

Embryology of a planktonic tunicate reveals traces of sessility

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A key problem in understanding deuterostome evolution has been the origin of the chordate body plan. A biphasic life cycle with a sessile adult and a free-swimming larva is traditionally considered ancestral in chordates with subsequent neotenic loss of the sessile adult stage. Molecular phylogenies challenged this view, suggesting that the primitive life cycle in chordates was entirely free-living as in modern day larvaceans. Here, we report the precise cell lineage and fate map in the normal embryo of the larvacean *Oikopleura dioica*, using 4D microscopy technique and transmission electron microscopy. We document the extraordinary rapidity of cleavage and morphogenetic events until hatching and demonstrate that—compared with ascidians—fate restriction occurs considerably earlier in *O. dioica* and that clonal organization of the cell lineage is more tightly coupled to tissue fate. We show that epidermal cells in the trunk migrate through 90°, reminiscent of events in ascidian metamorphosis and that the axis of bilateral symmetry in the tail rotates in relation to the trunk. We argue that part of the tail muscle cells are ectomesodermal, because they are more closely associated with prospective epidermis than with other tissues in the cell lineage. Cladistic comparison with other deuterostomes suggests that these traits are derived within tunicates strengthening the hypothesis that the last common ancestor of tunicates had a sessile adult and thus support traditional morphology-derived scenarios. Our results allow hypothesizing that molecular developmental mechanisms known from ascidian models are restricted to fewer, yet identifiable, cells in *O. dioica*.

heterochrony | larvacea | *Oikopleura dioica* | ontogeny

A key problem in understanding deuterostome evolution has been the origin of the tadpole-like chordate body plan. Ascidians are sessile tunicates with larval stages that possess chordate features such as notochord or dorsal nerve tube. Larvaceans are also tunicates but display these typical chordate features throughout their entirely planktonic life. Traditionally, a biphasic, ascidian-like life cycle with a free-swimming larva inherited from a deuterostome ancestor had been considered ancestral in chordates with subsequent neotenic loss of the sessile adult stage (1–5). Molecular phylogenies challenged this view and suggested that the primitive life cycle in chordates was entirely free-living as exemplified by modern day larvaceans (6–8). Thus, larvaceans are pivotal for the understanding of chordate evolution and figured prominently in discussions about the role neoteny might have played in the evolution of chordates (9–12). However, whereas cell lineage and molecular aspects of tissue restriction have been intensely studied in ascidians, culminating in the deciphering of the first metazoan whole-embryo gene regulatory network (13, 14), lack of comparable knowledge in larvaceans has hindered a deeper understanding of chordate ontogeny and evolution (2, 4, 15, 16). Here, we report the precise cell lineage and fate map in the embryo of the larvacean *Oikopleura dioica*, using 4D microscopy technique and transmission electron microscopy. We document the extraordinary rapidity of cleavage and morphogenetic events and demonstrate that—compared with ascidians—fate restriction occurs consid-

erably earlier in *O. dioica* and that clonal organization of the cell lineage is more tightly coupled to tissue fate. We show that epidermal cells in the trunk migrate through 90°, reminiscent of events in ascidian metamorphosis, and that the tail rotates through ontogeny in relation to the trunk. Comparison with other deuterostomes shows that cell lineage characteristics are derived within tunicates supporting the hypothesis that the last common ancestor of tunicates had a sessile adult stage. Our results allow us to hypothesize that molecular developmental mechanisms known from ascidians are restricted to fewer, yet identifiable, cells in *O. dioica*. The present study reveals the simplest cell lineage (in the sense of ref. 17) known from metazoans and amounts to a major step forward in tracing evolutionary changes in cellular mechanisms that will ultimately facilitate understanding molecular mechanisms correlated with drastic changes in life history strategies.

Results and Discussion

To facilitate comparisons among animals, we adopted the nomenclature established by Conklin (18) for tunicate ascidians instead of the one introduced later by Delsman (15) for appendicularians. See *Materials and Methods* for explanation.

Early Cleavage. Cleavage in *O. dioica* is stereotyped and characterized by rapid divisions, early fate determinations, and overall bilateral symmetry [Fig. 1, supporting information (SI) Fig. S1, and Movies S1 and S2]. The first cleavage, 15 min after fertilization, in principle separates left from right (Figs. 1 and 2; see next paragraph). The second cleavages, 5 min after the first, occur longitudinally at right angles to the first one (Fig. 2). The third round of cell divisions is at right angles to both previous planes and separates animal from vegetal (Figs. 2 and 3). After the initial three cleavages, a stereoblastula is formed. The blastula stage lasts until after the fifth round of cell divisions, when most adult tissue types can be assigned to specific fate-restricted cells (Fig. 4). At the ventral vegetal side of the embryo, an exceptionally asymmetric cell division from B5.2 (B5.2) to B6.3 and B6.4 (B6.3, B6.4) takes place, B6.3 (B6.3) being bigger than B6.4 (B6.4). At this stage embryos can easily be oriented.

Cell Divisions and Body Axes. The planes of the first three cleavages in *O. dioica* coincide with embryonic axes at least to late neurula stages. The first cleavage separates left from right (Figs. 1 and 2), but, during later stages, cells cross the bilateral symmetry axis

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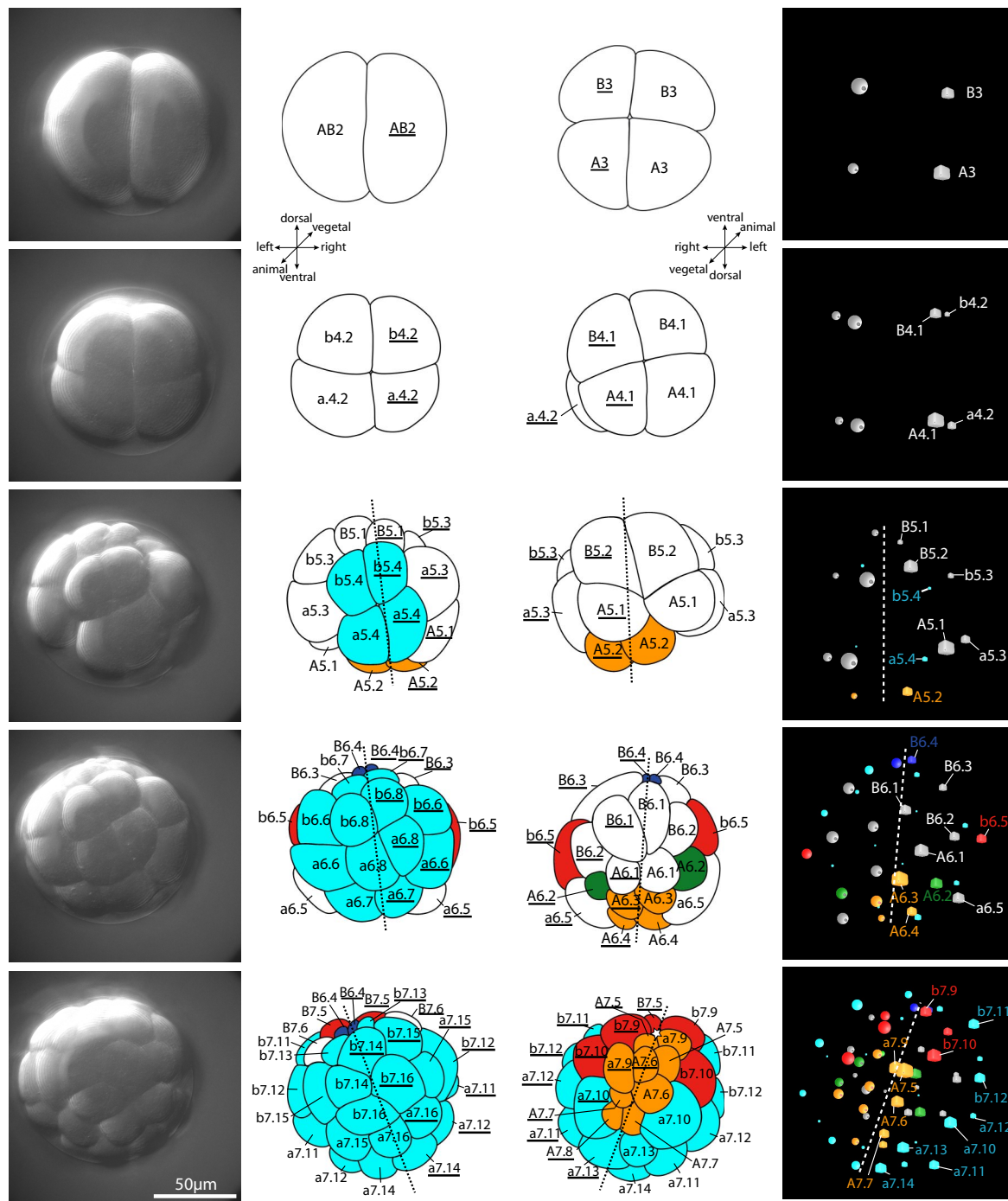


Fig. 2. First cleavages in the ontogeny of the tunicate *Oikopleura dioica*. (Left) Projections of the upper (animal) half of DIC-stacks recorded for 4D microscopy. (Line drawings in Left) View from animal pole. (Line drawings in Right) View from vegetal pole. Asterisk marks the position of the blastopore. (Right) 3D representation of the positions of the nuclei of each cell in the 4D microscopy software SIMI*BioCell as seen from the vegetal pole. Cubes derive from the left side (AB2), spheres from the right (AB2). Nomenclature and color code for tissues that are already restricted as in Fig. 1. Dashed line indicates axis of bilateral symmetry. Note bilateral symmetry in these early embryonic stages.

term “meso/endoderm.” Based on the detailed anatomical knowledge derived from the TEM investigation, we could assign almost every cell of the hatchling to a tissue fate that was then traced back by using 4D microscopy to result in the cell lineage fate map (Fig. 1, Fig. S1, and Movie S1) and 3D representation of nuclei in time (Figs. 2–4).

Clonal organization of the tissues is essentially invariant among individuals (Fig. S1), and fate restriction occurs as early

as the 16-cell stage, where $a5.4$, $b5.4$ and their counterparts $a5.4$, $b5.4$ are destined to become epidermis, and $A5.2$ and $A5.2$ are restricted to nervous fate. Most lineages achieve clonal restriction at the 62-cell stage, many already at the 32-cell stage. Epidermis, albeit being entirely animal in derivation, has a variety of clonal precursors ($a5.4$, $a6.6$, $a7.10$, $b5.4$, $b6.6$, and their counterparts). The majority of cells in the nervous system derive from vegetal cells $A5.2$ and $A5.2$; only a few derive from the

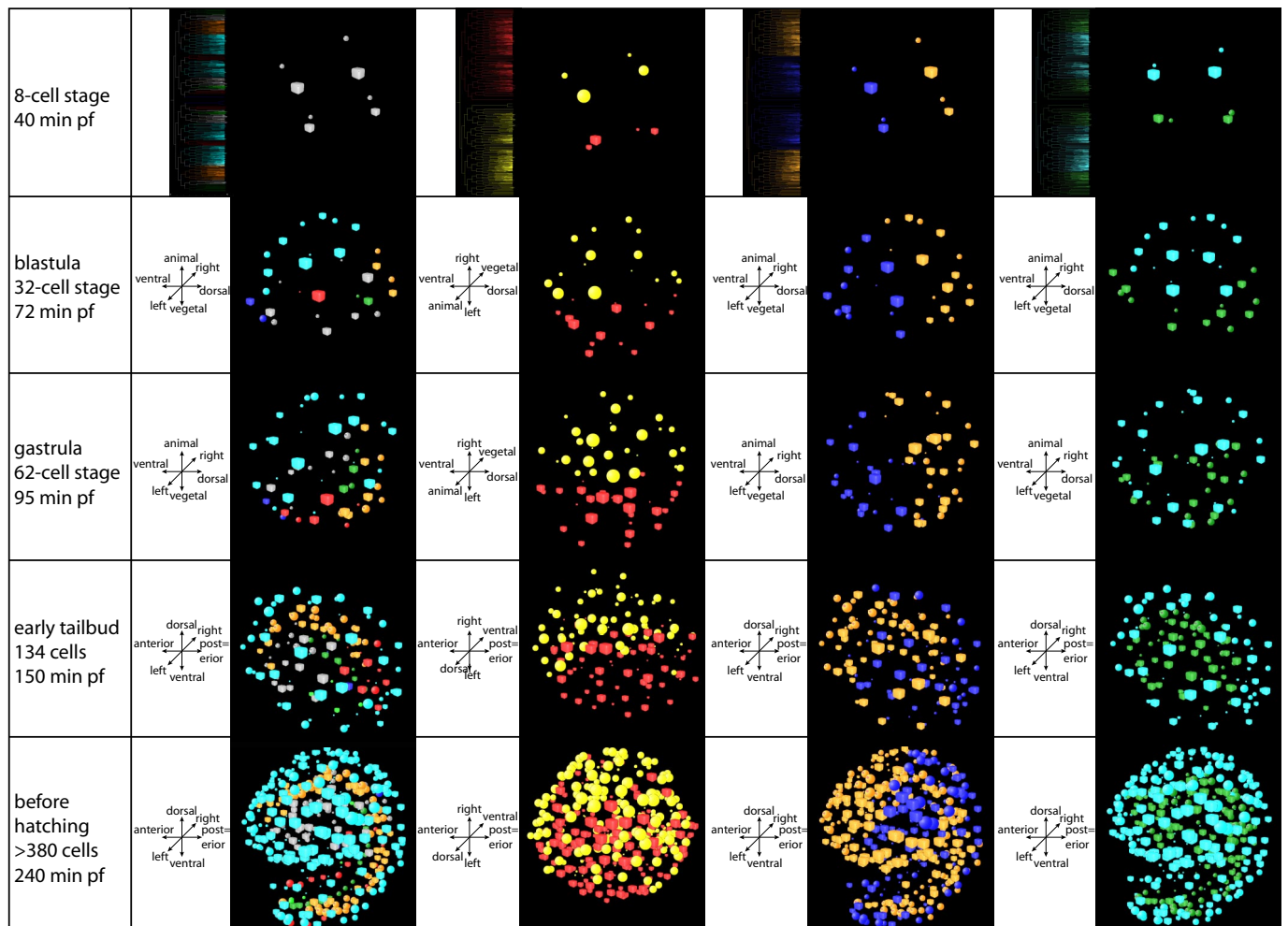


Fig. 3. Virtual cell tracing experiments, using the 3D representation of the positions of the nuclei of each cell implemented in the 4D microscopy software SIMI^oBioCell. Cubes derive from the left side (AB2); spheres derive from the right side (AB2). (Second column) Colors indicate tissues as in Fig. 1. (Third column) Red cubes derive from the left side (AB2); yellow spheres derive from the right side (AB2). Note that the orientation in this panel differs from the remaining ones to give a better view of the dorsal midline in the prehatching state. (Fourth column) Orange shapes derive from dorsal cells A3 and A3, traditionally labeled “anterior” (see text for details); blue shapes derive from ventral cells B3 and B3, traditionally labeled “posterior.” Note that the center of orange colored nuclei shifts from dorsal to anterior between gastrula and early tailbud stage. (Far right column) Blue shapes derive from the animal cells a4.2, b4.2, a4.2, and b4.2; green shapes derive from the vegetal cells A4.2, B4.2, A4.2, and B4.2.

animal a7.9 and a7.9. Similarly, tail muscle cells are in part vegetal (B7.5, B8.11, B7.5, and B8.11) and part animal (b6.5 and b6.5) in origin. Muscle cells and notochord cells display a stereotyped pattern of complex morphogenetic movements ending up in precise and invariant positions in the hatchling (Fig. S4). Vegetal muscle cells end up anteriorly in the tail and remain in their respective bilateral sides. Animal muscle cells become more posteriorly situated and—like many epidermal cells—partly cross the bilateral symmetry axis. What we termed meso-/endoderm is entirely vegetal in origin, but we were unable to assign a more specific fate to these cells. Part of the vegetal progeny, the lines originating in B7.6 and B7.6 can be assigned to the endodermal strand in the tail. B6.4 and B6.4 derive via a strongly asymmetric cleavage from B5.2 and B5.2 and arrest cleavage until hatching, characteristic for germ line cells (in one individual one division in the B6.4 line was seen 20 min before hatching). These conspicuous small cells are situated in the ventral midline in early stages and invaginate during gastrulation via the ventral blastopore lip. Delsman, in the classic description of the ontogeny of *Oikopleura dioica*, was unsure whether these cells were not merely plasmaprotusions (15). Later, these cells become situated in the posterior trunk and are impacted by the

tail rotation, because the right cell is more dorsally located than the left. On the dorsal vegetal side, eight notochord cells derive from A6.2 and A6.2, and, on the right side, an additional pair derive from A8. Thus, using 4D microscopy, we identified 18 notochord cells in the hatchling, 19 using TEM, and 20 that were described later as juveniles (15, 16, 26, 27). Notochord cells position themselves along the embryonic midline and start intercalating ≈ 2.5 h pf. Intercalation proceeds from anterior to posterior and is completed ≈ 3.5 h pf. It remains unclear where the two remaining cells needed to complete juvenile cell number originate.

The cell lineage and fate map of *O. dioica* share similarities with those of ascidians, such as rapid and precise determinative cleavage, general bilateral symmetry of cleavage pattern, and the derivation of nervous system and musculature from both vegetal and animal lineages (18–20). However, closer inspection reveals also major differences between larvaceans and ascidians. Fate restrictions occur considerably earlier, and clonal organization of the cell lineage is even more tightly coupled to tissue fate in *O. dioica*. For example, tail muscle cells derive from four distinct lineages in the ascidian embryo (A8.16, B7.4, B7.5, B7.8, and b8.17) (13, 19), whereas, in *O. dioica*, no A-cell line gives rise to musculature. Similarly, whereas the nervous system derives from

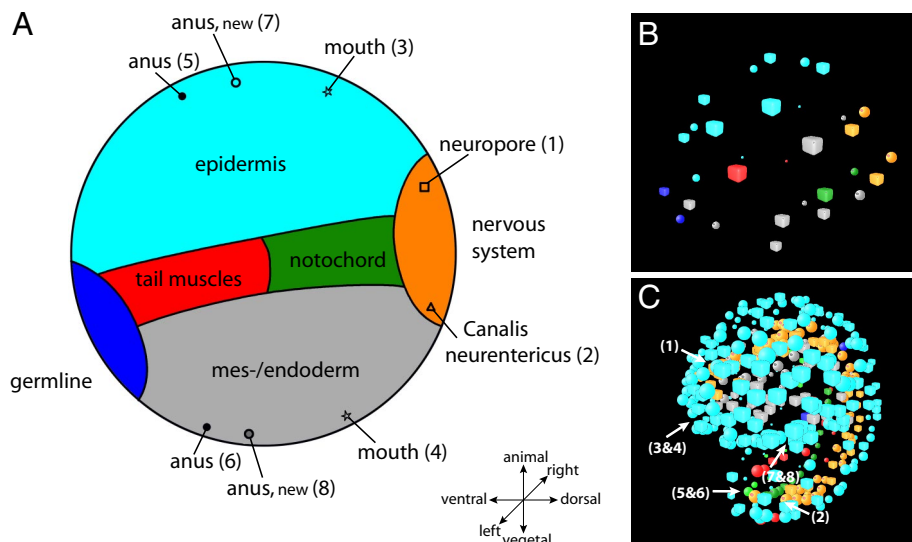


Fig. 4. Virtual tracing of chordate landmarks. (A) Schematic projection of the prospective fate in *Oikopleura dioica* as derived from the blastula (B). Positions of anatomic landmarks that are indicated in the larva (C) are also indicated in the schematic projection. (B) Representation of nuclei with colors coding for prospective tissue fates (see Fig. 1). (C) Representation of nuclei in the larva with positions of anatomic landmarks indicated that are depicted in the schematic fate map in A.

A-, a-, and b-lines in ascidians (19, 28, 29), in *O. dioica*, nervous tissue originates in the lineages A5.2 and a7.9, but no b-line contribution to the nervous tissue was found. Another example is seen in the development of the notochord; in ascidian embryos, the anterior 32 primary notochord cells derive from A7.3, A7.7, and the posterior eight secondary notochord cells derive from B8.6 (13, 30). In *O. dioica* notochord originates only in A-cell lines, indicating the absence of secondary notochordal cells. In *Ciona intestinalis*, notochordal cells in the A-line depend on activation of brachyury, which depends on ZicL and Fgf9/16/20 activation. ZicL is activated by FoxA-a and FoxD, and Fgf9/16/20 is mediated by phosphorylation of the ETS-containing transcription factor ets/pointed2 (13). Although we would predict that a similar network is responsible for the determination of A6.2, A6.2, and A8.1 in *O. dioica*, it will be interesting to see whether the slightly different circuit that results in secondary notochordal cells in *C. intestinalis* (13, 30) is missing or silenced in *O. dioica* like genes in the case of the tailless ascidian larva of *Molgula occulta* that is evolutionarily derived from a tailed ancestor (31, 32). Based on our results, it is now possible to target molecular networks for comparative analysis elucidating ontogenetic mechanisms in the evolution of drastically divergent life cycles.

Larvaceans have been used as proxies for the suggested role heterochrony might have played in the evolution of the chordate body plan (2, 4, 33), but Ruppert (34) used a cladistic approach to argue that heterochrony occurred in the stem lineage leading to tunicates and that developmental speed was even more derived in larvaceans. Comparison of our data to available data from other deuterostomes (Table S1 and Fig. S16) supports Ruppert's conclusions and strengthens the hypothesis that larvaceans are derived within tunicates. The observed rotation of the tail can be used as an informative phylogenetic character to place larvaceans more precisely as the sister group to aplousobranch ascidians (35, 36).

The observed acceleration in the development of larvaceans is not only achieved by shortening cell cycles or lowering number of cleavages at which specific events occur, but the lineage tree is altered. This finding is consistent with the hypothesis that selection for decreased cell lineage complexity played a major role in the evolution of metazoan ontogenies (17, 37). Although *Caenorhabditis elegans* had been considered a prime example of

mosaic development (38), modern investigations have demonstrated that its ontogeny is based on principles of regional specification rather than clonal organization of cell lineage (39, 40). Thus, although tunicates also show regulative cell interactions during ontogeny, their fate map is tightly coupled to cell lineage and in that sense remain extreme examples of deterministic (mosaic) development, with *Oikopleura dioica* as the most extreme example.

Materials and Methods

For recordings, two- or four-cell embryos were mounted on a microscope slide coated with 0.01% polylysine. Fundamentals of 4D microscopy are described by Schnabel *et al.* (40). A Zeiss Axioplan Imaging 2 microscope with internal focus drive was used to move the temperature-controlled stage to record a z series with a Hamamatsu Newvicon camera. Images were digitized with an Inspecta 3 frame grabber (Mikroton) and compressed with a wavelet function (Lurawave). The microscope is controlled with a software programmed by K. Schulz and R. Schnabel. Embryos were recorded at 15°C (measured on stage) until hatching. Recordings of three different embryos—all of which were developing normally and, after hatching, actively swimming—stemming from three different parental pairs were analyzed by using SIMI[®]BioCell software (SIMI). Z projections of the upper half of the DIC-stacks recorded at specific times were generated in ImageJ software (<http://rsb.info.nih.gov/ij>).

The anatomy of a hatched larva was analyzed by using TEM. Fixation was in 1.5% glutaraldehyde in 0.1 M phosphate buffer (pH 7.4) adjusted to 950 mosM with NaCl followed by a postfixation in 1% OsO₄. Araldite was used as embedding resin, and a complete series of ultrathin sections prepared. Analysis was on a Philips Biotwin CM 120 (FEI) and a virtual 3D reconstruction was prepared by using 3Ds Studio Max (Autodesk).

In this article, we used the standard nomenclature established for tunicate ascidians by Conklin (18). In a study of *Oikopleura dioica*, Delsman (15) introduced a different nomenclature. We provide a translation table for the two systems in Table S2 and compare the two systems in Fig. S1. We preferred the Conklin system, because by adopting Conklin's nomenclature the similarity of the fate maps becomes immediately obvious and allows detailed evolutionary comparisons.

Note. While this article was under review, a description of the early development up to the 64-cell stage of *O. dioica* was published by Fuji *et al.* (41). Note that these authors use Delsman's nomenclature as explained here and in Table S2.

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