

Aug 07, 2025

KAPP-Sen TMC: Xenium - Cell DIVE - H&E

DOI

<https://dx.doi.org/10.17504/protocols.io.n92ld6zj7g5b/v1>

Emily Soja¹, Shruti Bhargava¹, Santhosh Sivajothi¹, William F Flynn¹, Elise T Courtois¹

¹Single Cell Biology Lab, The Jackson Laboratory, USA

Cellular Senescence Net...



Sergii Domanskyi

The Jackson Laboratory for Genomic Medicine

Create & collaborate more with a free account

Edit and publish protocols, collaborate in communities, share insights through comments, and track progress with run records.

Create free account

OPEN  ACCESS



DOI: <https://dx.doi.org/10.17504/protocols.io.n92ld6zj7g5b/v1>

Protocol Citation: Emily Soja, Shruti Bhargava, Santhosh Sivajothi, William F Flynn, Elise T Courtois 2025. KAPP-Sen TMC: Xenium - Cell DIVE - H&E. [protocols.io https://dx.doi.org/10.17504/protocols.io.n92ld6zj7g5b/v1](https://dx.doi.org/10.17504/protocols.io.n92ld6zj7g5b/v1)

License: This is an open access protocol distributed under the terms of the [Creative Commons Attribution License](#), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited



Protocol status: Working

We use this protocol and it's working

Created: July 31, 2025

Last Modified: August 07, 2025

Protocol Integer ID: 223835

Keywords: SenNet, HuBMAP, 10x genomics xenium platform, resolution spatial gene expression profiling, detailed molecular characterization of tissue architecture, spatial gene expression profiling, situ spatial transcriptomic analysis, spatial transcriptomic analysis, tissue architecture, tissue section preparation, tissue section, tissue processing, xenium, cell dive, enabling detailed molecular characterization

Disclaimer

DISCLAIMER – FOR INFORMATIONAL PURPOSES ONLY; USE AT YOUR OWN RISK

The protocol content here is for informational purposes only and does not constitute legal, medical, clinical, or safety advice, or otherwise; content added to protocols.io is not peer reviewed and may not have undergone a formal approval of any kind. Information presented in this protocol should not substitute for independent professional judgment, advice, diagnosis, or treatment. Any action you take or refrain from taking using or relying upon the information presented here is strictly at your own risk. You agree that neither the Company nor any of the authors, contributors, administrators, or anyone else associated with protocols.io, can be held responsible for your use of the information contained in or linked to this protocol or any of our Sites/Apps and Services.

Abstract

This protocol describes tissue section preparation, in situ spatial transcriptomic analysis, antibody-based imaging and H&E staining of tissue sections using an integrated workflow. Standardized procedures were followed for tissue processing, slide preparation, and staining to ensure consistency and quality. Modifications were applied where appropriate to optimize sample integrity. The 10x Genomics Xenium platform was employed for high-resolution spatial gene expression profiling, enabling detailed molecular characterization of tissue architecture.

Tissue section preparation

- 1 Block QC and sectioning according to **CG000578 Rev E**.
- 2 Post sectioning, all slides are baked at 42°C for 3 hours.

Xenium

- 3 Deparaffinization and decrosslinking is performed according to **CG000580 Rev C or E**. Temperature and duration were as default in the protocol.
- 4 Probe Hybridization, Ligation & Amplification **CG000582 Rev F or G** were followed.
- 5 Xenium Analyzer is used according to **CG000584 Rev E, F or G** were followed.

Cell DIVE

- 6 **Materials:**
 - 1X PBS, pH 7.4 (Gibco; 10010023, ThermoFisher Scientific)
 - Glycerol (G5516, Sigma Aldrich)
 - DI water or Milli-Q water
 - Antigen retrieval buffer (pH9, Abcam ab93684)
 - SuperBlock Blocking Buffer (37580; ThermoFisher Scientific)
 - Antibody diluent (ab64211; Abcam)
 - Hydrogen peroxide, 30% (H1009-500ML; Sigma)
 - 0.5M Sodium bicarbonate (NaHCO₃) (pH= 10.9-11.3)
 - Antigen retrieval chamber (BioSB TintoRetriever)
- 7 **Tissue preparation:**
 - Retrieve the Xenium slides containing the FFPE tissue
 - Wash the slides three times in 1X PBS for 5 min each
 - Place the slides in a coplin jar filled with Tris-EDTA Antigen Retrieval Solution (pH 9.0) and loosely cover the jar with aluminum foil to prevent evaporation of the solution
 - Place the jar in the BioSB TintoRetriever (or any other pressure cooker) containing enough water to cover the foot of the coplin jar
 - Close the TintoRetriever and incubate at 95C (low pressure) for 20 min

- After the TintoRetriever beeps to signal completion of 20 min, switch it off and let it de-pressurize
- After depressurization, retrieve the coplin jar and leave it open on the bench till it cools to room temperature
- Wash the slides in 1X PBS for 5 min
- Permeabilize the tissue by incubating in PBS + 0.3% TX-100 for 10 min
- Wash the slides in 1X PBS for 5 min
- Incubate the slides with SuperBlock solution (ThermoFisher) and incubate for 1 hr at RT
- Wash the slides in 1X PBS for 5 min
- Label the nuclei by incubating with DAPI solution (1mg/ml) for 10 min at RT.
- Wash the slides in 1X PBS for 5 min
- Install the slides into the ClickWell
- Mount slides by adding 2 ml of mounting medium (50% glycerol in PBS) into the ClickWell chamber

8 **Acquisition of mIF image using Cell DIVE imager:**

The Mx Workflow and Cell DIVE acquisition software will be used to set up the imaging protocol and acquire images. Create the protocol appropriate for the study and create a batch by adding slides to be imaged.

9 **ROI selection and autofluorescence recording:**

Select tissue ROIs on the acquisition software and image the slides without antibodies to record the first background/autofluorescence round

10 **Antibody hybridization and biomarker image acquisition:**

- Wash the slides three times with 2 ml 1X PBS for 5 min each
- Prepare the antibody staining mix by adding the appropriate amount of antibody into antibody diluent to achieve the optimized dilution for each antibody in the cycle. Final volume of the staining mix (antibody + diluent) should be 350 µl for each slide. Centrifuge the mixture at 2,000 g for 60 sec to remove the antibody aggregates.
- Add the staining mix to each slide and incubate in the dark at RT for 1 hr
- After incubation, wash the slides three times with 2 ml 1X PBS for 5 min each
- Add 2 µl DAPI solution and incubate for 2 min
- Wash the slides with 2 ml 1X PBS for 5 min
- Mount slides by adding 2 µl of mounting medium. Slides are ready for imaging.
- Advance the workflow on the Mx Workflow software and load the ClickWell into the Cell Dive image to begin biomarker imaging.

11 **Dye inactivation:**

- After completing biomarker imaging, retrieve the ClickWells containing the slides
- Wash the slides three times with 2 ml 1X PBS for 5 min each. Hold the slides in the last PBS wash while preparing the dye inactivation solution
- Prepare the dye inactivation solution with the formula: 10 ml dye inactivation solution = 7 ml MilliQ water + 2 ml 0.5M sodium bicarbonate solution + 1 ml 30% hydrogen



peroxide. Prepare the solution fresh prior to each use and add the hydrogen peroxide immediately prior to adding the solution to tissue

- Remove the PBS and add 2 ml of dye inactivation solution. Incubate for 15 min at RT
- Wash the slides in 2 ml 1X PBS for 5 min
- Add 2 ml DAPI solution and incubate for 2 min
- Wash the slides in 2 ml 1X PBS for 5 min
- Mount slides by adding 2 ml of mounting medium. Slides are ready for imaging.
- After imaging, return to Step x for the next cycle of antibody hybridization
- Repeat this process till the final cycle of biomarker imaging is complete.

H&E

12 All slides are H&E stained according to **CG000613 Rev B**.