

1 **TITLE:**

2 Whole-cell cryo-electron tomography of cultured and primary eukaryotic cells on
3 micropatterned TEM grids
4

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26 **KEYWORDS:**

27 Cell culture, correlative light and electron microscopy (CLEM), cryo-electron microscopy (cryo-
28 EM), cryo-electron tomography (cryo-ET), fluorescence light microscopy (fLM), maskless
29 photopatterning, micropatterning, neurons, respiratory syncytial virus (RSV)
30

31 **ABSTRACT**

32 Whole-cell cryo-electron tomography (cryo-ET) is a powerful technique that can provide
33 nanometer-level resolution of biological structures within the cellular context and in a near-
34 native frozen-hydrated state. It remains a challenge to culture or adhere cells on TEM grids in a
35 manner that is suitable for tomography while preserving the physiological state of the cells.
36 Here, we demonstrate the versatility of micropatterning to direct and promote growth of both
37 cultured and primary eukaryotic cells on TEM grids. We show that micropatterning is
38 compatible with and can be used to enhance studies of host-pathogen interactions using
39 respiratory syncytial virus infected BEAS-2B cells as an example. We demonstrate the ability to
40 use whole-cell tomography of primary *Drosophila* neuronal cells to identify organelles and
41 cytoskeletal structures in cellular axons and the potential for micropatterning to dramatically
42 increase throughput for these studies. During micropatterning, cell growth is targeted by
43 depositing extra-cellular matrix (ECM) proteins within specified patterns and positions on the
44 foil of the TEM grid while the other areas remain coated with an anti-fouling layer. Flexibility in

45 the choice of surface coating and pattern design make micropatterning broadly applicable for a
46 wide range of cell types. Micropatterning is useful for studies of structures within individual
47 cells as well as more complex experimental systems such as host-pathogen interactions or
48 differentiated multi-cellular communities. Micropatterning may also be integrated into many
49 downstream whole-cell cryo-ET workflows including correlative light and electron microscopy
50 (cryo-CLEM) and focused-ion beam milling (FIB-SEM).

51

52 **INTRODUCTION:**

53 With the development, expansion, and versatility of cryo-electron microscopy (cryo-EM),
54 researchers have examined a wide-range of biological samples in a near-native state from
55 macromolecular (~1 nm) to high (~2 Å) resolution. Single-particle cryo-EM and electron
56 diffraction techniques are best applied to purified macromolecules in solution or in a crystalline
57 state, respectively ^{1,2}. Whereas, cryo-electron tomography (cryo-ET) is uniquely suited for near-
58 native structural and ultrastructural studies of large, heterologous objects such as bacteria,
59 pleomorphic viruses, and eukaryotic cells ³. In cryo-ET, three-dimensional (3D) information is
60 obtained by physically tilting the sample on the microscope stage and acquiring a series of
61 images through the sample at different angles. These images, or tilt-series, often cover a range
62 of +60/-60 degrees in one to three degree increments. The tilt-series can then be
63 computationally reconstructed into a 3D volume, also known as a tomogram ⁴.

64

65 All cryo-EM techniques require the sample to be embedded in a thin layer of amorphous, non-
66 crystalline, vitreous ice. One of the most commonly used cryo-fixation techniques is plunge
67 freezing, where the sample is applied to the EM grid, blotted, and rapidly plunged into liquid
68 ethane or a mixture of liquid ethane and propane. This technique is sufficient for the
69 vitrification of samples from <100 nm to ~10 µm in thickness including cultured human cells,
70 such as HeLa cells ^{5,6}. Thicker samples, such as mini-organoids or tissue biopsies, up to 200 µm
71 in thickness, can be vitrified by high-pressure freezing ⁷. However, due to increased electron
72 scattering of thicker samples, sample and ice thickness for cryo-ET is limited to ~0.5 – 1 µm in
73 300 kV transmission electron microscopes. Therefore, whole-cell cryo-ET of many eukaryotic
74 cells is limited to the cell periphery or extensions of cells unless additional sample preparation
75 steps are used such as cryo-sectioning ⁸ or focused-ion beam milling ⁹⁻¹¹.

76

77 A limitation of many whole-cell cryo-ET imaging experiments is data collection throughput ¹².
78 Unlike single particle cryo-EM, where thousands of isolated particles can often be imaged from
79 a single TEM grid square, cells are large, spread-out, and must be grown at low enough density
80 to allow for the cells to be preserved in a thin layer of vitreous ice. Often the region of interest
81 is limited to a particular feature or sub-area of the cell. Further limiting throughout is the
82 propensity of cells to grow on areas that are not amenable for TEM imaging, such as on or near
83 TEM grid bars. Due to unpredictable factors associated with cell culture on TEM grids,
84 technological developments are needed to improve sample accessibility and throughput for
85 data acquisition.

86

87 Substrate micropatterning with adherent extra-cellular matrix (ECM) proteins is a well-
88 established technique to direct the growth of cells on glass and other tissue culture substrates

89 ¹³. Such techniques have not only allowed for the precise positioning of cells, they have also
90 supported the creation of multicellular networks, such as patterned neural cell circuits ¹⁴.
91 Bringing micropatterning to cryo-ET will not only increase throughput, but it can also open up
92 new studies for exploring complex and dynamic cellular microenvironments.

93

94 Recently, we and others have begun using micropatterning techniques on TEM grids through
95 multiple approaches ¹⁵⁻¹⁷. Here, we describe the use of a maskless photopatterning technique
96 for TEM grids using the Alvéole PRIMO system. With the PRIMO process, an antifouling layer is
97 applied on top of the substrate, followed by application of a photocatalyst and ablation of the
98 antifouling layer in user-defined patterns with a UV laser. ECM proteins can then be added to
99 the patterns for the appropriate cell culture. This method has been used by several groups for
100 cryo-ET studies of RPE1, MDCKII, HFF, and endothelial cell lines ¹⁵⁻¹⁷. The PRIMO system is
101 compatible with multiple anti-fouling layer substrates as well as either a liquid or gel
102 photocatalyst reagent. A variety of ECM proteins can be selected from and adapted for the
103 specificity of the cell line, conferring versatility for the user.

104

105 We have successfully applied micropatterning to a number of projects within the lab. Here we
106 present results from our use of micropatterning for cryo-ET studies of cultured HeLa cells,
107 respiratory syncytial virus (RSV)-infected BEAS-2B cells, and primary larval *Drosophila*
108 *melanogaster* neurons ¹⁸. Significant findings include the identification of ECM proteins,
109 patterns, and other technological adaptations to allow for the micropatterning of the fragile
110 primary *Drosophila melanogaster* neurons. This is a valuable model system for a number of
111 reasons, including the ability to perform whole-cell tomography on neuronal extensions
112 without the need for downstream thinning techniques post-vitrification. We also show that
113 micropatterning can be applied to virus infected cells which remain competent for viral release
114 after targeted growth in micropatterned regions. Further, released virions remain in proximity
115 to infected cells in areas suitable for cryo-ET.

116

117 **RESULTS:**

118 This procedure was used to pattern EM grids for whole cell cryo-ET experiments. The entire
119 workflow presented in this study, including initial cell culture preparations, micropatterning (Fig
120 1 and Fig 2), and LM and cryo-EM imaging encompasses 3-7 days. In our protocol we use a two-
121 step procedure to generate the anti-fouling layer by applying PLL to the grid and subsequently
122 linking PEG by addition of the reactive PEG-SVA. The anti-fouling layer can also be applied in a
123 single step by adding PLL-g-PEG in one incubation. We used the PLPP gel as a catalyst for the UV
124 micropatterning, the catalyst is also available as a less concentrated liquid. The gel allows for
125 patterning at a significantly reduced dose compared to the liquid, which results in much faster
126 patterning. With our system, the actual patterning time of a full TEM grid was ~2 minutes. The
127 micropatterning workflow alone generally spans five to six hours and allows an individual to
128 pattern eight grids for standard cell-culture on TEM grids.

129

130 A number of the steps during the micropatterning process require long incubation times.
131 Conveniently, some of these steps, such as PLL passivation or PEG-SVA passivation may be
132 extended to an overnight incubation. Additionally, grids may be patterned in advance and

133 stored in a solution of the ECM protein or PBS for later use. In our study, these options were
134 valuable in instances where the timing of cell preparation and seeding is critical such as for
135 primary *Drosophila* neurons and RSV-infection of BEAS-2B cells.

136

137 We prepare grids in a general biosafety-level 2 (BSL-2) lab setting using clean tools, sterile
138 solutions, and include antibiotics/antimycotics in the growth media^{6,19-21}. For samples
139 particularly sensitive to microbial contamination, the anti-fouling layer and ECM can be applied
140 in a tissue culture hood or other sterile environment. Additionally, the grid could be washed in
141 ethanol between patterning and ECM application. If working with infectious agents, it is
142 important to adapt the procedure to comply with appropriate biosafety protocols.

143

144 This workflow and the procedures presented (Fig 1 and Fig 2) allowed HeLa cells (Fig 3) and
145 RSV-infected BEAS-2B cells (Figs 4 & 5), and primary *Drosophila* larval neurons (Figs 6 & 7) to be
146 seeded onto patterned EM grids to control for cell density and spatial positioning for optimal
147 cryo-ET data collection.

148

149 **Cultured HeLa cells adhere to and spread out over patterns**

150 We show that HeLa cells seeded onto micropatterned TEM grids remain viable as determined
151 by fluorescent staining using a calcein-AM and ethidium homodimer-1 based cell viability assay
152 (Fig 3A & 3B). Using a mixed collagen and fibrinogen ECM, HeLa cells readily adhere to patterns
153 across the grid (Fig 3A & 3C). The overall morphology of cells that expand along the pattern is
154 similar to that of cells on grown on unpatterned grids (Fig 3C & 3D). In the case of HeLa cells,
155 the total cell thickness remains $\sim < 10 \mu\text{m}$ with significantly thinner areas $\sim < 1 \mu\text{m}$ thick near the
156 cell periphery (Fig 3C).

157

158 **Virus infected cells on micropatterned areas remain competent for viral release**

159 For our RSV studies, we patterned entire grid squares using a gradient, with a low-dose
160 exposure on the edges and a higher dose pattern towards the center (Fig 4A). Gradient patterns
161 yielded better results when searching for released viruses present near the periphery of cells.
162 With these patterns, we find that cells preferentially adhere to the higher ECM concentration,
163 but are also able to adhere to and grow on the lower ECM concentrations. The relative dose
164 between areas will need to be optimized when using patterns that require multiple doses. If the
165 doses and thus ECM concentrations are too similar or too disparate to one another, the effect
166 of using multiple doses will be lost.

167

168 In Fig 4 we show a TEM grid that has been patterned and subsequently seeded with RSV
169 infected BEAS-2B cells and used for cryo-EM data collection. Fig 4A is a fluorescent image of
170 ECM patterned onto a TEM grid using a gradient pattern. Cell adhesion and growth along the
171 central region of the pattern can be seen in Fig 4B, a brightfield image of the cells 18 hours
172 post-seeding. In Fig 4C, fluorescent signal (red) from replication of RSV-A2mK+ is overlaid with
173 signal from the ECM. The majority of the infected cells are positioned along the higher density
174 central region of the gradient pattern. A low-mag TEM map of the grid post cryo-fixation
175 reveals a number of cells, including RSV-infected cells, positioned on the carbon foil near the
176 center of the grid squares. As previously shown for cells grown on standard TEM grids²⁰, we are

177 able to locate and collect tilt-series of RSV virions in close proximity to the periphery of infected
178 BEAS-2B cells grown on micropatterned grids (Fig 5A & 5B). Many of the RSV structural proteins
179 can be identified within the tomograms including nucleocapsid (N) and the viral fusion protein
180 (F) (Fig 5C).

181

182 **Micropatterning allows for optimized distribution and positioning of *Drosophila* neurons on** 183 **TEM grids**

184 For our primary *Drosophila* neuron studies, we found that the narrow pattern, near the
185 resolution limit offered by PRIMO (where the thickness of the pattern was 2 μm), allowed from
186 one to a few cells to be isolated within a grid square (Fig 6). The neuronal soma was able to
187 extend its neurites over a period of several days within the pattern. This allowed easy
188 identification and tilt series acquisition of the neurites compared to neurons cultured on
189 unpatterned grids (Fig 7). We also found that fluorescently-labeled concanavalin A, a lectin that
190 has been used as an ECM for *in vitro Drosophila* neuronal cultures^{18,22}, is amenable for PRIMO
191 patterning.

192

193 *Drosophila* neurons from third instar larvae were isolated according to previously published
194 protocols^{18,22,23}. The neuronal preparations were applied to micropatterned cryo-EM grids
195 where concanavalin A was deposited on the pattern to regulate cell placement, spreading, and
196 organization. The neurons on patterned or unpatterned grids were allowed to incubate for 72-
197 96 hours and the grids were then plunge frozen. A representative image of a micropatterned
198 EM grid with several *Drosophila* neurons distributed across the patterned regions is shown in
199 Fig 6A. These neurons, derived from a transgenic fly strain that has pan-neuronal GFP
200 expression in the membrane, can be easily tracked by light microscopy not only due to its
201 fluorescent labeling, but also because of its location within the micropatterns. While neurons
202 cultured on unpatterned grids can also be tracked through its GFP signaling by light microscopy
203 (Fig 7A, yellow circle), locating them in cryo-EM became substantially more difficult due to the
204 presence of cellular debris and contamination from the media (Fig 7B, yellow circle). Such
205 presence was lessened for neurons on patterned grids, likely due to the pattern being narrow
206 enough to allow neurons to attach to the grid while excluding undesired contaminants. Due to
207 the dimensions of the neuron cell body and the extended neurites (Fig 6A & 6B, yellow circle),
208 cryo-ET tilt series were collected along thinner regions of the cells (Fig 6C & 6D, red circle). The
209 neuronal cell membrane, a mitochondrion, microtubules, actin filaments, and vesicular
210 structures were well resolved in higher-magnification image montages and slices through the
211 3D tomogram (Fig 6E). While similar sub-cellular features can be seen from 3D tomograms of
212 unpatterned neurons (Fig 7E), the difficulty in locating viable cellular targets for data collection
213 decreased throughput substantially.

214

215 When first starting with micropatterning, there are a few potential pitfalls that are detrimental
216 to the final result. We have found that careful grid handling and sterile technique, a uniform
217 distribution of the PLPP gel, proper dose and focus during patterning, and maintenance of cell
218 viability prior to seeding are among the most important considerations for success. We have
219 assembled a list of some of the potential issues as well as solutions in Table 1. In Fig 8 we've
220 assembled representative images from grids with some of these issues to assist in their

221 identification and troubleshooting. Once optimal conditions are determined, micropatterning
222 with PRIMO is a reliable and reproducible method for the positioning of cells on grids for cryo-
223 TEM.

224

225 **DISCUSSION:**

226 Substrate micropatterning is a well-established technique for live-cell light microscopy^{13,24}
227 where investigators benefit from the ability to use rigid, durable, and optically transparent
228 surfaces such as glass coverslips. Micropatterning has also been done on soft and three-
229 dimensional surfaces. Here we present our application of micropatterning to extend this
230 technique for cryo-EM studies of multiple cell types by utilizing features such as high resolution
231 and contactless patterning of the PRIMO system to pattern TEM grids.

232

233 Modern, advanced electron microscopes and software packages now support streamlined
234 automated cryo-EM and cryo-ET data collection where hundreds to thousands of positions can
235 be targeted and imaged within a few days²⁵⁻²⁸. One significant limiting factor for whole-cell
236 cryo-ET workflows has been obtaining sufficient numbers of collectable targets per grid.

237 Recently, a number of groups have developed protocols for micropatterning grids for cryo-EM,
238 with one advantage being improved data collection efficiency¹⁵⁻¹⁷. Here we use
239 micropatterning to optimize cryo-ET studies of primary *Drosophila* neurons and cultured human
240 cell lines (uninfected or RSV-infected). The PRIMO system is versatile and many steps can be
241 optimized and tailored to fit specific experimental goals. A user with TEM and fluorescent
242 microscopy experience can quickly become skilled in grid preparation and micropatterning.
243 With careful practice, good results should be achievable after a few iterations. Below, we
244 discuss some of the options available, user considerations, potential benefits, and future
245 applications of micropatterning for cryo-EM.

246

247 One of the important considerations for whole cell cryo-ET is EM grid selection. EM grids are
248 composed of two parts: a mesh frame (or structural support) and the foil (or film), which is the
249 continuous or holey film surface on which cells will grow. Copper mesh grids are commonly
250 used for cryo-EM of proteins and isolated complexes. However, they are unsuitable for whole-
251 cell cryo-ET due to the cytotoxicity of copper. Instead, gold mesh is commonly used for cellular
252 tomography. Other options include nickel or titanium, which may provide benefits over gold
253 such as increased rigidity¹⁵. EM grids are available with different mesh dimensions to support a
254 range of applications. Larger mesh sizes provide more room for cells to grow between grid bars
255 and more areas that are amenable for tilt series collection, though at the cost of increased
256 overall specimen fragility. The most commonly used foil is perforated or holey amorphous
257 carbon, such as Quantifoils or C-flat grids. Biological targets can be imaged either through the
258 holes in the carbon or through the electron-translucent carbon. Grids such as R 2/1 or R 2/2,
259 where the holes are 2 μm wide that are spaced 1 and 2 μm apart respectively, provide a large
260 number of holes and thus a large number of potential areas for data collection. However, some
261 cells may grow and expand better on more uniform surfaces such as R 1.2/20 grids or
262 continuous carbon. For downstream sample processing by FIB-SEM, the foil is removed through
263 milling, reducing concerns over the continued presence of the underlying film. As with the
264 mesh, foils from other materials are also available, with the patterning protocol presented here

265 being equally suitable for SiO₂ grids. We commonly use gold Quantifoil, continuous carbon, or
266 SiO₂ film 200-mesh grids (~90 μm spacing between grids bars) for whole-cell cryo-ET.

267

268 There are a number of considerations when designing a pattern. A majority of these decisions
269 are guided by the cell type and purpose of experiment. A good starting point is to choose a
270 pattern that approximates the shape and dimensions of the cells in culture. Many studies have
271 demonstrated significant effects of pattern shape on cell growth and cytoskeletal arrangement
272 ^{13,29,30}. Special care should be taken during pattern design if this could alter the target of
273 interest. We tested several patterns for each cell type to determine which patterns promoted
274 cellular adhesion and growth. The flexibility of the PRIMO system permits testing of multiple
275 patterns on a single grid and changing patterns for different grids within a single experiment.
276 Larger patterns (~50-90 μm), such as those used here, increase the likelihood that multiple cells
277 adhere to a single region of the pattern and allow cells to expand and extend after adhesion.
278 More constrained patterns (20-30 μm) may be appropriate in experiments where cell isolation
279 is more critical than cell expansion, such as for FIB-SEM experiments. For tomography
280 applications, one may need to consider the impact of the tilt-axis. If a pattern is positioned such
281 that all cells grow parallel to one another in a single direction, it is possible that all of the cells
282 will be perpendicular to the tilt-axis when loaded onto the microscope stage, resulting in a
283 lower quality of data.

284

285 On unpatterned grids, cells often preferentially adhere to the grid bars, where they cannot be
286 imaged by TEM. Even on patterned grids, we often observe cells which are positioned in the
287 corners of grid squares partially on both the patterned carbon foil and grid bar. Recently,
288 micropatterning was used to intentionally position part of the cell over the grid bar ¹⁷. This
289 could be considered for experiments where it is not critical to have the entire cell periphery on
290 the foil. This can be especially important for cells that can grow larger than a single grid square,
291 such as primary neurons growing over multiple days.

292

293 There are many tools that can be used to design a pattern. Here, we limit the pattern to less
294 than 800 pixels in any dimension such that the pattern can be rotated to any angle and still fit
295 within the maximum area that can be patterned in a single projection by the PRIMO system. This
296 allows the user to rotate the pattern to be properly oriented with the grid regardless of the
297 orientation of the grid on the microscope. In our experiments, we divide the grid into six
298 patterning areas. Primarily, this allows us to adjust the focus between different regions of the
299 grid. Gold grids in particular are very malleable and may not laydown completely flat on the
300 glass. Proper focus is essential for clean, refined patterning results. By using segmented
301 patterns, we are able to make minor adjustments to the pattern position if the grid shifts
302 slightly during patterning process, this is usually not an issue when using the PLPP gel and PDMS
303 stencils. Finally, it allows us to keep the center of the grid unpatterned. Being able to clearly
304 identify the center of the grid is very useful for correlative-imaging experiments.

305

306 The PRIMO patterning software, Leonardo, also has more advanced features such as stitching
307 and the ability to import patterns as PDFs which are not described here. Leonardo also includes
308 microstructure detection and automated pattern positioning that can be used on TEM grids.

309 This feature is most useful when the grid is very flat and can be patterned without the need to
310 adjust focus between different areas

311

312 Selection of an ECM protein can have a significant impact on cell adhesion and expansion. Some
313 cells are known to undergo physiological changes when grown on specific substrates³¹. We
314 tested multiple ECM proteins and concentrations for any new cell type based on prior work
315 reported in the literature. Laminin, fibrinogen, fibronectin, and collagen are widely used for
316 cultured cells and can be used as a starting point if other data is not available. However, other
317 ECM proteins must also be considered if the commonly used ECM proteins fail to confer proper
318 adherence properties for the cells. This was particularly true for primary *Drosophila* neurons, as
319 a high-concentration of the plant lectin concanavalin A was necessary for proper cellular
320 adherence. The compatibility of cellular adhesion and growth with the ECM can be tested by
321 patterning on glass dishes or slides prior to transitioning to TEM grids. This pre-screening
322 approach is time and cost effective if a large number of combinations need to be examined. The
323 inclusion of a fluorescently conjugated ECM protein is valuable for assessing the success and
324 quality of patterning.

325

326 Cell seeding is one of the most important steps for whole cell cryo-ET, either with or without
327 micropatterning^{6,15,32}. For primary *Drosophila* or other neurons, which are fragile, unstable in
328 suspension, and may be limited in quantity, single seeding approaches are preferred over
329 monitored, sequential cell seeding. A single seeding step at an optimized cell density, as
330 described in the methods for *Drosophila* neurons, is a viable option for most cell types.
331 However, it is also possible to seed cells onto the substrate at a lower initial concentration and
332 the add more cells in a monitored fashion as described here and in other literature¹⁷. This
333 sequential seeding can provide more consistent results in some cases. Similar to standard cell
334 culture, care should always be taken to maintain cell viability and minimize cell clumping during
335 isolation.

336

337 Micropatterned grids can be used to help position cells to establish a consistent cell density
338 across the grid and to position regions of interest in areas suitable for tilt-series collection. The
339 placement and positioning of cells can be used as fiducial markers for correlation in cryo-CLEM
340 experiments, reducing the need for fragile finder-grids and fluorescent fiducial markers.
341 However, it should be noted that such fiducial markers may still be useful for sub-micron
342 accuracy correlation^{19,33}. Furthermore, an even distribution of isolated cells is also highly
343 beneficial for FIB-SEM experiments to maximize the number of cells from which lamella can be
344 cut¹⁵.

345

346 The addition of micropatterning to cryo-EM workflows will result in measurable improvements
347 in data throughput and potentially enable new experiments. As the technique is further
348 adopted and developed, more advanced applications of micropatterning including ECM
349 gradients, multiple ECM depositions, and microstructure assembly will further expand the
350 capabilities of cryo-ET to study biological targets and processes in full cellular context.

351

352 **MATERIALS AND METHODS:**

353 The methods and materials (supplementary) described here is a compilation of the cell culture,
354 micropatterning, and imaging methods used by the Wright lab and the Cryo-EM Research
355 Center at the University of Wisconsin, Madison. Additional training and instructional materials
356 are available at the following site: <https://cryoem.wisc.edu>

357

358 **Preparation of grids for patterning**

359 Prior to micropatterning 5-8 nm of carbon was evaporated onto the grids in a Leica ACE600
360 carbon evaporator. Carbon coated grids were stored in a low humidity environment such as a
361 vacuum desiccator. Within 15-30 minutes of use the grids are placed on either a grid prep
362 holder (see materials) or a piece of filter paper on a small petri dish and glow discharged for 60
363 s at 10 mA with an 80 mm working distance and vacuum pressure of $1.0e^{-3}$ mbar using a Leica
364 ACE600.

365

366 **Application of the anti-fouling layer**

367 Using proper sterile technique the grids were transferred to a clean glass slide or coverslip with
368 at least 1 cm of separation between the grids. The grids were incubated in 10 μ L of sterile 0.05
369 % Poly-L-lysine (PLL) for 30 minutes to overnight in a humid chamber, such as an enclosed
370 plastic box with moist paper towels. Each grid was washed three times with 15 μ L of 0.1 M
371 HEPES pH 8.5. For each wash, most of the liquid is removed from the grid with a pipet without
372 letting the grid dry. The grids are then incubated in 15 μ L fresh HEPES pH 8.5 for at least 30
373 seconds and left in the final wash.

374

375 10 μ L of 100 mg/mL Poly-ethylene glycol-succinimidyl valerate (PEG-SVA) in 0.1 M HEPES pH 8.5
376 was prepared for each grid immediately prior to use. PEG-SVA has a half-life of 10 minutes at
377 pH 8.5. It is important to avoid exposing the PEG-SVA stock to excessive moisture by storing in a
378 desiccator or dry environment and warming to room temperature before opening. PEG-SVA
379 dissolves quickly with gentle mixing resulting in a clear solution. The 15 μ L drop of HEPES pH 8.5
380 was removed from each grid followed by incubation in a 10 μ L drop of the PEG-SVA solution.
381 The grids are kept in a humid chamber to prevent drying during this incubation for one hour to
382 overnight. Following PEG-SVA coating, each grid is washed three times with 15 μ L sterile water.
383 For each wash. The grids are then stored in 15 μ L water in a humid chamber until the next step.

384

385 **Applying PLPP gel**

386 A clean microscope coverslip was prepared for each grid. A 1.0 μ L drop of water was placed in
387 the center of a new coverslip for each grid to assist in placing the grid on the coverslip and
388 keeping the grid wet. The grids are then carefully transferred from the 15 μ L water drop to the
389 1.0 μ L drop on the coverslip carbon side up. We then carefully place a PDMS stencil over the
390 grid, taking care to keep the grid centered and to minimize stencil contact with the carbon foil
391 of the grid. Next, we pipet 1.0 μ L of PLPP gel onto each grid and mix gently by pipetting. Finally,
392 the gel is allowed to dry in a dark environment for 15-30 minutes.

393

394 **Calibration and design of the micropattern**

395 The PRIMO system was calibrated using a glass coverslip with highlighter on one side. The slide
396 was placed on the microscope, highlighted sides down, and brought into focus. After setting up

397 the appropriate light path and starting up the PRIMO system, we followed the on screen
398 instructions to calibrate the system. During this calibration the UV laser is used to project an
399 image onto the slide which must be brought into focus before proceeding. Following
400 calibration the micrometer/pixel ($\mu\text{m}/\text{px}$) ratio is reported under Calibration data in the top left
401 window of Leonardo (Fig 2, area 1). This ratio is needed to determine the number of pixels to
402 use per micrometer when designing a pattern.

403
404 In order to measure the grid to determine pattern size, we loaded a prepared grid on a
405 coverslip (from above) onto the stage with the grid facing the objective lens. We then adjust
406 stage position and focus so that the grid is visible in the Leonardo software window. We used
407 the ruler function built-in to Leonardo activated by the button near the bottom left corner to
408 measure the grids (Fig 2, area 2). For example, the patterns used here for a 200 mesh grid were
409 measured to have $\sim 87 \times 87 \mu\text{m}$ grid squares and $\sim 36 \mu\text{m}$ grid bars. The Leonardo software
410 offers flexibility in resizing patterns on-the-fly, so minor inaccuracies in measurement can be
411 tolerated.

412
413 We then designed the patterns used in Figures 3-8 based on the measurements and ratios
414 reported in the steps above. Patterns can be designed in any image creation software. The
415 minimum feature size with a 20 \times objective is 1.2 μm . Patterns should be saved as
416 uncompressed 8-bit .tiff files. Be sure the software does not rescale images to a different pixel
417 size when saving. The pattern should fit within an 800 \times 800 pixel box, which is sufficient to
418 cover four grid squares. Pixels with a value of 255 (white) will be patterned at the highest
419 intensity (total dose of the laser) and pixels with a value of zero (black) will not be patterned.
420 Any pixels with an intermediate value will be patterned with a dose of approximately
421 $(X/255) \times \text{total dose}$. In Fig 4A, pixel values of 255 and 129 were used for the greyscale patterns.
422 Once the pattern is designed it can be saved and reused without modification.

423

424 **Micropatterning**

425 For our initial run we created a new 3000 μm ROI In Leonardo using the Add ROI function (not
426 shown, in the location of Fig 2 area 3). For subsequent runs we reloaded and modified this
427 initial template. We used the Add Pattern function (not shown, in the location of Fig 2 area 3)
428 to place six patterns on each grid which allow for independent focusing and positioning in each
429 region. An 8 \times 8 grid square region for each corner of the grid and a 2 \times 8 grid square region on
430 each side of the center, leaving the center four grid squares unpatterned (Fig 2, center image).
431 The replication options (Fig 2, area 4) were used to generate copies of the initial pattern to
432 reach the desired number of total copies of the pattern.

433

434 The angle, position, space between, and ratio (size) of the patterns were iteratively adjusted
435 using the exper options panel (Fig 2, area 4) until the patterns aligned with the grid squares.
436 Total dose was set to 30 or 45 mm/mJ^2 (see discussion) with 100% laser power for each
437 pattern. The template file was saved within Leonardo for use in future experiments (Fig 2, area
438 6, bar with up arrow icon in top toolbar).

439

440 Each of the six regions on the grid were patterned one at a time by selecting only a single
441 pattern in Action panel (Fig 2, area 5). Prior to patterning a region we navigate to that region
442 and focus on the carbon foil. Focusing on the area to be patterned is an essential step. After
443 patterning we remove the coverslip from the microscope and immediately pipet 10 μ L of sterile
444 PBS onto the grid. After 10 minutes the PDMS stencil is removed the grid is washed 3 \times with 15
445 μ L PBS and stored in PBS until the next step.

446 447 **Deposition of ECM proteins for cultured cells**

448 Each grid was incubated for one hour at room temperature or overnight at 4°C in 15 μ L of
449 freshly prepared ECM solution. For BEAS-2B cells a final concentration of 0.01 mg/mL bovine
450 fibronectin and 0.01 mg/mL fluorophore-conjugated fibrinogen in sterile PBS was used. For
451 HeLa cells we used 0.01 mg/mL bovine collagen I and 0.1 mg/mL fluorophore-conjugated
452 fibrinogen in sterile PBS. After incubation in ECM the grids are washed 5x with sterile PBS and
453 stored in PBS at 4°C. We have stored grids for up to a week in PBS at 4 °C with no observed
454 deterioration in quality.

455 456 **Deposition of ECM proteins for *Drosophila* neurons**

457 For primary *Drosophila* neurons, the patterned grids were first moved to a 30 mm glass bottom
458 dish containing sterile PBS. The PBS was then aspirated from the dish and replaced with 2 mL
459 of 0.5 mg/mL fluorescently conjugated concanavalin A. The grids were incubated in this
460 solution overnight at 25°C before 3x washes in 2 mL PBS. After the final wash the grids are
461 incubated at 25°C in 2 mL of freshly-prepared, sterile-filtered supplemented Schneider's
462 *Drosophila* media ²², containing 20% heat-inactivated FBS, 5 μ g/mL insulin, 100 μ g/mL penicillin,
463 100 μ g/mL streptomycin, and 10 μ g/mL tetracycline until the neurons are ready to be plated.

464 465 **Preparation of primary *Drosophila* cells prior to seeding**

466 All dissection dishes were sterilized with 70 % EtOH, then submerge the plate with 2-3 mL of
467 sterile-filtered 1 \times dissection saline (9.9 mM HEPES pH 7.5, 137 mM NaCl, 5.4 mM KCl, 0.17 mM
468 NaH₂PO₄, 0.22 KH₂PO₄, 3.3 mM glucose, 43.8 mM sucrose) ²².

469
470 Thirty to forty 3rd instar larvae were gently removed from the food using a pair of tweezers
471 and placed into a tube of PBS and transferred to a fresh tube of PBS. The larvae were then
472 transferred into a tube of 70% EtOH and a second fresh tube of 70% EtOH for 2-3 minutes to
473 sterilize the larvae. A final rinse was done in dissection saline by transferring the larvae through
474 two tubes of dissection saline. The larvae were then transferred to a dissecting dish containing
475 1 \times dissection saline. The brains were extracted with a pair of forceps and a dissection
476 microscope and transferred to a third tube with 1 \times dissection saline. The brains are washed
477 three times by centrifugation at 300 x g for 1 minute followed by discarding and replacing the
478 supernatant with 1 mL of fresh 1 \times dissection saline, leaving 200-250 μ L after the final wash.

479
480 To digest the tissue we added 20 μ L of 2.5 mg/mL Liberase in 1 x dissection saline to the
481 remaining 200-250 μ L volume and rotated the tubes for one hour at room temperature. The
482 solution was mixed by pipetting 25-30 times every ten minutes during this step. The solution
483 was centrifuged for 5 minutes at 300 x g and supernatant was replaced with 1 mL of

484 supplemented Schneider's *Drosophila* media. This was repeated 3 x, leaving 300 μ L of volume
485 after the final step. The cells were pipetted 30-40 times to mix.

486

487 **Culture and RSV infection of BEAS-2B and HeLa cells**

488 HeLa cells and BEAS-2B cells are maintained in T75 flasks at 37 °C and 5 % CO₂. Cells are
489 passaged every 3-4 days once reaching approximately 80 % confluency. HeLa cells are
490 maintained in DMEM + 10 % FBS + 1 \times Antibiotic-Antimycotic. BEAS-2B are maintained in RPMI
491 + 10 % FBS + 1 \times Antibiotic-Antimycotic^{6,20,34}. Prior to RSV infection 5 \times 10⁴ cells per well were
492 passaged into a 6-well plate (surface area \sim 9.6 cm²) with 2mL of growth media and incubated
493 overnight. The next day one well of cells was trypsinized and used for cell counting. Media was
494 aspirated from the well and washed with 2 mL sterile PBS without Mg²⁺ and Ca²⁺ to remove
495 residual media. The well was then incubated in 500 μ L 0.25 % trypsin solution at 37 °C for 5-10
496 min. Once the cells were released they were diluted with 1.5 mL culture media. The cells were
497 counted using a hemacytometer and trypan blue staining.

498

499 Media was then aspirated from the remaining wells and replaced with 750 μ L of media with
500 RSV-A2mK+³⁵ at a concentration calculated to achieve MOI 10. The MOI of RSV-A2mK+ can be
501 calculated from fluorescent focus units (FFU) titers of the stock (For example: for 1.0 \times 10⁵ cells
502 per well and an RSV stock of 1.0 \times 10⁸ FFU/mL, dilute the viral stock 1:75 to 1 \times 10⁶ FFU/750 μ L or
503 1.33 \times 10⁶ FFU/mL). The plate was incubated with rocking at room temperature for one hour.
504 After one hour the total volume per well was brought to 2 mL with growth media pre-warmed
505 to 37°C and place the plate in an incubator set to 37 °C with 5 % CO₂ for 6 hours. The cells were
506 trypsinized for seeding as described below. After seeding the grids were incubated for an
507 additional 18 hours before plunge freezing (for a total 24 hours post-infection).

508

509 **Cell seeding of cultured cells onto micropatterned grids**

510 The cells from each well were released with trypsin at a confluency of 60% or less to avoid
511 aggregation. The cells were counted using trypan blue staining and a hemacytometer and
512 diluted to 2 \times 10⁴ cells/mL in prewarmed media.

513

514 One μ L of media was placed in the center of a glass bottom dish and a micropatterned grid was
515 transferred to the dish. Ten μ L of cell solution was added to each grid and incubated at 37°C for
516 five minutes. Every five minutes the grids were checked with a brightfield microscope and an
517 additional 10 μ L of cells were added until all patterns were occupied or many patterns had
518 multiple cells. The grids were incubated for 2 hours (37 °C, 5 % CO₂) after the final addition of
519 cells before the addition of 2 mL of growth media and overnight incubation.

520

521 **Cell seeding of primary *Drosophila* neurons onto micropatterned grids**

522 Media was removed from the dish containing the micropatterned grids and the cells were
523 plated onto the grids. After a 30-60 minute incubation at 25°C to allow for cell attachment, the
524 dish was flooded with 2 mL of supplemented Schneider's *Drosophila* media. The neurons were
525 cultured for a minimum of 2-3 days in a 25 °C incubator before plunge-freezing.

526

527 **LM imaging, vitrification, and cryo-EM imaging of patterned grids**

528 All grids were checked before and after cell seeding using a Leica DMI8 with a 20X objective for
529 brightfield and fluorescent imaging. This ensures the grid quality is suitable for cryo-
530 preservation and data collection. Images were processed in the FIJI software package ³⁶.

531

532 Primary *Drosophila* neurons were prepared on a Leica EM-GP and BEAS-2B cells were prepared
533 using the Gatan CP3. Gold fiducials were applied to all samples to allow for proper alignment of
534 tilt series. Primary *Drosophila* neurons were blotted for 4 s from the backside. For HeLa and
535 BEAS-2B cells we used double sided blotting for 4-6 s. The frozen grids can be stored in liquid
536 nitrogen until further use.

537

538 Cryo-EM data was collected on a Titan Krios set at 300 kV with a direct electron detector
539 camera. Tilt-series were collected for each region of interest using SerialEM ³⁷. Tilt-series of
540 primary *Drosophila* neurons were collected from -60° to 60° bidirectionally at 2 ° increments
541 using a Falcon3 detector at -8 μm defocus with a pixel size of 4.628 Å for a total dose of 70-75 e⁻
542 /Å². Tilt-series of RSV-infected BEAS-2B were collected on a Gatan K3 with a BioQuantum
543 energy filter (20 eV slit) at -5 μm defocus with a pixel size of 4.603 Å and total dose of ~80 e⁻/Å².
544 The tilt series were aligned and reconstructed into tomograms using IMOD package ³⁸; lowpass
545 filtering was done using the EMAN2 software package ³⁹.

546 **FIGURE AND TABLE LEGENDS:**

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548 **Prepare grids**

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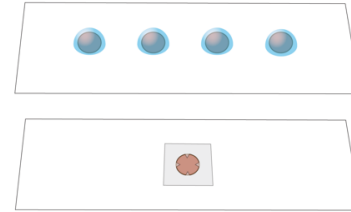
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Approx.
3 hours

Carbon coat and glow discharge grids
Incubate grid in PLL and wash
Incubate grid PEG-SVA and wash
Add PDMS stencil over grid on coverslip
Add PLPP gel and dry



558 **Micropatterning**

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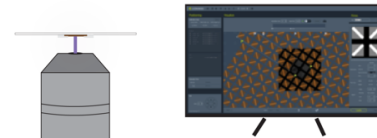
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Approx.
20 min/grid

Load grid onto microscope
Load/design pattern template
Position pattern on grid
Pattern grid
Wash grid in PBS



568 **ECM and cell seeding**

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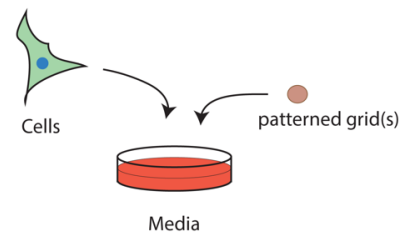
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2-3 hours

Incubate grid in ECM and wash
Prepare cells
Add cells to grid
Add media and grow in incubator



577 **Cryo-preparation and data-collection**

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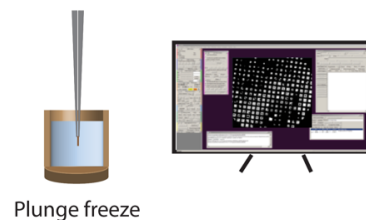
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Time varies
significantly
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workflow

Screen grids by light microscopy
Plunge-freeze grids
Cryo-CLEM or FIB-SEM (optional)
Cryo-electron tomography

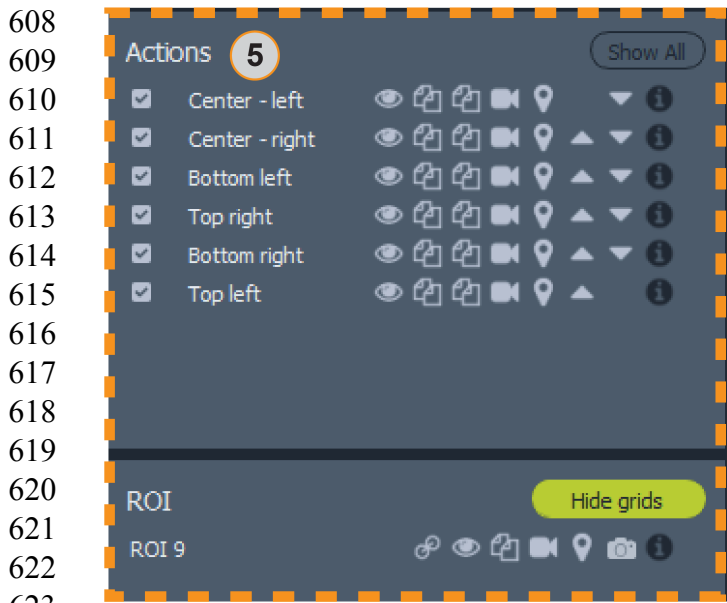
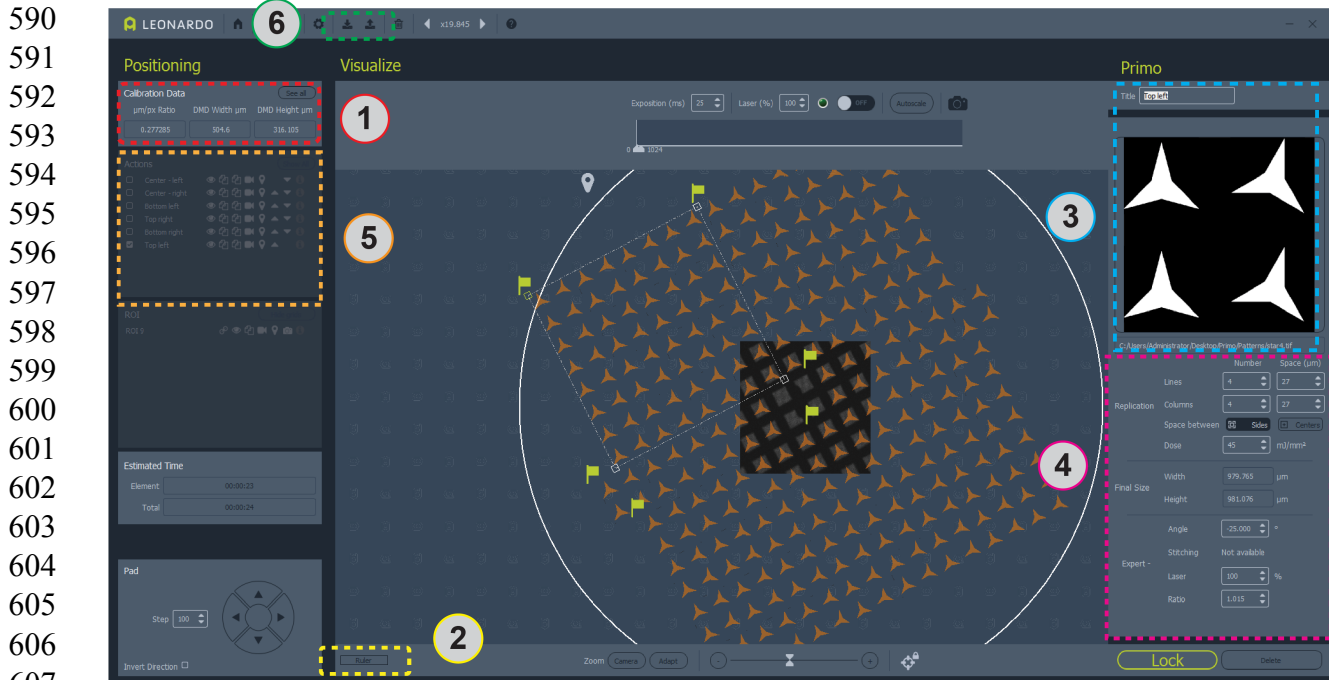


586 **Figure 1. General workflow of micropatterning for cryo-EM.** The workflow can be roughly

587 divided four parts: Grid preparation, micropatterning, ECM and cell seeding, and cryo-

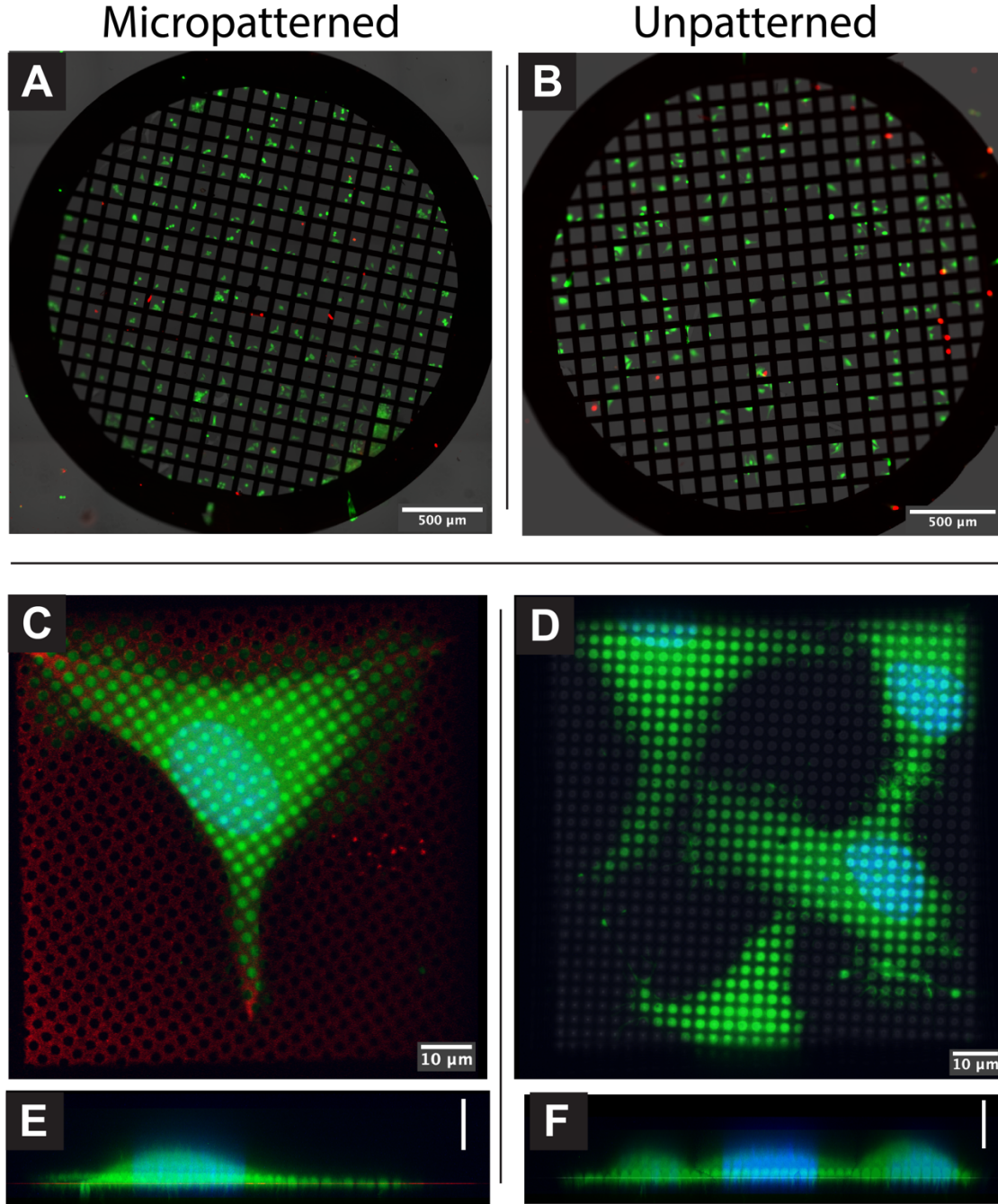
588 preparation and data collection. Major steps of each section are listed below the headings and

589 the approximate time to complete each section is shown to the left



627 **Figure 2. Screen shot of Leonardo software with pattern positioned on grid.** Area 1 contains
628 the $\mu\text{m}/\text{pix}$ ratio for pattern design. Area 2 is the ruler for measuring your grid. Area 3 is where
629 to add or change patterns and ROIs. Area 4 contains all of the information for pattern
630 positioning and dose. Area 5 contains options for patterns including toggling overlays, copying
631 or deleting patterns, and selecting patterns for micropatterning. Area 6 is where templates can
632 be saved and loaded. Larger views of areas 4 and 5 are shown below for clarity.
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669 **Figure 3. Live/Dead staining of patterned and unpatterned cells.** **A.** Fluorescent image of HeLa
670 cells grown on a patterned grid and stained with calcein-AM (live cell stain, green) and ethidium
671 homodimer-1 (dead cell stain, red). **B.** HeLa cells grown on an unpatterned grid and stained as
672 in **A.** **C.** Projection of confocal z-stacks of a HeLa cell on a patterned Quantifoil R2/2 grid with
673 0.01 mg/mL collagen and fibrinogen 647 ECM (red). Cell was stained with calcein-AM (green)
674 and Hoechst-33342 (blue). **D.** X,Z projection of **C.** **E.** HeLa cell on unpatterned grid incubated
675 with 0.01 mg/mL collagen and fibrinogen 647 ECM, incubated and stained with calcein-AM and
676 Hoechst-33342. The fluorescent images were merged with transmitted light (grayscale). **F.** X,Z
677 projection of **D.** Images are pseudocolored. All scale bars are 10 μm .

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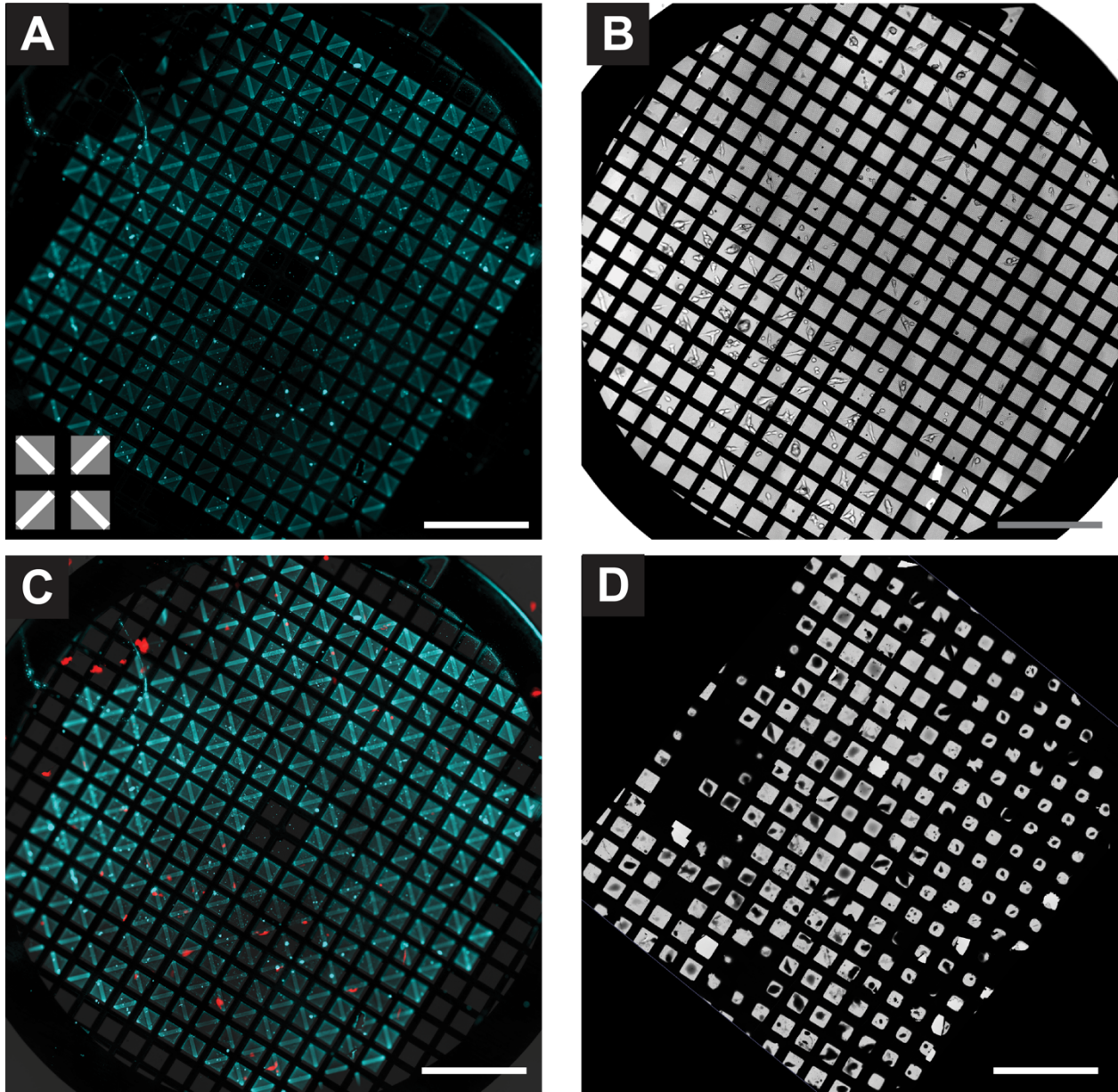


Figure 4. RSV infected BEAS-2B cells on patterned cryo-TEM grid. A. Fluorescent image of patterned grid after addition of fluorescently labelled ECM. The input pattern is shown in the lower left corner. **B.** Brightfield image of BEAS-2B cells grown on grid in A. **C.** Merge of image in A (cyan) and B (grey) with fluorescent image of RSV infected cells (red) immediately prior to plunge-freezing; infected cells express mKate-2. Scale bar 500 μm . Fluorescent images are pseudocolored. **D.** Low-magnification cryo-TEM map of grid in B after plunge-freezing.

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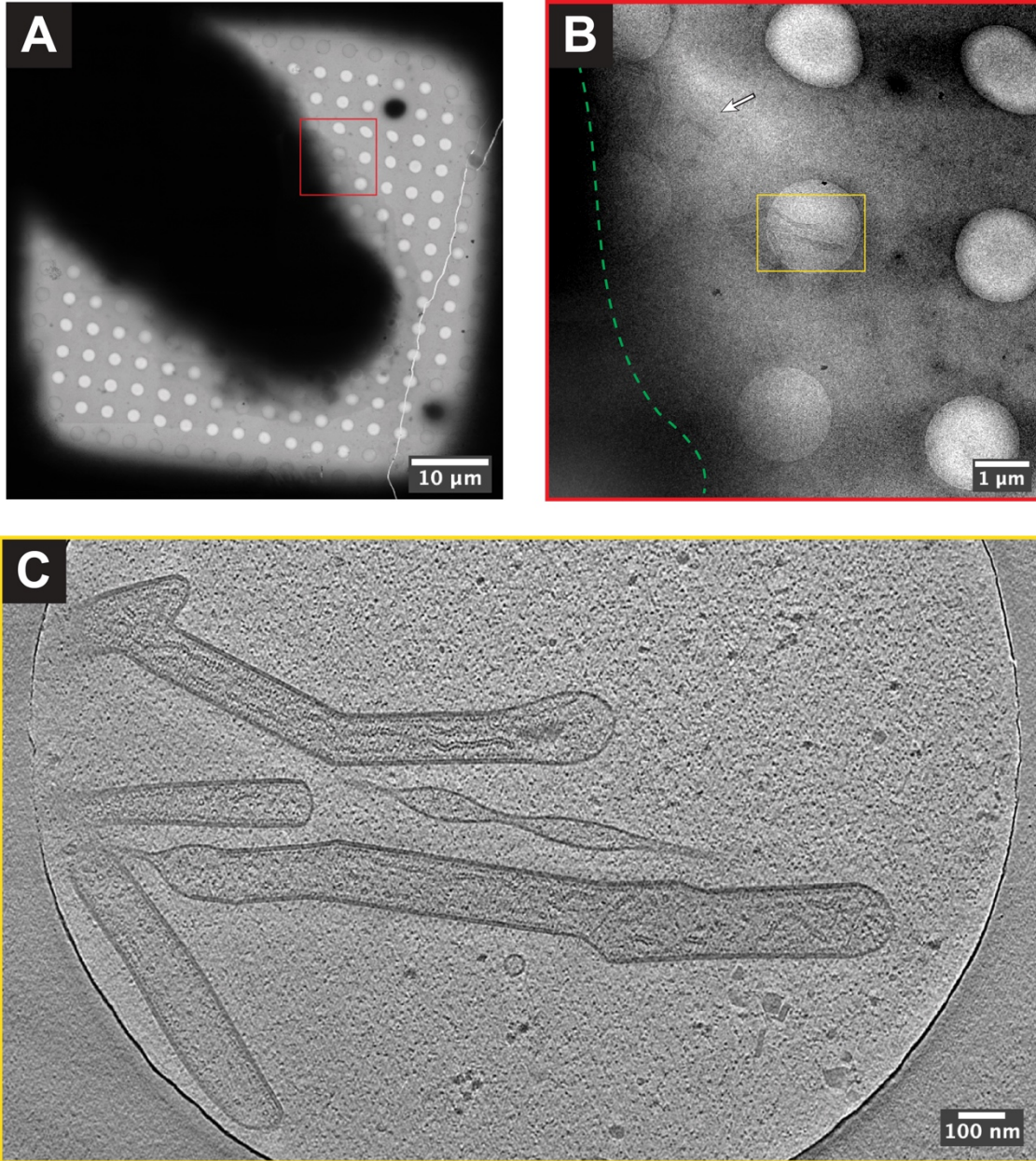


Figure 5. Cryo-ET of RSV infected BEAS-2B cell on patterned cryo-TEM grid. A. Cryo-EM grid square map of RSV infected BEAS-2B cell. **B.** Higher resolution image of area boxed in red in (A). Approximate cell boundary is indicated by dashed green line. RSV virions can be seen near the cell periphery (white arrow and yellow box). **C.** Single z-slice from tomogram collected in the area of the yellow box in (B). The scale bars in (A)-(C) are embedded in the image.

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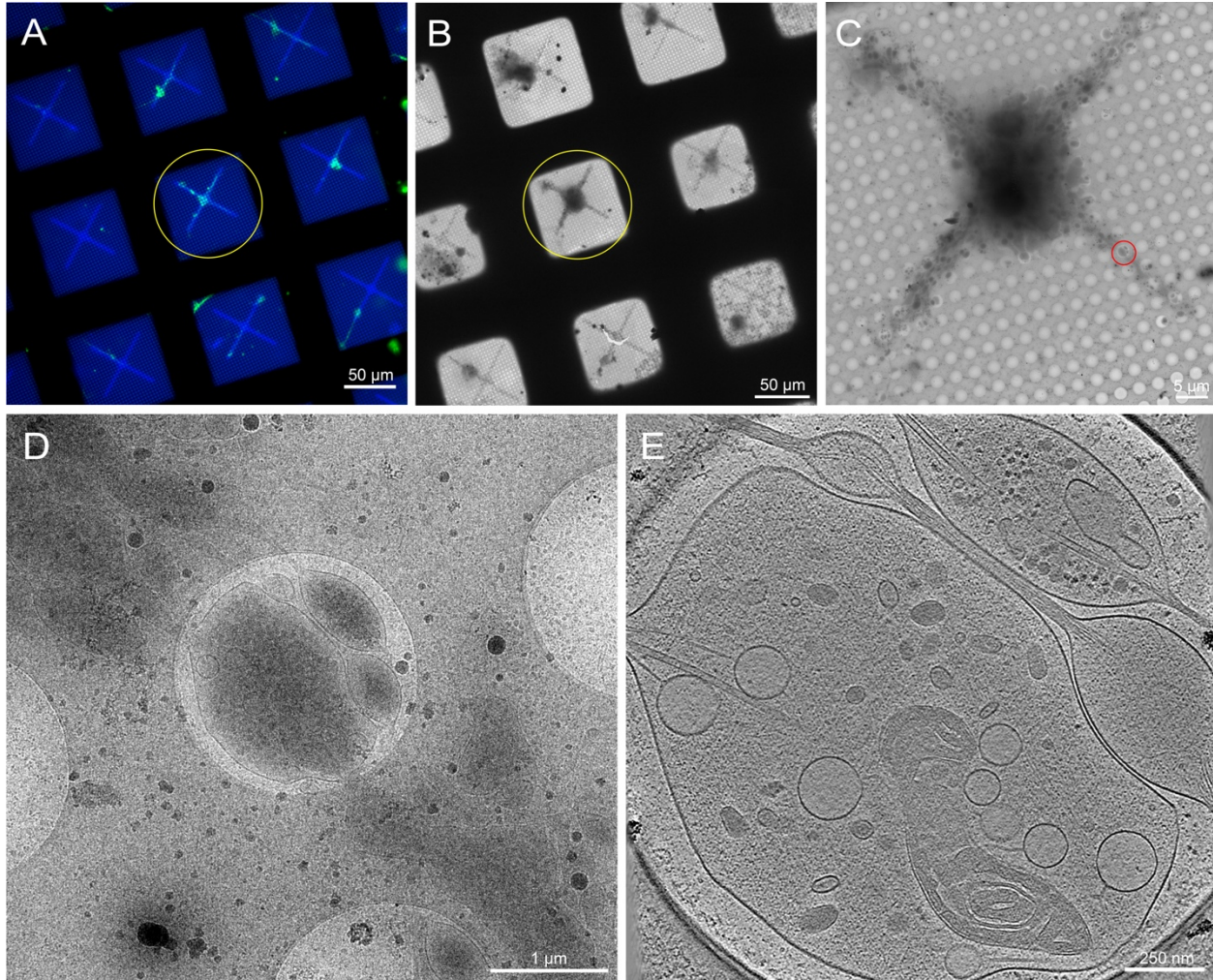


Figure 6. Primary neurons derived from the brains of 3rd instar *Drosophila melanogaster* larvae on patterned cryo-TEM grid. **A.** Overlaid live-cell fluorescent microscopy grid montage of *Drosophila* neurons expressing membrane-targeted GFP on patterned grid squares with 0.5 mg/mL fluorescent concanavalin A. Green: *Drosophila* neurons. Blue: Photopattern. **B.** Cryo-EM grid montage of the same grid in (A) after plunge-freezing. Yellow circle shows the same grid square as in (A). **C.** Magnified cryo-EM square montage of the yellow circle in (A) and (B). **D.** Magnified view of the red circle in (C), where a tilt series was collected on the neurites. **E.** 25 nm thick slice of a tomogram reconstructed from the tilt series that was acquired from the red circle in (C). Various organelles can be seen in this tomogram, such as the mitochondria, microtubules, dense core vesicles, light vesicles, the endoplasmic reticulum, and actin. Macromolecules, such as ribosomes, can also be seen in the upper right corner. Fluorescent images are pseudocolored. The scale bars in (A)-(E) are embedded in the image.

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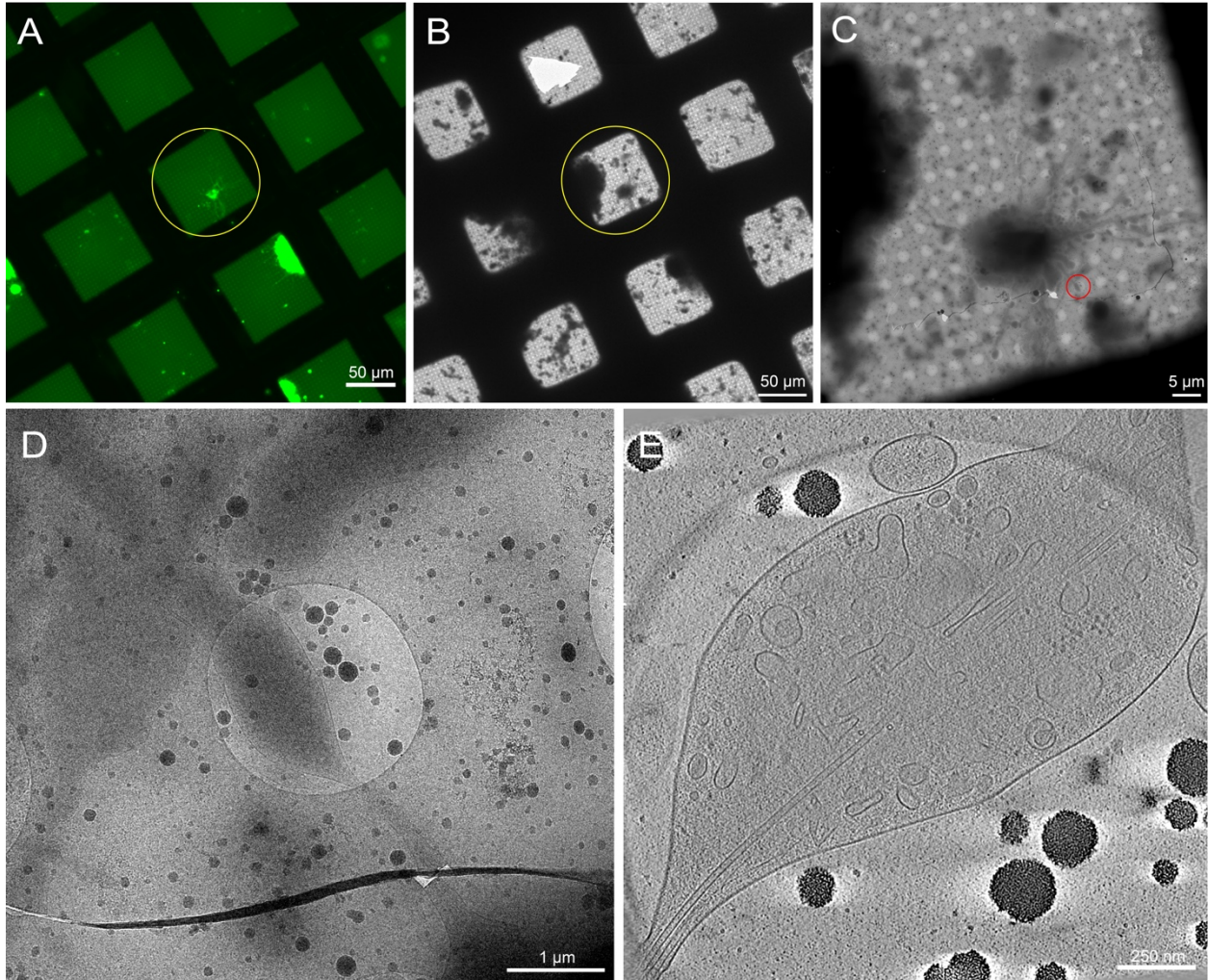


Figure 7. Primary neurons derived from the brains of 3rd instar *Drosophila melanogaster* larvae on unpatterned grids. A. Live-cell fluorescent microscopy grid montage of *Drosophila* neurons expressing membrane-targeted GFP on grid squares with 0.5 mg/mL concanavalin A. Green: *Drosophila* neurons. **B.** Cryo-EM grid montage of the same grid in (A) after plunge-freezing. Yellow circle shows the same grid square as in (A). Note the presence of cellular debris and media contamination, which made target identification difficult compared to patterned grids. **C.** Magnified cryo-EM square montage of the yellow circle in (A) and (B) maps. **D.** Magnified view of the red circle in (C), where a tilt series was collected on the neurites. **E.** 25 nm thick slice of a tomogram reconstructed from the tilt series that was acquired from the red circle in (C). Various organelles can be seen in this tomogram, such as microtubules and vesicles. Macromolecules, such as ribosomes, can also be seen. Fluorescent images are pseudocolored. The scale bars in (A)-(E) are embedded in the image.

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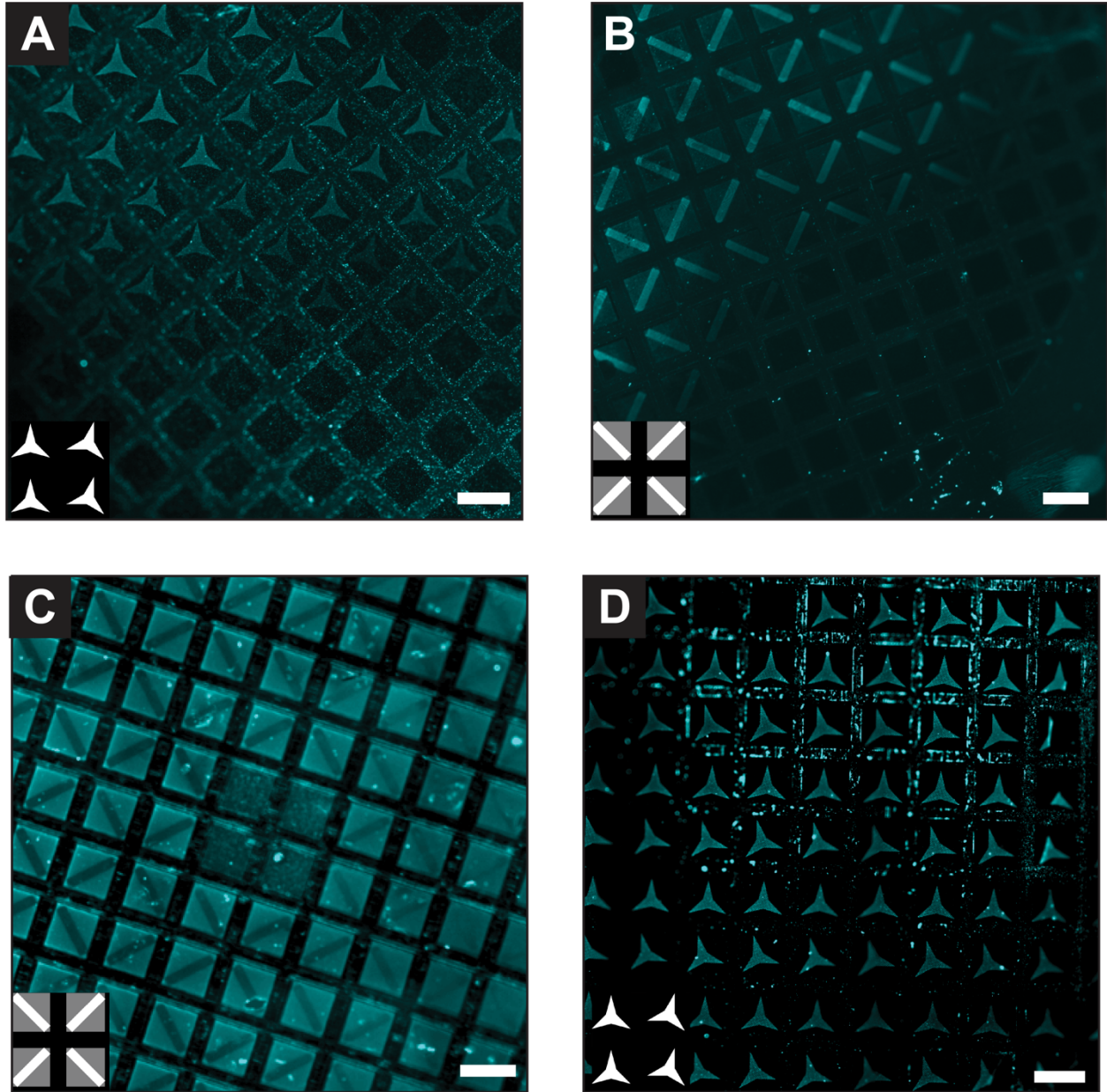


Figure 8. Examples of possible problems with patterning. Fluorescent images of labelled ECM deposited on micropatterned grids. **A.** Uneven patterning across the grid due to uneven distribution of PLPP gel. **B.** ECM cannot adhere to areas covered by the PDMS stencil during patterning. **C.** Saturated gradient pattern (right side) or inverted pattern (left) on a grid patterned with too high total dose. **D.** ECM is adhering to areas on the grid bars as well as patterned area due to reflections of the UV laser during patterning. Images are pseudocolored; input pattern is shown in lower left; scale bars are 100 μm .

Issue	Potential cause(s)	Troubleshooting
Micropatterning		
Cannot see illumination from PRIMO laser	<ul style="list-style-type: none"> • Light path is not set up correctly • PRIMO laser is not on or laser is interlocked 	<ul style="list-style-type: none"> • Check that the microscope light path is set up properly
Many broken grid squares	<ul style="list-style-type: none"> • Touching grid foil with tweezers or pipet while handling • Grid dried out during incubations or washing 	<ul style="list-style-type: none"> • Handle grids with care • Do not allow grid to dry during washes and incubations
Large unpatterned areas	<ul style="list-style-type: none"> • Insufficient gel coverage • Grid foil out of focus during patterning • Area covered by stencil 	<ul style="list-style-type: none"> • Ensure gel spreads evenly over grid while adding • Add an additional microliter of gel • Check focus before patterning each region • Carefully center grid in stencil
Saturated or inverted pattern	<ul style="list-style-type: none"> • Incorrect dose • Insufficient gel coverage 	<ul style="list-style-type: none"> • Try a range of total doses for pattern • Ensure grid is evenly covered with gel • Try different values for grayscale patterns
Blurry pattern	<ul style="list-style-type: none"> • Poor focus during patterning • Incorrect calibration 	<ul style="list-style-type: none"> • Repeat PRIMO calibration at same height as sample • Focus on grid foil before patterning • Divide pattern into additional regions for patterning
ECM adhering outside of pattern	<ul style="list-style-type: none"> • Reflections from gel or dust 	<ul style="list-style-type: none"> • Ensure gel is dry before patterning • Make sure coverslip and objective lens are clean
ECM not visible after patterning	<ul style="list-style-type: none"> • Photo bleaching • Incorrect dose during patterning • Insufficient ECM incubation time 	<ul style="list-style-type: none"> • Minimize light exposure to ECM prior to imaging • Try a range of total dose values for pattern • Increase incubation time for ECM
Cell seeding		
Cells clumping	<ul style="list-style-type: none"> • Over digestion • High cell density 	<ul style="list-style-type: none"> • Use lower percentage of trypsin or time for release of adherent cells • Passage and/or digest cells at lower confluency • Do not agitate cells during release • Gently pipet cell solution or use cell strainers
Cells not adhering to patterned areas	<ul style="list-style-type: none"> • ECM is not suitable for cell type • Cells viability is decreased prior to seeding 	<ul style="list-style-type: none"> • Try different ECM concentrations and composition • Ensure cell culture and cell release conditions are not damaging cells
Cells not expanding after adhesion	<ul style="list-style-type: none"> • ECM or pattern not suitable for cell type 	<ul style="list-style-type: none"> • Try different patterns and ECM • In some cases a more continuous foil (R1.2/20 vs R2/1) may promote cell expansion

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Table 1. Potential issues during micropatterning. This table describes some issues a user may experience during micropatterning or cell-seeding. Potential causes and troubleshooting are provided for each issue. Representative images of some problems can be seen in Figure 8.

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916

917 **DISCLOSURES:**

918

919 The authors have nothing to disclose.

920

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