## 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 BAZ1A in K562 cells Cell Line: K562 RNA ID: BAZ1A-BGKcLV17-3 ENCODE BIOSAMPLE ACCESSION: ENCSR825EPB

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	BAZ1A-human
gRNA ID	BGC0000453
gRNA sequence	CCGGCTCTTACTCGTAGTGG

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.

## 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 $\mu l$
Total volume	$60 \ \mu l$

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

If BsmBI digested, a  $\sim 2$ 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 $\mu$ M)	$1 \ \mu l$
Oligo 2 (100 $\mu$ 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 $\mu$ 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 0.8-1x10<sup>6</sup> 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 $\mu l$

- 2. Add 6  $\mu$ 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  $\mu$ 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  $\mu$ g/ml of puromycin.

#### Day 3

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

#### Day 5

1. Change to fresh media (2 ml) with 3  $\mu$ 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  $\mu$ l of chilled 1-Thioglycerol/Homogenization solution to the cell pellet and vortex until the pellet is dispersed.
- 3. Add 200  $\mu$ l of lysis buffer and votex vigorously for 15 sec to mix.
- 4. Transfer all 400  $\mu l$  lysate to well 1 of the Maxwell 16 LEV cartridge.
- 5. Add 5  $\mu$ l of DNase I solution to well 4 of the cartridge.
- 6. Put elution tubes with 40-50  $\mu l$  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).

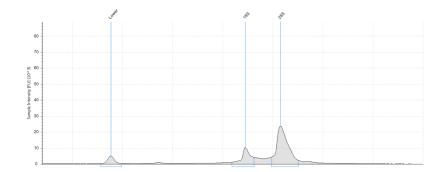


Figure 1: Agilent TapeStation image of 1  $\mu$ l of total RNA sample.

## 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 $(200 \text{ ng})$	$\mathbf{x} \ \mu \mathbf{l}$
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 $\mu$ 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 $\mu l$
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.

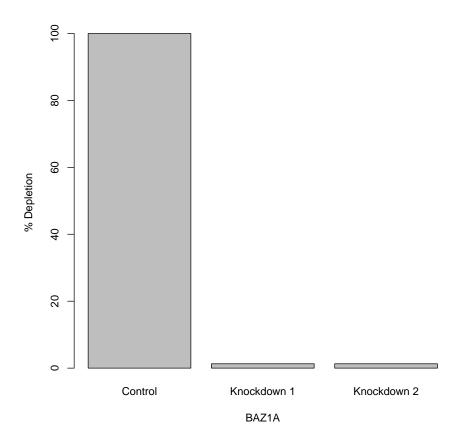


Figure 2: qRT-PCR analysis of depletion level of the target RNA binding protein in control and knockdown cells. The percent depletion was calculated in the RNA sample isolated from K562s transduced with a gRNA targeting BAZ1A (ENCODE Biosample ENCSR825EPB and ENCBS287JVX) in comparison to K562 cells transduced with a control non-target gRNA (ENCODE Biosample ENCSR516TLX and ENCBS813SYB). The efficiency of depletion is normalized using GAPDH 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: BAZ1A

Loading control primary antibody: GAPDH

## Wes Protocol

12-230 kDa Master kit with split Running Buffer from Proteinsimple

## A. PREPARE STANDARD PACK REAGENTS

- 1. Add 40  $\mu$ L deionized water to make a 400 mM solution of the DTT.
- 2. Add 20  $\mu L$  10X Sample buffer and 20  $\mu L$  prepared 400 mM DTT to make 5X Fluorescent Master Mix.
- 3. Add 16  $\mu$ L deionized water, 2  $\mu$ L 10X Sample Buffer and 2  $\mu$ L prepared 400 mM DTT solution to make Biotinylated Ladder.

#### **B. PREPARE YOUR SAMPLES**

- 1. Combine 1 part 5X Fluorescent Master Mix with 4 parts lysate in a microcentrifuge tube (If needed, dilute the lysate with 0.1X Sample Buffer).
- 2. Denature the samples and biotinylated ladder at 70 °C for 10 min.

## C. PREPARE REAGENTS FROM DETECTION MODULE

- 1. Dilute primary antibody with antibody diluent 2.
- 2. The supplied secondary antibody is ready to use without dilution.
- 3. Combine 150  $\mu$ L Luminol-S and 150  $\mu$ L Peroxid in a microcentrifuge tube. Gently pipette up and down to mix and store on ice.

### D. PIPETTE YOUR PLATE (IMMUNOASSAY)



For more consistent results, keep the lid on between reagent additions. Centrifuge the plate for 5 minutes at 2500 rpm at room temperature.



## E. START WES

- 1. Load the desired assay in Compass software v2.7 or higher.
- 2. Open Wes' door.
- 3. Insert a capillary cartridge into the cartridge holder. The interior light will change from orange to blue.
- 4. Remove the assay plate lid. Hold plate firmly on bench and carefully peel off evaporation seal. Pop any bubbles observed in the Separation Matrix wells with a pipette tip.
- 5. Place the assay plate on the plate holder
- 6. Close Wes' door.
- 7. Click the Start button in compass.
- 8. When the run is complete, discard the plate and cartridge.

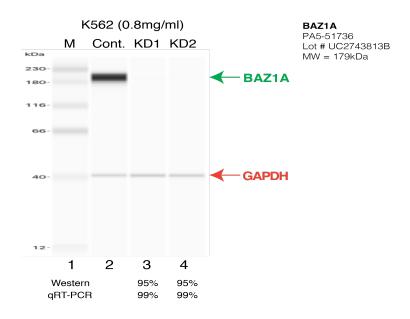


Figure 3: Western Blot Analysis of depletion level of the target RNA binding protein in control and knockdown cells. Lane 1: Molecular weight marker. Lane 2: 30  $\mu$ g of protein from K562 transduced with a control, non-target gRNA (ENCODE Biosample ENCSR516TLX and ENCBS813SYB). Lane 3: 30  $\mu$ g of protein from K562 transduced with a gRNA targeting BAZ1A (ENCODE Biosample ENCSR825EPB). Lane 4: 30  $\mu$ g of protein from K562 transduced with a gRNA targeting BAZ1A (ENCODE Biosample ENCBS287JVX). Samples were separated by SDS-PAGE, transferred to a membrane and blotted using antibodies against BAZ1A (ENCODE Antibody ENCAB850VOO) and GAPDH as controls.