

# TF ORF barcode dial out PCR (10X)

## Introduction

Get started by giving your protocol a name and editing this introduction.

## Materials

- › Forward primers
  - › ORF-perturb-NGS-F: AATGATAACGGCGACCACCGAGATCTACACTCTTCCCTACACGACGCTTCCGATCT
- › Reverse primers
  - › ORF-perturb-NGS-R1:  
CAAGCAGAACAGCGCATACGAGATTGCCTGGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTTAAGTAGAGTTGGCTTATA  
TATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R2:  
CAAGCAGAACAGCGCATACGAGATAGCGTCGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTATCATGCTTATTGGCTTATA  
TATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R3:  
CAAGCAGAACAGCGCATACGAGATGAAGAAGTGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTGACATCTTGGCTT  
ATATATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R4:  
CAAGCAGAACAGCGCATACGAGATATTCTAGGGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTCGATTGCTCGACTTGGCTT  
ATATATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R5:  
CAAGCAGAACAGCGCATACGAGATCGTACCGAGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTCGATAGCAATTCTGGCTT  
ATATATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R6:  
CAAGCAGAACAGCGCATACGAGATGTCTGATGGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTCGATAGTTGCTTGGCTT  
TTATATATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R7:  
CAAGCAGAACAGCGCATACGAGATTACGCACGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTGATCCAGTTAGTTGG  
CTTATATATCTTGAAAGGACGA
  - › ORF-perturb-NGS-R8:  
CAAGCAGAACAGCGCATACGAGATTGAATAGGTGAUTGGAGTTAGACGTGTGCTCTCCGATCTCGATCGATTGAGCCTTG  
GCTTATATATCTTGAAAGGACGA
- › NEBNext High Fidelity PCR Master Mix, 2× (New England BioLabs, cat. no. M0541)

## Procedure

### Library PCR for NGS

- We have provided NGS primers that amplify the TF ORF barcode region with Illumina adaptor sequences. Pool the 10 ORF-perturb-NGS-R primers at equimolar ratio to produce ORF-perturb-NGS-R (pooled). These primers stagger the resulting PCR amplicons and diversify the library.
- To prepare the TF ORF barcode library for NGS, set up a reaction using the ORF-pertrub-NGS-F and ORF-perturb-NGS-F (pooled) for each 10X channel according to the following table.

<b>NEBNext PCR reaction components</b>				
	<b>A</b>	<b>B</b>	<b>C</b>	<b>D</b>
1	<b>Component</b>	<b>Starting concentration</b>	<b>Amount per reaction (uL)</b>	<b>Final concentration</b>
2	<b>NEBNext PCR Master Mix</b>	2X	25	1X
3	<b>ORF-perturb-NGS-F</b>	10 uM	1.25	0.25 uM
4	<b>ORF-perturb-NGS-R (pooled)</b>	10 uM	1.25	0.25 uM
5	<b>WTA cDNA template</b>	10-20 ng/uL	2	0.4-0.8 ng/uL
6	<b>H2O</b>		20.5	
7			50	

- Perform a PCR using the following cycling conditions:

<b>NEBNext PCR Cycle Conditions</b>				
	<b>A</b>	<b>B</b>	<b>C</b>	<b>D</b>
1	Step	Temperature	Duration	Cycles
2	Initial denaturation	98 °C	3 min	1
3	Denaturation	98 °C	10 sec	22
4	Annealing	60 °C	10 sec	
5	Extension	72 °C	40 s	
6	Final extension	72 °C	2 min	1

- After the reaction is complete, proceed according to the "Next-generation sequencing of the amplified sgRNA library to determine sgRNA distribution" section in Joung, Nature Protocols 2017.