annealing and cloning into digested vector

1. Digest and dephosphorylate 5ug of the lentiviral CRISPR plasmid with BsmBI for 30 min at 37 $^{\circ}\mathrm{C}:$

| Reagent | Quantity |
|------------------------------|------------------|
| lentiCRISPRv2 | $5 \ \mu { m g}$ |
| FastDigest BsmBI (Fermentas) | $3 \ \mu l$ |
| FastAP (Fermentas) | $3 \ \mu l$ |
| 10X FastDigest Buffer | $6 \ \mu l$ |
| 100 mM DTT | $0.6 \ \mu l$ |
| ddH_2O (freshly prepared) | X μl |
| Total volume | $60 \ \mu l$ |
| | |

2. Gel purify digested plasmid using QIAquick Gel Extraction Kit and elute in EB.

If BsmBI digested, a ~ 2 kb filler piece should be present on the gel. Only gel purify the larger band. Leave the 2kb band.

3. Phosphorylate and anneal each pair of oligos:

| Reagent | Quantity |
|------------------------------|---------------|
| Oligo 1 (100 μ M) | $1 \ \mu l$ |
| Oligo 2 (100 μ M) | $1 \ \mu l$ |
| 10X T4 Ligation Buffer (NEB) | $1 \ \mu l$ |
| T4 PNK (NEB M0201S) | $0.5 \ \mu l$ |
| ddH_2O | $6.5 \ \mu l$ |
| Total volume | $10 \ \mu l$ |

Please use the T4 Ligation Buffer since the buffer supplied with the T4 PNK enzyme does not include ATP (or supplement to 1mM ATP).

Put the phosphorylation/annealing reaction in a thermocycler using the following parameters:

 $\begin{array}{rl} 37\,^{\circ}\mathrm{C} & 30\ \mathrm{min}\\ 95\,^{\circ}\mathrm{C} & 5\ \mathrm{min}\ \mathrm{and}\ \mathrm{then}\ \mathrm{ramp}\ \mathrm{down}\ \mathrm{to}\ 25\,^{\circ}\mathrm{C}\ \mathrm{at}\ 5\,^{\circ}\mathrm{C}/\mathrm{min} \end{array}$

- 4. Dilute annealed oligos from Step 3 at a 1:200 dilution into sterile water or EB.
- 5. Set up ligation reaction and incubate at room temperature for 10 min:

| Reagent | Quantity |
|--|--------------|
| BsmBI digested plasmid from Step 2 $(50\mu g)$ | X μl |
| diluted oligo duplex from Step 4 | $1 \ \mu l$ |
| 2X Quick Ligase Buffer (NEB) | $5 \ \mu l$ |
| ddH_2O | X μ l |
| Subotal volume | $10 \ \mu l$ |
| | |
| Quick Ligase | $1 \ \mu l$ |
| Total volume | $11 \ \mu l$ |

Also perform a negative control ligation (vector-only with water in place of oligos) and transformation.

6. Transformation into Stbl3 bacteria.

Lentiviral transfer plasmids contain Long-Terminal Repeats (LTRs) and must be transformed into recombination-deficient bacteria. We use homemade Stbl3 (propagated from Invitrogen C7373-03) and get excellent plasmid yields. Although other RecA-strains may work, we have found the most consistent transformations and yields using Stbl3.

Protocol for producing lentiCRISPRV2-gRNA particles

Day 1

- Plate 0.8-1x10⁶ 293 T cells (catalog number: CRL-11268, ATCC) in each well of 6-well plate with 10 % FBS (catalog number: 30-2020, ATCC) DMEM (catalog number: 11995-065, Life technologies) medium without penicillin and streptomycin.
- 2. Incubate overnight. Cells should be 70-80% confluent.

Day 2

1. In polypropylene tubes, make a cocktail for each transfection as follows:

| Reagent | Quantity |
|----------------------|---------------------|
| lentiCRISPRv2-gRNA | $1 \ \mu { m g}$ |
| psPAX2 Packaging DNA | $750 \ \mathrm{ng}$ |
| pCMV-VSV-G DNA | 250 ng |
| serum-free OPTI-MEM | to 100 μl |

- 2. Add 6 μ l of FuGENE HD Transfection reagent (Catalog number: E2311, Promega) to the tube (FuGENE:DNA=3:1)
- 3. Incubate for 20 minutes at room temperature.
- 4. Gently add the DNA mix dropwise to cells.
- 5. Incubate the cells at $37\,^{\rm o}{\rm C}$ for 12-15 hr.

Day 3

1. In the morning, change the media to remove the transfection reagent, wash with PBS once and add 1.5 ml fresh media +10% FBS + penicillin/streptomycin.

Day 4

- 1. Harvest media from cells, store at $4\,^{\circ}\mathrm{C}.$
- 2. Add 1.5 ml fresh media.

Day 5

- 1. Harvest the media from the cells and pool with the media collected on Day 4.
- 2. Spin the media at 1250 rpm for 5 min to remove cells.
- 3. Freeze the virus stock at $-40\,^{\circ}\mathrm{C}.$

Lentiviral CRISPR Protocol

For K562 cells

Source: ATCC CCL-243 (lot 59300853)

Growth Media for K562 cells:

500 ml RPMI-1640 with glutamine medium (Hyclone, SH30027.01) 50 ml Fetal Bovine Serum (FBS) (10% Final Concentration) (Hyclone, SH30071.03) 5 ml Pen-Strep (1% Final Concentration) (Invitrogen, 15140-163)

Culturing

- 1. Remove a frozen stock vial of K562 cells from liquid nitrogen.
- 2. Thaw it in a $37 \,^{\circ}$ C water bath.
- 3. As soon as it thawed, transfer the cells into the growth medium.
- 4. Centrifuge at 1000 rpm for 5 minutes.
- 5. Resuspend the cells in the growth medium such that the final cell density is about $1x10^{5}$ cells/ml.
- 6. Split the cells every 2 to 3 days.
- 7. Grow the cells to required numbers.

Prepare cells for transduction

Day 0

- 1. Plate 0.8-1ml of lentiCRISPRv2-gRNA particles in each well of 12-well plates.
- 2. Add 5-7x10^5 cells to appropriate wells with 8 μ g/ml of polybrene (Catalog Number H9268, Sigma-Aldrich), incubate at 37 °C for 2 hours, and then add 1 ml of complete medium.

Day 1

1. After 24 hrs, change to fresh media (2 ml) with 3 μ g/ml of puromycin.

Day 3

1. Change to fresh media (2 ml) with 3 μ g/ml of puromycin.

Day 5

1. Change to fresh media (2 ml) with 3 μ g/ml of puromycin.

Day 6

1. Harvest half of the cells to prepare RNA and half of the cells to prepare a protein lysate for western blotting.

RNA Isolation

RNA isolation is performed using a Promega Maxwell (R)16 Instrument and the Maxwell (R)16 LEV simply RNA Cells Kits (Catlog Number AS1270).

1. Pellet cells at 300 x g for 3 minutes and remove medium.

- 2. Add 200 μ l of chilled 1-Thioglycerol/Homogenization solution to the cell pellet and vortex until the pellet is dispersed.
- 3. Add 200 μ l of lysis buffer and votex vigorously for 15 sec to mix.
- 4. Transfer all 400 μl lysate to well 1 of the Maxwell 16 LEV cartridge.
- 5. Add 5 μ l of DNase I solution to well 4 of the cartridge.
- 6. Put elution tubes with 40-50 μl of nuclease-free water and LEV plungers in the cartridge.
- 7. Transfer the Maxwell 16 LEV cartridge rack containing prepared cartridges on the Maxwell 16 Instrument.
- 8. Push Run/Stop button to start run.

RNA Quality Control

The quality of the RNA is measured using an Agilent TapeStation Instrument with the RNA screen tape (Catlog Number 5067-5576).

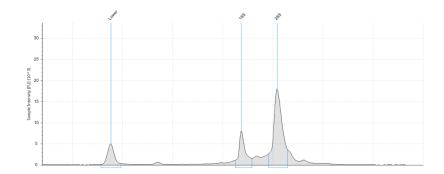


Figure 1: Agilent TapeStation image of 1 μ l of total RNA sample.

qRT-PCR Assay to Monitor mRNA Target Knockdown Efficiency

cDNA Synthesis

This assay uses the iScript cDNA Synthsis Kit from BIO-RAD (Catalog number: 170-8891)

1. Reaction Setup:

| Reagent | Quantity |
|---------------------------------|-------------------------------|
| 5x iScript reaction mix | $2 \ \mu l$ |
| iScript reverse transcriptase | $0.5 \ \mu l$ |
| Nuclease-free water | $\mathbf{x} \ \mu \mathbf{l}$ |
| RNA template (200 ng) | $\mathbf{x} \ \mu \mathbf{l}$ |
| Total volume | $10 \ \mu l$ |

2. Reaction Protocol:

| Time | Temperature |
|------------|-------------|
| 5 minutes | 25 °C |
| 30 minutes | 42 °C |
| 5 minutes | 85 °C |
| Hold | 4 °C |

qPCR Assay

This assay uses Phusion High-Fidelity DNA Polymerase from NEB (Catalog number: M0530L) and SYBR Green from Invitrogen (Catalog number: S7563)

1. Reaction setup:

| Reagent | Quantity |
|----------------------------------|---------------|
| 5X Phusion HF Buffer | $4 \ \mu l$ |
| 10 mM dNTPs | $0.4 \ \mu l$ |
| 10 μ M Forward Primer | $1 \ \mu l$ |
| $10 \ \mu M$ Reverse Primer | $1 \ \mu l$ |
| Template (1:20 of cDNA reaction) | $1 \ \mu l$ |
| Phusion DNA Polymerase | $0.2 \ \mu l$ |
| SYBR Green $(10,000 \text{ X})$ | $0.1 \ \mu l$ |
| Nuclease-free water | to 20 μl |
| Total volume | $20 \ \mu l$ |

2. Reaction Protocol:

| STEP | TEMP | TIME |
|----------------------|---------------------------|------------|
| Initial Denaturation | 98 °C | 30 seconds |
| 35 Cycles | 98 °C | 10 seconds |
| | $58-66^{\circ}\mathrm{C}$ | 15 seconds |

3. Data Analysis:

Data analysis is perfored using the $2-\Delta\Delta Ct$ Method.

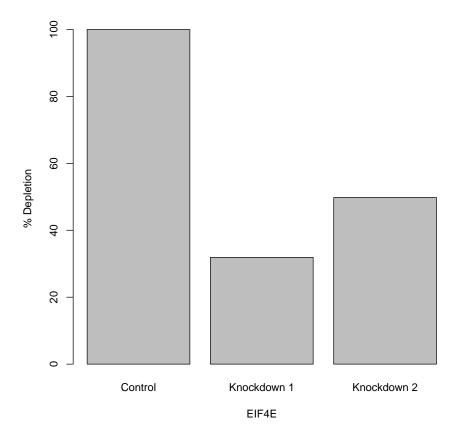


Figure 2: qRT-PCR analysis of depletion level of the target RNA binding protein in control and knockdown cells. The percent depletion was calculated in the RNA sample isolated from K562s transduced with a gRNA targeting EIF4E (ENCODE Biosample ENCSR033XYV and ENCBS485INP) in comparison to K562 cells transduced with a control non-target gRNA (ENCODE Biosample ENCSR404OHQ and ENCBS996SVS). The efficiency of depletion is normalized using Beta-actin as a control.

Western Blot Assay to Monitor Protein Target Knockdown Efficiency

A western blot is performed to determine the knockdown efficiency of the target RNA binding protein. For this biosample, the following antibodies were used:

RNA binding protein primary antibody: EIF4E

Loading control primary antibody: Beta-actin

1. Purify protein by resuspending cell pellets in 100 μ l Lysis buffer (50 mM

Tris-HCl, pH 7.4; 100 mM NaCl; 1% NP-40; 0.1% SDS; 0.5% sodium deoxycholate) with protease inhibitor (Roche cocktail).

- 2. Vortex vigorously and leave tubes on ice at least 30 min.
- Spin 18,000g, 20 min, 4 °C. Recover the supernatant in a new tube on ice. Discard the pellet.
- 4. Quantitate the protein concentration using the Pierce BCA kit.
- 5. Load 30–60 μ g of protein per sample depending on expression levels of the protein of interest in the cell type.
- 6. Dilute the protein samples with 4X sample buffer and 10X reducing agent (Invitrogen NuPage reagents) and heat at 70 °C for 10 min.
- 7. While samples are heating, prepare the gel: 4-12% Bis–Tris gel for proteins 10-220 kDa, or 8% Tris–glycine for proteins over 220 kDa. Dilute MOPS running buffer to 1X and pour into buffer tank. Add 500 μ l of antioxidant to the inner chamber of the buffer tank. Wash out all wells of the gel with buffer.
- 8. Load 3 μ l of Licor Odyssey prestained molecular weight marker (928–4000) and protein samples. Run the gel at 200 V for about an hour, until the dye just runs off the gel (less for small molecular weight proteins).
- 9. Transfer to PVDF for 30 min in transfer buffer, with methanol and antioxidant, using BioRad Semi-dry transfer apparatus.
- 10. After transfer, block membrane in 5-10 mls of Licor Blocking buffer for 1 hour at room temperature.
- 11. Incubate with the RNA binding protein primary antibody $(0.2 \ \mu g/ml)$ and the loading control primary antibody (mfg recommended dilution) diluted in Licor block with 0.1% tween 20 on a rocker at 4 °C overnight.
- 12. The next day, wash the blot 4 x 5 min each in TBST.
- 13. Prepare the secondary antibody (Rockland Fluorescent TrueBlot antirabbit IgG IRDye800 (Catalog number 18-3216-32) for the RBP and Licor IRDye680 secondary antibody for the loading control). Dilute the antibodies according to the mfg instructions in Licor blocking buffer with 0.1% Tween 20 and 0.01% SDS. Incubate on a rocker for 30-60 min, at room temperature.
- 14. Wash 4 x 5 min each in TBST. Rinse 1x in TBS (no T) and scan on the Licor Odyssey instrument.

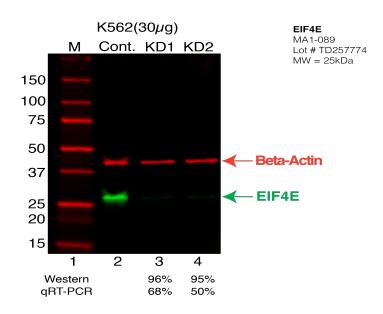


Figure 3: Western Blot Analysis of depletion level of the target RNA binding protein in control and knockdown cells. Lane 1: Molecular weight marker. Lane 2: 30 μ g of protein from K562 transduced with a control, non-target gRNA (ENCODE Biosample ENCSR404OHQ and ENCBS996SVS). Lane 3: 30 μ g of protein from K562 transduced with a gRNA targeting EIF4E (ENCODE Biosample ENCSR033XYV). Lane 4: 30 μ g of protein from K562 transduced with a gRNA targeting EIF4E (ENCODE Biosample ENCSR033XYV). Lane 4: 30 μ g of protein from K562 transduced with a gRNA targeting EIF4E (ENCODE Biosample ENCBS485INP). Samples were separated by SDS-PAGE, transferred to a membrane and blotted using antibodies against EIF4E (ENCODE Antibody ENCAB864VOP) and Beta-actin as controls.