

**!!! RNase-free working environment is required for the whole procedure!!!**

## sgRNA preparation

### . Amplify the template for *in vitro* sgRNA transcription

1) Oligos need to be ordered:

Template: TAATACGACTCACTATAGGGGCCACTAGGGACAGGATGTTTT  
 AGAGCTAGAAATAGCAAGTTAAAATAAGGCTAGTCCGTTATCAACTTGAAAA  
 AGTGGCACCGAGTCGGTGTCTTTTTT  
Fw: ggcactTAATACGACTCACTATAGG  
Rv: cggagcGAAAAAGCACCGACTC

2) PCR reaction mix:

**!! 8-16 reactions will be pooled together.**

Template (10ng/μL)		1μL
Fw (10μM)		1μL
Rv (10μM)		1μL
5x HF buffer		4μL
dNTP (10mM)	0.4μL	
DMSO		0.6μL
Phusion		0.2μL
H <sub>2</sub> O		11.8μL

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Total reaction volume 20μL

3) PCR cycles:

98°C	20s	
98°C	10s	
60°C	25s	x34
72°C	5s	
72°C	3m	
4°C	∞	

4) PCR purification

! Use one column to bind DNA.

! Elute with 20μL Nuclease-free water.

**!! The expected concentration should be more than 300ng/μL.**

! Adjust the concentration to 250ng/μL as the template of the sgRNA synthesis.

### . *In vitro* sgRNA synthesis using MEGAscript™ T7 Kit

Thaw the T7 10X Reaction Buffer at room temperature by vortexing.

**CRITICAL!!! Make sure there is NO DTT crystal remaining in the T7 10X Reaction Buffer.**

When a new kit is opened, thaw the T7 nTP Solutions, transfer 50μL of each T7 nTP Solution into a new tube, well-mixed and labeled as 'T7 nTP Mix'

**!! Assemble the reaction mix (in PCR tube) at room temperature, in the following order:**

**!! 3 reactions will be pooled together.**

Nuclease-free water	7μL
T7 10X Reaction Buffer	2μL
T7 nTP Mix	8μL

Template DNA (250ng/ $\mu$ L)	1 $\mu$ L
T7 Enzyme Mix	2 $\mu$ L
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Total reaction volume	20 $\mu$ L

Mix thoroughly by pipetting or flicking the tube, then centrifuge the tube briefly.  
Incubate in a thermocycler at 37°C for 4 hours.

#### . DNase treatment

sgRNA synthesis reaction	20 $\mu$ L
TURBO DNase	1 $\mu$ L

Mix well by pipetting or flicking. Incubate at 37°C for 15min

! After DNase treatment, proceed immediately to the purification procedure. No DNase inactivation needed!

#### . sgRNA purification using MEGAclean™ Kit

Set the heating block to 95°C, pre-heat elution buffer in a 1.5mL eppendorf.

1. Pool 3 DNase-treated samples together (in total 60 $\mu$ L), add 40 $\mu$ L of Elution Solution (4°C, **NOT the pre-heated!!!**). Vortex briefly to mix thoroughly. Briefly spin the tube.
2. Add 350 $\mu$ L of Binding Solution Concentration (4°C) to the sample. Mix gently by pipetting up and down immediately after adding.
3. Take 100% Ethanol from -20°C. Add 250 $\mu$ L of cold 100% ethanol to the sample. Invert the tube few times to mix thoroughly. Stand on ice for 1min, and invert again for few times.

**NOTE: This is the critical step, directly affecting the final yield!!!**

1). The cold ethanol is very important for this step, to precipitate the RNA!!! So, take your 100% ethanol from -20°C just before you start on this step

2) Because ethanol precipitates the RNA, after thoroughly mixing, the solution can become opaque suspension. **DO NOT SPIN THE SAMPLES AFTER MIX!!!**

4. Load the sample to a filter with a collection tube.
5. Centrifuge for 30s at 12000xg at r/t. Remove the flow-through.
6. Wash the filter 2 times with 500 $\mu$ L of Washing Solution (4°C). Centrifuge for 30s at 12000xg at r/t. Discard the flow-through.
7. Centrifuge again for 1min at 12000xg to completely remove the Washing Solution.
8. Remove the filter to a new collection tube.
9. Apply 50 $\mu$ L of pre-heated Elution Solution to the center of the filter. Centrifuge for 1min at 12000xg at r/t.

**NOTE: pipetting the pre-heated elution buffer at the heating block, in stead of taking the pre-heated elution solution to the RNA hood.**

10. Apply another 50 $\mu$ L of pre-heated Elution Solution to the same filter. Centrifuge for 1min at 12000xg at r/t.
11. Add 10 $\mu$ L of 5M NH<sub>4</sub>Ac to 100 $\mu$ L of the eluted sample. Vortex to mix well and spin down briefly.
12. Add 275 $\mu$ L of cold 100% ethanol (from -20°C). Mix well by inverting the tube for few times.
13. Stand the tube at -20°C for at least 30min. (Normally I do it for 45 min)
14. Centrifuge at maximum speed for 15min at 4°C.
15. Carefully remove the supernatant by pipetting, without disturbing the pellet.
16. Add 500 $\mu$ L of cold 70% ethanol (from -20°C). Centrifuge at maximum speed for 5min at 4°C.

17. Remove most of the supernatant by pipetting. Centrifuge again, at maximum speed for 1min at 4°C. Carefully remove the rest of supernatant by pipetting.
18. Air-dry the pellet, 5-8 mins.  
**!!! DO NOT over-dry the pellet!!!** When the pellet is dried, the body of the pellet should remain white, whereas the edge of the pellet turns transparent.
19. Dissolve the pellet in 40µL (varies depending on the size of your pellet) Nuclease free water.
20. Store sgRNA samples at -80°C, after checking the concentration\*.

\*For checking the concentration with Nanodrop:

sgRNA	2µL
H <sub>2</sub> O	38µL
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In total	40µL

Mix thoroughly, and spin down briefly.

Check the concentration with Nanodrop (RNA-40).

**!!! From 3 reactions, the expected yield should be more than 150µg, or with the minimum concentration above 3.5µg/µL.**

**. Prepare sgRNA for the assay**

Dilute the sgRNA to 1µM (1pmol/µL, 31ng/µL).

To cut 100fmol substrate with 1pmole CAS9, 3-4pmol of sgRNA is recommended.

**NOTE: repeatedly freeze and thaw the sgRNA samples can sharply decrease quality. We suggest to aliquot you sgRNA samples in working concentration and store at -80°C.**