Kit v1 - Manual v1.0

CUTANA[™] CUT&TAG Kit

QUICK-START CARD



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DAY 1

Section I: Buffer Prep (~30 min)

1. Prepare CUT&Tag buffers (see Table). Buffers contain 20% excess volume - no additional overage is needed.

BUFFER	COMPONENTS	1X	8X	16X	STORAGE
Nuclear Extraction Buffer	Pre-Nuclear Extraction Buffer	235 µL	1.9 mL	3.8 mL	lce for use on Day 1
	25X Protease Inhibitor	9.8 µL	78.4 µL	157 µL	
	1 M Spermidine	0.13 µL	1.0 µL	2.0 µL	
Wash Buffer 1	Pre-Wash Buffer	1.3 mL	10.4 mL	20.8 mL	4°C for use on Day 2
	25X Protease Inhibitor	56 µL	448 µL	896 µL	
	1 M Spermidine	0.7 µL	5.6 µL	11.2 µL	
	5% Digitonin	2.8 µL	22.4 µL	44.8 µL	
	Wash Buffer 1	600 µL	4.8 mL	9.6 mL	4°C for use on Day 2
	4.5 M NaCl	20.7 µL	166 µL	331 µL	
Antibody Buffer	Wash Buffer 1	60 µL	480 µL	960 µL	lce for use on Day 1
	0.5 M EDTA	0.25 µL	2 µL	4 µL	

Section II: Bead Activation (~30 min)

- Resuspend ConA beads and transfer 11 µL/reaction to a 1.5 mL tube. Place tube on a magnet, allow slurry to clear, and pipette to remove supernatant.
- Resuspend beads in 100 µL/reaction cold Bead Activation Buffer. Return to magnet, allow slurry to clear, and remove supernatant. Repeat one time.
- 4. Resuspend beads in 11 µL/reaction cold Bead Activation Buffer.
- 5. Aliquot 10 µL/reaction of bead slurry into 8-strip tubes. Place on ice.

Section III: Nuclei Prep and Binding to Beads (~30 min)

- 6. Harvest 100,000 cells/reaction in a 1.5 mL tube. Spin 600 x g for 3 min at room temperature (RT).
- 7. Remove supernatant and resuspend cells in 100 µL/reaction cold Nuclear Extraction Buffer.
- 8. Incubate 10 min on ice. Spin 600 x g for 3 min at 4°C. Pipette to remove supernatant.
- Resuspend nuclei in 105 µL/reaction cold Nuclear Extraction Buffer. Add 100 µL nuclei to 10 µL ConA beads in 8-strip tubes. Gently vortex to mix and quick spin to collect bead slurry.
- 10. Incubate 10 min at RT to adsorb cells to beads.
- 11. Place tubes on magnet, allow slurry to clear, remove supernatant. Take tubes off magnet and resuspend in 50 µL/reaction cold **Antibody Buffer**.

Section IV: Primary Antibody Binding (~30 min + overnight)

- Quick spin the K-MetStat Panel stock and mix by pipetting (do NOT vortex). To reactions designated for H3K27me3 & IgG control antibodies, add 2 μL K-MetStat Panel and vortex to mix. If using <100,000 nuclei, decrease K-MetStat Panel amount as per the manual instructions.
- 13. Add 0.5 µg primary antibody to each reaction. For designated control reactions, add 1µL H3K27me3 positive control antibody & 1 µL IgG negative control antibody. Vortex to mix.
- 14. Incubate overnight on a nutator at 4°C, caps elevated. Do NOT rotate tubes.



DAY 2

Section V: Secondary Antibody Binding (~1 hr)

- 15. Prepare Tagmentation Buffer: Transfer 60 μL/reaction Wash Buffer 2 to a 1.5 mL tube. Add 1 M MgCl₂ at a 1:100 dilution (10 mM final concentration). Place on ice. Recipe includes 20% excess volume.
- 16. Quick spin tubes, place on a magnet, and allow slurry to clear. Remove supernatant and resuspend in 50 µL/reaction cold **Wash Buffer 1**.
- 17. Add 0.5 μg/reaction secondary antibody. For control antibodies and other rabbit primary antibodies, use 0.5 μL **anti-rabbit secondary antibody**. Gently vortex to mix.
- 18. Incubate on nutator, caps elevated, for 30 min at RT.
- 19. Quick spin tubes, place on a magnet, and allow slurry to clear. Remove supernatant.
- 20. On magnet, add 200 µL/reaction cold Wash Buffer 1. Remove supernatant. Repeat one time.
- 21. Remove tubes from magnet and resuspend in 50 µL/reaction cold Wash Buffer 2.

Section VI: pAG-Tn5 Binding & Targeted Chromatin Tagmentation (~4 hrs)

- 22. Add 2.5 µL/reaction pAG-Tn5 and gently vortex to mix. Incubate on nutator, caps elevated, for 1 hr at RT.
- 23. Quick spin tubes, place on a magnet, and allow slurry to clear. Remove supernatant.
- 24. Resuspend in 200 μL/reaction cold **Wash Buffer 2**. Return tubes to magnet, allow slurry to clear, and remove supernatant. Repeat one time.
- 25. Resuspend in 50 µL/reaction cold Tagmentation Buffer. Incubate 1 hr in a thermocycler with block set to 37°C and heated lid set to 47°C.
- 26. Quick spin tubes, place on magnet, and allow slurry to clear. Remove supernatant.
- 27. Add 50 μL/reaction **TAPS Buffer** and gently vortex to resuspend (do **NOT** pipette-mix after adding TAPS Buffer). Place on magnet, allow slurry to clear, and remove supernatant.
- 28. Add 5 µL/reaction SDS Release Buffer, vortex to mix (do NOT pipette), and quick spin.
- 29. Incubate 1 hr in a thermocycler with block set to 58°C and heated lid set to 68°C.
- 30. Quick spin tubes. Add 15 µL/reaction SDS Quench Buffer, vortex to mix (do NOT pipette), and quick spin.

Section VII: Indexing PCR & Library Cleanup (~1 hr)

- Assign a unique pair of i5 & i7 indexing primers to each reaction. See kit manual, Appendix 3.
- 32. To the entire reaction mixture (including ConA beads) add: 2 μL i5 primer, 2 μL i7 primer, and 25 μL Non-Hot Start 2X PCR Master Mix. Vortex to mix and quick spin to collect liquid.
- Perform PCR in a thermocycler with a heated lid (105°C) using the parameters in the table. During PCR, prepare 80% EtOH.
- 34. After PCR, quick spin tubes. Resuspend **SPRIselect** reagent (manufactured by Beckman Coulter, Inc.*) and slowly add 65 μ L/reaction. Vortex to mix and quick spin.
- 35. Incubate 5 min at RT. Place tubes on magnet for 2 min. Pipette to remove supernatant.
- 36. On magnet, add 200 µL/reaction 80% EtOH. Pipette to remove supernatant. Repeat one time.
- 37. Quick spin and return tubes to magnet. Remove residual EtOH.
- 38. Remove tubes from magnet and air-dry 2 min at RT, caps open.
- 39. Add 17 µL/reaction 0.1X TE Buffer to elute libraries. Vortex to resuspend beads and incubate 2 min at RT.
- 40. Place tubes on magnet for 2 min. Transfer 15 µL eluted CUT&Tag libraries to new 8-strip tubes.

Section VIII: Analysis of Library Fragment Size (~1 hr)

- 41. Quantify libraries using the Qubit[™] fluorometer and 2X dsDNA HS Assay Kit.
- 42. Confirm fragment size distribution (~300 bp) on the Agilent BioAnalyzer® or TapeStation®.
- 43. Store prepared libraries at -20°C. See manual for guidance on sample pooling and Illumina® sequencing.

EpiCypher.
Bringing Epigenetics to Life

ТЕМР

58°C

72°C

98°C

98°C

60°C

72°C

4°C

TIME

5 min

5 min

45 sec

15 sec

10 sec

1 min

m

1

1

1

14-21

1

1

STEP #

1

2

3

Δ

5

6

7

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