### Chromium Total: single-nucleus total RNA sequencing via in situ polyA-tailing

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#### Materials/Reagents

- Yeast PAP
- 5X Yeast PAP reaction buffer
- 1M Tris-HCl, pH 7.5
- 5M NaCl
- 1M MgCl<sub>2</sub>

- 1M Sucrose
- Nuclease-Free H2O
- 10X PBS
- 10% BSA
- 20U/ul SUPERase in RNAse Inhibitor

#### <u>Buffers</u>

#### Yeast PAP Enzyme Mix:

	Final Conc.	Volume to add		
		1 RXN	5 RXNs	
Yeast PAP	0.25M	2ul	10ul	
25mM ATP	1%	1ul	5ul	
SUPERase		2ul	10ul	
Total Volume		5ul	25ul	

# Yeast Reaction Buffer:

	Final Conc.	Volume to add			
5X Yeast PAP reaction buffer	1X	50ul	20ul		
SUPERase	0.2U/ul	2.5ul	20ul		
H2O		197.5ul			
Total Volume		250ul	1000ul		

# **Nuclei Buffer** (from sciRNAseq3):

	Final Conc.	Volume to add		
1M Tris-HCl, pH 7.5	10mM	500µl		
5M NaCl	10mM	100μΙ		
1M MgCl <sub>2</sub>	3mM	150µl		
Nuclease-Free H <sub>2</sub> O		49.25ml		
Total Volume		50ml		

## Nuclei Suspension Buffer (NSB) (from sciRNAseq3):

	Final Conc.	Volume to add			
Nuclei Buffer (NB)		4.8ml			
*20U/µl SUPERase	1% v/v	50 μΙ			
10% BSA	1% w/v	100µl			
Total Volume		5ml			

<sup>\*</sup>Add just before use

# **Protocol**

- 1. Set hotplate/thermomixer to 37°C and centrifuge to 4°C.
- 2. Prepare appropriate volumes of Yeast PAP Enzyme Mix and Yeast Reaction Buffer.
- 3. Thaw, resuspend, and pellet nuclei according to the methanol fixation protocol (5min, 1000 x g, 4°C).
- 4. Resuspend nuclei in **200ul** of **Yeast Reaction Buffer**. Pellet nuclei by spinning (5min, 1000 x g, 4°C).
- 5. Gently resuspend in 45ul of Yeast Reaction Buffer.
- 6. Add 5ul of Yeast PAP Enzyme Mix.
- 7. Incubate at 37°C for 25min without shaking (shaking causes clumping).
- 8. Wash nuclei in 500ul of NSB.
- 9. Resuspend in appropriate volume for downstream applications (10x Chromium, drop-seq, etc)