Formation of the vulva in *Caenorhabditis elegans*: a paradigm for organogenesis

Ranjana Sharma-Kishore¹, John G. White²*, Eileen Southgate² and Benjamin Podbilewicz¹,²‡

¹Department of Biology, Technion-Israel Institute of Technology, Haifa, 32000, Israel
²MRC Laboratory of Molecular Biology, Hills Road, Cambridge, UK
*Present address: Laboratory of Molecular Biology and Department of Anatomy, University of Wisconsin, Madison, WI 53704, USA
‡Author for correspondence (e-mail: podbilew@tx.technion.ac.il)

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SUMMARY

The genes involved in the inductive interactions that specify cell fates in the vulva of *Caenorhabditis elegans* are known in some detail. However, little is known about the morphogenesis of this organ. Using a combination of cell biological and anatomical approaches, we have determined a complete morphogenetic pathway of cellular events that lead to the formation of the vulva. These events include reproducible cell divisions, migrations, remodeling of adherens junctions, cell fusions and muscle attachments. In the course of these events, an epithelial channel comprising a stack of 7 toroidal cells is formed that connects the internal epithelium of the uterus with the external body epithelium, forming the vulva. Vulval muscles attach to the epithelial channel and the whole structure everts during the final molt. The mature vulva has rotational, two-fold symmetry. Using laser microsurgery, we found that the two halves of the vulva develop autonomously.

Key words: Morphogenesis, Vulva, *Caenorhabditis elegans*, Cell fusion, Cell migration, Adherens junctions, Cell fate

INTRODUCTION

Morphogenesis is one of the most striking manifestations of the development of a multicellular organism, but one that is as yet poorly understood. Organismal morphogenesis is often modular, producing discrete, non-overlapping organs such as the eye or kidney. Focused studies of organogenesis allow knowledge of development to be acquired piecemeal. In vertebrates, the development of organs such as the kidney (Saxen and Sariola, 1987), bone (Reddi, 1994), tooth (Thesleff and Nieminen, 1996) and placenta (Cross et al., 1994) have been subjected to considerable scrutiny. These studies have revealed insights into the process of tissue patterning and are beginning to provide information about mechanisms that specify cell identity. Developmental genetic studies of organisms such as *Drosophila melanogaster* or *Caenorhabditis elegans* have revealed signal transduction pathways that specify cell identity (Freeman, 1997; Kornfeld, 1997). In particular, studies directed towards the development of the eye in *Drosophila* (Banerjee and Zipursky, 1990) and the vulva in *C. elegans* (Horvitz and Sternberg, 1991) have revealed the inductive interactions and the signaling pathways that are used to generate the diversity of component cells that go to make up these organs (Greenwald and Rubin, 1992). Organogenesis in animals involves the remodeling of epithelia into organs (Trinkaus, 1984). In this study, we have determined the precise remodeling of 22 epithelial cells into an organ. We have endeavored to complement the considerable genetic studies that have been made on the *C. elegans* vulva by determining the cellular basis of organogenesis of the vulva.

Background

Vulval development involves very few cells, occurs over a period of 20 hours, can be observed in living animals and can be analyzed by genetic and molecular methods. During the first larval stage, a linear sequence of 12 ventral hypodermal cells P(1-12).p is generated. In wild-type hermaphrodites, P(1-2).p and P(9-11).p fuse with the surrounding syncytial hypodermal cell (hyp7) (Sulston and Horvitz, 1977). Six vulval precursor cells (VPCs; P(3-8).p) are competent to generate vulval cells (Sulston and White, 1980). A somatic gonadal cell, the anchor cell (AC), induces the nearby P5.p, P6.p and P7.p to adopt vulval cell fates (Kimble, 1981) and the remaining three VPCs usually divide once and fuse with hyp7 (3° sublineage) (Kimble, 1981; Sulston and White, 1980). A somatic gonadal cell, the anchor cell (AC), induces the nearby P5.p, P6.p and P7.p to adopt vulval cell fates (Kimble, 1981) and the remaining three VPCs usually divide once and fuse with hyp7 (3° sublineage) (Kimble, 1981; Sulston and White, 1980). VPCs induced to adopt a vulval fate signal laterally to each other causing P5.p and P7.p to adopt a 2° sublineage and P6.p to adopt a 1° sublineage (Greenwald, 1997; Sternberg and Horvitz, 1989). Each sublineage (1°, 2° and 3°) consists of a distinct pattern of cell divisions generating a characteristic number and types of cells. Without the inductive signal from the AC, all VPCs fuse to hyp7 and no vulva is formed (Kimble, 1981; Sulston and White, 1980). There might be an independent signal from hyp7 dependent on the products of lin-15 locus that represses a vulval fate or promotes a hyp7 3° (fused) fate (Herman and Hedgecock, 1990). The 2° sublineages require the product of...
lin-12 gene in order to occur while the 1° sublineage is independent of lin-12 activity (Greenwald et al., 1983).

Classic and molecular genetic analyses have shown that the inductive signal from the anchor cell is a molecule similar to mammalian growth factors encoded by the gene lin-3. A receptor tyrosine kinase (LET-23) on the surface of the VPCs is proposed to bind LIN-3 and activate a signal transduction cascade involving LET-60 Ras and MPK-1/SUR-1 MAP kinase. MAPK and its targets in C. elegans appear to be required for the decision between vulval and epidermal fates (for review, see (Greenwald, 1997). Epidermal fates in nematodes involve cell fusion as an important developmental decision during the morphogenesis of many organs (Baird et al., 1991; Podbilewicz and White, 1994; Sommer, 1997; Wang et al., 1993).

MATERIALS AND METHODS

Nematode strains, growth in liquid culture and synchronization of populations

Wild-type (N2) C. elegans was cultured as described (Brenner, 1974). A one-liter liquid culture of nematodes was grown and collected by sucrose floatation. Eggs were obtained by digesting this population with a final concentration of 1-2% alkaline hypochlorite and allowed to hatch overnight in liquid medium without food. The L1s so generated were collected by centrifugation, washed in M9 and allowed to grow in liquid medium with food. Aliquots were collected at 2 hour intervals (from L3 to the adult stage) and the specific populations were collected by sucrose floatation (Sulston and Hodgkin, 1988).

Permeabilization and fixation of worms

The different worm aliquots were resuspended in fixation buffer (Finney and Ruvkun, 1990) and 20% para-formaldehyde (Sigma) was added to a final concentration of 1%. All permeabilization, fixation and subsequent antibody-staining reactions were performed in 1 ml Eppendorf tubes (Finney and Ruvkun, 1990). A fraction of the worms were fixed using methanol-acetone, with the same results (Podbilewicz, 1996; Podbilewicz and White, 1994).

Immunofluorescence and confocal microscopy

Aliquots of the different fixed worm populations were washed with PBS-BSA (0.1%)-Triton X-100 (0.5%). Cell adherens junctions were visualized by staining with the monoclonal antibody MH27 (Francis and Waterston, 1991) and nuclei were stained with propidium iodide (Podbilewicz, 1996; Podbilewicz and White, 1994). The specimens were analyzed using a MRC-1024, laser confocal scanning microscope (Bio-Rad, Hempstead, UK) with the objective Nikon Plan Apo 60x/1.40. Projections were obtained from 10-20 serial optical sections collected every 0.5 μm, in the z-axis. Figures are oriented with the anterior of the worm on the left-hand side of the page and are ventral views unless specified. Developmental timings were estimated from the size of animal and from studying the developing vulva with Nomarski optics using a Nikon Eclipse E-800 microscope. For all descriptions, at least 20 animals were observed and all developed as described, except where noted. Cell lineages were followed using Nomarski optics (Sulston and Horvitz, 1977). Serial EM reconstructions and laser microsurgery were performed as described (Sulston and White, 1980; White et al., 1986).

RESULTS

Formation of the vulva

Morphogenesis of the vulva in the wild-type hermaphrodite

The process of vulva formation can conveniently be split into three phases. In the first phase (execution and short-range migrations), 22 epithelial cells are produced and migrate relative to their neighbors to take up their final configuration and form a tube containing 7 rings (toroidal cells) that join the endoderm of the uterus to the external epithelium (Fig. 1). The second phase is characterized by cell fusions within the 7 rings, vulval muscle attachments at defined regions and specific vulval cell attachments to lateral epithelial cells. The second phase covers most of the L4 stage. In the third phase (eversion), the vulva partially evets (turns inside out) at the L4 to adult molt and the lumen closes to block the transit of eggs until it is opened by activation of the vulval muscles.

To study developing organs, we obtained 3-dimensional reconstructions of confocal optical sections of vulvae stained with an antibody that recognizes the adherens junctions (apical surfaces and boundaries between cells) together with serial section electron microscopy. This approach reveals the construction of the organ involving short-range cell migrations and cell fusions (Podbilewicz and White, 1994).

First phase of vulva formation (execution and short-range migrations)

To study cell migrations, cell fusions and formation of cell junctions with the hypodermis, we reconstructed intermediates in the early stages of vulva morphogenesis. The first phase covers the division of the vulval precursor cells (VPCs) to the termination of the vulval lineages at the end of the L3 stage (Figs 2, 3).

P3.p, P4.p and P8.p are ventral epithelial cells that have the potential to generate vulval cells (Sulston and White, 1980) but in wild-type animals they divide and fuse with hyp7, the larger hypodermal syncytial cell that surrounds the VPCs. Induced P5.p, P6.p and P7.p undergo two rounds of division and form a linear row of 12 epithelial cells situated on the ventral midline (Fig. 1). EM and confocal reconstructed vulval primordia after the first and second rounds of VPC division exhibited no signs of differentiation. Fig. 2 shows confocal reconstructions of the apical domains of vulval precursors stained with the MH27 monoclonal antibody (Francis and Waterston, 1991) that binds at, or close to, adherens junctions during (Fig. 2A) and after (Fig. 2B,C) the second round of longitudinal division. In summary, during the first two divisions all cells divide longitudinally and there is an apparent bilateral symmetry; however, there is some degree of asymmetry between the anterior and posterior halves of the vulvae (e.g. Fig. 2B). During the terminal (third) round of divisions, six cells divide transversely, two cells do not divide and four divide longitudinally (Fig. 1). Fig. 2 shows examples of VPC granddaughters during the third round of division. The precursors of vulE (P6.ppp and P6.paa) are the first to divide transversely (Fig. 2D) and in general the anterior half of the vulvae develops faster than the posterior half (Fig. 2E).

One EM reconstruction of a vulval primordium at the intermediate stage (not shown) in which the first terminal divisions of the VPCs had just occurred shows four daughter cells that migrate towards the center of symmetry of the vulva (where the anchor cell is situated). The four cells form a square configuration (precursor to vulE) with a hole in the middle displacing the central two cells dorsally (vulF). Few signs of morphogenesis were detected in the other vulval cells in this EM reconstruction and in similar intermediates reconstructed with the confocal microscope (Fig. 2F). Fig. 3 shows ventral views...
of intermediates in different stages of early morphogenesis and ring formation. Fig. 3B shows an intermediate vulva reconstructed with the confocal microscope that has a central ring precursor containing four cells (vulF), four cells that surrounded the central ring form the next ring (vulE) and a third ring with two cells (vulD). The next two cells on each side (vulC precursors) have divided transversely and are sending processes towards the center of symmetry of the vulva surrounding vulD. The outer cells on each side are the precursors of vulA that fused (see Fig. 3A before cell fusions) and will be the last to migrate. A second EM reconstruction of a vulva just after the terminal round of divisions revealed intermediate stages of the short-range migrations (Fig. 4A,B ventral and lateral views respectively). The cells that will form vulC and vulD had sent processes that displaced the adherens junctions that were between the central vulval cells (the precursors of rings vulE and vulF) and the hypodermal cell (hyp7). Precursors to the more distal vulval rings (vulA, vulB1 and vulB2) were further behind developmentally with vulB2 sending out only a small extension and vulA showing no morphological changes (see also Fig. 3B). The innermost cells (the daughters of P6.pap and P6.ppa), making up the vulF ring formed a rectangle with the anchor cell nestling into a pocket in the middle (Fig. 4B). At slightly later times, the anchor cell breaks completely through the center of the group of four cells in the middle of the primordium thus forming the vulF ring. This second EM reconstruction and confocal reconstructions (e.g. Fig. 3) revealed how the series of stacked epithelial rings that make up the vulva is formed out of a row of 22 epithelial cells. These observations show that short-range cell migrations in the vulva primordium involve rearrangements of adherens junctions, as shown for other migrations in <i>C. elegans</i> (Podbilewicz and White, 1994), sea urchin (Hardin, 1989), <i>Drosophila</i> (Grawe et al., 1996) and other organisms (Yap et al., 1997).

In a temporal sequence, starting in the center i.e. proximal to the anchor cell, vulval primordial cells (VPC grand and great-granddaughters; Fig. 1) send out unidirectional extensions towards the center of the primordium that displace the adherens junction between hyp7 and the inner, neighboring precursor cell. This process begins with the vulE cells, each of which sends out two lateral extensions towards the vulval midline. Towards the center of the primordium, these extensions become stacked up, with the processes from the outer cells running ventral to those of their inner neighbors. These process-like extensions interdigitate between the two displaced cells, forming their own adherens junctions giving rise to a ‘Y’ configuration of junctions at the apex (Figs 3, 4). This structure is situated right out at the tip of the extending process suggesting that the act of opening up and reforming the junctions may be part of the driving force for process extension.

In summary, organogenesis begins with the final round of

**Fig. 1.** Schematic representation of the cellular events that occur during the morphogenesis of the vulva in *C. elegans*. The top diagram (lateral view) represents a larva in the L3 stage. A somatic gonadal cell, the anchor cell (AC), through inductive signaling induces three (P5.p, P6.p and P7.p) of the six vulval precursor cells (VPCs), P3.p to P8.p, to adopt vulval fates. The non-vulval cells - P3.p, P4.p and P8.p undergo a single proliferation step and fuse with the underlying syncytial cell. The remaining three VPCs signal laterally to each other to adopt fates that involve distinct patterns of cell divisions and generate specific numbers and types of cells. The cells of the vulva primordium perform short-range migrations from the outer regions towards the central midline, in a symmetrical fashion and also undergo fusions with specific partners. This involves the cells adjacent to the innermost cells forming processes and migrating around them (i.e., precursor cells of the ring vulE send processes around the precursor cells of the inner ring vulF). This process is repeated in a temporal sequence by the next outer cells that send processes around their inner neighbours, which meet at the midline. This results in the formation of a stack of 7 toroids (vulA to vulF) at the center. This is followed by fusion between the component cells of each ring, in a specific and temporal order. The resultant structure is an epithelial tube that connects the uterus of the worm to the outside. The anchor cell (black circle in the center) eventually penetrates the inner ring (vulF) and fuses with a uterine cell (open white circle at the center). Diagrams are ventral views, the anterior of the worm on the left-hand side of the page and represent the apical domains of cells unless specified. The same color code for the different cell types and the vulval rings has been maintained for all the figures. The vulva in the L4 stage (last diagram; lateral view) appears like an invagination from the ventral side of the worm. Times given are hours from hatching at 20°C (Sulston and Horvitz, 1977).
division when symmetrical groups of cells on either side of the four central cells (precursors of vulF; Fig. 3) send out processes between hyp7 and their inner neighbors. The precursors of vulE (the daughters of P6.paa and P6.ppp; Fig. 5) migrate ventrally and laterally towards the midline of the vulva where they form a ring of 4 cells connected by adherens junctions. The next ring of epithelium is formed when precursors of vulD migrate surrounding the first ring. In a similar fashion, more concentric rings of epithelium are formed. The process is sequential from inner to outer groups until 7 rings have formed.

Second phase of vulva formation (cell fusions within the 7 rings)
The anchor cell invades the center containing 4 cells making a hole in the middle that results in the formation of a central ring (called vulF); this ring attaches to the uterus via adherens junctions to the uv1 cells. The anchor cell then fuses (Fig. 4C) with the precursors of the utse syncytium of the gonad (Newman et al., 1996). Cell fusions occur among cells within the rings (Figs 6, 7). The order in which the cells within the rings fuse is vulD, vulA, vulC, vulF and vulE. vulD is a binucleate cell and the pair of cells within vulB1 and vulB2 remain unfused, the other rings are tetranucleate. These 7 concentric rings are stacked (Fig. 5). The ring vulE sends processes to the seam lateral epithelial cells, vulval muscles attach to specific sites and vulA makes junctions with the syncytial hypodermal cell hyp7 (Figs 8, 9).

The confocal reconstructions in Figs 6 and 7 show intermediates containing 7 rings that have completed the migrations and are in different stages of the cell fusions within each concentric toroidal ring. For example, the most external vulval ring adjacent to the external epithelium (vulA) was formed by the migration of the most external cells in Fig. 3. These cells fuse in pairs before they migrate (Fig. 3B), their processes meet in the center and form two adherens junctions between them (Fig. 6A). In an intermediate step between Fig. 6B and C, the two junctions disappear because the two binucleate cells fuse at two points forming the syncytial ring vulA, a cell containing four nuclei.

The epithelial rings of the vulva are attached to each other by means of circular adherens junctions (Figs 6, 7). These junctions, in common with those described in other epithelia (Podbilewicz and White, 1994), demarcate the apical and basal domains of epithelial cells. The organization of these junctions was reconstructed from EMs of serial sections (Fig. 4C-E) and in the light microscope by immunofluorescent staining with the MH27 monoclonal antibody (Fig. 6). The apical domains of the toroids are situated on their inside surfaces and together make up the internal luminal surface of the vulva.

The toroid at the dorsal end of the stack (vulF) is attached to the uterus (Newman et al., 1996). Four small, extended epithelial cells (uv1) arranged in a rhomboid configuration attach to vulF. The four uv1 cells in turn are attached to a large syncytial uterine cell (utse) that has nine nuclei. The bulk of this cell is situated on each side of the uterus and appears to be used to attach the uterus to the lateral epithelial cells (Fig. 8).

Fig. 2. Proliferation of vulval precursor cells. Confocal micrographs of early L3 vulvae stained with the nuclear stain propidium iodide (red) and MH27 mAb (green) that reveals apical domains of cells (Francis and Waterston, 1991; Podbilewicz, 1996; Podbilewicz and White, 1994). Pictures were obtained by projecting 5-10 serial confocal sections of 0.5 μm. All estimated times are in hours after hatching. (A) Ventral view of vulval primordium showing the 12 resultant grand-daughters of the VPCs, P5.p, P6.p and P7.p, after two rounds of cell division; grand-daughters of P5.p and P7.p are still undergoing cytokinesis; a, b1, b2, c, d, e and f are the precursor cells of the future vulval rings – vulA to vulF, respectively (approximately 31.5 hours). (B) Oblique view of an asymmetric vulva. The posterior precursor of E (P6.ppp) is twice as long as the anterior precursor of E (P6.paa). (C) Symmetric primordium with the 12 granddaughters of the VPCs. (D) Oblique view of 14 cell intermediate; precursors of vulE have divided, resulting in four E cells. (E) Ventral view of asymmetric primordium with precursors of vulA and vulBx having divided in the anterior half, resulting in the cells B1 and B2, and two A cells. (F) Ventral view of symmetrical 18 cell primordium where the terminal divisions of A, B1, B2 and E are completed (32.5 hours), before the terminal divisions of vulC and vulF precursors. Arrowhead, centre of symmetry. Scale bar, 10 μm.
L4s and early adults show that cuticular material is secreted in a polarized way to the lumen of the vulva and the vulval labia are formed. Confocal reconstructions of evert ing vulvae show that vulB1 and vulB2 remain unfused \((n=5; \text{ data not shown})\). After vulva eversion, the lumen closes preventing the transit of eggs until it is opened by the action of the vulval muscles. The structure of the adult vulva is accomplished without major relative cell movements. The final structure is reached by plastic deformations of the immature L4 vulva from the second phase.

In summary, the vulva in the fourth larval stage (L4) of C. elegans consists of a tube made of a stack of 7 toroidal cells with the anchor cell in the apex of this tube connecting to the uterus (Fig. 5). During the L4, the anchor cell fuses to the utse uterine multinucleated cell causing the opening of the tube (Figs 4C-E, 7). The vulva then everts to form a mature adult vulva through which eggs can be laid.

**Laser microsurgery shows that each side of the vulva develops autonomously**

The developing vulva has a mirror-symmetric structure. As we have described above, during vulval morphogenesis cells from each side of the center line send two extensions towards the midline until they meet their symmetrical partners on the other side, thereby forming the stack of rings. It appeared as if each cell could be attracted towards its symmetric partner. However, we observed that 27\% \((n=55\) vulvae from the first phase) of the vulvae analyzed had developing rings containing asymmetric divisions and migrations (e.g. Figs 2B,E, 3B). In general, the anterior half of the vulva developed faster suggesting that the two halves of each ring precursor develop autonomously. We tested this possibility by removing one half of the cells of a developing primordium with a laser microbeam. We killed the posterior three cells of a six-cell primordium (just after the first division of the VPCs). The half vulva appeared to develop normally except that the terminal division of the P6.pap cell (giving rise to the cells of the vulF ring) was longitudinal instead of transverse. To analyze the cell migrations and cell fusions within the semi-vulva, this animal was reconstructed from electron micrographs of serial sections. We found that the two components of the vulF cell looked normal in that the anchor cell was deeply embedded in them and they made adherens junctions to uterine cells. Thus, the cells seemed to have acquired the same differentiated state in spite of being daughters of a longitudinal division instead of the normal transverse division. In addition, all the cells had sent out two extensions in the normal way, to what would have been the mid-line of the vulva (Fig. 4F) However, instead of contacting their symmetrical partners, they had grown around the primordium until they contacted themselves (Fig. 9C). In several toroids, these processes had actually fused forming a complete ring (Fig. 9D). Therefore, we propose that, in the absence of a corresponding ring precursor partner on the opposite side, the two processes can continue migrating until they contact each other. This can prevent their further migration and eventually promotes 'self' fusion. In summary, the events that can occur in the absence of half the precursors are cell migrations, stacking of 7 rings, fusions within the rings and generation of a functional semivulva. Taken together, the resulting semivulva and the high incidence of asymmetric vulvae suggest a model in which at least two positive signals, one originating from the adjacent dorsally located ring and the other coming from the ventrally located hypodermal
syncytium, attract the growing processes to open the junctions between them. The physical interaction between the opposing tips of the growing processes constitutes a third signal in our model for the arrest of further migration.

Summary of sequence of events in vulva formation

Our studies, along with previous genetic and cell ablation studies (reviewed in Greenwald, 1997), suggest that the following sequence of events occurs during the organogenesis of the vulva:

(1) An EGF-like signal (LIN-3) from the anchor cell defines three adjacent epithelia cells, P5.p, P6.p and P7.p (or their six daughters) as vulval precursors (possibly by inhibiting fusion). P3.p, P4.p and P8.p, which are more distal from the AC, divide once and their daughters fuse with the hypodermal syncytium hyp7.

(2) Lateral signaling among the vulval precursors using a LIN-12 signal (and possibly other unidentified signals) establishes a set of six cell fates (A, B, C, D, E and F). The cell fates are established in a palindromic sequence along the single row of vulva precursor cells (Figs 1, 5).

(3) The terminal divisions of C, E and F are transverse, this being orthogonal to the previous divisions of the precursors. This event marks the start of vulva morphogenesis. The anchor cell is positioned in the center of the four ventral vulF cells and starts to open up a central pore marking the beginning of lumen formation (Figs 1, 5).

(4) In a proximal-distal sequence, cells produced by the vulval lineage migrate towards the center of the vulva primordium wrapping themselves around their proximal neighbors until they meet their symmetrical homologues on the other side of the vulva. In so doing, the central apex of the vulval primordium moves inwards (i.e. dorsally) to make a conical structure (Fig. 5). We have shown...
that this process occurs independently in each half of the vulva. Furthermore, EM studies of the multivulva mutant lin-15(e1763) have demonstrated that this process does not require the presence of the anchor cell (unpublished observations). It therefore seems as if a given cell in a sequence is programmed to wrap around its proximal neighbor in the sequence. This is the central process that transforms a linear array of epithelial cells into a hollow, conical vulva primordium.

(5) Concomitantly with the cell migrations described above, a stereotyped sequence of homotypic cell fusions occurs transforming the vulva primordium into a sequence of toroidal syncytia (Figs 1, 7).

(6) The anchor cell undergoes a heterotypic cell fusion to a large uterine syncytium (utse) opening up the lumen of the vulva to the uterus (although a thin membrane of utse separates the vulval from the uterine lumens until the first eggs are laid). At this time, the attachment of the vulva primordium to the uterus becomes consolidated by means of the uterine uv1 and utse cells.

(7) Vulval muscles attach to specific regions of the vulva and the vulE cell makes attachments to the lateral hypodermal (seam cells) (Fig. 8). The inner cuticular lining of the vulva and cuticle on the labia start to be secreted at this time. The morphogenesis of the vulva is now essentially complete.

(8) The last phase of development (eversion) occurs during the final molt and involves plastic deformations of the structure to form a valve that is opened by the contraction of the vulval and uterine muscles to allow eggs to be laid.

**DISCUSSION**

**The cellular basis of pattern formation**

We have defined a developmental pathway responsible for the formation of a simple organ. In summary, we have characterized all the cellular migrations and fusions required for the assembly...
of a functional vulva. Previous work has characterized the molecular genetic basis of the fates of the six VPC cells (Greenwald, 1997; Kornfeld, 1997; Sternberg and Horvitz, 1989). Here we establish the cellular basis of vulval morphogenesis that provides a necessary developmental foundation for molecular genetic analyses of organ and pattern formation. We have shown that each half of the vulva develops autonomously implicating that interactions between the anterior and the posterior halves are not necessary for the execution of short-range migrations, toroid formation and intratorialid fusions.

Role of cell fusion in vulval morphogenesis

Cell fusion is a process involved in entry of viruses into cells, fertilization, conjugation, and formation of muscle, bone and placenta (reviewed in Hernandez et al., 1996). In *C. elegans*, 30% of all somatic nuclei are part of syncytia formed by cell fusion. We have previously characterized embryonic and postembryonic fusions in the hypoderm (Podbielwicz and White, 1994). In the development of the vulva, we observe six distinct types of topological cell fusion events:

(1) Heterotypic fusions of the daughters of P3.p, P4.p and P8.p with the syncytial hypodermis hyp7 (Fs) (Fig. 1). Experiments have shown that these cells are capable of acquiring vulval fates but fuse with hyp7 because they are beyond the range of the inductive influence of the anchor cell (Sternberg and Horvitz, 1986; Sulston and White, 1980).

(2) Fusions between two pairs of anterior-posterior cells before toroid formation (FIL: longitudinal fusions between vulA precursors Fig. 3).

(3) Homotypic transverse fusions (FT) between two pairs of vulC precursors (Figs 3, 5).

(4) Homotypic intratorialid longitudinal fusions (FIL) within vulA, vulC, vulD and vulF.

(5) Homotypic intratorialid transverse fusions (FIT) within vulE and vulF (Fig. 7).

(6) Heterotypic fusion of the anchor cell with utse (FAC) opening up a channel between the uterus and vulva (Figs 4, 7).

One of the functions of heterotypic cell fusion in *C. elegans* development seems to be to suppress the expression of specific differentiated states (Podbielwicz and White, 1994). This is likely to be the reason for the Fs fusions in (1). The FAC fusion in (6) could also be of this type only, in this case, the anchor cell fate has been expressed and may be only suppressed when the function of that cell is no longer required. Interestingly, evolutionary studies have shown that, in the development of the vulva in the nematode *P. pacificus*, cell death appears to be used as an alternative strategy to cell fusion (Eizinger and Sommer, 1997).

We conclude that AC and the descendants of the VPCs participate in the formation of the vulva resulting in:

(1) Specification of eight cell classes derived from the VPCs: vulA, vulB1, vulB2, vulC, vulD, vulE, vulF and hyp7.

(2) Formation of 7 syncytia: hyp7, vulA, vulC, vulD, vulE, vulF and utse.

The strategies used in the construction of the vulva in *C. elegans* are probably used in pattern formation during development of other organs. Very little is known about the mechanisms of migrations and fusions during the formation of intratorialid fusions.
different tissues in biological systems (Hernandez et al., 1996; Podbilewicz and White, 1994). Our results provide ideas for the analysis of organ formation and a way of correlating the lineage with the execution of cell migrations and cell fusions. Many pathological conditions in a variety of tissues contain syncytial cells most probably formed by cell fusion (Cross et al., 1994; Huang et al., 1993; Sitar et al., 1994). Gross vulval morphological defects have been identified in many mutants in C. elegans. The defective phenotypes have been described as reduced invagination at the mid-L4 stage, abnormal eversion of the vulva, ‘squashed’ vulva or development of abnormal vulval protrusion (Bettinger et al., 1997; Greenwald, 1997; Herman and Horvitz, 1997; Maloof and Kenyon, 1998; Seydoux et al., 1993). The exact nature of these defects in terms of cell-cell recognitions, interactions, fusions, migrations and attachments is not known. The detailed knowledge of short-range migrations, changes in cell shapes, cell fusions and the anatomical organization of the vulva allows a more precise description of the phenotypes of mutations that disrupt pattern formation in the vulva (unpublished observations).

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REFERENCES


