

Immunoisolation of Centrosomes from *Drosophila melanogaster*

The isolation of centrosomes from cells has been, for a long time, a challenging task. The main reason is that this small organelle (~0.3 μm in diameter in mammalian cells) is present only in a single copy in the cell and therefore represents only a minuscule part of the whole cell proteome. Another technical difficulty is that the density of this organelle ($\rho > 1.32 \text{ g/ml}$) is too high for it to be purified in a continuous gradient using isopycnic density centrifugation. Hence, step-gradient centrifugation has been the method of choice for most of the isolation approaches used to date. Both Ficoll and sucrose density gradients, as well as a combination of the two, have been employed. Most recent methods use sucrose gradient centrifugation only. Density gradient centrifugation is followed by immunofluorescence assays, microtubule nucleation assays, or electron microscopy to evaluate enrichment of cell organelles in particular fractions of the gradient.

Currently, isolation of centrosomes from *Drosophila* embryos, although relatively inefficient (i.e., only ~10% of total centrosomes present in starting material are isolated), results in some of the best total yields compared to isolation of centrosomes from other organisms. In short, this method permits production of excellent quantities for many biochemical, functional, and structural studies.

The method described here focuses on the isolation of centrosomes from the syncytial stages of the early *Drosophila* embryo (0 to 3.5 hr). This has the particular advantage that cell organelles are not bounded by cellular membranes and that lysis can be performed on a relatively small scale, keeping the resulting volume low. This is an essential factor for subsequent gradient centrifugation steps, because ultracentrifugation of large buffer volumes is impractical. The abundance of pericentriolar material of these centrosomes isolated from the highly mitotic embryos produces excellent total protein yields. Moreover, maintenance of large fly populations that produce grams of starting material within a few hours is relatively cost efficient.

Basic Protocol 1 describes isolation of centrosomes from early-syncytial-stage *Drosophila* embryos by sucrose step gradient centrifugation. Basic Protocol 2 details an immunomagnetic isolation procedure for further purification of the centrosomes isolated in Basic Protocol 1. Support Protocol 1 details an immunofluorescence microscopy procedure for quantitatively tracking the gradient purification of centrosomes, and, finally, Support Protocol 2 describes fluorescence microscopy to assess the purification of the immunomagnetically isolated centrosome preparations.

ISOLATION OF CENTROSOMES FROM THE EARLY SYNCYTIAL STAGES OF THE *DROSOPHILA* EMBRYO USING SUCROSE STEP GRADIENT CENTRIFUGATION

This protocol outlines the production of *Drosophila* embryo extract as starting material for the isolation of centrosomes. Embryos are homogenized in homogenization buffer (HB), centrifuged at low speed, and filtered twice to remove large aggregates of membranes, debris, and other cell organelles. The two consecutive low-speed centrifugations also eliminate the large quantities of lipids that float on top of the embryo extract; these lipids need to be carefully removed to avoid contamination in subsequent isolation steps. The first two steps of density centrifugation serve to pool centrosomes for the subsequent immunoisolation, rather than to enrich cell organelles in a particular fraction. The

BASIC PROTOCOL 1

Subcellular Fractionation and Isolation of Organelles

3.17.1

Contributed by Verena Lehmann, Hannah Müller, and Bodo M.H. Lange

Current Protocols in Cell Biology (2005) 3.17.1-3.17.13

Copyright © 2005 by John Wiley & Sons, Inc.

affinity-isolation step then removes contaminants that copurify in the sucrose gradient to obtain a high degree of purity.

This protocol was adapted from Moritz et al. (1995) and Moritz and Alberts (1999) for use with the immunoisolation protocols developed by the authors of this unit.

Materials

Large population of *Drosophila* (e.g., ~250,000 strain W118 flies) maintained in large environmental incubator or temperature- and humidity-controlled room with fixed day (12 hr) and night (12 hr) cycles (Greenspan, 1997; Bonte and Becker, 1999; Sullivan et al., 2000)

Apple juice/molasses agar plates (see recipe)

Yeast paste: ordinary baker's yeast dissolved in warm water to form a paste

Embryo wash: 0.7% (w/v) NaCl/0.04% (v/v) Triton X-100

3% (v/v) sodium hypochlorite in embryo wash (see above)

Homogenization buffer (HB, see recipe)

Liquid N₂

5× BRB80 with 100 mM KCl (see recipe), diluted to 1×

25% (v/v) Triton X-100

100× protease inhibitor mix (PIM, see recipe)

55% and 70% (w/v) sucrose solutions (see recipes)

Fine strainer

Filter unit fitted onto vacuum flask

Motor-driven Wheaton homogenizer with tight-fitting Teflon pestle (60-ml volume)

50-ml conical polypropylene centrifuge tubes

Refrigerated low-speed centrifuge

Miracloth (Calbiochem)

15-ml snap-cap polypropylene tubes for freezing and storage of supernatants

39-ml thin-walled polyallomer ultracentrifuge tubes (e.g., Beckman)

Large-volume ultracentrifuge (e.g., Beckman) with swinging-bucket rotor (e.g., Beckman SW 32)

Additional reagents and equipment for assessing purify of centrosome fractions from gradient (see Support Protocol 1)

Homogenize embryo material

1. Collect 30 to 50 g of 0- to 3.5-hr-old *Drosophila* embryos by placing a fresh apple juice/molasses agar plate with a lump of yeast paste on the surface into the population cage every 3 hr.

Depending on the size of the fly population, three collections of 3 hr each will be needed in order to obtain a total of ~30 to 50 g of embryo material.

2. After the 3-hr collection period for each plate, wash embryos off the plate with embryo wash using a paint brush to aid in removing the embryos, collecting them in a fine strainer, and store the embryos in ice-cold embryo wash until all embryos have been collected.
3. Remove the chorion layer by stirring embryos for 3 min in 250 ml of 3% sodium hypochlorite solution.
4. Intensively wash the embryos with a forceful stream of cold tap water, then three times with a total volume of 1 liter of ice-cold distilled, deionized water, on a filter unit fitted onto a vacuum flask. Determine the dry weight after removing all excess water.
5. Resuspend dechorionated embryos in HB at a ratio of 1:5 (w/v).

6. Homogenize material on ice using five strokes of the motor-driven Wheaton homogenizer at a speed of 1500 rpm in a cold room.

CAUTION: *Wear protective gloves and goggles when using the homogenizer.*

Remove lipids and prepare/store HSS

7. Transfer homogenate to 50-ml conical polypropylene centrifuge tubes. Centrifuge 10 min at $1800 \times g$, 4°C . Remove floating lipids by aspirating with a suction pump and filter supernatant through Miracloth. Repeat the centrifugation, lipid removal, and filtration a second time.
8. Divide the resulting supernatant (referred to as HSS: i.e., “homogenate, supernatant, supernatant”) into 10-ml aliquots in 15-ml conical polypropylene tubes, snap-freeze in liquid nitrogen, and store at -80°C until further use.

The extract can be conveniently frozen at this stage if the isolation procedure must be interrupted. Freezing also gives the opportunity to produce stockpiles of embryo extracts without affecting the functional and structural integrity of the centrosome.

Perform sucrose-gradient purifications

9. Prepare six 50-ml conical polypropylene centrifuge tubes, each containing:
 - 20 ml 70% (w/v) sucrose
 - 1 ml $1 \times$ BRB80 with 100 mM KCl
 - 640 μl 25% Triton X-100
 - 400 μl $100 \times$ PIM.
10. Drop the contents of each 15-ml tube containing a frozen 10-ml HSS aliquot into one of the 50-ml tubes containing the mixture prepared in the previous step, and thaw on ice. Collect a 30- μl aliquot for immunofluorescence microscopy (see below).
11. For each of the extract mixtures prepared in the previous step, pour a sucrose gradient in a 39-ml thin-walled polyallomer ultracentrifuge tube by carefully overlaying 3 ml of 70% sucrose solution with 4 ml of 55% sucrose, and, finally, at the top, the HSS mixture from the preceding step. Mark the upper margin of the 55% sucrose layer of the sucrose gradient on the plastic wall of the tube.
12. Centrifuge samples for 1.5 hr at $100,000 \times g$ in a SW 32 rotor with slow acceleration and slow braking.

Slow acceleration and braking correspond to “setting 4” on the Beckman instrument.
13. Aspirate the supernatant above the marked line with a vacuum aspirator. Pool the gradients (all of the sucrose solutions below the mark; Pool 1) and discard the pellets. Take a 30- μl sample for immunofluorescence microscopy.
14. Dilute the ~ 35 to 40 ml Pool 1 to 70 ml with $1 \times$ BRB80 containing 100 mM KCl and add $100 \times$ PIM to a final concentration of $1 \times$.
15. Prepare two sucrose gradients in 39-ml thin-walled polyallomer tubes ultracentrifuge tubes by carefully overlaying 3 ml of 70% sucrose solution with 1 ml of 55% sucrose solution. Split the diluted Pool 1 (step 14) into two equal portions, and carefully overlay each of these portions over the 55% sucrose layer in each of the two respective tubes. Mark the upper margin of the 55% sucrose layer of the sucrose gradient on the plastic wall of the tube.
16. Ultracentrifuge the gradients 1 hr at $100,000 \times g$, 4°C . Remove the supernatant from each tube with a vacuum aspirator and discard the pellets. Pool the sucrose gradients (resulting in Pool 2). Take a 30- μl sample for immunofluorescence microscopy.

**SUPPORT
PROTOCOL 1**

17. Assess the sample aliquots from each purification stage by immunofluorescence microscopy (see Support Protocol 1). Use the remainder of Pool 2 for further purification by immunoisolation (Basic Protocol 2).

In principle, these enriched preparations can also be used for functional in vitro microtubule nucleation assays and for standard immunofluorescence colocalization studies. However, the preparations are too crude to be used for biochemical analysis of centrosome composition. For convenience, samples can also be snap-frozen in liquid nitrogen until used in Basic Protocol 2.

IMMUNOFLUORESCENCE MICROSCOPY OF ISOLATED CENTROSOMES

This protocol is used to assess the enrichment and yield of aliquots from the intermediate steps in preparation of centrosomes.

Materials

- 30- μ l aliquots from Pool 1 and Pool 2 (see Basic Protocol 1)
- PBS-T: phosphate-buffered saline (PBS; APPENDIX 2A), pH 7.4, containing 0.003% (v/v) Triton X-100
- Methanol, -20°C
- Primary antibody: anti- γ -tubulin (Sigma)
- Secondary antibody: fluorochrome-conjugated antibody specific for IgG of the species from which the primary anti- γ -tubulin antibody was obtained
- Mounting medium (e.g., Mowiol from Calbiochem, or 50% v/v glycerol)
- 20 mg/ml (10 \times) *p*-phenylenediamine in H₂O
- Nail polish
- Plastic inserts to support a 11-mm round glass coverslip (Evans et al., 1985)
- 15-ml Corex tubes
- 11-mm round glass coverslips (grade 1)
- Refrigerated centrifuge with swinging-bucket rotor (e.g., Sorvall HB-6 or Beckman JS 13.1, or equivalent) and adapters
- 24-well tissue culture plates
- Spatula bent at the tip for removing the plastic insert
- Needle with bent tip
- Forceps
- Microscope slides
- Epifluorescence microscope equipped with immunofluorescence filters and appropriate optics

NOTE: To avoid staining artifacts, never allow the coverslips to dry out at any point during the following protocol.

Centrifuge centrosome sample onto coverslip

1. Resuspend the 30- μ l aliquot of *Drosophila* centrosomes from HSS mix Pool 1 or Pool 2 in 11 ml PBS-T.
Keep samples on ice until centrifugation.
2. Place a pair of plastic inserts into a 15 ml-Corex glass tube to support a 11-mm round glass coverslip.
3. Transfer the 11-ml diluted centrosome sample into a 15-ml Corex tube and centrifuge onto the coverslips for 15 min at 15,000 \times g, 4 $^{\circ}\text{C}$.
4. Remove the coverslip by lifting the top insert out of the Corex tube with a spatula bent at the tip. Transfer the coverslip with a pair of forceps immediately into -20°C methanol in a well of a 24-well tissue culture plate and incubate at least 5 min at -20°C to fix the sample.

5. Aspirate methanol and rehydrate sample in the well by covering with 1.5 ml PBS-T and incubating 15 min at room temperature. Aspirate PBS-T.

Stain with antibodies

6. Add ~80 μ l of the anti- γ -tubulin primary antibody, appropriately diluted in PBS-T (e.g., 1:400 for polyclonal antisera, depending on antibody used), to each coverslip-containing well. Incubate 1 hr at room temperature in a moist environment.
7. Wash four times, each time by filling the sample-containing well(s) with an excess (e.g., 1.5 ml) of PBS-T, incubating 2 min at room temperature, and aspirating the solution.

Apply the PBS along the wall of each well and not directly onto the sample, to avoid rinsing the centrosomes away.

8. Add ~80 μ l of the fluorophore-labeled secondary antibody diluted in PBS-T according to the manufacturer's recommendations to each coverslip-containing well. Incubate 45 min at room temperature in the dark in a moist environment.

The antibody dilution will vary; e.g., anti-mouse IgG whole-molecule dilutions will range between 1:400 and 1:2000.

Mount coverslip on slide and examine centrosome preparation

9. Remove coverslip(s) from well(s) with a bent needle and a pair of forceps and dip each coverslip briefly in water to rinse off excess PBS.

Coverslips can more easily be removed from the plastic wells using a needle with a slightly bent tip.

10. Add 20-mg/ml *p*-phenylenediamine to the mounting medium (e.g., Mowiol or 50% glycerol) for a final concentration of 1 mg/ml. Mount each coverslip on a microscope slide with 6 μ l of the phenylenediamine-containing mounting medium. Carefully remove excess mounting medium with tissue, and allow to set for 20 min at room temperature in the dark. Seal with nail polish.
11. Check preparations under a fluorescence microscope or store at 4°C in the dark until samples are viewed.

The centrosomes should be homogenous in size and shape, and should be enriched through each step of the isolation procedure.

IMMUNOPURIFICATION OF CENTROSOMES WITH MAGNETIC BEADS

Prepare magnetic beads cross-linked with affinity-purified anti- γ -tubulin antibody. For negative control, magnetic beads are cross-linked with the relevant preimmune antibody. Centrosomes should be well enriched and quite similar in size. A ratio of 1:6 (v/v) between beads and centrosomes is recommended.

Materials

- Protein G magnetic bead suspension (Dynal)
- Phosphate-buffered saline (PBS; APPENDIX 2A)
- 0.2 M triethanolamine, pH 8.2, with and without 20 mM dimethyl pimelidate dihydrochloride (DMP); prepare fresh
- Anti- γ -tubulin antibody (anti-peptide antibody; not commercially available; affinity-purified; Tavosanis et al., 1997)
- Tris-Cl, pH 7.5 (APPENDIX 2A)
- PBS (APPENDIX 2A) containing 0.1% (v/v) Tween 20
- PBS (APPENDIX 2A) containing 0.1% (v/v) Tween 20 and 0.02% (w/v) thimerosal
- Pool 2 of centrosomes (see Basic Protocol 1)
- Dilution buffer (see recipe), 4°C

BASIC PROTOCOL 2

Subcellular Fractionation and Isolation of Organelles

3.17.5

PBS-T: phosphate-buffered saline (PBS; *APPENDIX 2A*), pH 7.4, containing 0.003% (v/v) Triton X-100, 4°C

Methanol, -20°C

SDS sample buffer (*UNIT 6.1*)

Magnetic particle collector (magnet and magnetic stand) suitable for 1.5-ml microcentrifuge tubes (e.g., Dynal)

End-over-end rotator

Additional reagents and equipment for SDS-PAGE (*UNIT 6.1*) and determining purity of centrosome preparations from immunomagnetic purification (see Support Protocols 1 and 2)

Prepare the antibody-coated magnetic beads

1. Place 120 μ l of protein G magnetic bead suspension in a 1.5-ml microcentrifuge tube. Place the tube on the magnetic stand to immobilize the beads with the magnet according to the manufacturer's documentation, and remove the supernatant. Add 100 μ l of PBS, resuspend the beads, and immobilize the beads on the magnet again. Repeat this wash four times.
2. Remove the supernatant and add 100 μ g of anti- γ -tubulin antibody to the washed protein G beads. Incubate 45 min on an end-over-end rotator at room temperature. Immobilize beads and remove supernatant.
3. Wash the beads three times, each time with 1 ml of 0.2 M triethanolamine, pH 8.2, by immobilizing the beads on the magnet.
4. Cross-link by incubating the antibody-coupled beads with 1 ml of freshly prepared 0.2 M triethanolamine, pH 8.2, containing 20 mM DMP, on an end-over-end rotator at room temperature for 30 min.
5. Place tube on the magnetic stand and remove supernatant. Add 1 ml of 50 mM Tris-Cl, pH 7.5, and incubate on an end-over-end rotator at room temperature for 15 min.
6. Remove supernatant and wash beads three times, each time with 1 ml PBS containing 0.1% Tween 20, by immobilizing the beads on the magnet. Store antibody-coupled beads at 4°C in the original bead-suspension volume of PBS containing 0.1% Tween 20 and 0.02% thimerosal.

*Antibody binding and cross-linking efficiency can be checked by SDS-PAGE (*UNIT 6.1*) of 20- μ l samples of beads before and after cross-linking.*

Perform immunomagnetic separation

7. Take 120 μ l antibody-coupled beads suspension and wash twice, each time with 1 ml wash buffer using the technique described above. Take a sample of 20 μ l for later analysis by SDS-PAGE (*UNIT 6.1*).
8. Thaw 0.5 ml of centrosome Pool 2 on ice, then mix with 0.5 ml dilution buffer. Take a 20- μ l sample for further analysis (see Support Protocol 1).
9. Add the 1 ml of diluted centrosomes to 100 μ l washed, cross-linked beads. Mix gently by inverting the tube. Incubate 1 hr at 4°C on an end-over-end rotator. Immobilize the beads on the magnetic collector and carefully remove the supernatant. Keep the supernatant on ice for later analysis.
10. Wash beads three times, each time with 1 ml cold wash buffer, using the technique described above.
11. Wash beads twice, each time with 1 ml cold PBS-T, using the technique described above. Resuspend in 80 μ l (original bead volume) of PBS-T. Remove an aliquot of 10 μ l for immunofluorescence microscopy and an aliquot of 20 μ l for SDS-PAGE.

12. Remove the PBS-T from the bead suspension with bound centrosomes (50 μ l), resuspend in 55% sucrose solution, snap-freeze in liquid nitrogen, and store at -80°C .

The purified centrosome preparation is used subsequently in various assays. These include, for example, functional tests such as microtubule nucleation assay, electron microscopic study of centrosome morphology, antigen localization, or antigen for the generation of monoclonal or polyclonal antibodies, and biochemical assays.

13. Fix the 10- μ l aliquot of beads reserved for immunofluorescence microscopy for at least 5 min in 1 ml of -20°C methanol and store at -20°C until analyzed according to Support Protocol 2.

Assess purity

14. Immobilize beads and remove supernatant as described above. Add SDS sample buffer, and boil 5 min. Perform magnetic collection to remove beads and run the supernatant on a 10% SDS-PAGE gel (UNIT 6.1).
15. Assay diluted centrosome fractions before (see Support Protocol 1) and after immunomagnetic separation (see Support Protocol 2) by fluorescence microscopy.

IMMUNOFLUORESCENCE MICROSCOPY OF IMMUNOPURIFIED CENTROSOMES ON MAGNETIC BEADS

SUPPORT PROTOCOL 2

Materials

Centrosomes immunopurified on magnetic beads (see Basic Protocol 2)
Methanol, -20°C
PBS-T: phosphate-buffered saline (PBS; APPENDIX 2A), pH 7.4, containing 0.003% (v/v) Triton X-100
Primary antibody: anti- γ -tubulin (Sigma)
PBS-T containing 0.1% (w/v) bovine serum albumin (BSA)
Secondary antibody: fluorochrome-conjugated antibody specific for IgG of the species from which the primary anti- γ -tubulin antibody was obtained
Mounting medium: Mowiol (Calbiochem)
20 mg/ml (10 \times) *p*-phenylenediamine in H_2O
Clear nail polish
Magnetic particle collector (magnet and magnet stand) suitable for 1.5-ml microcentrifuge tubes (e.g., Dynal)
End-over-end rotator
Microscope slides
Round coverslips (grade 1, 11 mm diameter)
Epifluorescence microscope equipped with immunofluorescence filters and appropriate optics

NOTE: Handle samples with care and avoid vigorous mixing or vortexing.

1. Fix a 10- μ l magnetic bead-centrosome sample in 1 ml -20°C methanol by incubating for at least 5 min at -20°C . Place the test tube containing the beads on the magnetic stand, immobilize the beads with the magnet as described in the manufacturer's documentation, and remove the supernatant.
2. Add 1 ml PBS-T to the beads in the tube. Mix gently by inverting the tube, then incubate 15 min on an end-over-end rotator. Collect beads and remove supernatant as described above.
3. Dilute the anti- γ -tubulin primary antibody 1:500 in PBS-T containing 0.1% BSA. Add 300 μ l of the diluted primary antibody to the beads in the tube and mix gently

Subcellular Fractionation and Isolation of Organelles

3.17.7

by inversion. Incubate 1 hr at room temperature with gentle rotation. Collect beads and remove supernatant as described above.

4. Wash beads four times, each time by immobilizing the beads on the magnet, removing the supernatant, resuspending the collected beads in 1 ml PBS-T, then removing the supernatant.
5. Dilute the fluorochrome-conjugated secondary antibody appropriately in PBS-T containing 0.1% BSA. Add 300 μ l of the diluted secondary antibody to the beads in the tube and mix gently by inverting the tube. Incubate 45 min at room temperature on an end-over-end rotator in the dark (cover tubes with aluminum foil). Collect beads and remove supernatant as described above.
6. Wash beads four times, each time with 1 ml PBS-T using the technique described in step 4.
7. Add 20 mg/ml *p*-phenylenediamine to Mowiol mounting medium for a final concentration of 2 mg/ml. Add \sim 30 μ l of this medium to the tube and mix gently by inverting the tube (solution is viscous).
8. Place 6 μ l of bead/Mowiol-mix on a microscope slide and carefully mount a coverslip on top. Let set for \sim 20 min at room temperature in the dark. Seal the edges of the coverslip with nail polish.
9. Examine preparation under the epifluorescence microscope or store at 4°C in the dark until samples are viewed.

REAGENTS AND SOLUTIONS

Use deionized or distilled water in all recipes and protocol steps. For common stock solutions, see APPENDIX 2A; for suppliers, see SUPPLIERS APPENDIX.

Apple juice/molasses agar plates

For 200 plates:

11.5 liters H₂O

5 liters apple juice

750 ml molasses

500 g agar

420 ml 10% (w/v) Nipagin (Fluka) in absolute ethanol

BRB80, 5 \times

To 100 ml H₂O add:

24.2 g PIPES (400 mM final)

2 ml 0.5 M EGTA (5 mM final)

1 ml 1 M MgCl₂ (5 mM final)

Adjust pH to 6.8 with KOH pellets.

Add H₂O to 200 ml

Store up to 2 months at 4°C

Dilution buffer

To 200 ml phosphate-buffered saline (PBS; APPENDIX 2A) add:

0.4 g bovine serum albumin (BSA, fraction V; 0.2% w/v final)

100 \times protease inhibitor mix (see recipe) to 2 \times final

20 mg/ml stock of DNase I, RNase free to final concentration of 20 μ g/ml (stock stored at -20° C)

Prepare fresh just prior to use

Homogenization buffer

To 140 ml H₂O add:

- 40 ml 5× BRB80 (1× final)
 - 20 ml 1 M KCl (100 mM final)
 - 28 g sucrose (14% w/v final)
 - 100× protease inhibitor mix (see recipe) to 2× final
- Prepare fresh

Protease inhibitor mix (PIM), 100×

To 3 ml of methanol add:

- 4 ml of 1 mg/ml aprotinin (dissolved in H₂O)
 - 4 ml of 1 mg/ml leupeptin (dissolved in H₂O)
 - 4 ml of 1 mg/ml pepstatin A (dissolved in methanol)
 - 5 ml of 100 mg/ml Pefabloc SC (Roche; dissolved in H₂O)
- Store up to 2 weeks at –20°C

Sucrose solution, 55% (w/v)

To 140 ml H₂O add:

- 40 ml 5× BRB80 (1× final)
 - 20 ml 1 M KCl (100 mM final)
 - 110 g sucrose (55% w/v final)
- Store up to 2 months at 4°C

Sucrose solution, 70% (w/v)

- 40 ml 5× BRB80 buffer (1× final)
 - 20 ml 1 M KCl (100 mM final)
 - 140 g sucrose (70% w/v final)
- H₂O to 200 ml
Heat without boiling to dissolve
Store up to 2 months at 4°C

Wash buffer

To 200 ml phosphate-buffered saline (PBS; APPENDIX 2A), add:

- 0.2 g bovine serum albumin (BSA, fraction V; 0.1% w/v final)
 - 100× protease inhibitor mix (see recipe) to 1× final
 - 20 mg/ml stock of DNase I, RNase free to final concentration of 20 µg/ml (stock stored at –20°C)
- Prepare fresh just prior to use

Yeast Paste

For 90 plates:

- 220 ml H₂O
- 150 g baker's yeast
- 1.4 ml propionic acid (Merck)

COMMENTARY

Background Information

Protocols for the isolation of centrosomes from higher eukaryotic cells are classically based on cell organelle enrichment via gradient centrifugation. Various successful protocols have been described that isolate centrosomes from mammalian tissue cul-

ture cells (Mitchison and Kirschner, 1986; Bornens et al., 1987; Blomberg-Wirschell and Doxsey, 1998; Bornens and Moudjou, 1999), tissue (Komesli et al., 1989; Lange and Gull, 1995), clam oocytes (Palazzo and Vogel, 1999), *Drosophila* (Moritz and Alberts, 1999; Lange et al., 2000), and yeast

(Wigge et al., 1998), to mention only some of the more frequently used sources. Most of the employed methods involve a lysis step in a low-ionic-strength buffer followed by density gradient centrifugation. The material thus produced is used subsequently in various assays. These include functional tests such as microtubule nucleation assay, electron microscopic study of centrosome morphology, antigen localization, or antigen production for the generation of monoclonal or polyclonal antibodies. Furthermore, centrosomal preparations have been used for the characterization of protein content of the organelle (protein complex); this application requires preparations in which the centrosomes are both highly enriched and present in biochemically meaningful quantities—i.e., micrograms of centrosomes for subsequent gel electrophoresis and mass spectrometry analysis (Wigge et al., 1998; Lange et al., 2000; Andersen et al., 2003).

Critical Parameters and Troubleshooting

Lysis and homogenization

Correct lysis and homogenization are critical for yield as well as for purity of centrosome preparations. Alternative methods using either tissue or mammalian tissue culture cells as starting material carry out the lysis in a low-ionic-strength buffer. Pretreatment of cells with anti-actin (e.g., cytochalasin B) and anti-microtubule drugs (e.g., nocodazole) has been employed in order to release the nuclear-centrosome interaction; this technique was initially proposed to efficiently dissociate the centrosome from the cytoskeleton and nuclei. However, cytochalasin treatment might not be an essential prerequisite for cytoskeletal dissociation (Bornens and Moudjou, 1999). Here, for *Drosophila* lysis and homogenization, buffer conditions are derived from the classical PIPES-based microtubule reassembly buffer (BRB80) but are modified (Moritz and Alberts, 1999) to include additional salt and sucrose, probably bringing them closer to isotonic conditions. An additional condition determining the outcome of the preparations is the efficient removal of the outer protective chorion layer of *Drosophila* embryos using washes with sodium hypochlorite solution. Incomplete removal can cause blockage or overheating of the tight-fitting Teflon pestle and render the material produced nearly useless.

Isolation of centrosomes by density gradient centrifugation

In the protocol developed by Moritz and Alberts (1999) for the isolation of *Drosophila* centrosomes, cell organelles are enriched in a single sucrose step gradient. The isolated centrosomes have been used in structural assays (electron microscopy) and functional assays (in vitro microtubule nucleation). Here centrifugation is carried out in two consecutive rounds, which represent concentration steps rather than purifications. These steps are then followed by affinity immunoisolation of centrosomes to remove co-migrating contaminants. Other groups employ, albeit for mammalian centrosome isolation, Ficoll gradients in a concentration step (Michison and Kirschner, 1986) or a rapid isolation step (Blomberg-Wirschell and Doxsey, 1998). This step can be followed by a purifying sucrose gradient centrifugation. The gradient is then fractionated and assayed by immunofluorescence microscopy using an anti- γ -tubulin antibody (Support Protocol 1) to select for the most enriched fractions for subsequent experiments. In the protocol in this unit, this assay is used to calculate the yield and determine the quality of isolated organelles. To perform this calculation, the number of centrosomes per relative area of the coverslip are counted, i.e., a given volume of the preparation is centrifuged down onto the coverslip, the count is performed, and the results extrapolated to the total volume of the preparation. A similar assay, though based on electron microscopy negative staining, can also be employed for rapidly checking the quality and structural integrity of centrosome preparations (Lange and Gull, 1996).

Purification of centrosomes by immunoaffinity purification

Because the avenue of density banding of centrosomes is blocked due to the high density of the organelle, the purity of most of the current centrosome isolation protocols is limited. Therefore, the authors of this unit developed a protocol for the affinity purification of centrosomes. The critical parameter here is the quality of the antibody that is used for the immunoprecipitation. Each antibody will need to be tested to determine if the antibody will or will not work under certain buffer or detergent concentrations and antibody concentrations, and protocols will have to be adapted accordingly. The initial protocol for affinity isolation

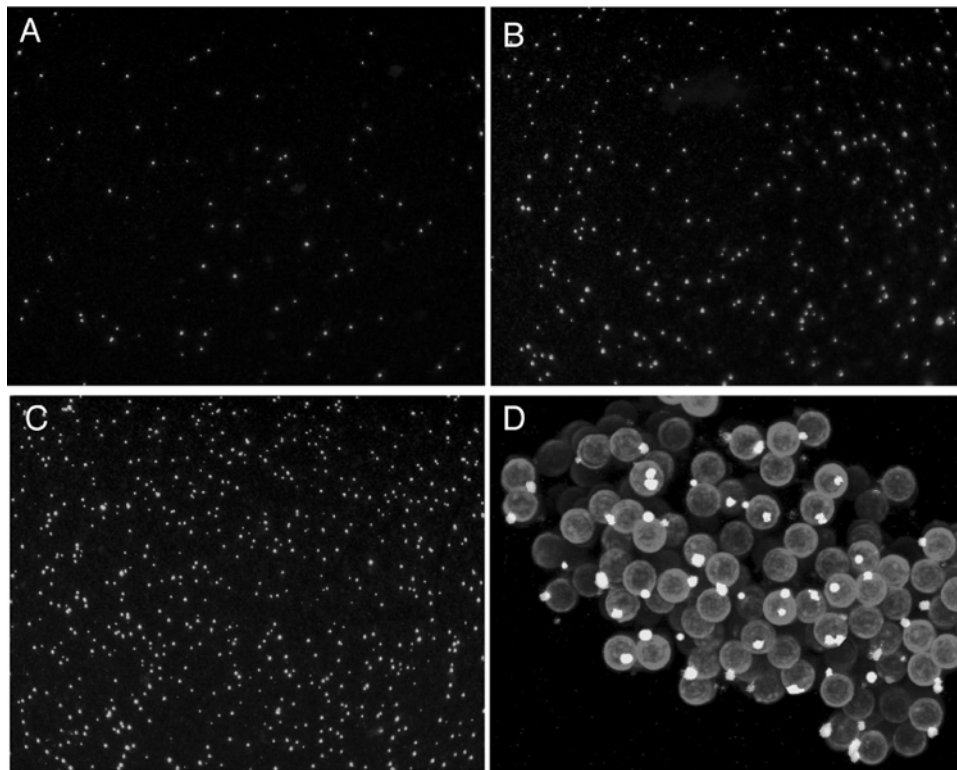


Figure 3.17.1 These images show the consecutive isolation steps assayed by immunofluorescence microscopy (Support Protocols 1 and 2) using an anti- γ -tubulin antibody to evaluate number and integrity of isolated centrosomes. **(A)** Centrosomes in embryo homogenate; **(B)** Pool 1; **(C)** Pool 2; **(D)** immunopurified centrosomes. Centrosomes are shown in yellow, beads are shown in red. For the color version of this figure go to <http://www.currentprotocols.com>.

of centrosomes was based on an indirect affinity method (Lange et al., 2000). In this technique, centrosomes were first coated with antibody, then anti-immunoglobulin beads were employed to isolate the centrosomes in a second step. In the protocol described in this unit, the authors use direct coupling of antibodies to magnetic beads to affinity purify centrosomes from enriched sucrose gradient fractions. This reduces the number of steps involved and consequently improves the yield.

Anticipated Results

With the described method microgram quantities of centrosomal protein can easily be obtained from ~ 10 g of embryo extract. Centrosomes should be intact as assayed by immunofluorescence microscopy (Support Protocols 1 and 2) and electron microscopy (Lange and Gull, 1996). Immunofluorescence images of centrosomes from consecutive isolations steps labeled with an anti- γ -tubulin antibody (see Fig. 3.17.1, panels A to D) are

quantified to control enrichment and yield. In the Coomassie-stained SDS-PAGE analysis (see Fig. 3.17.2) of the immunopurified preparations, a major band should be CNN, a centrosomal protein of about 130 kD (see Fig. 3.17.2, asterisk) indicating the level of enrichment of the preparations.

Time Considerations

In well maintained fly populations, large quantities of embryo material can be obtained almost on a daily basis. Collection of material and production of embryo extract can be streamlined so that extracts are stockpiled for future use. The actual preparation of centrosomes by density gradient centrifugation can be easily achieved within 4 hr, and samples can be safely frozen without losing microtubule nucleation activity. Affinity purification, including all of the washing and coupling steps, will consume ~ 4 hr, not taking into account buffer and antibody preparation time. Assays monitoring the quality of the

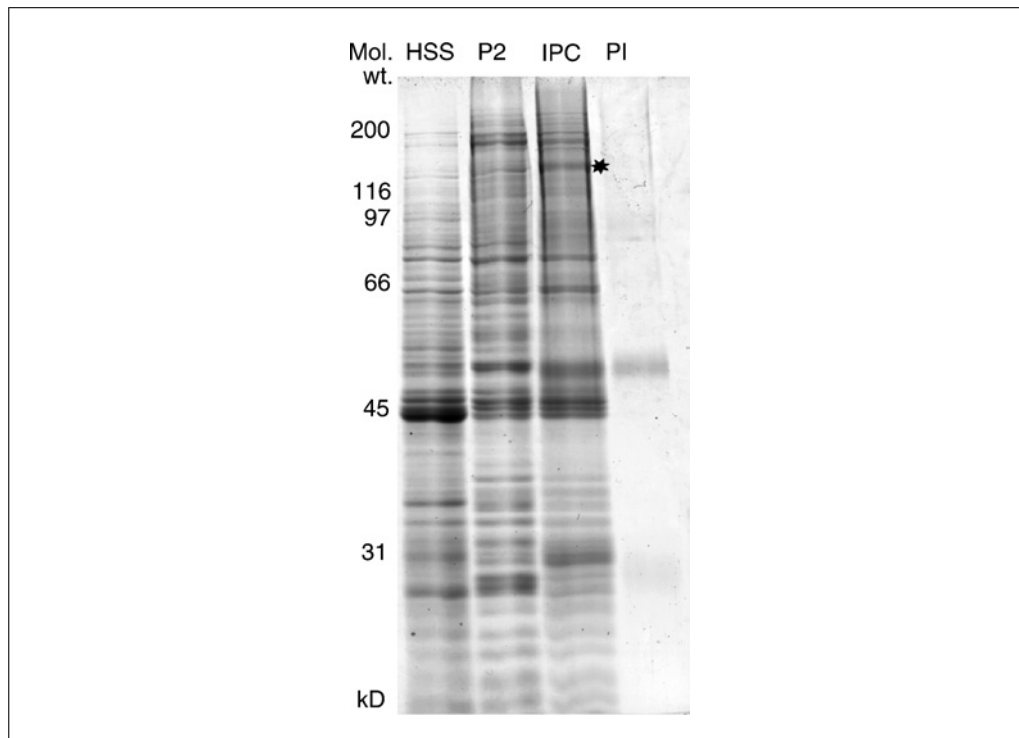


Figure 3.17.2 Isolation profile showing the protein composition of samples from the major consecutive steps in a Coomassie-stained gel. Abbreviations: HSS, homogenate supernatant supernatant; P2, Pool 2; IPC, immunopurified centrosomes; PI, preimmune isolation negative control. The asterisk at ~130 kDa labels the position of one of the major centrosome components (CNN) that is enriched throughout the isolation process.

preparation using immunofluorescence microscopy and SDS-PAGE are performed within 3 to 4 hr, not including microscope viewing time. Most solutions can be made ahead of time and stored either at 4°C or -20°C as indicated. Some solutions, such as proteinase inhibitors and DNase, must be added fresh from stock solutions to the solutions prior to use.

Literature Cited

- Andersen, J.S., Wilkinson, C.J., Mayor, T., Mortensen, P., Nigg, E.A., and Mann, M. 2003. Characterization of the human centrosome by protein correlation profiling. *Nature* 426:570-574.
- Blomberg-Wirschell, M. and Doxsey, S.J. 1998. Rapid isolation of centrosomes. *Methods Enzymol.* 298:228-238.
- Bonte, E. and Becker, P.B. 1999. Preparation of chromatin assembly extracts from preblastoderm *Drosophila* embryos. *Methods Mol. Biol.* 119:187-194.
- Bornens, M., Paintrand, M., Berges, J., Marty, M.C., and Karsenti, E. 1987. Structural and chemical characterization of isolated centrosomes. *Cell Motil. Cytoskel.* 8:238-249.
- Bornens, M. and Moudjou, M. 1999. Studying the composition and function of centrosomes in vertebrates. *Methods Cell Biol.* 61:13-34.
- Evans, L., Mitchison, T., and Kirschner, M. 1985. Influence of the centrosome on the structure of nucleated microtubules. *J. Cell Biol.* 100:1185-1191.
- Greenspan, R.J. 1997. *Fly Pushing: The Theory and Practice of Drosophila Genetics*. Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y.
- Komesli, S., Tournier, F., Paintrand, M., Margolis, R.L., Job, D., and Bornens, M. 1989. Mass isolation of calf thymus centrosomes: Identification of a specific configuration. *J. Cell Biol.* 109:2869-2878.
- Lange, B.M.H. and Gull, K. 1995. A molecular marker for centriole maturation in the mammalian cell cycle. *J. Cell Biol.* 130:919-927.
- Lange, B.M.H. and Gull, K. 1996. A structural study of isolated mammalian centrioles using negative staining electron microscopy. *J. Struct. Biol.* 117:222-226.
- Lange, B.M.H., Bachi, A., Wilm, M., and Gonzalez, C. 2000. Hsp90 is a core centrosomal component and is required at different stages of the centrosome cycle in *Drosophila* and vertebrates. *EMBO J.* 19:1252-1262.
- Mitchison, T.J. and Kirschner, M.W. 1986. Isolation of mammalian centrosomes. *Methods Enzymol.* 134:261-268.
- Moritz, M. and Alberts, B.M. 1999. Isolation of centrosomes from *Drosophila* embryos. *Methods Cell Biol.* 61:1-12.

Moritz, M., Braunfeld, M.B., Fung, J.C., Sedat, J.W., Alberts, B.M., and Agard, D.A. 1995. Three-dimensional structural characterization of centrosomes from early *Drosophila* embryos. *J. Cell Biol.* 130:1149-1159.

Palazzo, R.E. and Vogel, J.M. 1999. Isolation of centrosomes from *Spisula solidissima* oocytes. *Methods Cell Biol.* 61:35-56.

Sullivan, W., Ashburner, M., and Hawley, R.S. (eds.) 2000. *Drosophila* Protocols. Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y.

Tavosanis, G., Llamazares, S., Goulielmos, G., and Gonzalez, C. 1997. Essential role for gamma-tubulin in the acentriolar female meiotic spindle of *Drosophila*. *EMBO J.* 16:1809-1819.

Wigge, P.A., Jensen, O.N., Holmes, S., Soues, S., Mann, M., and Kilmartin, J.V. 1998. Analysis of the *Saccharomyces* spindle pole by matrix-assisted laser desorption/ionization (MALDI) mass spectrometry. *J. Cell Biol.* 141:967-977.

Contributed by Verena Lehmann,
Hannah Müller, and Bodo M.H. Lange
Max Planck Institute for Molecular
Genetics
Berlin, Germany