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Sequential immunofluorescence and immunohistochemistry on cryosectioned zebrafish embryos --Manuscript Draft--

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Dear Editors:

We are submitting our manuscript entitled “**Sequential immunofluorescence and immunohistochemistry on cryosectioned zebrafish embryos**” for consideration as an original methods article in *JoVE*. Zebrafish are a widely used animal model in biomedical research, but the availability of standardized protocols and reagents for scientific experimentation is limited in comparison to mammalian models. In particular, the relatively small number of commercially available antibodies that are validated for use in zebrafish is a hindrance for antibody-based detection methods. Here we describe a protocol that we developed for the sequential application of immunofluorescence (IF) and immunohistochemical (IHC) assays on a single slide containing cryosectioned zebrafish embryos. This method is designed for precise co-localization of two proteins of interest at the single-cell level in the same tissue section. Our method provides a significant advantage over sequential protocols that are limited to only one methodology (dual IF or dual IHC). This is especially important for studies with zebrafish, as antibodies that have been validated for use in this species are often compatible with either IF or IHC assays, but not both. While designed for use in cryosectioned zebrafish embryos, this protocol could easily be adapted for use with other model organisms.

This manuscript has not been published nor is under consideration elsewhere, including the internet.

Sincerely,

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TITLE:

Sequential Immunofluorescence and Immunohistochemistry on Cryosectioned Zebrafish Embryos

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KEYWORDS:

Zebrafish, embryos, immunohistochemistry, immunofluorescence, colocalization, intercellular interaction

SUMMARY:

This protocol demonstrates sequential immunofluorescence and immunohistochemistry on cryosections from early-stage zebrafish embryos enabling precise colocalization analyses in specific cell populations.

ABSTRACT:

Investigation of intercellular interactions often requires discrete labeling of specific cell populations and precise protein localization. The zebrafish embryo is an excellent tool for examining such interactions with an in vivo model. Whole-mount immunohistochemical and immunofluorescence assays are frequently applied in zebrafish embryos to assess protein expression. However, it can be difficult to achieve accurate mapping of co-localized proteins in three-dimensional space. In addition, some studies may require the use of two antibodies that are not compatible with the same technique (e.g., antibody 1 is only suitable for immunohistochemistry and antibody 2 is only suitable for immunofluorescence). The purpose of the method described herein is to perform sequential immunofluorescence and/or immunohistochemistry on individual cryosections derived from early-stage zebrafish embryos. Here we describe the use of sequential rounds of immunofluorescence, imaging, immunohistochemistry, imaging for a single cryosection in order to achieve precise identification of protein expression at the single-cell level. This methodology is suitable for any study in early-stage zebrafish embryos that requires accurate identification of multiple protein targets in individual cells.

INTRODUCTION:

The zebrafish is an extremely robust model organism that is currently used across a wide variety of disciplines in biomedical research. In particular, the rapid external development and translucency of zebrafish embryos provide an excellent tool for in vivo studies. Herein, we describe a method for sequential immunofluorescence (IF) and immunohistochemical (IHC) analyses of cryosectioned zebrafish embryos. This novel procedure uses sequential application of two antibodies on a single slide, enabling accurate identification of colocalized proteins at the cellular level while conserving tissue sections. This protocol is particularly useful for studies with the zebrafish model, since a relatively small number of antibodies have been validated for use in IF and/or IHC applications in zebrafish compared to mouse models.

The observation of intercellular interactions is an essential element in many studies, and may provide key insights into molecular mechanisms functioning at the cellular level that underlie phenotypes at the organismal level. Additionally, protein expression can provide information about cellular function, especially when examining expression of multiple proteins simultaneously within the cell (co-localization). Although whole-mount IHC and IF are commonly used techniques for analyzing protein expression in zebrafish embryos¹⁻⁵, whole mount procedures can be problematic for achieving precise colocalization data. In our experience, it can be difficult to differentiate between layers of tissue and visualize protein expression at the single-cell level in whole-mount specimens. Imaging software programs may generally be unable to distinguish between surface staining versus deeper staining. Non-surface level protein expressions can be obscured by more brightly expressed surface level expression, leading to inaccuracies in quantification. Additionally, most traditional zebrafish clearing methods are quite toxic⁶ and thus less desirable for use.

Antibody-based techniques, such as IF and IHC, are often used to detect protein expression in sectioned material, simplifying the identification of discrete cell populations that express a particular protein within complex tissues. IHC is commonly used for colocalization, most frequently by using two different antibodies conjugated in different host species and visualized with different colored chromogens^{4,7-10}. However, using multiple chromogens can lead to nonspecific background staining or incompatibility of colors^{11,12}.

We developed a novel protocol for detection of multiple proteins by sequential IF and IHC on cryosectioned early-stage zebrafish embryos. Cryosectioning is particularly well-suited to delicate tissues such as zebrafish embryos, and cryosections are superior to paraffin-embedded sections for fluorescence-based assays^{13,14}. We chose to optimize combined IF and IHC rather than dual-color IF or IHC, to circumvent the problem of antibody incompatibility for a single assay type. These problems are particularly relevant for research involving zebrafish due to the limited number of commercially available antibodies that are validated for use in zebrafish. In fact, a study of four large companies showed that commercially available antibodies for use in mouse was about 112,000 versus about 5,300 for use in zebrafish¹⁵. Finally, we chose to develop a protocol that could be performed on a single cryosection, which is essential when working with small or limited tissue samples such as are obtained from zebrafish embryos.

This protocol was designed to assess the proliferative behavior of donor cells in 48 h post-fertilization chimeric zebrafish embryos that were generated by blastula-to-blastula transplant as described by Carmany-Rampey and Moens¹⁶. Donor embryos were injected at the one-cell stage with a fluorescently labeled dextran conjugate prior to transplant of donor cells into recipient embryos. We used immunofluorescence for Ser 10 phosphorylated Histone H3 (pH3) to detect proliferating cells followed by immunohistochemistry for the labeled dextran to detect donor cells in chimeric zebrafish embryos. Sequential detection of pH3 and the labeled dextran within a single cryosection enabled us to identify and quantify individual cells that expressed both markers.

This sequential IF/IHC protocol for cryosectioned zebrafish will provide a useful tool for zebrafish researchers who desire a colocalization protocol for protein expression. The problems that this protocol was designed to address, such as small tissue specimens and limited antibody availability, are not unique to the zebrafish model. This method may therefore be of use to any researcher desiring to perform sequential IF/IHC.

ETHICS STATEMENT:

All animal studies were approved by the Institutional Animal Care and Use Committee, North Carolina State University, Raleigh, North Carolina, USA.

PROTOCOL:

1. Embryo preparation

1.1. Fix 48 h post-fertilization (hpf) chimeric zebrafish embryos generated by blastula-to-blastula transplant between AB wild-type zebrafish embryos in 4% paraformaldehyde (PFA) overnight with rocking at 4 °C. Perform two 5 min washes with rocking at room temperature using 500 µL of 1x phosphate buffered saline containing 0.1% of a non-ionic surfactant (1x PBSt; see the **Table of Materials**)

NOTE: Paraformaldehyde is toxic and a carcinogen and must be properly disposed of per institutional regulations. Work with paraformaldehyde should be performed in a chemical hood, and appropriate personal protective equipment (gloves, lab coat, and safety glasses) should always be used.

1.2. Dehydrate embryos by washing sequentially with 500 µL of 30%, 50%, and 70% methanol (MeOH) diluted with 1x PBSt for 10 min each at room temperature with rocking. Incubate embryos in 500 µL of 100% MeOH at -20 °C for at least 14 h.

NOTE: Methanol is toxic and must be properly disposed of per institutional regulations. Work with methanol should be performed in a chemical hood, and appropriate personal protective equipment (gloves, lab coat, and safety glasses) should always be used.

130 1.3. Rehydrate embryos by washing sequentially with 500 μ L of 70%, 50%, and 30% MeOH diluted
131 with 1x PBSt for 10 min each at room temperature with rocking. Perform two 5 min washes with
132 500 μ L of 1x PBSt at room temperature with rocking.

133
134 1.4. Remove the 1x PBSt and incubate in 500 μ L of 30% sucrose diluted with deionized water at
135 room temperature with rocking until the embryos sink to the bottom of the tube (approximately
136 1 h). Replace sucrose with 500 μ L of 15% fish gelatin/25% sucrose (15/25; see the **Table of**
137 **Materials**) and incubate at room temperature with rocking overnight.

138
139 1.5. Replace approximately half of the volume of 15/25 with optimal cutting temperature
140 medium (OCT medium) and incubate at room temperature with rocking until the embryos sink
141 to the bottom of the tube (approximately 1 h).

142
143 1.6. Replace approximately half of the volume with OCT medium and incubate at room
144 temperature with rocking for 1 h. Repeat this step one time.

145
146 1.7. During incubation steps with 15/25 and OCT medium, invert or flick the tube as needed to
147 combine these reagents.

148 149 **2. Embryo embedding and preparation of cryosections**

150
151 2.1. Transfer embryos to a plastic mold (see the **Table of Materials**) with forceps, minimizing any
152 transfer of the 15/25-OCT mixture. Fill the mold approximately half full with OCT medium and
153 gently mix the embryos in OCT medium.

154
155 2.2. Prepare labeled plastic molds and transfer desired embryos (usually 1–3 embryos) into the
156 empty, labeled plastic molds, minimizing carry-over of OCT medium. Once the desired embryos
157 are in the plastic mold, gently fill with OCT medium to the top of the mold.

158
159 2.3. Use forceps or 25 G needles to arrange embryos to the desired orientation, using a light
160 stereomicroscope for visualization (**Figure 1A,B**).

161
162 2.4. Freeze the prepared molds on dry ice in an insulated container with a metal platform. Place
163 an ice bucket or foam cooler gently over the metal platform to create a cold chamber (**Figure**
164 **1C,D**)

165
166 2.5. Prepare 10–12 μ m thick cryosections using a cryostat set at -20 $^{\circ}$ C.

167
168 2.5.1. Set up one block at a time and use OCT medium to freeze the block onto the disc/chuck
169 with the bottom of the block facing toward the blade (**Figure 2A**). Ensure that the block is
170 completely frozen to the disc (the OCT medium will turn white) before beginning sectioning
171 (**Figure 2B**).

2.5.2. Place cryosections on charged glass slides (**Figure 2C,D**) and air-dry the slides overnight at room temperature.

3. Immunofluorescence for pH3

3.1. Perform three 5 min washes of the slides in 1x PBS in an appropriate container, such as a Coplin jar. If tissues are not well adhered to the slide, perform washes by laying slides on a flat surface and gently pipetting 500 μ L of 1x PBS onto the surface. Pour off 1x PBS between washes by gently tipping the slides.

3.2. Outline the sections with a barrier pen or wax pencil to keep liquid on the slides (**Figure 3**).

3.3. Lay the slides on a flat surface in a humid chamber and pipette 200 μ L of block buffer per section onto the slides. Incubate in block buffer for 2 h at room temperature or overnight at 4 °C.

3.4. Prepare the primary antibody dilution (rabbit anti-pH3 antibody, 1:200; see **Table of Materials**) in block buffer and mix well by pipetting. Gently tip the slides to drain off the block buffer, return to the humid chamber, and pipette 200 μ L of primary antibody solution per section onto the slides. For the secondary-only control, pipette 200 μ L of block buffer onto the appropriate sections.

3.5. Incubate the slides at 4 °C overnight in a humid chamber filled with deionized water. Seal the edges of the humid chamber to help retain moisture.

3.6. Perform three 5 min washes of the slides in 1x PBS in an appropriate container, such as a Coplin jar. During wash steps, prepare the secondary antibody dilution (anti-rabbit fluorescent secondary antibody conjugate, 1:2,000; see **Table of Materials**) in block buffer and mix well by pipetting.

3.7. Lay the slides on a flat surface in a humid chamber and pipette 200 μ L of secondary antibody solution per section onto the slides. Incubate the slides in secondary antibody solution(s) at room temperature, shielded from light, for 30 min.

3.8. Perform three 5 min washes of the slides in 1X PBS in an appropriate container, such as a Coplin jar, shielded from light. During wash steps, prepare a nuclear staining solution (see the **Table of Materials**).

3.9. Place the slides on a flat surface and pipette 200–500 μ L (depending on the size of sample) of the nuclear staining solution onto each section. Incubate the slides in the nuclear staining solution for 10 min, shielded from light.

3.10. Drain off the nuclear staining solution and mount the slides with non-hardening fluorescent mounting media and a glass coverslip. Maintain the slides in the dark at 4 °C until imaging is performed.

NOTE: The best results are obtained when the slides are imaged on the same day or the next day.

3.11. Following IF, visualize and image the slides with a confocal fluorescence microscope and digital camera using 555 nm (red) and 645 nm (far-red) emission filters at 100x magnification (10x ocular magnification and 10x objective magnification). Keep the laser power consistent throughout imaging.

3.12. After imaging, place the slides in individual containers of 1x PBS and store flat at 4 °C overnight to prepare for coverslip removal. When placing the slides in 1x PBS, gently agitate the slides to assist in removal of the coverslip.

NOTE: Do not press down on the coverslip or attempt to remove the coverslip manually, as this can compromise quality of the sections. The coverslips must be removed horizontally with minimal forced movement.

4. Immunohistochemistry for labeled dextran

4.1. After the coverslips have been removed, gently transfer the slides into new 1x PBS in an appropriate container, such as a Coplin jar. Remove the 1x PBS and perform three 5 min incubations of the slides in 1x Tris-buffered saline with 0.1% of a non-ionic surfactant (1x TBSt) at room temperature.

4.2. Prepare 250 mL of 3% hydrogen peroxide (H_2O_2) diluted in deionized water in an appropriately sized container. Incubate the slides in 3% H_2O_2 solution at room temperature for 15 min.

NOTE: Hydrogen peroxide is corrosive and must be properly disposed of per institutional regulations. Appropriate personal protective equipment (gloves, lab coat, and safety glasses) should always be used.

4.3. Quickly rinse the slides with deionized water. Perform three 5 min washes of the slides in 1x TBSt. Carefully dry the area around the tissue sections and place the slides flat in a humid chamber. Outline the tissue sections with a barrier pen or wax pencil to keep liquid on the sections if needed.

4.4. Apply a ready-to-use 2.5% serum blocking solution provided with a secondary antibody labeling kit (see the **Table of Materials**) dropwise to each section. Incubate the slides in a humid chamber at room temperature for 20 min.

4.5. Prepare the primary antibody diluted in antibody diluent (rabbit anti-labeled dextran, 1:7500; see the **Table of Materials**) and mix by gentle pipetting. Gently tip the slides to drain off the block buffer, dry the area around the sections, and place the slides flat in a humid chamber.

4.6. Do not wash before applying the primary antibody solution. Pipette 200 μ L of primary antibody solution per section onto the slides and incubate in a humid chamber at 4 °C overnight. For the secondary-only control, pipette 200 μ L of antibody diluent onto the appropriate sections.

4.7. Gently tip the slides to drain off the primary antibody solution and place in 1x TBSt in an appropriate container, such as a Coplin jar. Perform two 5 min washes of the slides in 1x TBSt. Dry the area around the sections and place flat in a humid chamber.

4.8. Apply a ready-to-use background-reducing blocking reagent (see the **Table of Materials**) dropwise to each section. Incubate the slides in a humid chamber at room temperature for 20 min.

4.9. Gently tip the slides to drain off the block buffer, carefully dry the area around the sections, and place the slides flat in a humid chamber. Do not wash before applying the secondary antibody solution.

4.10. Apply a ready-to-use secondary antibody solution (anti-rabbit horseradish peroxidase (HRP) polymer secondary antibody; see the **Table of Materials**) to each section dropwise and incubate in a humid chamber at room temperature for 30 min.

4.11. Gently tip the slides to drain off the secondary antibody solution and place in 1x TBSt in an appropriate container, such as a Coplin jar. Perform two 5 min washes of the slides in 1x TBSt.

4.12. Prepare the HRP chromogenic substrate immediately before use according to the manufacturer's protocol (see the **Table of Materials**).

4.13. Dry the area around the sections, place the slides on a flat surface, and apply 200 μ L of the HRP chromogenic substrate to each section. Incubate at room temperature for 6 min, starting a timer when the substrate has been applied to the first slide (**Figure 4**).

4.14. Drain off the substrate solution and rinse the slides briefly in a Coplin jar containing 1x PBS. Place the slides in deionized water in an appropriate container, such as a Coplin jar, and perform two 5 min washes in deionized water with gentle agitation.

4.15. Counterstain the slides by placing them in hematoxylin staining solution for up to 30 s.

NOTE: Hematoxylin is toxic and must be disposed of per institutional regulations. Appropriate personal protective equipment (gloves, lab coat, and safety glasses) should always be used.

4.16. Place the slides in deionized water in an appropriate container, such as a Coplin jar, and perform three 5 min washes in deionized water. Transfer the slides to a container containing Scott's tap water (see the **Table of Materials**) and incubate for 1 min.

4.17. Place the slides in deionized water in an appropriate container, such as a Coplin jar, and perform three 5 min washes in deionized water.

4.18. Dehydrate the slides through a series of ethanol (EtOH) grades (diluted in deionized water) and xylene in a chemical hood. Perform one 2 min incubation in 70% EtOH, two 1 min incubations in 95% EtOH, two 1 min incubations in 100% EtOH, and three 1 min incubations in xylene. Remove the slides from xylene and place a coverslip while wet with a toluene-based mounting medium in a chemical hood.

NOTE: Xylene and toluene are toxic and flammable and must be properly disposed of per institutional regulations. Work with xylene and toluene should be performed in a chemical hood, and appropriate personal protective equipment (gloves, lab coat, and safety glasses) should always be used.

4.19. Following IHC, visualize and image the slides with a compound light microscope and digital camera at 100x magnification (10x ocular magnification and 10x objective magnification). (Figure 5).

REPRESENTATIVE RESULTS:

We developed this protocol in order to identify and analyze protein expression in individual donor cells in 48 h post-fertilization chimeric zebrafish embryos that were generated by blastula-to-blastula transplant between AB wild-type zebrafish embryos. Successful analysis of protein expression at the one-cell level required preparation of cryosections with appropriately oriented embryos (Figure 1 and Figure 2) and careful, sequential application of IF and IHC techniques to these cryosections (Figure 3 and Figure 4). Use of specific antibodies to simultaneously detect proliferating cells (anti-pH3, Figure 5A,B) and donor cells (anti-labeled dextran, Figure 5D) can be used to identify and quantify donor cells that are actively proliferating (Figure 5E). It is essential to generate high-quality images after both IF (Figure 2A,B) and IHC (Figure 2D) in order to accurately identify individual cells. An image analysis program should be used to perform image overlay (Figure 5E). Depending on the image analysis program used, it may be possible to perform automated counts of labeled cells should quantification be desired.

FIGURE AND TABLE LEGENDS:

Figure 1. Preparation of frozen OCT blocks. (A) 50x microscopic view of embryos in OCT in a cryogenic mold prepared for freezing. Red arrow indicates head of zebrafish embryo position against the bottom surface of the mold; black arrow indicates tail pointing up towards the user. (B) Plastic molds with embryos and OCT placed on chilled metal platform. (C) Foam ice bucket placed over plastic molds to create freezing chamber. (D) OCT turns white from transparent once the block is fully frozen.

Figure 2. Cryosectioning of frozen OCT blocks. (A), Application of fresh OCT onto cryostat disc and placement of frozen block on OCT, rotated 180° from the freezing orientation of the block.

(B) Side view of block frozen to sectioning disc. (C) View of sectioning block with roll plate in position. (D) Pick-up of sections using a charged slide from the metal platform of the cryostat.

Figure 3. View of prepared sections after placement on a slide. Arrow indicates hydrophobic barrier, and dotted line delineates the zone containing sections within barrier perimeter.

Figure 4. Preparation for chromogenic substrate application during IHC. The slide is placed on a flat surface covered with plastic wrap to allow uniform application of the chromogenic substrate while containing any spillage of hazardous material.

Figure 5. Representative IF and IHC labeling of cryosectioned 48 hpf chimeric zebrafish embryos generated by blastula-to-blastula transplant between AB wild-type zebrafish embryos. (A) IF to detect Ser 10 phosphorylated Histone H3 (anti-pH3, 1:200) expression in 48 hpf chimeric zebrafish embryo. Red = pH3-positive cells; blue = nuclei. Yellow arrows indicate examples of positive cells. (B) Subtraction of the blue nuclear stain in the digital image (ImageJ software) enhances visualization of pH3-positive cells for quantification and image overlay. (C) Negative control (secondary antibody only) for IF assay in cryosectioned 48 hpf chimeric zebrafish embryo. Scale bar represents 50 μ m (applicable to all panels). (D) IHC to detect labeled dextran (anti-labeled dextran, 1:7500) in chimeric 48 hpf zebrafish embryo generated by blastula-to-blastula transplant between AB wild-type zebrafish embryos. Donor embryos were injected with a fluorescently labeled dextran conjugate at the one-cell stage prior to use for blastula-to-blastula transplant. Red indicates cells labeled with a dextran conjugate; blue indicates nuclei. Yellow arrows indicate examples of positive cells. (E) Overlay of panel B (IF for pH3) and panel C (IHC for labeled dextran) showing colocalization of pH3 expression and dextran labeling in individual cells. Yellow arrows indicate examples of double-positive cells. (F) Negative control (secondary antibody only) for IHC assay in cryosectioned 48 hpf chimeric zebrafish embryo.

DISCUSSION:

We have presented a novel method for combined immunofluorescence and immunohistochemistry that represents an important step forward in performing colocalization experiments on cryosectioned zebrafish embryos. There is a critical lack of existing colocalization protocols that are optimized for use with small embryos and cryosectioned material^{17,18}, both of which are otherwise commonly used in molecular studies¹⁴. Existing colocalization protocols mainly focus on simultaneous visualization of two fluorophores^{17,18}. While these protocols can work well, their usefulness is limited by the availability of antibodies that (i) can be used in zebrafish and (ii) are compatible with IF.

We feel that use of cryosectioning for zebrafish embryos provides a significant advantage in terms of preserved tissue morphology. As IHC is more commonly performed on paraffin sections than cryosections^{7,8}, further exploration of cryosection-based IHC methods for detection of protein expression is warranted. The use of whole mount IF is a common method for visualizing expression levels in zebrafish embryos, and is more commonly described in the literature than IF using cryosections^{1-4,19}. However, whole mount protocols have limitations, including the lack of accurate localization of proteins expressed in deep tissues and the potential for surface level

expression to obscure protein expression in deeper tissues. We have consistently observed excellent maintenance of cellular morphology following cryopreservation, sectioning, and sequential IF and IHC. For applications that require precise localization of protein expression at the single-cell level, there is a significant advantage to section-based procedures as we have described in this protocol. Resin embedding provides an alternative option for performing IF or IHC assays on sections from early-stage zebrafish embryos²⁰. Resin sections provide superior preservation of tissue morphology compared to cryosections, and relatively thinner tissue sections can be prepared. However, manufacturers generally do not recommend the use of resin sections for immuno-based assays, as some components of the resins cannot be removed from the sections and may mask antibody binding sites²¹.

While the whole of the protocol is essential for accurate and successful analysis of expression patterns, a few specific steps are critical for experimental success. The first critical step is the handling of embryos during embedding in OCT^{13,14}. It can be challenging to orient each embryo in the same transverse plane so that sections are cut from approximately the same area for all embryos. We have found that using small gauge needles to perform small, deliberate movements through the viscous OCT medium for embryo orientation is superior to using forceps, which are too large and displace too much OCT medium. A second critical step occurs during sectioning of frozen blocks, as it can be difficult to obtain high-quality sections that contain the desired tissue(s) without significant training and experience. We thus recommend use of test samples until the technique is mastered. A third critical step is coverslip removal, which has the potential to disturb or strip off tissue samples. We have found that a combination of gentle agitation to loosen the coverslip and slow removal of the slide from the PBS container is the most effective approach for removing coverslips. A gentle and steady hand is best; researchers may find that some trial and error is necessary to learn how to remove coverslips effectively.

Additional important steps in the protocol include antibody optimization (primary and secondary antibodies for both IF and IHC assays) and exposure of slides to the chromogenic substrate and counterstain. Thorough research regarding primary antibody specificity and target concentration is essential; generally, it is best to collect sufficient non-precious samples for at least two separate experiments for IF and IHC that will be performed to optimize primary antibody concentrations before beginning the IF/IHC combination. Including a known positive and negative control for each primary antibody is essential in each experiment to ensure that both IF and IHC assays are performing appropriately. Adjustments to secondary antibody concentration may also be necessary. Optimization of exposure of tissue sections to the chromogenic substrate and counterstain for the IHC assay is required, since manufacturer guidelines often describe a wide range of possible exposure times. Not all chromogens may be compatible with counterstains and endogenous tissue pigmentation, requiring careful planning with regard to chromogen selection.

There are numerous potential modifications that can be applied to the described protocol, and we have successfully performed this protocol with other combinations of antibodies that could not both be used for IF. As alluded to above, chromogenic substances have distinct compatibility with specific counterstains. We have found that these components are easily modifiable in the protocol. Additionally, the parameters for antibody incubation are quite flexible, and can be

increased or decreased depending on desired staining intensity. The embedding process can also be modified. While we have focused on transverse sections, embryos can easily be oriented in any direction to address particular tissues of interest. Finally, there are multiple possible pause points in this protocol. Dehydrated embryos can be stored in 100% MeOH at -20 °C for a few months; frozen blocks can be stored at -80 °C for up to three months; and prepared slides can be stored at -80 °C for up to nine months in a slide box.

Due to the relative complexity of this combined IF/IHC protocol, there are multiple possible sources of variation or error that may require troubleshooting, ranging from tissue quality to researcher experience to sample handling. Most of the critical steps described above are considered critical because they either require optimization at the individual user level or they require particular dexterity, skill, and experience. However, we have found that nonspecific background staining and low signal intensity are the most significant issues requiring optimization. As with any IF or IHC protocol, addressing these issues can be time consuming and labor intensive, and can be compounded with this method since the two techniques are combined. For this reason, we highly recommend optimizing IF and IHC separately before proceeding with the combined protocol.

While we predict that this method will be useful for a broad range of experiments, there are potential limitations. Successful performance of this procedure is dependent on maintaining intact tissue samples through two sequential experiments that include numerous wash steps. For this reason, any problems that arise during sectioning or tissue handling, including use of older slides that may have degraded in quality, will significantly limit researchers' ability to perform this protocol successfully. Although slides can potentially be stored at -80 °C for up to nine months, tissue adherence to slides declines over time and we had the best success with slides that are used within a month of preparation or ideally, the next day. A second limitation is the small footprint of zebrafish embryos. While we have found this experiment to be useful in visualizing proteins that are relatively abundantly expressed in early-stage embryos, proteins that are expressed inconsistently or at low levels may be very difficult to capture in a section. Finally, since the protocol uses 10–12 µm cryosections, it is best suited for evaluation of protein expression in larger structures, such as nuclei, rather than smaller sub-cellular components.

In summary, our combined IF/IHC protocol will be advantageous for a wide variety of studies in early-stage zebrafish embryos, and represents an important innovation in precise protein expression analysis in such specimens. Our method, shown successfully in **Figure 5**, will allow researchers to preserve the delicate morphology of early-stage zebrafish embryos in cryosections while working with the somewhat limited range of currently available primary antibodies that are validated for use in IF or IHC in this species. This protocol will likely be useful for studies in other fish and amphibian species (e.g., *Medaka* and *Xenopus* spp.), which are hampered by similar limitations to antibody availability, and may be applicable for traditional mammalian models as well.

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DISCLOSURES:

The authors have nothing to disclose.

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Figure 1

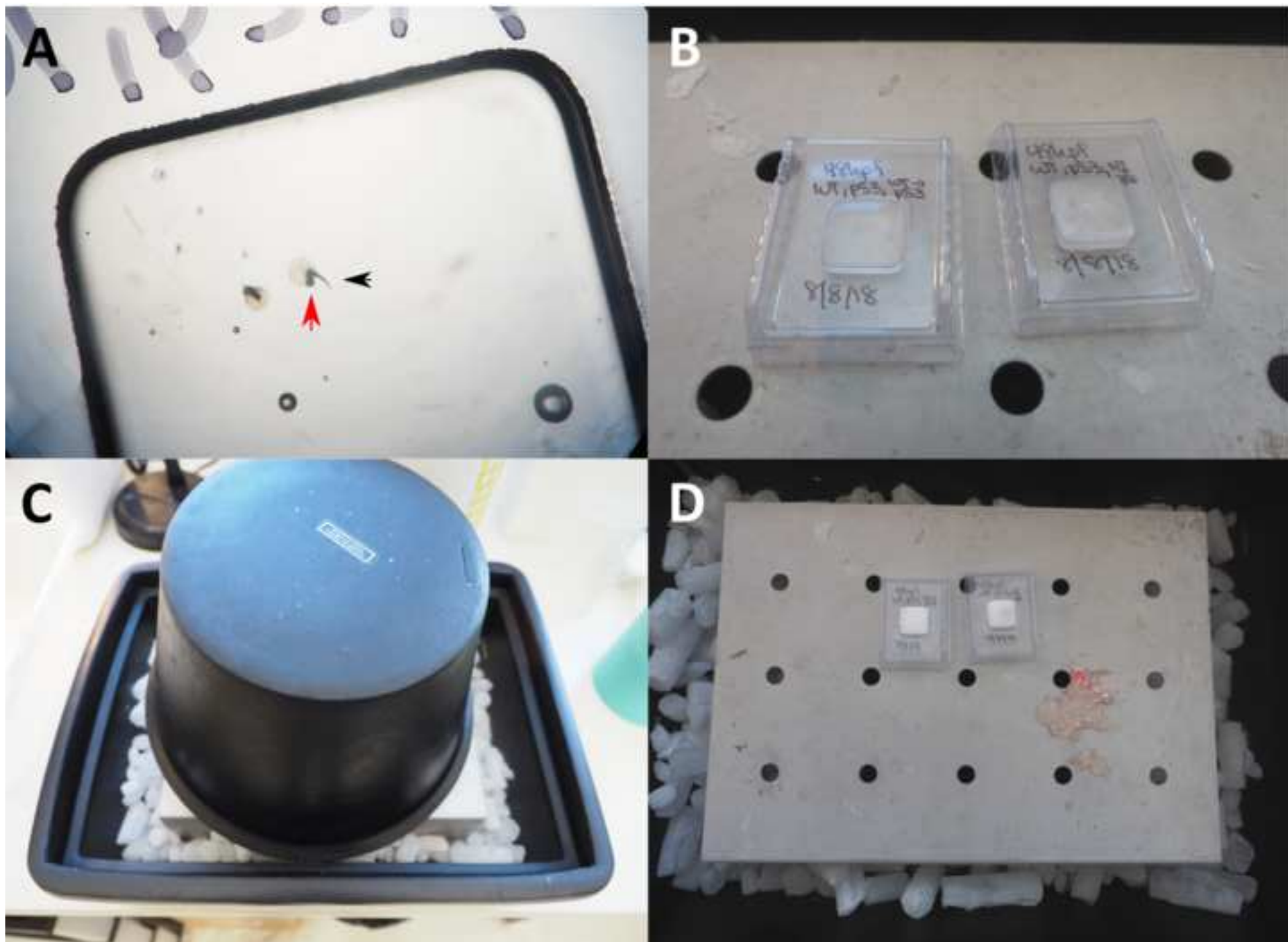


Figure 2

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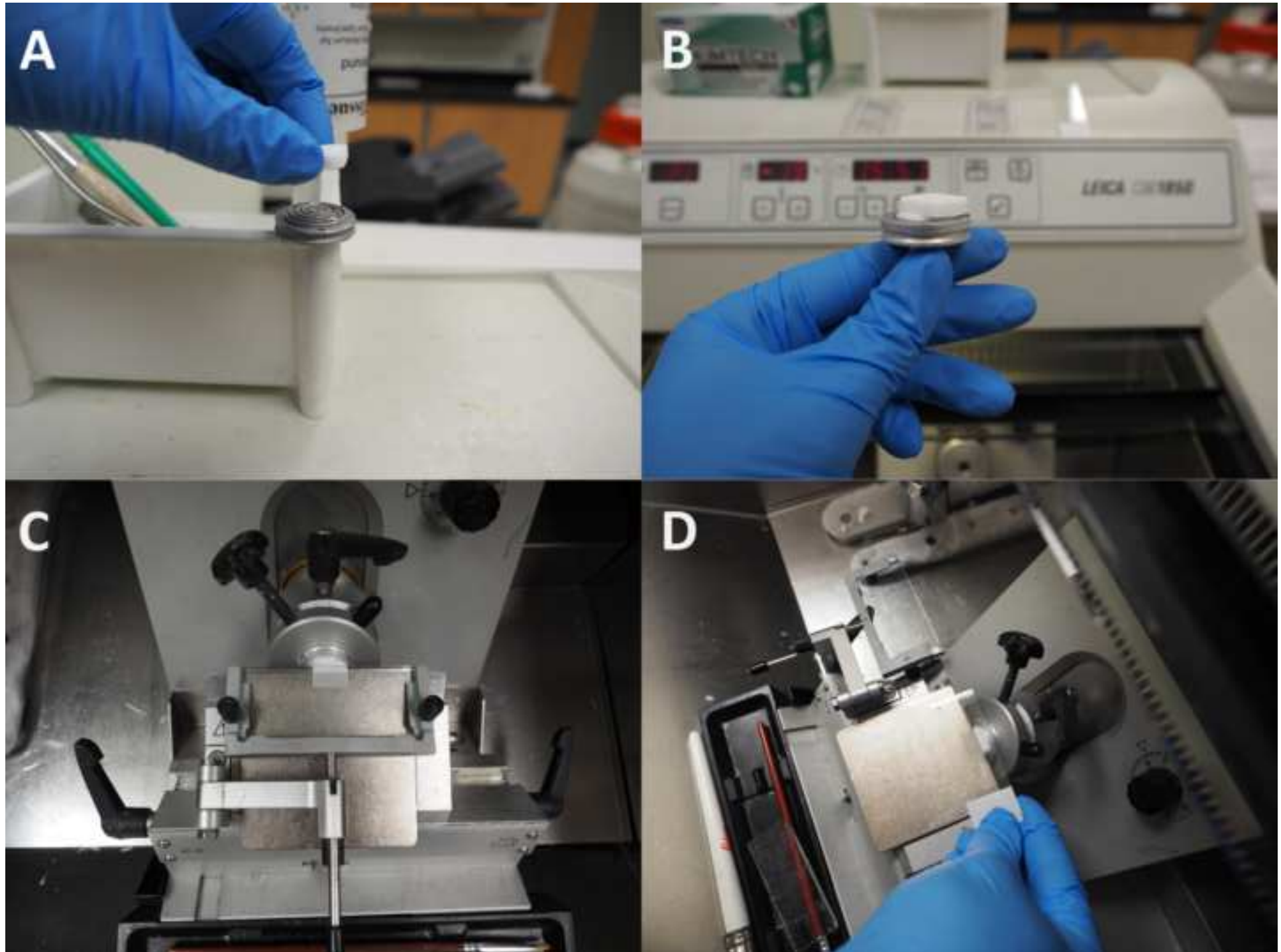


Figure 3

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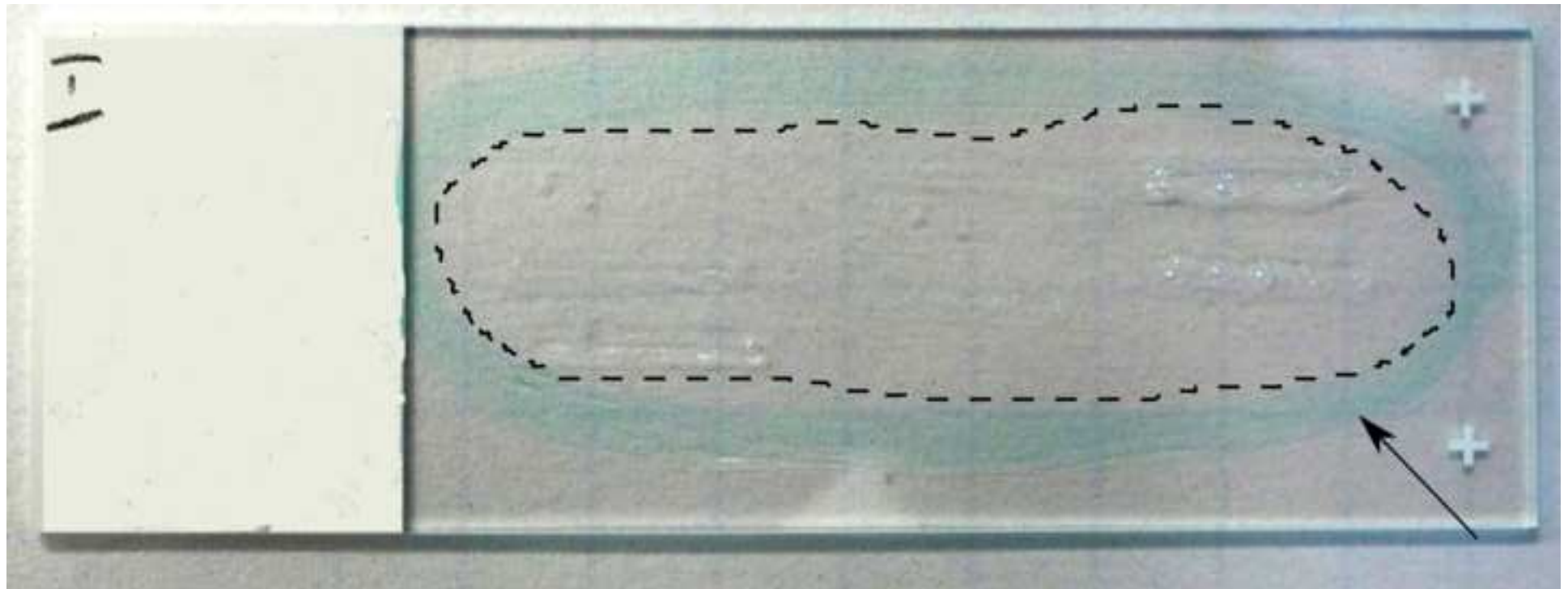
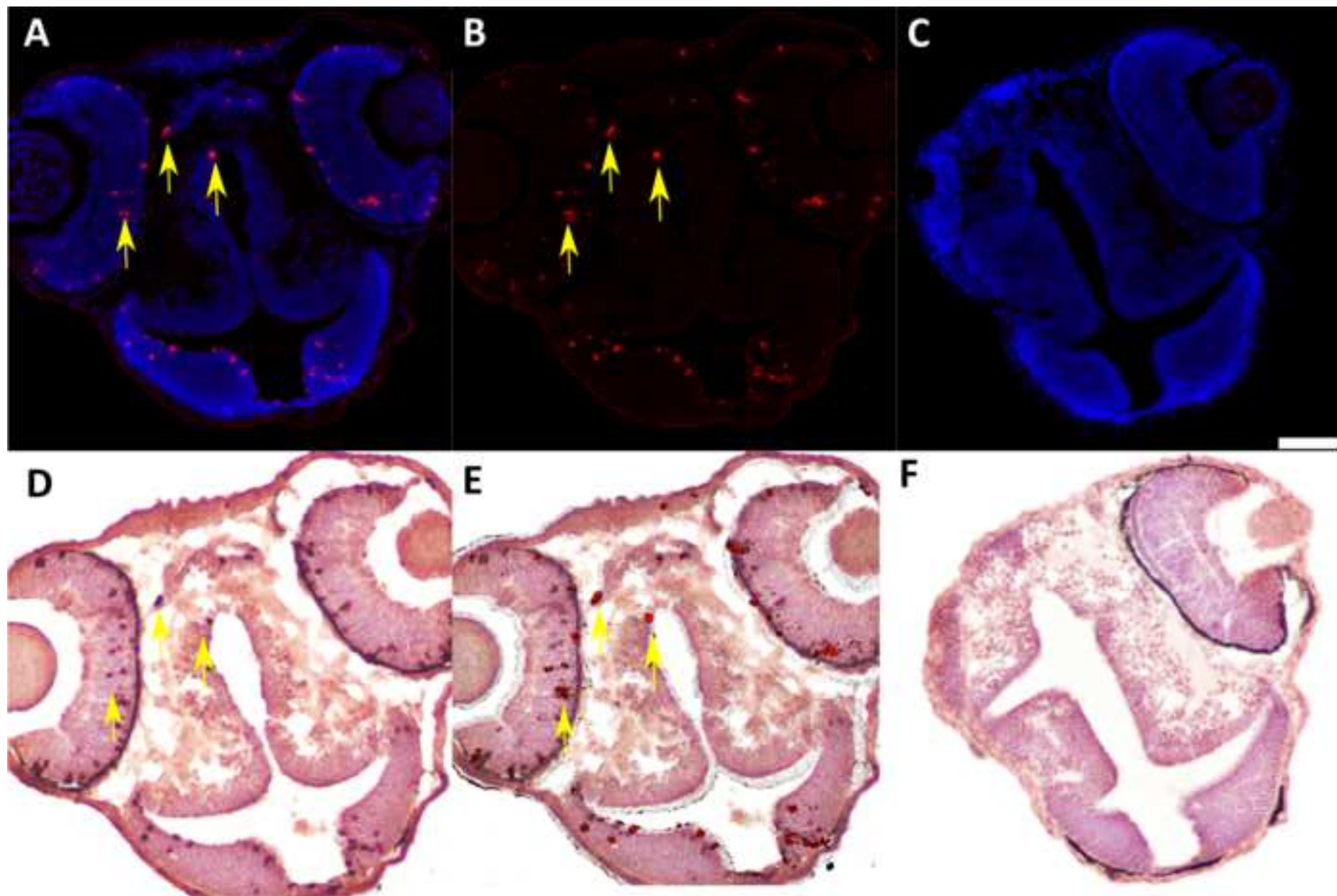


Figure 4

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Figure 5



Name of Material/ Equipment	Company	Catalog Number	Comments/Description
15/25	N/A	N/A	15% fish gelatin/25% sucrose w/v, made in dH ₂ O
Alexa 555 secondary antibody	Invitrogen	A21429	used at 1:2000 dilution in block buffer for IF
Antibody Diluent	Dako	S3022	
Background Buster	Innovex	NB306	
block buffer (IF)	N/A	N/A	
cellSens imaging software	Olympus	cellSens	5% normal goat serum, 0.1% Triton X-100 in 1X F
chimeric zebrafish embryos	N/A	N/A	imaging software used with compound light micr
Compound light microscope	Olympus	BX51	AB wild-type chimera zebrafish embryos analyze
Confocal fluorescence microscope	Leica	DM 2500	used with digital camera and cellSens imaging so
			used with digital camera and Leica Application Si
disposable plastic molds, 15 mm x 15 mm x 5 mm	Ted Pella	27147-2	
Digital camera, light microscope	Olympus	DP27	
Digital camera, fluorescence microscope	Leica	TCS SPE	
Ethanol	Koptec	V1001	
fish gelatin	Sigma	G7041	
Glass slides	Fisher	12-550-15	
Hydrogen peroxide, 30%	Fisher	H325-100	
ImmPRESS HRP Polymer detection kit	Vector	MP-7401	anti-rabbit secondary antibody, HRP polymer
Leica Application Suite Advanced Fluorescence imaging software	Leica	LAS AF 2.3.6	imaging software used with confocal fluorescenc
Methanol	Fisher	A452-4	
Modified Meyer's Hematoxylin	Thermo	72804	
Normal Goat Serum	MP Biomedical	191356	
OCT medium	Tissue-Tek/Fisher	4583/4585	

anti-Oregon Green antibody	Molecular Probes/Life Technologies	A889	used at 1:7500 dilution for IHC
Oregon Green Dextran	Invitrogen	P7171	used at 2.5% in 0.2M KCl
Paraformaldehyde, 4%	Acros Organics	416780030	made in lab in 1x PBS
Permunt	Fisher	SP15	
1X phosphate-buffered saline	made in lab	N/A	50 mL 10x PBS, up to 500 mL volume with dH ₂ O
10X phoshpate-buffered saline	made in lab	N/A	use Cold Spring Harbor protocol
1X PBSt	made in lab	N/A	50 mL 10x PBS, 1 mL 10% Tween 20, up to 500 mL
anti-Ser10 phosphorylated	Santa Cruz	sc-8656-R	used at 1:500 dilution for IF
Histone H3 antibody			
Scott's Tap Water	Electron Microscopy Sciences	26070-06	have used other brands successfully in addition to
sucrose	Amresco		335
Tween-20	Fisher	BP337	
TO-PRO3	Invitrogen	T3605	used at 1:1000 in 1x PBS for IF
1X Tris-buffered saline	made in lab	N/A	50 mL 10x TBS, up to 500 mL volume with dH ₂ O
10x Tris-buffered saline	made in lab	N/A	85 g NaCl, 61 g Tris base, up to 1 L volume with dH ₂ O
1X TBSt	made in lab	N/A	50 mL 10x TBS, 1 mL 10% Tween 20, up to 500 mL
Triton X-100	Acros Organics	327371000	
Vectashield	Vector	H-1000	non-hardening mounting media
Vector NovaRed substrate	Vector	SK-4800	HRP substrate
Xylene	Fisher	X3P-1GAL	

)

BS

roscope

d at 48 hpf

oftware for image capture after IHC

uite Advanced Fluorescence imaging software for image capture after IF

ce microscope

1L volume with dH₂O

to EMS

dH₂O

1L volume with dH₂O



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Author(s):

Jordan L. Ferguson, Heather R. Shive

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Institution:	North Carolina State University	
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RE: Response to review of manuscript ID JoVE59344

We appreciate that the editor and reviewers found our manuscript to be of interest. We have revised our manuscript according to the comments and recommendations provided by the editor and reviewers, and feel that these changes have improved the quality and clarity of this work.

Editorial comments:

Changes to be made by the author(s) regarding the manuscript:

1. Please take this opportunity to thoroughly proofread the manuscript to ensure that there are no spelling or grammar issues.

We have proofread the document as requested.

2. Please define all abbreviations before use.

We have defined abbreviations as requested.

3. Please use SI abbreviations for all units: L, mL, μ L, h, min, s, etc.

We have made this change.

4. Please include a space between all numerical values and their corresponding units: 15 mL, 37 °C, 60 s; etc.

We have made this change.

5. Please adjust the numbering of the Protocol to follow the JoVE Instructions for Authors. For example, 1 should be followed by 1.1 and then 1.1.1 and 1.1.2 if necessary. Please refrain from using bullets, dashes, or indentations.

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We have made this change.

7. Please revise the protocol to contain only action items that direct the reader to do something (e.g., “Do this,” “Ensure that,” etc.). The actions should be described in the imperative tense in complete sentences wherever possible. Avoid usage of phrases such as “could be,” “should be,” and “would be” throughout the Protocol. Any text that cannot be written in the imperative tense may be added as a “Note.” Please include all safety procedures and use of hoods, etc. However, notes should be used sparingly and actions should be described in the imperative tense wherever possible. Please move the discussion about the protocol to the Discussion.

We have made these changes.

8. Please add more details to your protocol steps. There should be enough detail in each step to supplement the actions seen in the video so that viewers can easily replicate the protocol. Please ensure you answer the “how” question, i.e., how is the step performed? Alternatively, add references to published material specifying how to perform the protocol action. See examples below.

We have made these changes.

9. Line 101: Please provide the composition of PBSt.

We have made this change.

10. Lines 170, 179: Please specify the primary/secondary antibody used. What is the dilution factor?

We have made this change.

11. Lines 180, 186: Please specify incubation time.

We have made this change.

12. Line 202: Please specify incubation temperature.

We have made this change.

13. Line 203: Please list an approximate volume to prepare.

We have made this change.

14. Please combine some of the shorter Protocol steps so that individual steps contain 2-3 actions and maximum of 4 sentences per step.

We have made this change.

15. Please include single-line spaces between all paragraphs, headings, steps, etc.

We have made this change.

16. After you have made all the recommended changes to your protocol (listed above), please highlight 2.75 pages or less of the Protocol (including headings and spacing) that identifies the essential steps of the protocol for the video, i.e., the steps that should be visualized to tell the most cohesive story of the Protocol.

We have made this change.

17. Please highlight complete sentences (not parts of sentences). Please ensure that the highlighted part of the step includes at least one action that is written in imperative tense.

We have made this change.

18. Please include all relevant details that are required to perform the step in the highlighting. For example: If step 2.5 is highlighted for filming and the details of how to perform the step are given in steps 2.5.1 and 2.5.2, then the sub-steps where the details are provided must be highlighted.

We have made this change.

19. Figure 5: The scale bars in panels A and B are difficult to read. Please revise. The unit should be μm instead of μM .

We have made this change.

20. Please include at least one paragraph of text to explain the Representative Results in the context of the technique you have described, e.g., how do these results show the technique, suggestions about how to analyze the outcome, etc. The paragraph text should refer to all of the figures. However for figures showing the experimental set-up, please reference them in the Protocol. Data from both successful and sub-optimal experiments can be included.

We have made this change.

21. References: Please do not abbreviate journal titles.

We have made this change.

22. Table of Materials: Please sort the items in alphabetical order according to the name of material/equipment.

We have made this change.

Reviewers' comments:

Reviewer #1:

Manuscript Summary:

The manuscript is clearly and concisely written, but with sufficient detail to allow replication of this rather complex protocol. The use of fish gelatin in embedding embryos for frozen sections is an advance over the standard protocol provided in the Zebrafish Book. Producing excellent frozen sections of zebrafish embryos is not trivial. Given the wide use of this model organism for many types of biomedical research and the limited number of antibodies available as research tools, the optimization of such procedures is valuable to the community.

Major Concerns:

No major concerns

Minor Concerns:

Negative controls are needed for primary antibody such as nonimmune serum from same species for immunofluorescence as well as immunohistochem. Did do secondary only control.

We appreciate the reviewer's comments and suggestions. We have added images from our negative controls (secondary only) for IF and IHC to Figure 5 as requested.

Reviewer #2:

Manuscript Summary:

The methods paper by Ferguson and Shive describes a method for sequential immunofluorescence and immunohistochemistry on cryosectioned zebrafish embryos. It is

useful technique that will help in specific experimental situations to show co-localization of specific antigens.

Minor Concerns:

The paper is generally clear. However, there are some clarification that might help the reader and additional limitations for the applicability of the method that could be addressed.

1. The paper discusses the use of IHC but it is not clear what immunohistochemistry they are using and why. The paper simply says "Immunohistochemistry was performed for 298 Oregon Green to detect donor cells". While there is nothing in the actual text about what histochemistry is associated with the secondary used to detect Oregon Green, the legend for Figure 5 mentions Nova Red and the reagent list mentions HRP. This was the only clue in the paper that the secondary antibody used to detect the Oregon Green Primary was associated with HRP. The text should contain a simple description of the reagents used sequentially in the immunofluorescence and immunohistochemistry protocols to make the actual protocols and choice of reagents easier to understand.

We appreciate the reviewer's comments and suggestions. We have provided more clarification with regard to the purpose of the IF and IHC we performed in the Introduction and Discussion. We have described the reagents used more clearly in the Protocol as recommended.

2. The paper illustrates the use of this technique for co-localization with co-labelled nuclei. These are relatively large cellular structures well suited to this protocol. However, I was concerned about how easy registration of sequentially collected images is how easily it might be effective in determining whether two structures are close or overlapping. What is the practical resolution of this approach?

We have discussed the resolution of this technique in the Discussion.

3. A concern with cryosections is the degree to which cellular morphology is retained. I was wondering how well cell membranes are preserved ? How does this approach compare with sections made with JB4 for example. In which situations would it be optimal to use one versus the other technique for embedding?

We have addressed these questions in the Discussion.