Zebrafish otic vesicle and mouse epididymis as model systems for studying columnar epithelial cell division

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ABSTRACT

Epithelial cell division maintains tissue architecture through coordinated nuclear migration, cell shape changes, and spindle orientation. In columnar epithelia, interkinetic nuclear migration (INM) involves apical nuclear translocation in G2 and basal return post-mitosis, yet its regulation and physiological significance remain understudied due to limited live imaging *in vivo* models. We adapted the zebrafish embryonic otic vesicle as an *in vivo* model to study epithelial division dynamics using high-resolution live imaging and genetic tools. We find that apical INM initiates in mid-to-late G2 and is driven by dynein, not myosin II. Mitotic rounding is achieved via actomyosin-mediated basolateral constriction while maintaining basal attachment. Inhibiting myosin II impairs rounding and planar division, causing apical retention of daughter cells, suggesting planar division ensures proper integration. We extend our analysis to the mouse epididymis epithelium, comparing nuclear migration, cell shape, and spindle orientation across species. Our work introduces optimized *in vivo* models and reveals conserved and tissue-specific mechanisms maintaining structure and function of columnar epithelia.

INTRODUCTION

Epithelial cell division is a tightly regulated process that preserves tissue architecture and function by coordinating cell shape changes, polarity maintenance, and spindle orientation. A defining feature of epithelial tissues is their apical-basal polarity, which is crucial for barrier function, directional transport, and homeostasis (Rodriguez-Boulan and Macara, 2014). Apical membranes form tissue lumens, basal membranes adhere to the basal lamina, and lateral membranes connect neighboring cells through tight and adherens junctions. During mitosis, epithelial cells undergo morphological remodeling while maintaining tissue integrity through dynamic rearrangements of cortical actin and cell junctions. Proper spindle orientation ensures that daughter cells integrate correctly into the epithelium, preventing disruptions that can lead to developmental defects and disease (Morin and Bellaiche, 2011; Godard and Heisenberg, 2019).

Columnar epithelium can be categorized into pseudostratified and simple columnar epithelium. In simple columnar epithelium, such as that in the stomach, intestine, airway epithelia, otic vesicle, epididymis, and prostate, cells are column-shaped with nuclei positioned near the basal lamina (Fujita, 1960; Lee et al., 1994; Tsue et al., 1994; Robaire and Hinton, 2002; Fleming et al., 2007; Bagley et al., 2024). In contrast, pseudostratified epithelium, such as neuroepithelium, exhibits a "pearl-on-a-string" morphology, where cells have thin apical and basal processes, and their nuclei appear multilayered (Baye and Link, 2008; Taverna and Huttner, 2010). A key feature of these columnar epithelia is interkinetic nuclear migration (INM)—nuclear movement toward the apical surface in G2 phase and back to the basal side post-mitosis (Sauer, 1935; Fujita, 1960; Fujita, 1962; Baye and Link, 2008; Taverna and Huttner, 2010). Additionally, cells in these tissues undergo division in specific orientations, typically parallel to the apical membrane, due to polarized mitotic spindle orientation (Fujita, 1960; Ahringer, 2003; Baye and Link, 2008; Taverna and Huttner, 2010; Nakajima et al., 2019). Most mechanistic studies on INM focus on stem or progenitor cells in the neuroepithelium of developing retina or neural

tube. Both microtubules (MTs) and filamentous actin (F-actin) are generally believed to be critical for INM (Baye and Link, 2008; Taverna and Huttner, 2010; Kosodo, 2012). Discrepancies, however, exist, concerning the actual contributions of each cytoskeletal element, possibly due to differences in experimental systems or methods. In rodent neuroepithelia, cytoplasmic dynein, a MT-bonded minus end-directed motor protein, facilitates basal-to-apical nuclear movement, supported by apically anchored centrosomes and nuclear tethering proteins (Gambello et al., 2003; Xie et al., 2007; Zhang et al., 2009; Ge et al., 2010; Taverna and Huttner, 2010; Tsai et al., 2010). Kinesin-3, a MT-bonded plus end-directed motor, drives apical-to-basal movement, while myosin II appears dispensable (Tsai et al., 2010). However, some studies suggest a critical role for myosin II in apical-to-basal migration (Schenk et al., 2009). In zebrafish retina, INM in both directions depends mainly on myosin II (Norden et al., 2009).

In pseudostratified epithelia like neuroepithelium, nuclei navigate through narrow cellular processes (Baye and Link, 2008; Taverna and Huttner, 2010; Kosodo, 2012), whereas in simple columnar epithelia, nuclear migration in G2 may be less constrained, but mitotic rounding requires cytoplasmic displacement toward the apical side. Thus, while INM and division share core mechanisms across epithelial subtypes, distinct structural constraints may influence their regulation. Recent studies analyzing fixed *in vivo* samples and live imaging studies in intestinal organoids and chicken neural tube slices have provided new insights into epithelial division (Spear and Erickson, 2012; Carroll et al., 2017; Saleh et al., 2023). However, key knowledge gaps persist, particularly in how cells integrate mechanical and biochemical cues to drive nuclear migration and position the mitotic spindle, orient division, and maintain tissue integrity. *In vitro* models often fail to capture the complexity of *in vivo* epithelial division, emphasizing the need for physiologically relevant systems of columnar epithelia. Although *in vivo* live imaging has been widely applied to study epithelial cell division in various systems, including mouse skin, zebrafish embryos, and the Drosophila wing disc (e.g., (Campinho et al., 2013;

Bergstralh et al., 2016; Ipponjima et al., 2016; Moore et al., 2023; Xin et al., 2024)),

columnar epithelia remain comparatively underexplored in live contexts.

Building on previous work by the Megason group and others (Megason, 2009;

Hoijman et al., 2015; Mosaliganti et al., 2019), we adapted the zebrafish embryonic otic

vesicle as an optimized *in vivo* model for studying columnar epithelial cell division

dynamics, an area that has not been explored in this context, to address the questions

outlined above. The zebrafish otic vesicle epithelium offers several advantages, including

optical accessibility for high-resolution live imaging and genetic tractability, making it an

ideal system for visualizing columnar epithelial cell division in its native context. We also

validated some of our findings in the mouse epididymis model, revealing conserved

cellular mechanisms underlying epithelial cell division.

RESULTS

INM in zebrafish otic vesicle epithelium begins in late G2

To investigate nuclear behavior during cell division in zebrafish otic vesicle epithelium, we

labeled nuclei with H2B-GFP and cell membranes with RFP-F (Hancock et al., 1991;

Aronheim et al., 1994) by microinjecting *in vitro*-transcribed mRNAs into fertilized

zebrafish eggs. We focused on the ventral side of the otic vesicle, where epithelial cells

maintain a tall columnar shape between 22 and 28 hpf (Fig. 1A, B). As expected, while

most nuclei remained at the basal side, occasional INM events interspersed with mitosis

were observed (Fig. 1C; Videos S1 and S2). The entire migration cycle lasted an average

of 70.4 ± 7.7 minutes (n = 10).

We then analyzed nuclear dynamics before and after mitosis. The prophase-

prometaphase and anaphase-telophase transitions were easily identifiable in time-lapse

image sequences (Fig. 1C; Videos S1 and S2), allowing us to use these as reference

points for timing and duration analyses of INM (Fig. 1D-F). Apically directed INM occurred

27.8 ± 3.5 minutes before the prophase-prometaphase transition, with an average velocity

of 0.38 ± 0.05 μm/min (n = 10, Fig. 1D, F). Basally directed INM lasted 26.2 ± 6.4 minutes

after telophase, with an average velocity of 0.46 ± 0.09 μm/min (n = 10) (Fig. 1E, F).

Although apically directed INM is generally associated with G2 phase of the cell cycle

(Baye and Link, 2008; Taverna and Huttner, 2010; Kosodo, 2012), the precise timing of

its initiation within G2 remains unclear. To address this, we measured G2 phase duration

using BrdU (5-Bromo-2'-deoxyuridine) labeling. Fish embryos were exposed to BrdU at

24 hpf, followed by fixation at various time points. Mitotic cells were identified using

phospho-histone H3 (pH3) staining (Hendzel et al., 1997), which allowed clear recognition

of prophase cells. No BrdU-positive cells were observed entering M phase at 40-, 60-, or

80-minutes post-labeling (Fig. 1G). However, BrdU-positive mitotic cells appeared in all

embryos at 100, 120, and 140 minutes, indicating that G2 phase lasts between 80 and

100 minutes. Since apically directed INM initiates approximately 27.8 minutes before the

prophase-prometaphase transition (Fig. 1D), this migration occurs in late G2 phase.

INM in mouse epididymis resembles that in zebrafish otic vesicle

To validate our findings in the zebrafish otic vesicle epithelium and determine the timing

of basal-to-apical INM, we examined the murine epididymis, a tubular organ composed

of simple columnar epithelial cells. Although spatial distribution of mitotic cells along the

epididymal tubule at the tissue level has been explored, polarized epithelial cell division

has not been studied in the tissue (Fig. 2A)(Robaire and Hinton, 2002; Hirashima and

Adachi, 2015). A survey of epididymal cryosections stained for chromatin DNA and pH3

revealed that mitotic events peaked around the fifth week postpartum and occurred

primarily in the initial segment (Fig. 2A, B). G2-phase nuclei, identified by punctate pH3-

positive foci (Hendzel et al., 1997), were observed in both the basal and middle regions

(Fig. 2C, panels 1, 3, 4). In contrast, prophase nuclei, marked by chromosome

condensation and strong pH3 staining (Hendzel et al., 1997), were found exclusively in

the middle-to-apical region (Fig. 2C, panel 2). Prometaphase and metaphase

chromosomes clustered apically (Fig. 2C, panels 2-3), while telophase nuclei were

observed in the middle region, suggesting a return migration (Fig. 2C, panel 4).

Additionally, mitotic cells underwent planar divisions at the apical side (Fig. 2C, panels 2,

4), confirming that epithelial cells in the mouse epididymis exhibit characteristic INM and

apical division patterns.

To further investigate the timing of apically directed INM, mice were injected

intraperitoneally with BrdU to label S-phase cells and sacrificed 1-hour post-injection.

Among 750 BrdU-positive cells examined, all S-phase nuclei, which were pH3-negative,

were located at the basal side (Fig. 2D, 2E). The majority of G2-phase nuclei, identified

by dual BrdU and pH3 labeling, were found in the basal region, with a smaller fraction in

the middle region (Fig. 2D-E). These findings suggest that INM begins after G2, but not

immediately upon, G2 entry.

To trace nuclear dynamics during cell cycle progression, we performed a pulse-chase

experiment by sacrificing mice at 1, 5, 12, and 24 hours post-BrdU injection. At 1 hour,

89.6% of BrdU-positive nuclei were concentrated in the basal region. By 5 and 12 hours,

they were distributed across all regions (Fig. 2F-G). At 24 hours, BrdU-positive nuclei

peaked again in the basal region (86.4%), with a marked reduction in other regions (Fig.

2F-G). Furthermore, BrdU-positive nuclei frequently appeared as pairs at the basal side

at 24 hours (Fig. 2F), indicating that daughter cells remained adjacent post-division.

These findings suggest that epithelial cells in both the mouse epididymis and the

zebrafish otic vesicle exhibit the INM characteristic of proliferative simple columnar

epithelia.

Cell shape dynamics and basal lamina attachment during INM and mitosis

To gain insights into the mechanisms of INM, we investigated cell shape dynamics during

INM and subsequent cell division. In the epididymis, lateral adherens junctions visualized

through F-actin or E-cadherin staining (Yap et al., 1997; Baum and Georgiou, 2011)

revealed that columnar cell morphology persisted from interphase to prophase (Fig. 3A,

B). In prometaphase, cells began to round up while retaining a narrow basal protrusion

attached to the basal lamina (Fig. 3A, B, arrowheads), which largely remained throughout

cell division (Fig. 3A). Although E-cadherin staining intensity on the lateral cell membrane

remained unchanged (Fig. 3B), signals for F-actin, phospho-E-cadherin, and β-catenin

were markedly increased in prometaphase and metaphase cells (Fig. 3A-C), suggesting

enhanced lateral membrane adhesion between mitotic and neighboring interphase cells

(Lickert et al., 2000; Navarro et al., 2005; Bertocchi et al., 2012).

To further validate these observations, we examined zebrafish otic vesicle cells

labeled with H2B-GFP and RFP-F. As shown in Fig. 1C and Video S2, live imaging

revealed that columnar cell morphology was largely maintained during basal-to-apical

INM, consistent with observations in the epididymis. From prometaphase, the lateral cell

membrane at the basal region gradually collapsed, forming a stalk-like process that

remained attached to the basal lamina as the cell body rounded up (Fig. 1C, 32-48',

arrowheads). Moreover, cleavage furrow ingression initiated at the basal side of the cell

(Fig. 1C, 50', yellow arrow), reaching the apical surface within approximately 2 minutes

(Fig. 1C, 52').

To further examine cell shape and F-actin dynamics, we expressed GFP-tagged

hUtrCH, a fragment of human Utrophin that binds specifically to F-actin without perturbing

actin dynamics (Burkel et al., 2007; Norden et al., 2009). Live imaging showed GFP-

hUtrCH enrichment at the cell cortex in otic vesicle epithelial cells (Fig. 3D, Video S3),

confirming proper F-actin labeling in zebrafish. We observed similar cell shape changes

in terms of cell rounding timing (30'-56'), basal lamina attachment (50'-62'), and cleavage

furrow ingression (58'-62') (Fig. 3D, arrowheads). Additionally, we detected increased

GFP signal intensity during cell rounding and cleavage furrow ingression (Fig. 3D, bottom),

indicating the involvement of actomyosin activity in these processes. These findings

suggest that columnar epithelial cells round up at prometaphase and undergo mitosis

while maintaining a cellular protrusion anchored to the basal lamina, with cleavage furrow ingression initiating from the basal side of the cell body.

Myosin II activity regulates mitotic cell rounding, spindle orientation, and epithelial Integrity

Our live imaging of F-actin dynamics suggested a role for actomyosin in epithelial cell division. To further investigate the contribution of myosin II-mediated F-actin contraction, we examined its role in zebrafish epithelial cells. In non-muscle cells, myosin II is a well-established actin-associated motor protein that generates contractile forces and has been implicated in INM of neuroepithelia (Norden et al., 2009; Schenk et al., 2009). To assess myosin II function, we treated zebrafish embryos with 100 µM blebbistatin, a small-molecule inhibitor of myosin II activity (Straight et al., 2003). Consistent with previous finding(Norden et al., 2009), blebbistatin treatment did not affect actin polymerization, as GFP-hUtrCH localization remained unchanged compared to controls (Fig. 4A; Videos S4 and S5). The treatment was effective, however, as evidenced by the pronounced cytokinesis defects observed throughout the embryos (Fig. 4A; Videos S4 and S5)(Straight et al., 2003).

Blebbistatin treatment did not alter the frequency of basal-to-apical INM or the average migration velocity (0.38 ± 0.06 µm/min for 16 blebbistatin-treated cells vs. 0.4 ± 0.08 µm/min for 20 vehicle-treated cells) in the otic vesicle (Fig. 4B, C), suggesting that myosin II activity is dispensable for apically directed INM. However, several defects were noted in blebbistatin-treated cells. Firstly, the treated cells failed to round up during mitosis. While 98.3% of vehicle-treated cells (58/59) successfully rounded up after entering prometaphase (Fig. 4D; Video S4), blebbistatin-treated cells largely failed to do so. Instead, 73.1% (38/52) of treated cells maintained a diamond-shaped morphology in prometaphase and metaphase (Fig. 4D; Video S5).

Secondly, although mitosis still occurred at the apical surface, spindle orientation was

frequently disrupted. In control cells, 97.6% (41/42) divided at angles between 60°-90°,

indicating proper planar spindle orientation (Fig. 4D-F). In contrast, only 39.2% (20/51) of

blebbistatin-treated cells maintained this orientation, while another 39.2% divided along

the basal-apical axis (0°-30°), demonstrating a severe spindle misorientation defect (Fig.

4D-F).

Thirdly, in control mitotic cells, both daughter nuclei migrated back to the basal side

within ~30 min after anaphase onset (Fig. 4A, D; Video S4). However, in 70.6% of

blebbistatin-treated mitotic cells (36/51), one daughter nucleus remained at the apical

side for over 50 min (Fig. 4A, D; Video S5). Among these, 33 of the 36 retained nuclei

persisted at the apical surface (Fig. 4F, red dots) throughout the imaging period, while

only 3 eventually migrated basally. GFP-hUtrCH fluorescence revealed that daughter

cells with retained nuclei remained round, indicating a failure to reintegrate into the

epithelium (Fig. 4A, D; Video S5). Further analysis revealed a correlation between division

angles and nuclear retention. Among blebbistatin-treated mitotic cells with division angles

of 0°-60°, 96.8% (30/31) exhibited apical nuclear retention (Fig. 4F, red dots), whereas

only 30% (6/20) of those with division angles between 60°-90° retained nuclei apically.

Cells positioned closer to the lumen were more likely to be retained (Fig. 4D). These

findings suggest that planar division is essential for proper daughter cell integration into

the epithelium. Together, our results demonstrate that while cell rounding is not required

for M-phase entry, it is crucial for proper spindle orientation and epithelial integrity.

Basal-to-apical INM requires microtubules

Both MTs and F-actin have been reported to play essential roles in INM (Baye and Link,

2008; Taverna and Huttner, 2010; Kosodo, 2012). To examine the specific contributions

of MTs in our INM models, we performed immunostaining against α-tubulin in mouse

epididymal epithelium. This revealed MTs oriented along the apicobasal axis of the cell

(Fig. 5A). These MTs were largely stable and surrounded migrating nuclei, as indicated

by acetylated tubulin (Ace-tu) staining (Fig. 5B, C). Similarly, live imaging of zebrafish otic

vesicle epithelium using EMTB-GFP, a live-cell marker for MTs (Bulinski et al., 1999; Faire

et al., 1999; Wuhr et al., 2011), showed a comparable MT organization (Fig. 5D; Video

S6).

To functionally validate these findings, we treated zebrafish embryos with 0.5 µg/ml

nocodazole to disrupt MTs. To confirm the drug efficacy, embryos were fixed 120 minutes

after treatment, revealing the absence or fragmentation of mitotic spindles in zebrafish

otic vesicle mitotic cells, whereas normal spindles were observed in vehicle (DMSO)-

treated controls (Fig. 5E, arrows). MTs in interphase cells were also largely disrupted

following nocodazole treatment (Fig. 5E).

Live imaging from ~20 minutes after nocodazole treatment showed mitotic cells

arrested in mitosis (Fig. 5F, arrowhead), confirming drug effectiveness. Time-lapse

microscopy further revealed a significant reduction in INM events (Fig. 5F, G; Videos S7

and S8). As nocodazole treatment progressively thinned the epithelium, we focused our

analysis on the thickest region, located in the anterior-ventral area, from 30 to 236 minutes

post-treatment, when epithelial cells remained columnar (Fig. 5F; Video S8). Compared

to the vehicle treatment, nocodazole treatment resulted in an average 7.5-fold reduction

in INM events (Fig. 5G). These results demonstrate that basal-to-apical INM relies on

MTs.

Cytoplasmic dynein activity is critical for basal-to-apical INM

Studies in rodent neuroepithelia suggest that cytoplasmic dynein, a microtubule-based

minus-end-directed motor protein, plays a major role in mediating basal-to-apical nuclear

movement. This function is supported by proteins localized both at the apically anchored

centrosome (such as Cep120, TACCs, Hook3, and PCM1), which help maintain

microtubule arrays (Gambello et al., 2003; Xie et al., 2007; Ge et al., 2010; Tsai et al.,

2010), and at the nuclear membrane (such as Syne1/2 and Sun1/2), which tether the

nucleus to dynein (Zhang et al., 2009; Taverna and Huttner, 2010). To assess the role of

dynein in INM in the zebrafish otic vesicle epithelium, we overexpressed RFP-tagged

human p50, a known inhibitor of dynein activity (Echeverri et al., 1996; Burkhardt et al.,

1997; Liang et al., 2007). To minimize potential developmental defects caused by dynein

inactivation, we microinjected zebrafish embryos with a plasmid encoding RFP-p50 under

the control of the zebrafish hsp70 promoter, followed by heat shock treatment at 20 hpf

for 1 hour to induce RFP-p50 expression (Asakawa et al., 2008). Live-cell imaging was

performed 1.5 hours after heat shock, when RFP fluorescence became visible. Compared

to RFP-expressing controls, RFP-p50 overexpression significantly suppressed apically

directed INM (Fig. 5H, 5I; Videos S9, S10). Over a 300-minute observation period, 55 ±

10% of RFP-positive otic vesicle cells underwent apical INM, whereas only 21 ± 5% of

RFP-p50-expressing cells did (Fig. 5I). These results suggest that cytoplasmic dynein is

required for MT-mediated basal-to-apical INM.

Nuclear migration is essential for M phase onset

Interestingly, in long columnar cells treated with nocodazole, mitotic entry was observed

only at the apical side, after cells had completed INM (Fig. 5F; Video S8). By contrast, in

the thinner epithelium of the posterior region, where nuclei are close to the apical

membrane, cells frequently entered M phase (Video S8). This difference implies a link

between nuclear position and mitotic entry. To investigate this further, immunostaining of

mouse epididymal epithelium revealed that centrosomes localized near the apical surface

during nuclear migration, and centrioles only moved to opposite sides of the nucleus after

it reached the apical side (Fig. 6A). Live imaging using EMTB-GFP to label MTs and MT

organizing centers (MTOCs) confirmed this dynamic process in the zebrafish otic vesicle

epithelium. The MTOCs remained apically localized until the nuclei reached the apical

side (Fig. 6B; Video S6). To explore the molecular mechanisms further, we stained for

cyclin B1, a marker of late G2 and M-phase cells. In the mouse epididymis epithelium,

Cyclin B1 was localized in the cytoplasm of migrating nuclei and only entered the nuclei

once they reached the apex (Fig. 6C, D). this further confirmed that the timing of mitosis

coincides with the apical localization of nuclei (Gavet and Pines, 2010; Santos et al.,

2012).

DISCUSSION

Recent studies using intestinal organoids, neural tube slices, and other cultured models

have provided insights into columnar epithelial cell division (Spear and Erickson, 2012;

Carroll et al., 2017; Saleh et al., 2023), but these systems often lack the spatial

organization and/or biomechanical context of native tissues. *In vivo* models, meanwhile,

are typically limited to fixed samples, which precludes dynamic analysis. Although

previous studies have characterized the morphogenesis of the zebrafish otic vesicle

(Megason, 2009; Hoijman et al., 2015; Mosaliganti et al., 2019), we adapted this system

as a live, physiological, and genetically tractable model for real-time visualization of

epithelial nuclear dynamics and mitosis, thereby addressing key limitations of earlier

approaches. Furthermore, we validated the translatability of our findings using a

complementary mammalian model, the mouse epididymal epithelium.

Basal-to-apical INM does not start immediately after G2 entry

Our results suggest that basal-to-apical INM begins in the late G2 phase in the zebrafish

otic vesicle epithelium (Fig. 1, 7). Similarly, in the mouse epididymis, INM predominantly

occurs after G2 initiation (Fig. 2). Interestingly, while nuclear oscillations are frequently

observed in neuroepithelia, a rapid, directional basal-to-apical movement occurs just prior

to mitosis (Norden et al., 2009; Tsai et al., 2010; Kosodo et al., 2011). This movement

appears to initiate in mid-to-late G2 across different epithelial systems. For example,

directional INM lasts 5-30 minutes in the developing zebrafish retina (Norden et al., 2009)

and approximately 60 minutes in the mouse neural tube[14, 45]. Thus, the late onset of INM

during G2 may be a conserved feature across both simple columnar and pseudostratified

epithelia.

Basal-to-apical INM requires MTs and dynein, but not myosin II

We show that basal-to-apical INM in the zebrafish otic vesicle is dependent on MTs and

dynein (Fig. 7). Both p50 overexpression and nocodazole treatment significantly

suppressed INM (Fig. 5), whereas myosin II inhibition had little effect (Fig. 4). This is in

contrast to neuroepithelia, where MTs, dynein, and actomyosin all contribute to INM

(Taverna and Huttner, 2010; Kosodo, 2012). This difference may be attributed to cell

shape. In pseudostratified epithelia, nuclei migrate along narrow processes, requiring

both MT motor-generated pulling forces and actomyosin-mediated squeezing forces

exerted by lateral membrane constriction (Taverna and Huttner, 2010; Kosodo, 2012).

However, in columnar epithelia, where cell shape remains unchanged during INM,

actomyosin-generated forces become dispensable. Because dynein inhibition or MT

disassembly blocked basal-to-apical INM, we could not assess the role of the MT system

in basally directed INM in the zebrafish otic vesicle. Similarly, blebbistatin-treated

embryos exhibited impaired planar cell division (Fig. 4), preventing us from determining

the role of myosin II in apical-to-basal nuclear movement. However, since one or both

daughter nuclei were still able to move back to the basal side (Fig. 4), myosin II may be

dispensable for apical-to-basal nuclear migration.

Mitotic cell rounding and planar division depend on actomyosin

We clarify that mitotic cell rounding in both zebrafish otic vesicle and mouse epididymal

epithelium occurs during prometaphase (Fig. 3, 4, and 7). Similar basal adhesion

maintenance has been reported in mitotic epithelial cells of the mouse intestine and

kidney, suggesting a shared feature in both simple columnar and pseudostratified

epithelia (Fleming et al., 2007; Baye and Link, 2008; Taverna and Huttner, 2010; Kosodo,

2012; Packard et al., 2013). We show that myosin II is the primary driver of mitotic

rounding (Fig. 4). The increased signals of F-actin, phospho-E-cadherin, and β-catenin at

lateral membranes (Fig. 3) likely strengthen actomyosin forces, facilitating lateral

membrane contraction while maintaining tight adhesion to neighboring interphase cells

during mitotic shape remodeling (Fig. 7) (Lickert et al., 2000; Navarro et al., 2005; Qin et

al., 2005). Understanding how INM and mitotic rounding are coupled to specific cell cycle

stages remains an intriguing avenue for future research. Planar cell division also depends

on myosin II activity (Fig. 4), likely because failure of mitotic rounding disrupts spindle

orientation. Spindle orientation is biased toward the long axis of the cell (O'Connell and

Wang, 2000; Saleh et al., 2023) and requires interactions between the spindle and the

cell cortex (Siller and Doe, 2009; Lu and Johnston, 2013). Thus, mitotic rounding may

provide the cortical geometry needed for proper spindle alignment and planar division

(Campinho et al., 2013; Godard and Heisenberg, 2019; Saleh et al., 2023).

Planar cell division facilitates daughter cell integration into the epithelium

In zebrafish otic vesicles treated with blebbistatin, perpendicular divisions (0-30° angles)

frequently resulted in apical retention of one daughter cell near the lumen, while the

basally positioned daughter nucleus returned to the basal side (Fig. 4). By contrast, planar

divisions (60-90° angles) were associated with a significantly reduced incidence of

nuclear retention. One possible explanation is that planar division preserves both apical

tight junctions(Packard et al., 2013) and basal lamina attachment, maintaining basal-

apical polarity of daughter cells (Fig. 7). In contrast, perpendicular division disrupts basal

lamina contact, preventing efficient epithelial reintegration. Such defects could

compromise otic vesicle structure and function, and the mechanism may also apply to

other simple columnar epithelia, where planar division failure disrupts tissue integrity.

Mitosis is coupled to INM in the zebrafish otic vesicle epithelium

In neuroepithelia, INM is not required for cell cycle progression. Blocking INM via
cytochalasin B treatment or disrupting Lis1/Dynein function allows mitosis to occur
throughout the neuroepithelium without halting the cell cycle (Murciano et al., 2002;
Gambello et al., 2003; Tsai et al., 2005; Del Bene et al., 2008). However, INM is
dependent on cell cycle progression, as cell cycle arrest (e.g., by 5-azacytidine and
cyclophosphamide) halts INM (Ueno et al., 2006). In our system, mitosis occurs
exclusively at the apical side. Disrupting MTs or dynein activity led to a significant
reduction in mitoses, with no mitotic events observed at the basal side in the tall columnar
epithelium. This finding aligns with our observation that mitotic spindle formation and
nuclear entry of cyclin B occur when nuclei approach the apical side. Future research
should investigate whether apically localized signals, such as centrosome-associated
components, contribute to M-phase onset.

Limitations of the study

Here, we demonstrated the use of the zebrafish embryonic otic vesicle epithelium as a
model for investigating epithelial cell division in a physiological context with a focus on
nuclear migration, cell shape dynamics, and spindle orientation. However, several
technical limitations should be considered. First, we utilized microinjection of *in vitro*
synthesized mRNA to label histones, F-actin, MTs to demonstrate the feasibility of this
approach. To enhance consistency and minimize potential toxicity from ectopic transgene
expression, available stable transgenic lines could be used. For example, the
h2afv:h2afv-mCherry (Knopf et al., 2011) or *hsp70:H2B-RFP* (Dzafic et al., 2015) line can
serve as a histone marker, while *actb2:mCherry-UTRN* (Compagnon et al., 2014) or
actb1:lifeact-GFP (Behrndt et al., 2012) can be used for F-actin labeling. Additionally,
Fluorescent Ubiquitin-Mediated Cell Cycle Indicator (FUCCI) reporters (Sakaue-Sawano
et al., 2008; Bouldin et al., 2014) provide a powerful tool for tracking cell cycle progression.
Various inducible transgenic tools are also widely available for conditional gene

manipulation in zebrafish embryos. Second, our live imaging experiments were performed

using an inverted confocal microscope, requiring careful orientation of the embryos so

that the otic vesicle is positioned close to the glass bottom of the mounting dish for optimal

imaging quality. The use of an upright microscope equipped with high-NA water

immersion objectives would simplify sample mounting and improve resolution.

Additionally, spinning-disk microscopy could further enhance spatiotemporal resolution.

Third, the morphology of the zebrafish embryonic otic vesicle evolves rapidly, with a

limited window of approximately six hours (22–28 hpf) during which the ventral side

maintains a columnar epithelial structure. This constraint makes it less suitable for studies

requiring extended observation periods. Nonetheless, this live imaging system provides

an unprecedented opportunity to explore epithelial cell division in a natural physiological

context, offering valuable insights into cellular and molecular dynamics.

Materials and Methods

Animals

All animal procedures were conducted in accordance with protocols approved by the Institutional Animal Care and Use Committee (IACUC). Zebrafish (Ekkwill (EK) and AB strains) were raised under standard conditions and staged in hours post-fertilization (hpf) as previously described (Kimmel et al., 1995; Westerfield, 2000). Male C57BL/6J mice 3-12 weeks old were obtained from Jackson Laboratories and maintained under a 12-hour light/dark cycle at a constant temperature of 22°C.

Plasmid constructs

To express RFP-F, the open reading frame (ORF) of monomeric RFP (mRFP) from pmRFP-C1 (Li et al., 2007) was fused to the c-Ha-Ras farnesylation signal (Hancock et al., 1991; Aronheim et al., 1994) via PCR. For GFP-UtrCH expression, the first 783 base pairs of the human utrophin coding sequence (Burkel et al., 2007; Norden et al., 2009) were PCR-amplified and inserted in-frame into the BspEI-XhoI sites of pEGFP-C1. For *in vitro* mRNA transcription, the ORFs for H2B-GFP (Liang et al., 2007), H2B-RFP (Li et al., 2007), RFP-F, GFP-hUtrCH, and EMTB-GFP (Wuhr et al., 2011) were subcloned into pCS2$^+$. For heat shock-inducible expression of RFP or RFP-p50, the ORFs were PCR-amplified and inserted into the BamHI-NheI sites of pT2KhspGFF (Asakawa et al., 2008). pT2KhspGFF and pCS-TP, which encodes a transposase required for stable integration of pT2KhspGFF into genomic DNA (Kawakami et al., 2004), were generously provided by Dr. K. Kawakami (Graduate University for Advanced Studies, Japan). All plasmids containing PCR-amplified fragments were confirmed by sequencing.

Zebrafish microinjection

In vitro mRNA synthesis was performed using the mMessage mMachine SP6 kit (Thermo Fisher, AM1340). For yolk injections, 5 nL of transcribed mRNA was injected, while for

cytoplasmic injections, 1 nL of a plasmid-mRNA mixture was delivered into fertilized eggs.

Phenol red (0.02%) was included as a tracer. For yolk injections, mRNAs encoding H2B-

GFP, H2B-RFP, GFP-hUtrCH, or EMTB-GFP were used at a concentration of 50 ng/µL,

while RFP-F mRNA was injected at 75 ng/µL. For cytoplasmic injections, a mixture

containing H2B-GFP mRNA (20 ng/µL), TP mRNA (30 ng/µL), and a heat shock-inducible

plasmid (30 ng/µL) was used. To induce gene expression, heat shock was applied for one

hour at 39°C in a water bath starting at approximately 20 hpf.

BrdU incorporation

For mouse experiments, C57BL/6 mice received intraperitoneal injections of BrdU at a

dose of 100 µg per gram of body weight, followed by sacrifice at different time points. The

epididymides were dissected and fixed overnight in 4% paraformaldehyde (PFA) in PBS

at 4°C. BrdU labeling in zebrafish was performed as previously described(Shepard et al.,

2004). Briefly, at 24 hpf, embryos were dechorionated and incubated in 10 mM BrdU

(Millipore Sigma, B5002) in embryo medium for 20 minutes on ice. They were then

transferred to warm medium and incubated for an additional 40 to 140 minutes at 28.5°C.

After incubation, embryos were fixed overnight in PFA and subsequently dehydrated in

graded methanol in PBT (PBS containing 0.1% Tween-20).

Immunofluorescence microscopy

Mouse epididymal cryosections were prepared at a thickness of 10 µm and processed for

staining as previously described (Wang et al., 2008). Briefly, tissue samples were blocked

in PBS containing 10% goat serum and 0.5% Triton X-100 for one hour, followed by

overnight incubation with primary antibodies at 4°C. After extensive washing, secondary

antibodies were applied for 1.5 hours at room temperature. For BrdU staining,

cryosections were denatured in 2 M HCl for 15 minutes at 37°C, neutralized in 0.1 M

$Na_2B_4O_7$ for 10 minutes at room temperature, and rinsed three times in PBS before

incubation with an anti-BrdU antibody at a 1:1000 dilution. Whole-mount immunostaining

of zebrafish embryos followed established protocols (Wloga et al., 2009). BrdU staining

was performed by digesting rehydrated embryos with proteinase K (10 µg/mL, Fisher

Scientific, BP1700-100) for 10 minutes at room temperature, followed by fixation in PFA

for 20 minutes. Embryos were incubated in 2 M HCl for one hour, washed three times in

PBT, and then subjected to immunostaining (Shepard et al., 2004). The primary

antibodies used in this study included mouse monoclonal antibodies against α-tubulin

(Millipore Sigma, T6199) and BrdU (Millipore Sigma, SAB4700630), rabbit antibodies

against phospho-histone H3 (Ser10) (Cell Signaling Technology, 9701), phospho-E-

cadherin (Ser838/840; Novus Biologicals, NBP3-12922), and β-catenin (Millipore Sigma,

C2206), as well as a rat monoclonal antibody against E-cadherin (Abcam, ab11512).

Secondary antibodies (ThermoFisher) used in this study were Alexa Fluor 488 goat anti-

rabbit (A11034), goat anti-rat (A11006) and goat anti-mouse (A11029), and Alexa Fluor

546 goat anti-rabbit (A11035) and goat anti-mouse (A11030). Phalloidin-TRITC (P1951)

and DAPI (D9542) were from Millipore Sigma. Fixed samples were imaged using Leica

TCS SP5 or Zeiss LSM 800 confocal microscope.

Live cell imaging

Embryos were first screened at 22 hpf under a Zeiss Axio Zoom V16 fluorescent

dissecting microscope to identify those with GFP or RFP expression in the otic vesicles.

Selected embryos were dechorionated, anesthetized with tricaine (Sigma-Aldrich,

E10521), mounted in 1% low-melting point agarose (Fisher Scientific, BP1360-100) within

a glass-bottom Petri dish (MatTek Life Sciences, P35G-1.5-14-C), and covered with E2

embryo medium containing tricaine and 0.003% phenylthiourea (Millipore Sigma, P7629).

For experiments involving drug treatments, blebbistatin (100 µM, Millipore Sigma, B0560),

nocodazole (0.5 µg/ml, Millipore Sigma, M1404), or DMSO (Millipore Sigma, D8418) as

the solvent control was added to both the medium and agarose. Embryos were positioned

so that one otic vesicle was in close proximity to the glass bottom. Time-lapse imaging

was conducted at 2-minute intervals over a period of 216 to 430 minutes using a 20× NA

0.7 objective on a Leica TCS SP5 or Zeiss LSM 800 inverted confocal microscope

equipped with a heating hood.

Statistical analysis of nuclear migration

Nuclear displacement was quantified as the distance between the bottom edge of the

nucleus and the basal membrane(Norden et al., 2009). Measurements were taken at 2-

minute intervals using ImageJ software. Instantaneous velocity (v_i) was generate using

formula $v_i = (p_{i+1}-p_i)/(t_{i+1}-t_i)$ (Norden et al., 2009). where p represents position, t represents

time, and i denotes the time point. Positive values indicated apical migration, while

negative values represented basal migration. Quantification values are reported as mean

± S.D. Student's t-tests (two-tailed) were applied when normality and equal variance tests

were passed.

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Author contributions

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Investigation: Y.X., B.P., A.G.C.Y., and M.Q.

Resources: J.C.

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Funding Acquisition: J.C.

Declaration of interest

The authors declare no competing interests.

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 Figure legends

Figure 1. INM in zebrafish otic vesicle epithelium begins in late G2.

(A) Schematic of the zebrafish otic vesicle at approximately 22 hpf.

(B) Morphology of the otic vesicle epithelium at 22 hpf and 28 hpf. Embryos were

microinjected with mRNA to express H2B-GFP (chromatin, green) and RFP-F (cell

membrane, red). The boxed regions indicate areas used for INM analyses. Scale bar, 20

μm.

(C) Selected frames from Video S2. H2B-GFP (green) and RFP-F (red) label chromatin

and the plasma membrane, respectively. Single-channel images of RFP-F are shown in

the middle. White arrows indicate the nucleus or chromosomes of the tracked cell, white

arrowheads highlight its basal protrusion and plasma membrane, and yellow arrows

denote the cleavage furrow. The cell boundary and nucleolus of the tracked cell are

outlined below. Scale bar, 20 μm.

(D, E) Nuclear trajectories before prometaphase (D) and after telophase (E). For daughter

nuclei, only one that could be clearly tracked was analyzed.

(F) Velocity of pooled nuclear movements.

(G) Estimation of G2 length. Embryos at 24 hpf were labeled with BrdU, fixed at the

indicated time points, and stained to visualize BrdU (red), pH3 (green), and DAPI (blue).

Representative S-phase cells (arrowheads) and mitotic cells (pH3-positive, arrows) are

shown. Yellow dashed lines indicate the basal membrane. Scale bar, 10 μm.

Figure 2. INM in the mouse epididymis resembles that in the zebrafish otic vesicle.

(A) Schematic of the anatomical structure of the mouse epididymis.

(B) Quantification of mitotic events per epididymal tubular cross-section. Mitotic cells were

identified based on chromosome morphology and/or strong pH3 staining (see C). The

number of analyzed tubular cross-sections is indicated above each bar.

(C) Section images showing nuclear positioning at different cell cycle stages. Insets

display pH3 (green) or DAPI (red) staining of the indicated nuclei. Yellow lines mark the

apical surface. Scale bar, 20 μm.

(D, E) Section images and quantification (E) showing S-phase and G2-phase nuclei 1

hour after BrdU labeling. Scale bar, 20 μm.

(F, G) Section images and quantification showing the distribution of BrdU-positive nuclei

or chromosome clusters at the indicated time points. Arrows in (F) indicate representative

nuclei or chromosome clusters. Scale bar, 20 μm.

Figure 3. Cell shape dynamics and basal lamina attachment during INM and mitosis.

(A-C) Section images of mouse epididymis epithelium showing cell-cell interfaces labeled

with the indicated markers (green). Insets show pH3 staining (magenta) in G2 or

prophase cells. Arrows indicate G2 or M-phase cells, and arrowheads point to the narrow

basal protrusions of mitotic cells. DAPI staining is shown in red. Fluorescent intensity of

F-action signals in (A) is shown at the bottom. Scale bars, 10 μm.

(D) Selected frames from Video S3 showing GFP-hUtCH (green) and H2B-RFP (red).

Single-channel GFP-hUtCH images are shown in grayscale at the bottom. Arrows indicate

the nucleus or chromosomes of the tracked cell, while arrowheads highlight its basal

process and plasma membrane. Fluorescent intensity of GFP signals is shown at the

bottom. Scale bar, 10 μm.

Figure 4. Myosin II activity regulates mitotic cell rounding, spindle orientation, and

epithelial Integrity.

(A) Selected frames from Videos S4 and S5 showing INM events. H2B-RFP (red) and

GFP-hUtrCH (green) are shown. Time is indicated relative to the addition of DMSO or

blebbistatin. Each number represents a nuclear migration event and marks a mitotic cell.

Scale bar, 20 μm.

(B) Quantification of INM events during a 230-min window while epithelial cells remained

columnar, starting 20 or 25 min after drug addition. Student's *t*-test.

(C) Quantification of migration speed of pooled apically directed INM. Student's *t*-test.

(D) Representative images from Videos S4 and S5 showing mitotic rounding, planar

division, and daughter cell insertion after treatment. The locations of mitotic cells in the

otic vesicle are indicated in (A) by the corresponding numbers. Arrows indicate parental

or daughter nuclei. Scale bar, 10 µm.

(E) Representative frames showing spindle orientation with GFP-EMTB (green) and H2B-

RFP (red). Scale bar, 10 µm.

(F) Statistical analysis of cell division angles. Each mitotic cell is represented as a dot in

the radial histogram. Cells with apically retained daughter nuclei are highlighted in red.

Figure 5. INM in zebrafish otic vesicle requires MTs and dynein.

(A, B) Section images of epididymal epithelium stained for α-tubulin (A, green), acetylated

tubulin (Ace-tu, B, green), pH3 (red), and DAPI (blue). Arrows indicate pH3$^+$ cells, and

arrowheads highlight microtubule bundles. Scale bars, 20 µm.

(C) Z-stack images of the same cell as in (B), showing Ace-tu (green) and DAPI (red)

staining.

(D) Video frame showing EMTB-GFP (green) and H2B-RFP (red). Scale bar, 20 µm.

(E) Nocodazole-induced microtubule disassembly. Zebrafish otic vesicle was stained for

α-tubulin (green) and pH3 (red). DAPI (blue) staining is also shown. Single-channel α-

tubulin images are displayed in grayscale at the bottom. Scale bar, 20 µm.

(F) Images from Videos S7 and S8 showing INM events (arrows) in the otic vesicle after

nocodazole or vehicle (DMSO) treatment. H2B-GFP (green) and RFP-F (red) are shown.

Time is indicated relative to drug addition. The arrowhead marks a cell arrested in M

phase from the beginning of imaging. Scale bar, 20 µm.

(G) Quantification of INM events during a 206-min window, starting 30 min after drug

addition. Data from three embryos per condition. Student's *t*-test.

(H) Images from Videos S9 and S10 showing INM events (arrows) in the otic vesicle. RFP or RFP-p50 was overexpressed under the hsp70 promoter (red). H2B-GFP was expressed to label chromatin (green). Each number represents an INM event in RFP-positive cells. Scale bar, 20 μm.

(I) Quantification of INM frequency over 300 min of imaging. Data from three embryos per condition. Student's *t*-test. Scale bar, 20 μm.

Figure 6. Nuclear migration is essential for M phase onset.

(A) Sections of epididymal epithelium stained for γ-tubulin (green), pH3 (red), and DAPI (blue). Centrosomes (arrowheads) localize near the apical surface during nuclear migration, while centrioles move to opposite sides of the nucleus (arrows) only after it reaches the apical side.

(B) Frames from Video S6. Nuclei labeled with H2B-RFP (red) and stable microtubules labeled with EMTB-GFP (green) show MTOC/centrosomes (arrowheads) migrating to opposite sides of the nucleus (arrows) after apical migration to form the spindle.

(C) Sections of epididymal epithelium stained for cyclin B1 (red), pH3 (green), and DAPI (blue). Scale bar, 20 μm.

(D) Quantification for images shown in (C). Subcellular localization of Cyclin B1 grouped by position of nuclei in the basal, middle and apical parts of the cell. Blue = cytoplasmic, red = nuclear Cyclin B1 expression. n = 299 cells from three mice.

Fig. 7. Summary of INM and cell division in simple columnar epithelium.

A schematic summarizing apically directed INM, which is driven by dynein in late G2. During prometaphase, actomyosin mediates mitotic rounding by constricting the basolateral membrane, while lateral adhesions to neighboring interphase cells are strengthened. Mitotic cells maintain attachment to the basal lamina during planar division.

Supplemental Information

Legends for supplemental Videos

Video S1. Nuclear migration and mitosis in zebrafish otic vesicle. Fertilized embryos were microinjected with *in vitro* transcribed mRNA to express H2B-GFP, labeling chromatin. Imaging of the otic vesicle began at approximately 22 hpf and continued for 170 minutes at 2-minute intervals.

Video S2. Morphological changes of a mitotic cell in the zebrafish otic vesicle. H2B-GFP (green) labels chromatin, while RFP-F (red or gray) labels the plasma membrane. The arrow highlights the nucleus, and arrowheads indicate the lateral plasma membrane.

Video S3. F-actin dynamics in a mitotic cell of the zebrafish otic vesicle. H2B-RFP (red) labels chromatin, and GFP-hUtCH (green or gray) labels F-actin. The arrow marks the nucleus, while arrowheads point to the lateral plasma membrane.

Video S4. Nuclear migration and mitosis in the zebrafish otic vesicle following DMSO treatment. Embryos were microinjected with mRNA to express H2B-RFP (red) and GFP-hUtrCH (green). Imaging started approximately 20 minutes after DMSO was added at 22 hpf.

Video S5. Nuclear migration and mitosis in the zebrafish otic vesicle following Blebbistatin treatment. Embryos were microinjected with mRNA to express H2B-RFP (red) and GFP-hUtrCH (green). Imaging began approximately 25 minutes after Blebbistatin was added at 22 hpf.

Video S6. Microtubule dynamics in an otic vesicle cell entering the cell cycle. H2B-

RFP (red) labels chromatin, and EMTB-GFP (green) labels stable microtubules.

Video S7. Control otic vesicle treated with DMSO. Embryos were microinjected with

mRNA to express H2B-GFP (green) and RFP-F (red). Imaging commenced

approximately 30 minutes after DMSO was added at 22 hpf.

Video S8. Otic vesicle morphology following nocodazole treatment. Embryos were

microinjected with mRNA to express H2B-GFP (green) and RFP-F (red). Imaging started

approximately 30 minutes after nocodazole was added at 22 hpf.

Video S9. A control otic vesicle overexpressing RFP. The zebrafish embryo was

microinjected with both mRNA to express H2B-GFP (green) and a plasmid to overexpress

RFP (red) in a heat shock-inducible manner. The heat shock treatment was performed at

20 hpf for 1hr. The otic vesicle was imaged at 1.5 hr after the heat shock was done.

Video 10. A representative otic vesicle overexpressing RFP-p50. The zebrafish

embryo was microinjected with both mRNA to express H2B-GFP (green) and a plasmid

to overexpress RFP-p50 (red) in a heat shock-inducible manner. The heat shock

treatment was performed at 20 hpf for 1hr. The otic vesicle was imaged at 1.5 hr after the

heat shock was done.

Figure 1. INM in zebrafish otic vesicle epithelium begins in late G2.

(A) Schematic of the zebrafish otic vesicle at approximately 22 hpf.

(B) Morphology of the otic vesicle epithelium at 22 hpf and 28 hpf. Embryos were microinject-ed with mRNA to express H2B-GFP (chromatin, green) and RFP-F (cell membrane, red). The boxed regions indicate areas used for INM analyses. Scale bar, 20 µm.

(C) Selected frames from Video S2. H2B-GFP (green) and RFP-F (red) label chromatin and the plasma membrane, respectively. Single-channel images of RFP-F are shown in the middle. White arrows indicate the nucleus or chromosomes of the tracked cell, white arrow-heads highlight its basal protrusion and plasma membrane, and yellow arrows denote the cleavage furrow. The cell boundary and nucleolus of the tracked cell are outlined below. Scale bar, 20 µm.

(D, E) Nuclear trajectories before prometaphase (D) and after telophase (E). For daughter nuclei, only one that could be clearly tracked was analyzed.

(F) Velocity of pooled nuclear movements.

(G) Estimation of G2 length. Embryos at 24 hpf were labeled with BrdU, fixed at the indicated time points, and stained to visualize BrdU (red), pH3 (green), and DAPI (blue). Representative S-phase cells (arrowheads) and mitotic cells (pH3-positive, arrows) are shown. Yellow dashed lines indicate the basal membrane. Scale bar, 10 µm.

Fig. 2

Figure 2. INM in the mouse epididymis resembles that in the zebrafish otic vesicle.

(A) Schematic of the anatomical structure of the mouse epididymis.

(B) Quantification of mitotic events per epididymal tubular cross-section. Mitotic cells were identified based on chromosome morphology and/or strong pH3 staining (see C). The number of analyzed tubular cross-sections is indicated above each histogram.

(C) Section images showing nuclear positioning at different cell cycle stages. Insets display pH3 (green) or DAPI (red) staining of the indicated nuclei. Yellow lines mark the apical surface. Scale bar, 20 μm.

(D, E) Section images and quantification (E) showing S-phase and G2-phase nuclei 1 hour after BrdU labeling. Scale bar, 20 μm.

(F, G) Section images and quantification showing the distribution of BrdU-positive nuclei or chromosome clusters at the indicated time points. Arrows in (F) indicate representative nuclei or chromosome clusters. Scale bar, 20 μm.

Figure 3. Cell shape dynamics and basal lamina attachment during INM and mitosis.
(A-C) Section images of mouse epididymis epithelium showing cell-cell interfaces labeled with the indicated markers (green). Insets show pH3 staining (magenta) in G2 or prophase cells. Arrows indicate G2 or M-phase cells, and arrowheads point to the narrow basal protrusions of mitotic cells. DAPI staining is shown in red. Fluorescent intensity of F-action signals in (A) is shown at the bottom. Scale bars, 10 μm.
(D) Selected frames from Video S3 showing GFP-hUtCH (green) and H2B-RFP (red). Single-channel GFP-hUtCH images are shown in grayscale at the bottom. Arrows indicate the nucleus or chromosomes of the tracked cell, while arrowheads highlight its basal process and plasma membrane. Fluorescent intensity of GFP signals is shown at the bottom. Scale bar, 10 μm.

Fig. 4

Figure 4. Myosin II activity regulates mitotic cell rounding, spindle orientation, and epithelial Integrity.
(A) Selected frames from Videos S4 and S5 showing INM events. H2B-RFP (red) and GFP-hUtrCH (green) are shown. Time is indicated relative to the addition of DMSO or blebbistatin. Each number represents a nuclear migration event and marks a mitotic cell. Scale bar, 20 μm. **(B)** Quantification of INM events during a 230-min window while epithelial cells remained columnar, starting 20 or 25 min after drug addition. Student's *t*-test. **(C)** Quantification of migration speed of pooled apically directed INM. Student's *t*-test. **(D)** Representative images from Videos S4 and S5 showing mitotic rounding, planar division, and daughter cell insertion after treatment. The locations of mitotic cells in the inner ear are indicated in (A) by the corresponding numbers. Arrows indicate parental or daughter nuclei. Scale bar, 10 μm. **(E)** Representative frames showing spindle orientation with GFP-EMTB (green) and H2B-RFP (red). Scale bar, 10 μm. **(F)** Statistical analysis of cell division angles. Each mitotic cell is represented as a dot in the radial histogram. Cells with apically retained daughter nuclei are highlighted in red.

Fig. 5

A α-tu/pH3/DAPI   **B** Ace-tu/pH3/DAPI   **C** z-stacks Ace-tu/DAPI   **D** →anterior EMTB-GFP H2B-RFP ↓ventral 20 μm

20 μm

E α-tubulin pH3 DAPI

DMSO   Noc

20 μm

G P = 0.0006

INM events

DMSO Noc
n = 38   5

I P = 0.0061

% cells with INM

RFP RFP-p50
n = 48   38

F   30'   40'   58'   80'   118'   152'   186'   234' anterior

dorsal

DMSO — H2B-GFP RFP-F

20 μm

Noc

H   0'   52'   92'   118'   126'   174'   184'   208'   248'   262'   288'   300'

RFP

H2B-GFP

RFP-p50

20 μm

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