Staining Protocol for Zhang et al., 2019

Ginty Lab Updated: 3/14/2021

This is the staining protocol for: Multiplexed peroxidase-based electron microscopy labeling enables simultaneous visualization of multiple cell types. *Nature Neuroscience* **22**, 828-839 (2019).

Treat all samples and solutions as light-sensitive (especially the ones indicated in the list of reagents) and cover plates and vials with aluminum foils at all times. Most organic reagents are moisture sensitive and therefore containers should be kept closed as much as possible and in desiccators. Most reagents used in this protocol are highly toxic, and therefore appropriate personal protective equipment should be worn at all times and exposures should be minimized by using a fume hood.

DAB staining:

All steps are done at room temperature (r.t.) unless otherwise specified.

- 1. Sacrifice the animal 2-3 weeks after AAV injection by transcardial perfusion with warm (37 °C) Ames' medium with heparin (equilibrated with carbogen: 95% O₂, 5% CO₂) to clear blood, followed by warm (37 °C) 2.5% glutaraldehyde, 2% paraformaldehyde in cacodylate buffer. Post-fix the sample using the same fixative at 4 °C overnight. If transcardial perfusion is known to not fix your particular type of samples well, fix your samples with an alternative protocol, but use the fixative described above. Note if no blood clearing is done you will see strong endogenous peroxidase labeling of red blood cells, which may interfere with the observation of peroxidase staining in LM but does not usually cause any issue in EM analysis.
- 2. Wash the sample with cacodylate buffer, and section the sample with a vibratome at 100-200 μ m in cacodylate buffer. Use low melting point agarose for embedding if possible to minimize thermal denaturation of the enzyme. Trim the sections to include only the area expected to have staining. This and the following DAB staining steps should be done as soon as the post-fixation is done, as we have noticed loss of enzyme activity over time in storage.
- 3. Use a shaker or rotator for all staining steps to ensure even staining. Place one section per well in 12-well or 24-well plates. Wash the sections for 2×10 min with 50 mM glycine in cacodylate buffer to quench aldehyde fixatives, and then wash the sections for 1×10 min with cacodylate buffer.
- 4. Incubate the sections in 1 mL of 0.3 mg/mL 3,3'-diaminobenzidine tetrahydrochloride hydrate (DAB, corresponds to ~ 0.16 mg/mL of the free base, assuming 9.5% water content) in cacodylate buffer for 30 min. Add 10 μL of 0.3% H_2O_2 in cacodylate buffer directly into the DAB solution (to a final H_2O_2 concentration of 0.003%) and mix well to initiate the peroxidase staining reaction. Allow the reaction to proceed for 1 hr. If higher staining solution volume is needed, scale up volumes of reagents proportionally.
- 5. Wash the sections with cacodylate buffer for 4×10 min. Post-fix in 3% glutaraldehyde in cacodylate buffer at $4\,^{\circ}\text{C}$ overnight.
- 6. Wash sections in cacodylate buffer for 1×10 min, and then in 50 mM glycine in cacodylate buffer for 1×10 min, and finally in cacodylate buffer for 2×10 min. Samples can be stored at 4 °C for a few days before further processing without apparent issues.

EM sample preparation:

Use a shaker or rotator for all staining steps to ensure even staining. Use glass scintillation vials for EM preparation.

For typical TEM applications, **1)** Normal-contrast TEM protocol is recommended. If high sample electron contrast is required, such as imaging using SEM or certain TEM platforms, follow **2)** SEM or High-contrast TEM protocol.

1) Normal-contrast TEM protocol:

Day 1:

- 1. Stain in 1% osmium tetroxide/1.5% potassium ferrocyanide in cacodylate buffer at r.t. for 1 hr. Place filter paper and weights on top of sections for the first 15 min to flatten the sections if necessary.
- 2. Wash sections in ddH_2O for 4×5 min.
- 3. Stain in 1% uranyl acetate in 0.05 M maleate buffer at 4 °C overnight.

Day 2:

- 1. Wash sections in ddH_2O for 4×5 min.
- 2. Dehydrate the sections with 50 % EtOH at 4 °C for 5 min.
- 3. Dehydrate the sections with 70 % EtOH at 4 °C for 5 min.
- 4. Dehydrate the sections with 90 % EtOH at 4 °C for 5 min.
- 5. Dehydrate the sections with 95 % EtOH for 3×10 min, with the first wash warm to r.t. from 4 °C, and second and third washes at r.t.
- 6. Dehydrate the sections with 100 % EtOH at r.t. for 3×20 min.
- 7. Exchange the sections into propylene oxide for 2×30 min.
- 8. Infiltrate the sections with 20% resin mix in propylene oxide for 1 hr.
- 9. Infiltrate the sections with 40% resin mix in propylene oxide for 1 hr.
- 10. Infiltrate the sections with 50% resin mix in propylene oxide overnight.

Day 3:

- 1. Infiltrate the sections with 60% resin mix in propylene oxide for 1 hr.
- 2. Infiltrate the sections with 80% resin mix in propylene oxide for 1 hr.
- 3. Infiltrate the sections with 100% resin mix for 1 hr.
- 4. De-gas the samples under vacuum for 1 hr.
- 5. Embed the sections in 100% resin mix and cure at 60 °C for 48-72 hrs. Samples can be stored indefinitely at r.t. after curing.

Sample imaging considerations:

- 1. Section at 30-40 nm for best resolution.
- 2. We pick up sections on Formvar or Pioloform films coated with carbon. Other types of sample support may work but may have different properties under the electron beam.
- 3. Do not stain with lead citrate or uranyl acetate on ultrathin sections, as extra staining could obscure the DAB signal. Samples should have sufficient contrast when imaged with a digital camera.

2) SEM or High-contrast TEM protocol (modified from Hua et al., 2015):

Day 1:

- 1. Stain in 2% osmium tetroxide in cacodylate buffer at r.t. for 1 hr. Place filter paper and weights on top of sections for the first 15 min to flatten the sections if necessary.
- 2. Stain in 2.5% potassium ferrocyanide in cacodylate buffer at r.t. for 1 hr.
- 3. Wash sections in ddH_2O for 4×5 min.
- 4. Stain in 1% thiocarbohydrazide in ddH₂O at 40 °C for 15 min.
- 5. Wash sections in ddH_2O for 4×5 min.
- 6. Stain in 2% osmium tetroxide in ddH₂O at r.t. for 1 hr.
- 7. Wash sections in ddH_2O for 4×5 min.
- 8. Stain in 1% uranyl acetate in 0.05 M maleate buffer at 4 °C overnight.

Day 2:

- 1. Heat sections in uranyl acetate solution to 50 °C for 2 hrs.
- 2. Wash sections in ddH₂O for 4×5 min.
- 3. Dehydrate the sections with 50 % EtOH at 4 °C for 5 min.
- 4. Dehydrate the sections with 70 % EtOH at 4 °C for 5 min.
- 5. Dehydrate the sections with 90 % EtOH at 4 °C for 5 min.
- 6. Dehydrate the sections with 95 % EtOH for 3×10 min, with the first wash warm to r.t. from 4 °C, and second and third washes at r.t.
- 7. Dehydrate the sections with 100 % EtOH at r.t. for 3×20 min.
- 8. Exchange the sections into propylene oxide for 2×30 min.
- 9. Infiltrate the sections with 20% resin mix in propylene oxide for 1 hr.
- 10. Infiltrate the sections with 40% resin mix in propylene oxide for 1 hr.
- 11. Infiltrate the sections with 50% resin mix in propylene oxide overnight.

Day 3:

- 1. Infiltrate the sections with 60% resin mix in propylene oxide for 1 hr.
- 2. Infiltrate the sections with 80% resin mix in propylene oxide for 1 hr.
- 3. Infiltrate the sections with 100% resin mix for 1 hr.
- 4. De-gas the samples under vacuum for 1 hr.
- 5. Embed the sections in 100% resin mix and cure at 60 °C for 48-72 hrs. Samples can be stored indefinitely at r.t. after curing.

Special considerations for using Matrix-dAPEX2 and IMS-dAPEX2:

- 1. This protocol leads to strong staining in the IMS for mitochondria labeled with Matrix-dAPEX2 (see **Supplementary Figure 5**). This is preferred because this IMS staining pattern is usually not obscured with heavy counterstaining required for SEM. Do not use IMS-dAPEX2 with this protocol, as it has a smaller range of optimal expression level than Matrix-dAPEX2 and often not distinguishable from Matrix-dAPEX2 under this staining condition.
- 2. Consider examining ultrathin sections with TEM first to determine the percentage of mitochondria with IMS staining when using Matrix-dAPEX2 with SEM. If the percentage is too low, you can increase the DAB concentration to 0.5 mg/mL and see if IMS staining becomes more prevalent.

Sample imaging considerations:

- 1. Section at 30-40 nm for best resolution. We use the same sectioning and pickup setup as in Kasthuri *et al.*, 2014.
- 2. Stain with lead citrate and uranyl acetate on sections to increase contrast.
- 3. Ultrathin sections can be imaged with SEM using either backscattered electrons or secondary electrons. Postsynaptic densities tend to be more prominent using backscattered electrons, but imaging using secondary electrons can be substantially faster.

Catalog numbers for reagents:

Please check expiration dates and stability of chemicals and avoid using degraded chemicals.

Ames' medium: Sigma A1420

Sodium bicarbonate: Sigma-Aldrich 792519

Heparin: Sigma-Aldrich H3393

Glutaraldehyde: Electron Microscopy Sciences 16316 Paraformaldehyde: Electron Microscopy Sciences 15712 Sodium cacodylate: Electron Microscopy Sciences 12310

Hydrochloric acid: Fisher A144SI-212 Calcium chloride: Sigma-Aldrich 793639

Magnesium chloride hexahydrate: Sigma-Aldrich M9272

Glycine: Sigma-Aldrich 410225

3,3'-diaminobenzidine tetrahydrochloride hydrate: Sigma D5637, **light sensitive**

Hydrogen peroxide: Sigma-Aldrich 216763, light sensitive

Osmium tetroxide: Electron Microscopy Sciences 19190, light sensitive

Potassium ferrocyanide: Sigma-Aldrich P3289, light sensitive

Uranyl acetate: Electron Microscopy Sciences 22400, light sensitive, radioactive

Maleic acid: Sigma-Aldrich M0375 Sodium hydroxide: Fisher SS266-1 Ethanol: Sigma-Aldrich 459836

Propylene oxide: Electron Microscopy Sciences 20411

LX-112: Ladd Research 21310

DDSA: Electron Microscopy Sciences 13710 NMA: Electron Microscopy Sciences 19000 DMP-30: Electron Microscopy Sciences 13600

SEM or High-contrast TEM protocol only:

Thiocarbohydrazide: Electron Microscopy Sciences 21900, light sensitive

Protocols for solution preparation:

Ames' medium with heparin (10 U/mL)

Dissolve the Ames' medium powder in appropriate amounts of ddH_2O . Gas the solution with 100% CO_2 , and then slowly add sodium bicarbonate at 1.9 g/L. Make sure no precipitates are formed. Dissolve heparin in ddH_2O to make a 10 kU/mL solution. Add 1 mL of 10 kU/mL heparin solution per 1 L of Ames' medium. Filter the solution with a 0.22 μ m filter and store at 4 °C.

Cacodylate buffer (0.15 M sodium cacodylate, 4 mM CaCl₂, 4 mM MgCl₂, pH 7.4)

Dissolve sodium cacodylate (64.21 g for 1 L), calcium chloride (0.89 g for 1 L) and magnesium chloride hexahydrate (1.63 g for 1 L) in ddH_2O to make a 0.3 M sodium cacodylate, 8 mM calcium chloride, 8 mM magnesium chloride solution. Adjust pH to 7.4 with 12 M hydrochloric acid. Filter the solution with a 0.22 μ m filter and store at 4 °C. Mix equal volumes of 0.3 M sodium cacodylate, 8 mM calcium chloride, 8 mM magnesium chloride solution (pH 7.4) and ddH_2O to make cacodylate buffer. Store at 4 °C.

DAB stock solution

Dissolve 3,3'-diaminobenzidine tetrahydrochloride hydrate in ddH_2O to make a 50 mg/mL solution. Aliquot and store the solution at -20 °C. Thaw the solution just before use and vortex to dissolve all solids. The solution can be refrozen for repeated uses but should be discarded when significant color appears in the solution.

1% uranvl acetate in 0.05 M maleate buffer (pH 5.15)

Dissolve 11.6~g of maleic acid with 100~mL 1~N sodium hydroxide solution and bring up the volume to 500~mL with ddH_2O to make a 0.2~M sodium maleate solution. Combine 100~mL of 0.2~M sodium maleate solution, 10.76~mL 1~N sodium hydroxide solution, bring up the volume to 400~mL with ddH_2O and verify that pH is 6.0, and then add 4~g of uranyl acetate to make 1% uranyl acetate in 0.05~M maleate buffer. Filter the solution with a $0.22~\mu m$ filter and store at 4~°C.

Resin mix

	~ 60 g total	~ 30 g total
Resin (WPE = 143)	30.00 g	15.00 g
DDSA	16.38 g	8.19 g
NMA	15.21 g	7.61 g
DMP-30	0.78 g	0.39 g

Mix the first three components together thoroughly for 5 min by hand. Then add DMP-30 and mix for another 5 min. Use the freshly prepared resin mix immediately for infiltration and do not store and reuse the resin mix, as polymerization occurs immediately after all the components are mixed in leading to increasing viscosity, even at r.t.

This resin mix is equivalent to the 5 mL/5 mL Mixture A/Mixture B combination in Luft, 1961. We used LX-112 as the resin. We have not tried other types of resins, but they may also give comparable results. Different infiltration/embedding schedules may need to be used if using other types of resins or processing hard to infiltrate sample types.

For consistent results, adjust the amounts of components using the WPE specific to your batch of resin. To calculate the amounts of components, multiply the amounts of DDSA and NMA in the table above by 143 / WPE. Then, for each gram of increase/decrease of total anhydrides (DDSA + NMA), increase/decrease the amount of DMP-30 by 0.0132 g. For example, if the WPE of your resin is 160, then for 30.00 g of resin, you will need 16.38 * 143 / 160 = 14.64 g of DDSA, 15.21 * 143 / 160 = 13.59 g of NMA, and 0.78 - (1.74 + 1.62) * 0.0132 = 0.74 g of DMP-30.

Normal-contrast TEM protocol only:

1% osmium tetroxide/1.5% potassium ferrocyanide in cacodylate buffer

Mix equal volumes of 4% osmium tetroxide and ddH_2O to make a 2% osmium tetroxide solution. Dissolve potassium ferrocyanide in 0.3 M sodium cacodylate, 8 mM calcium chloride, 8 mM magnesium chloride solution (pH 7.4) to make a 3% potassium ferrocyanide solution. Add equal

volumes of the 3% potassium ferrocyanide solution dropwise to the 2% osmium tetroxide solution, and then mix well to make 1% osmium tetroxide/1.5% potassium ferrocyanide in cacodylate buffer.

SEM or High-contrast TEM protocol only:

2% osmium tetroxide in cacodylate buffer

Mix equal volumes of 4% osmium tetroxide and 0.3 M sodium cacodylate, 8 mM calcium chloride, 8 mM magnesium chloride solution (pH 7.4) to make 2% osmium tetroxide in cacodylate buffer.

2.5% potassium ferrocyanide in cacodylate buffer

Dissolve potassium ferrocyanide in cacodylate buffer to make 2.5% potassium ferrocyanide in cacodylate buffer.

1% thiocarbohydrazide solution

This solution needs to be freshly made every time. Add thiocarbohydrazide to ddH_2O for a 1% solution. Warm up the mixture to 60 °C and vortex to dissolve thiocarbohydrazide, then cool it down to 40 °C and filter it with a 0.22 μ m filter before use.

Troubleshooting:

1) Low staining intensity observed in light microscopy

- 1. Inadequate enzyme expression level: check virus injection quality, virus concentration, suitability of viral delivery route, recombinase activity, etc.
- 2. Degraded enzyme: conduct DAB staining immediately after post-fixation and do not store the samples, avoid exposing tissues to high temperatures.
- 3. Degraded chemicals: replace chemical stocks, especially hydrogen peroxide and DAB.
- 4. Insufficient/inappropriate mixing of chemicals: mix chemicals well.
- 5. Using SV-HRP construct: this construct does not usually give strong staining observable in light microscopy, proceed to conduct electron microscopy.

2) Low staining intensity observed in electron microscopy

- 1. Check issues and solutions in 1).
- 2. Lead citrate and uranyl acetate staining used on ultrathin sections with *Normal-contrast TEM protocol*: do not use additional staining.
- 3. Poor ultrathin section: image other ultrathin sections from the same sample.
- 4. Insufficient staining penetration in the center of the sample: reduce sample thickness.

3) Abnormal organelle ultrastructure

1. Excessive enzyme expression level: most typically seen with IMS-dAPEX2, reduce virus volume and/or concentration.

4) Poor tissue ultrastructure

- 1. Poor fixation: prepare new samples with better fixation.
- 2. Physical injury from virus injections: use finer needles, inject more slowly, inject lower volumes, etc.