

Immunofluorescence in *Pleurobrachia pileus* adult tissues

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– PROTOCOL –

Day 1 – Sample preparation

In each step, samples can be rocked on a nutator or on an orbital (horizontal) shaker.

1. Dissect the desired adult tissue in 100% methanol.
2. Wash samples for 15 min each with a graded series of 500 µL (or 1 mL) of **methanol (60 and 30%) in PBSTx** at RT.
3. Wash samples for 15 min with 500 µL (or 1 mL) of **PBSTx** at RT.
4. (optional) Rock samples for 2 min with 500 µL (or 1 mL) of **2.6 µg/mL proteinase K** at RT.
5. (optional) Wash samples for two times without incubation with 500 µL (or 1 mL) of **PBST** at RT.
6. (optional) Post-fix samples for 30 min with 500 µL (or 1 mL) of **3.7% PFA** at RT.
7. (optional) Wash samples for 2 × 10 min with 500 µL (or 1 mL) of **PBSTx** at RT.
8. Incubate samples for 2 h with 500 µL (or 1 mL) of **PBSTx** at RT.
9. Incubate samples for >12 h (overnight) with 500 µL (or 1 mL) of **a blocking solution of 6% goat serum in PBSTx** at RT.

Day 2 – Primary antibody reaction

1. Incubate samples for 24–48 h with 500 µL of **1:40 primary antibody in 6% goat serum solution** at 4 °C.

Day 3 – Primary antibody reaction

1. *ongoing incubation with primary antibody solution*

Day 4 – Secondary antibody reaction

1. Wash samples for 6 × 30 min with 500 µL (or 1 mL) of **TBSTx** at RT.
2. optional Incubate samples for 1 h with 500 µL (or 1 mL) of **a blocking solution of 6% goat serum in PBSTx** at RT.
3. Incubate samples for >12 h (overnight) with 500 µL of **1:20 secondary antibody in 6% goat serum** at RT.

Day 5 – IF conclusion

1. Wash samples for 2×5 min, 5×15 min, and 1×5 min with 500 μ L (or 1 mL) of **PBSTx** at RT.
2. Incubate samples for 30–60 min with 1 mL of **50 % glycerol in 1 \times PBS** at RT.
3. Incubate samples for 30–60 min with 1 mL of **75 % glycerol in 1 \times PBS** at RT.
4. Collect samples from the 75 % glycerol solution and place them on a cleaned (bridged, if necessary) slide.
5. Add ~ 15 – 20 μ L of mounting medium (or VECTASHIELD® PLUS Antifade Mounting Medium with DAPI [H-2000]) directly on samples.

Adjust the amount of mounting medium according to the cover glass dimensions, the quantity of samples, and the thickness of the bridge.

6. Seal the slide with nail polish and store in the dark at 4 °C.

– RECIPES –**1× PBS with 0.01 % Triton X-100 (PBSTx or PTx)**

Compound	Quantity	Molar mass (g/mol)	Final concentration (for 50 mL)
Triton X-100	5.0 µL	647.00	0.01 %
20× PBS	5.0 mL	–	1×
Ultrapure water	–	–	–

1. Add 5 mL of 20× PBS to a graduated cylinder.
2. Add 5 µL of Triton X-100.
3. Fill up to 50 mL with ultrapure water.

6 % goat serum in PBSTx

Compound	Quantity	Molar mass (g/mol)	Final concentration (for 5 mL)
Normal goat serum	300.0 µL	–	6 %
Triton X-100	0.5 µL	647.00	0.01 %
20× PBS	250.0 µL	–	1×
Ultrapure water	–	–	–

1. Add 250 µL of 20× PBS to a beaker.
2. Add 300 µL of normal goat serum.
3. Add 0.5 µL of Triton X-100.
4. Fill up to 5 mL with ultrapure water.

Proteinase K working solution

Compound	Quantity	Molar mass (g/mol)	Final concentration (for 10 mL)
20 mg/mL proteinase K stock solution	1.30 µL	–	2.6 µg/mL
PBST	–	–	–

1. Add 1.3 µL of 20 mg/mL proteinase K stock solution to 8 mL of PBST.
2. Fill up to 10 mL with PBST.

1:500 DAPI staining solution

Compound	Quantity	Molar mass (g/mol)	Final concentration (for 500 µL)
DAPI	1.0 µL	–	1:500
1× PBS	499.0 µL	–	–

– RESOURCES –

1. IHC in various ctenophore species.

- Burkhardt, P., Colgren, J., Medhus, A., Digel, L., Naumann, B., Soto-Angel, J. J., ... & Kittelmann, M. (2023). Syncytial nerve net in a ctenophore adds insights on the evolution of nervous systems. *Science*, 380(6642), 293-297. [10.1126/science.ade5645](https://doi.org/10.1126/science.ade5645)
- Jager, M., Chiori, R., Alié, A., Dayraud, C., Quéinnec, E., & Manuel, M. (2011). New insights on ctenophore neural anatomy: immunofluorescence study in *Pleurobrachia pileus* (Müller, 1776). *Journal of Experimental Zoology Part B: Molecular and Developmental Evolution*, 316(3), 171-187. [10.1002/jez.b.21386](https://doi.org/10.1002/jez.b.21386)
- Moroz, L. L., Kocot, K. M., Citarella, M. R., Dosung, S., Norekian, T. P., Povolotskaya, I. S., ... & Kohn, A. B. (2014). The ctenophore genome and the evolutionary origins of neural systems. *Nature*, 510(7503), 109-114. [10.1038/nature13400](https://doi.org/10.1038/nature13400)
- Sachkova, M. Y., Nordmann, E. L., Soto-Àngel, J. J., Meeda, Y., Górski, B., Naumann, B., ... & Burkhardt, P. (2021). Neuropeptide repertoire and 3D anatomy of the ctenophore nervous system. *Current Biology*, 31(23), 5274-5285. [10.1016/j.cub.2021.09.005](https://doi.org/10.1016/j.cub.2021.09.005)