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Calcium signaling in tunicate development

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A comparative overview is provided of Ca²⁺ signaling and its potential mechanistic roles during development in tunicates. As background, the review presents an introduction to tunicate taxonomy, and then a general overview of Ca²⁺ signaling and methods for recording and measuring Ca²⁺ signals. It then covers the dynamics and implicated mechanisms of Ca²⁺ signals during different phases of development from oocyte to larva. These include signals arising in the unfertilized oocyte, signals associated with fertilization and meiosis, intercellular signals occurring from early cleavage stages through gastrulation, intercellular signals during organogenesis, and signals associated with early behavior. Comparisons are made among different tunicate species and where relevant to other chordate species. In many tunicate species, Ca²⁺ currents across the oocyte membrane are present prior to fertilization, and in the appendicularian *Oikopleura dioica* regular Ca²⁺ transients have been recorded optically prior to fertilization. Ca²⁺ signals at this stage have been implicated in pre-fertilization oocyte maturation events. The fertilization transient is the most well-studied Ca²⁺ signal and is triggered by factors from the sperm, including pivotally a phospholipase C (PLC) isoform that catalyzes the generation of IP₃, which elicits release of Ca²⁺ from the endoplasmic reticulum. Post-fertilization signals are similarly dependent on IP₃ signaling and are regulated by cyclin-dependent kinase 1 (Cdk1), and thereby linked to the meiotic divisions required for zygote formation. Ca²⁺ signals associated with early cleavages through gastrulation arise in blastomeres of the muscle lineage and spread from these in a coordinated fashion to other blastomeres through gap junctions. Post-gastrulation Ca²⁺ signals begin to show tissue-specificity in their temporal pattern as organogenesis proceeds, likely associated with loss of general gap junction transmission. Once neurulation has occurred, Ca²⁺ signals arise first in the nervous system and are transmitted synaptically to muscle, while Ca²⁺ signals arising spontaneously in the epidermis follow a separate temporal pattern. Species differences in the spatiotemporal characteristics of pre- and postgastrulation Ca²⁺ signals are discussed.

KEYWORDS

oocyte, blastula, gastrula, neurulation, *Oikopleura*, *Ciona*, ascidian, appendicularian

Introduction

Calcium signaling is a universal feature of animal development and influences many developmental processes starting from oocyte maturation and fertilization and extending through organogenesis and cell differentiation. It has been investigated extensively in several chordate species at various development stages (reviewed in Webb and Miller, 2003; Whitaker, 2008).

As the invertebrate chordate clade most closely related phylogenetically to vertebrates (Delsuc et al., 2006, 2008), and with a relatively small number of constituent cells, tunicates provide a favorable model for studies of calcium signaling during development (McDougall et al., 2012). Indeed, some of the early studies of calcium signaling during development were carried out in tunicates (Block and Moody, 1987; Bosma and Moody, 1990; Speksnijder et al., 1990; Dale et al., 1991; Coombs et al., 1992; Arnoult and Villaz, 1994). This review provides an overview of current knowledge about the developmental dynamics and function of calcium signaling in tunicates.

Tunicate diversity

Tunicates are highly diverse in morphology, ecology and behavior, and encompass both sessile and pelagic forms (reviewed in Holland, 2016). They are commonly categorized into three classes, the sessile Ascidiacea and the pelagic Thaliacea and Larvacea (the latter also termed Appendicularia), although more robust cladistic groupings based on large-scale genomic analyses are now recognized (Appendicularia, Stolidobranchia, and Thaliacea + Phlebobranchia +

Aplousobranchia, with the Ascidiacea being paraphyletic; Delsuc et al., 2018). Pending a more thorough resolution of evolutionary relationships, we use here the traditional (but admittedly outdated) taxonomic designations to illustrate general variations in developmental features. In ascidian, appendicularian and some thaliacean species, early development generates a free-swimming larval stage that exhibits the typical chordate body plan with a motile tail (some thaliaceans also employ sexual reproduction or an asexual budding reproductive mode that lacks a free-swimming larval stage). In appendicularians, the larval body plan is maintained in the adult (hence the name Larvacea), which remains individually mobile throughout life. Thaliaceans generally form pelagic colonies of tail-less adults, whereas ascidian larvae metamorphose into sessile bottom-dwelling adults with a sac-like body plan. Of relevance to the Ca²⁺ signaling events to be presented below, model tunicate species across taxa exhibit invariant cell lineages, most have a well-developed notochord, and a muscular tail is the main mode of locomotion.

Sources of Ca²⁺ signals

Intracellular Ca²⁺ signals (reviewed in Clapham, 2007) derive from two sources: 1) influx through the cell membrane mediated by Ca²⁺ channels, driven by the large electrochemical gradient for Ca²⁺, and 2) release from intracellular Ca²⁺ stores, principally the endoplasmic/sarcoplasmic reticulum but extending to other compartments such as mitochondria and lysosomes (Figure 1). Cell membrane influx can be mediated by voltage-gated Ca²⁺ channels (VGCs), mechanosensitive cation channels (MCCs), which pass both Ca²⁺ and other cations), ligand-gated Ca²⁺

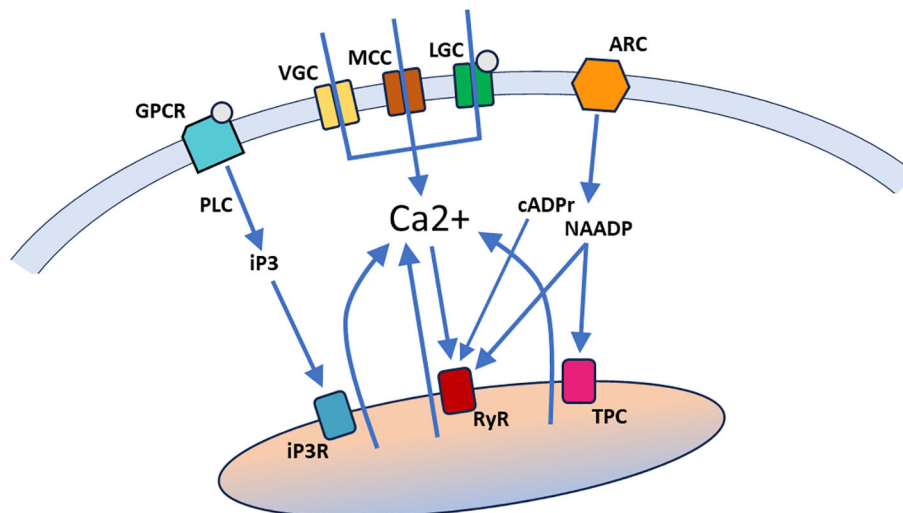


FIGURE 1

A general overview of Ca²⁺ signaling pathways. Ca²⁺ can enter the cell through ion channels in the cell membrane, including voltage-gated Ca²⁺ channels (VGC), mechanosensitive cation channels (MCC), and ligand-gated Ca²⁺ channels (LGC). Ca²⁺ can also be released from intracellular stores (indicated by a single organelle here that encompasses the various Ca²⁺ storing organelles), through binding of second messengers (iP3, cADPr, NAADP and Ca²⁺ itself) to specific receptors in the organelle membrane. Ligand binding by G-protein coupled receptors in the cell membrane (GPCR) activate phospholipase C (PLC) to generate iP3, which can bind to iP3 receptors (iP3R) in the endoplasmic/sarcoplasmic reticulum. Activation of ADP-ribosyl cyclases (ARC) in the cell membrane generates cADPr and NAADP, both of which bind to ryanodine receptors (RyR) in the endoplasmic/sarcoplasmic reticulum. NAADP also binds to two-pore channels (TPC) in acidic lysosomes.

channels (LGCs), and second messenger-activated Ca²⁺ channels, of which G-protein coupled receptors (GPCR) and calcium-activated Ca²⁺ channels are subcategories. Voltage-gated, mechanosensitive and ligand-gated Ca²⁺ channels have fast opening kinetics and generate inward Ca²⁺ currents of abrupt onset, variable duration (depending on the channel closing mechanisms) and localized action (since the Ca²⁺ concentration falls markedly within a few tens of nanometers from the inner surface of the cell membrane). Second messenger-activated Ca²⁺ channels are slower to open and close and typically generate more protracted Ca²⁺ signals.

Release from intracellular Ca²⁺ stores depends on second messenger systems and thus generates Ca²⁺ signals with a more gradual onset and relatively long durations. In addition, these signals can spread through a larger cytoplasmic volume due to the extent of the storage organelle(s) and the amplifying nature of the second messenger systems. The principal second messengers are inositol trisphosphate (IP₃), generated by different isoforms of phospholipase C (PLC), cyclic ADP-ribose (cADPr) and nicotinic acid adenine dinucleotide phosphate (NAADP), generated by ADP-ribosyl cyclases (ARCs), and Ca²⁺ itself (Galione, 2019; Shah et al., 2022; Kim, 2022; Guse, 2023). To effect Ca²⁺ release, IP₃ binds to IP₃ receptors (IP₃Rs) on the endoplasmic/sarcoplasmic reticulum, cADPr, NAADP and Ca²⁺ bind to ryanodine receptors (RYRs) on the endoplasmic/sarcoplasmic reticulum, and NAADP also binds to two-pore channels on acidic lysosomes (Figure 1). The full picture of regulated release from intracellular Ca²⁺ stores is more complex, but the general overview in Figure 1 provides sufficient context for the discussion of Ca²⁺ signal mechanisms in tunicate embryos below.

Methods for studying calcium signaling

Calcium signals involve the movement of Ca²⁺ ions across organelle and cell membranes by dint of calcium channels, pumps and antiporters, the diffusion of Ca²⁺ ions through gap junctions and over limited distances within the cytoplasm, and the binding of Ca²⁺ ions to specific calcium buffer and signaling proteins. Because Ca²⁺ is a universal intra- and intercellular signaling molecule, involved in a myriad of cell functions including cytoskeletal dynamics, cell adhesion, synaptic transmission and muscle contraction, its movement and distribution is highly regulated.

Methods for registering and measuring calcium signals take advantage of either the ionic charge of Ca²⁺ or the binding of Ca²⁺ to specific calcium-binding fluorophores or fluorescent proteins. In the former case, electrodes can be used to measure Ca²⁺ currents and/or resultant membrane potential changes with very high temporal resolution (limited by the electronics used, typically at sub-microsecond levels). A commonly used technique is patch clamp recording, which can measure Ca²⁺ currents through large populations of Ca²⁺ channels (as in whole cell recordings) and down to single channels (if only a single or a very few channels are opened or closed within the patch of membrane recorded from) (Cuomo et al., 2006; Gallo et al., 2013; Carvacho et al., 2018). Other types of electrode-based recording can also be used. For example, a

vibrating probe system was developed to measure Ca²⁺ current densities across cell membranes (Jaffe and Nuccitelli, 1974), including associated with the activation current following fertilization of fish oocytes (Nuccitelli, 1987).

Calcium binding to calcium-sensitive fluorophores and fluorescent proteins has been utilized to transform Ca²⁺ signals into fluorescent signals that can be captured optically (Paredes et al., 2008; de Melo Reis et al., 2020). Calcium-sensitive fluorophores such as Fura-2, Fluo-4 and Calcium Green are organic molecules which, when photo-excited, emit fluorescence of specific wavelengths in a manner that is calcium-dependent. The kinetics of their response to changes in Ca²⁺ levels vary over a domain spanning milliseconds, limiting the temporal resolution obtainable relative to electrophysiological measurements, but still capable of discriminating signals such as broad calcium action potentials or protracted synaptic potentials. Most calcium-sensitive fluorophores (and fluorescent proteins, see below) only report relative changes in Ca²⁺ concentration. Fura-2 is an example of a calcium-sensitive fluorophore with a bimodal excitation spectrum, which permits ratiometric measurements that can be used to quantitate the concentration of Ca²⁺ ions at a given time point (Paredes et al., 2008).

Calcium-sensitive fluorescent proteins exist in nature and are intrinsically fluorescent or bioluminescent with an intensity that is dependent on Ca²⁺ concentration. Aequorin was the first such protein to be used for imaging Ca²⁺ signals, particularly in connection with the fertilization of oocytes (Ridgway et al., 1977; Steinhardt et al., 1977). More recently, genetic engineering has been used to create specially designed calcium-sensitive fluorescent proteins through combination of calcium-binding protein domains with an intrinsically fluorescent protein (Zhang et al., 2002). The advantage of engineered calcium-sensitive fluorescent proteins is that their expression can be genetically targeted to specific cell types. The GCaMP family of calcium-sensitive fluorescent proteins has become a particularly popular tool for imaging Ca²⁺ signals in a variety of organisms and cells (Kettunen, 2020; Shemesh et al., 2020) and encompasses variants of different sensitivities and kinetics. In particular, GCaMP has been used to assess Ca²⁺ signals during early embryogenesis in both vertebrates (Chen et al., 2017) and tunicates (Akahoshi et al., 2017; Mikhaleva et al., 2019; Tolstenkov et al., 2022). As is the case for calcium-sensitive fluorophores, calcium-sensitive fluorescent proteins have much slower kinetics than electrophysiological measurements but nevertheless can in many cases discriminate individual Ca²⁺ signals, especially when combined with temporal deconvolution methods (de Melo Reis et al., 2020).

Calcium signaling from oocyte to first cleavage

Pre-fertilization

Calcium is known to play an important role in oocyte maturation and the progression through meiotic arrest (Whitaker, 2008; Gallo et al., 2013). In line with this, Ca²⁺ currents and signals have been detected in ascidian and appendicularian oocytes prior to

fertilization (Block and Moody, 1987; Moody and Bosma, 1989; Bosma and Moody, 1990; Dale et al., 1991; Coombs et al., 1992; Arnoult and Villaz, 1994; Silvestre et al., 2009; Tosti et al., 2011; Gallo et al., 2013; Mikhaleva et al., 2019). All but the most recent of these studies employed electrophysiological measurements to demonstrate the presence in the oocyte membrane of either mechanosensitive non-selective cation channels (which pass Ca^{2+} in addition to Na^{+} and K^{+}) or voltage-sensitive Ca^{2+} channels, as well as other voltage-sensitive monovalent cation channels. These studies characterized channel types and kinetics and showed that Ca^{2+} currents depended on extracellular Ca^{2+} but did not describe the developmental dynamics of pre-fertilization calcium signals. Mikhaleva et al. (2019) employed Fluo-4 or GCaMP6 to visualize these signals in unfertilized oocytes of the appendicularian *Oikopleura dioica* and showed that regularly repetitive Ca^{2+} transients occur at a frequency of about one every 2-3 minutes. The transients have a stereotypical double-peaked waveform consisting of a minor peak lasting about 3 sec followed after a delay of 10-15 sec by a 20-fold higher amplitude major peak lasting 8-10 sec. They are always initiated at a focal point near the inner surface of the oocyte and spread from there throughout the oocyte as a wave.

It is unclear what initiates these pre-fertilization signals in *Oikopleura*, in particular whether the trigger is intrinsic or extrinsic, such as mechanical or chemical stimulation associated with spawning. In this context it is of interest to compare how appendicularian and ascidian oocytes are obtained experimentally prior to calcium imaging. In *Oikopleura*, natural spawning in free-swimming adults involves wholesale rupture of the gonads, a traumatic and life-terminating event. In the lab this can be induced by mechanical agitation. In both cases, oocytes are violently released en masse, likely resulting in substantial

mechanical stimulation, although some protection of the oocyte membrane is afforded by the chorion. By contrast, in ascidians, natural and laboratory-based spawning involves a gradual, enzyme-mediated release of individual follicles from the ovary of sessile adults (Matsubara et al., 2019), which likely has a much milder mechanical impact on the oocytes. Mikhaleva et al. (2019) discuss the possibility that mechanical stimulation of oocytes arising from induced spawning, impalement with the electrodes used to inject GCaMP6 mRNA or the dechoriation necessary to allow loading with Fluo-4 might be stimuli that initiate pre-fertilization signals in *Oikopleura*. Irrespective of how these signals are initiated in the experimental context, it is clear that they do not depend on fertilization and exhibit regular, clock-like dynamics and a stereotypical waveform that are not disrupted by the fertilization transient (although there may be a resetting of the clock; Figure 2; Mikhaleva et al., 2019).

Fertilization transient

Calcium plays a major role in polyspermy block at fertilization, by dint of the generation of a large, global Ca^{2+} transient that arises shortly after sperm contact, which initiates a chain of events that creates a barrier to penetration by later-arriving sperm. The fertilization transient is arguably the most studied Ca^{2+} signal in development and has been characterized in many invertebrate and vertebrate species (reviewed in Stricker, 1999; Nixon et al., 2000; Runft et al., 2002; Whitaker, 2006; Carvacho et al., 2018). In tunicates, the fertilization transient has been studied extensively in ascidians, including *Ciona robusta* (*Ciona intestinalis* type A), *Ciona intestinalis* (*Ciona intestinalis* type B), *Ciona savignyi*, *Ascidella aspersa* and *Phallusia mamillata* (Hirano and

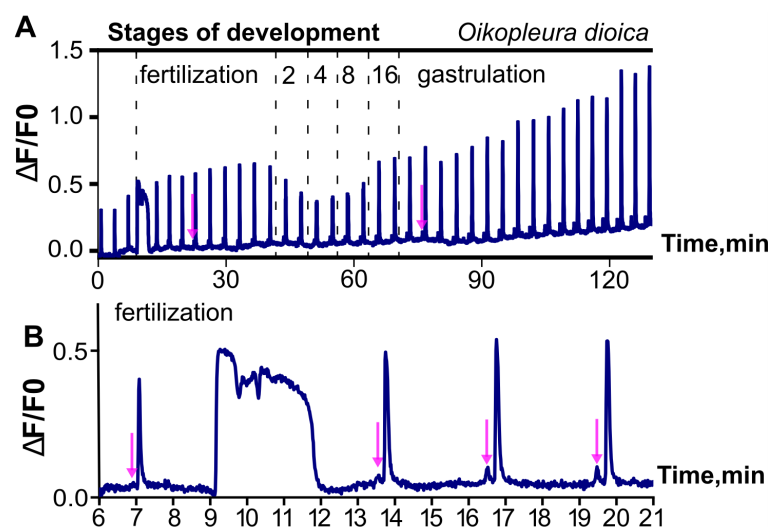


FIGURE 2

Illustration of ongoing, regular Ca^{2+} signals from pre-fertilization to gastrula stages of *Oikopleura dioica*. Ca^{2+} signals are represented as fractional changes of fluorescence (F) over a baseline ($\Delta F/F_0$). (A) Note the highly regular, repetitive nature of the signals, which begin in the unfertilized oocyte and continue as coordinated intercellular signals as the embryo develops, interrupted only by the fertilization transient. (B) Higher magnification highlights the much longer time course of the fertilization transient, and the double-peak waveform of the regular calcium signals, which begin with a small "mini-peak" (indicated by arrows in both A, B) followed by a main peak. Modified from Mikhaleva et al. (2019).

Takahashi, 1984; Dale and DeFelice, 1984; Speksnijder et al., 1989; Brownlee and Dale, 1990; Speksnijder, 1992; Goudeau and Goudeau, 1993; Goudeau et al., 1994; Lambert et al., 1994; McDougall et al., 1995; Cuomo et al., 2006; Tosti et al., 2011; reviewed in Dumollard et al., 2004). It has the following general features conserved across tunicate species: 1) it is initiated at the site of sperm entry; 2) it involves multiple sources of Ca²⁺, including influx of extracellular Ca²⁺ through voltage-gated and potentially other types of Ca²⁺ channels in the oocyte membrane and a primary IP₃-mediated release of Ca²⁺ from the endoplasmic reticulum triggered by sperm-derived factors introduced into the egg cytoplasm; 3) it lasts for about 3–5 min (reviewed in Dumollard et al., 2004; Whitaker, 2008). The rise of Ca²⁺ concentration during the fertilization transient is necessary for subsequent events that prevent polyspermy, such as the release of N-acetylglycosaminidase, which inactivates sperm receptors in the vitelline coat (Lambert et al., 1994; McDougall et al., 1995). In addition, ion depletion experiments show that an inward Na⁺ current contributes to the fertilization transient in *Ciona*, and that when Na⁺ is removed extracellularly the transient is diminished and development is defective (Tosti et al., 2011).

As in other species (both vertebrate and invertebrate), the fertilization transient in ascidians is triggered by the PLC-mediated generation of IP₃ and binding of IP₃ to IP₃R_s on the endoplasmic reticulum (reviewed in Runft et al., 2002; Dumollard et al., 2004). In mouse and human, injection of the PLC zeta 1 isoform (PLCζ1), triggers the fertilization transient, and genetic knockout of PLCζ1 in mouse sperm prevents it (reviewed in Carvacho et al., 2018). It has been speculated that PLCζ1 is also the relevant sperm factor in tunicates (Dale et al., 2010), but this has not yet been confirmed. In ascidian oocytes, injection of IP₃ is sufficient to initiate a fertilization transient, and specific blockade of PLC gamma (PLCγ) or Src protein tyrosine kinases prevents the fertilization transient, implicating these enzymes in the IP₃ transduction signal (Runft and Jaffe, 2000). By contrast, inhibition of RyR does not prevent the fertilization transient (reviewed in Runft et al., 2002). Thus, the primacy of the IP₃ pathway evidently is conserved in tunicates and non-tunicate species, although the PLC isoforms involved differ.

Post-fertilization signals prior to first cleavage

Most studies that have characterized the fertilization transient in tunicates note that it is followed by a series of highly regular Ca²⁺ signals during ensuing meiosis and cleavage (Figure 3, Speksnijder et al., 1989; Speksnijder, 1992; Mikhaleva et al., 2019; Matsuo et al., 2020). In the appendicularian *Oikopleura*, the regular post-fertilization signals are indistinguishable in frequency and waveform from the pre-fertilization signals (Figure 1; Mikhaleva et al., 2019). Moreover, in ascidians, the post-fertilization signals are similar to the pre- and post-fertilization signals in *Oikopleura* in that they are regular and have a much shorter time-course than the fertilization transient (reviewed in Dumollard et al., 2004).

However, they appear to lack the doublet waveform seen in *Oikopleura* (Mikhaleva et al., 2019; an absence that may be artefactual depending on the temporal resolution of the Ca²⁺ recording used). The similarity of pre- and post-fertilization transients across tunicate classes suggests that they represent the same phenomenon and may occur independently of the fertilization transient rather than being necessarily triggered by it. Indeed, it is clear in *Oikopleura* that the fertilization transient arises on a background of regular signals that spans from pre- to post-

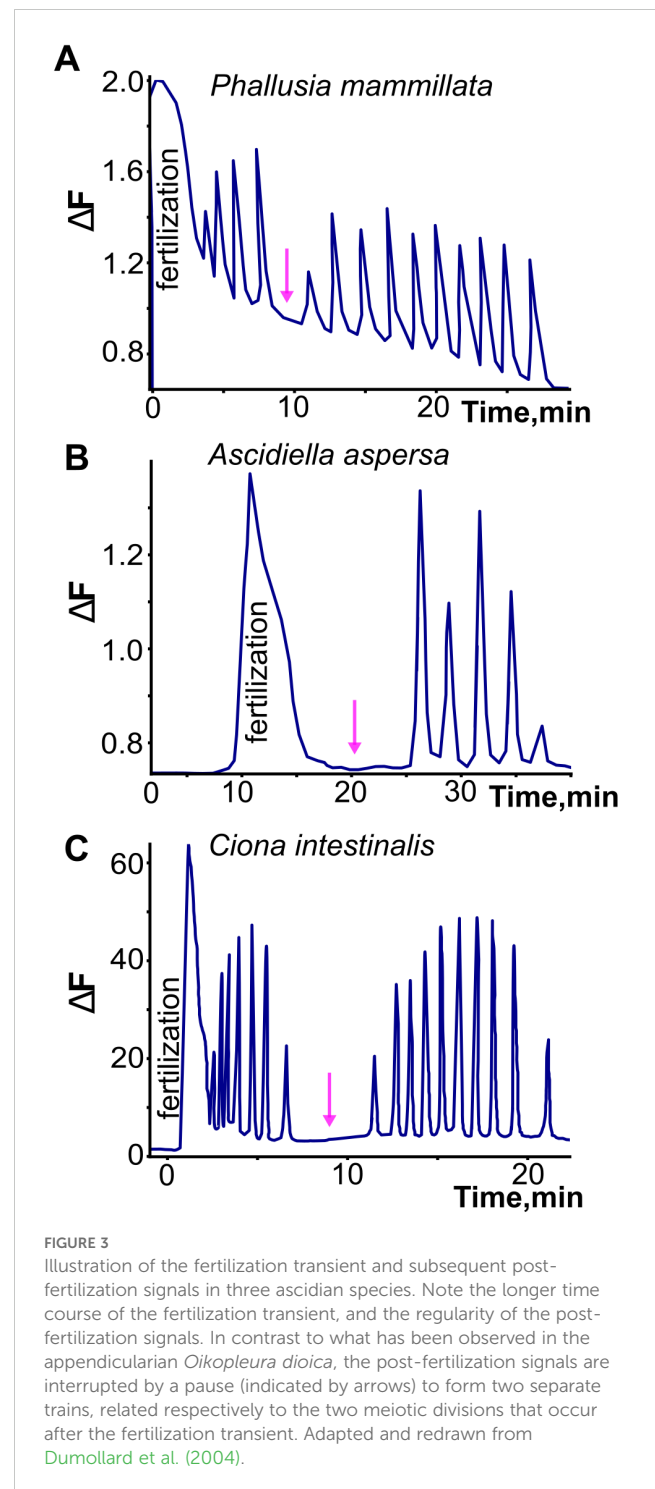


FIGURE 3
 Illustration of the fertilization transient and subsequent post-fertilization signals in three ascidian species. Note the longer time course of the fertilization transient, and the regularity of the post-fertilization signals. In contrast to what has been observed in the appendicularian *Oikopleura dioica*, the post-fertilization signals are interrupted by a pause (indicated by arrows) to form two separate trains, related respectively to the two meiotic divisions that occur after the fertilization transient. Adapted and redrawn from Dumollard et al. (2004).

fertilization, and neither initiates these nor alters their character once it has occurred (Mikhaleva et al., 2019).

In ascidians, the post-fertilization signals have been discriminated into two different series separated by a pause, with different pacemakers, related respectively to the first and second meiotic divisions (reviewed extensively in Dumollard et al., 2004). They are reported to cease after the second meiotic division. These temporal features differ from the unabated continuation of the post-fertilization signals into subsequent cleavage stages in the appendicularian *Oikopleura* (Mikhaleva et al., 2019, see below). Whereas the first series of post-fertilization signals in ascidians originate like the fertilization transient from the sperm entry site, the second series shifts to a different origin, within the cortex of the vegetal contraction pole. This region is rich in mitochondria and contains the so-called myoplasm, where maternal determinants of muscle and endoderm are concentrated. Both origins contain endoplasmic reticulum that provide the source of Ca²⁺ released by IP₃ (reviewed in Dumollard et al., 2004).

The molecular mechanisms underlying the post-fertilization signals that regulate meiosis have been studied extensively in ascidians (McDougall and Sardet, 1995; Albrieux et al., 1997, 1998; Albrieux and Villaz, 2000; Levasseur and McDougall, 2000; Wilding et al., 2000; Carroll et al., 2003; Levasseur et al., 2007, 2013; Lambert, 2011), less so in appendicularians (Matsuo et al., 2020). Pharmacological block of IP₃R using the competitive inhibitor heparin prevents the post-fertilization signals, and UV-stimulated release of caged IP₃ is sufficient to trigger similar series of signals and the associated meiotic divisions, with a single large UV pulse sufficient for the first series and additional pulses necessary for the second (McDougall and Sardet, 1995). Direct injection of IP₃ through a patch pipette also elicits similar series of signals (Albrieux et al., 1997), whereas injection of NAADP can inhibit them, presumably through depletion of the NAADP-dependent Ca²⁺ store (Albrieux et al., 1998). Neither injection of ryanodine nor of cADPr elicits the series, corroborating the primacy of IP₃ signaling underlying them (Albrieux et al., 1997). The durations of the series are correlated with and regulated by cyclin-dependent kinase 1 (Cdk1) activity, underscoring their tight relationship to the meiotic cell cycle, and possibly extending that relationship to mitotic divisions (McDougall and Levasseur, 1998; Levasseur and McDougall, 2000). Inhibiting Cdk1 eliminates the Ca²⁺ signals, but not the ability of IP₃R agonists to elicit them (Levasseur et al., 2007). Evidently, Cdk1 promotes IP₃ formation rather than acting directly on IP₃R-mediated Ca²⁺ release.

The results of these mechanistic studies have in several instances informed similar studies in vertebrates. The study of Ca²⁺ signaling in tunicates have thus contributed significantly to the overall view of the role of calcium in the early embryo (reviewed in Whitaker, 2008).

Calcium signaling from first cleavage to gastrulation

Although post-fertilization Ca²⁺ signals have been reported to cease after the second meiosis in several studies of ascidians, Ca²⁺ is known to play a pivotal role in mitosis (reviewed in Nugues et al., 2022), and calcium signaling elements, including voltage-sensitive

Ca²⁺ channels and ryanodine receptor-sensitive and IP₃-sensitive internal release, have been demonstrated in 2-cell embryos of the ascidians *Boltenia villosa* and *Phallusia mammillata* (Coombs et al., 1992; Albrieux and Villaz, 2000). Oscillating Ca²⁺ currents carried by T-type Ca²⁺ channels in the cell membranes have been recorded during early cleavage stages of *Styela plicata*, and pharmacological blockade of these channels reduces the rate of cleavage and disrupts normal embryonic development (Gallo et al., 2013). As noted above, in the appendicularian *Oikopleura dioica* post-fertilization calcium signals continue throughout the mitotic divisions that generate the blastula and gastrula (Mikhaleva et al., 2019). These are not linked temporally to cleavages but follow rather the distinctly different clock already established prior to fertilization (once every 2-3 min, as opposed to mitotic cleavages occurring once every 13 min). They exhibit a stereotyped, sequential pattern related to stage and tissue lineage (Figure 4). At the 2-cell stage, a Ca²⁺ signal is first generated in one blastomere and then spreads to the other. The order is not obligatory, as the two blastomeres can switch in taking the lead. At the 4-cell stage, a Ca²⁺ signal is first generated in one blastomere before spreading into a neighboring blastomere (either a sister or a non-sister blastomere), before spreading from these into the other pair of blastomeres in sequence. At the 8-cell stage, Ca²⁺ signals arise in two of the posterior blastomeres and spread from these in a coordinated manner to the other blastomeres. By the 16-cell stage, this temporal pattern achieves a clear tissue type relationship, as the initiation is always in the blastomeres giving rise to the muscle cell lineage (Mikhaleva et al., 2019). This may be related to the shift in post-fertilization Ca²⁺ signal origin to the myoplasm of the uncleaved ascidian oocyte, suggesting a link between early Ca²⁺ signaling and body pattern that begins already when ooplasm is reorganized prior to the first cleavage. During subsequent divisions up to gastrulation, the muscle lineage-specific origin is maintained in *Oikopleura* (Mikhaleva et al., 2019). The muscle cell lineage is also where Ca²⁺ + signals are initiated in gastrulae of *Ciona* (Akaoshi et al., 2017), is where a coupling of voltage-gated Ca²⁺ channels to ryanodine receptors has been characterized in *Halocynthia roretzi* (Nakajo et al., 1999) and is the specific site of voltage-gated Ca²⁺ currents recorded electrophysiologically after gastrulation in *Boltenia villosa* (Simoncini et al., 1988). Thus, the muscle cell lineage as the origin of whole-embryo Ca²⁺ signals through gastrulation appears to be a conserved feature in tunicates.

Contrasting with the description in *Oikopleura*, Ca²⁺ signals in gastrulae of *Ciona* were not reported to follow a patterned sequence, but rather to arise in separate, independent cells (Akaoshi et al., 2017). This is likely due to a lower temporal resolution in the *Ciona* study (0.5-1 frames/second, as opposed to 4 frames/second used by Mikhaleva et al., 2019).

Throughout development to gastrulation in *Oikopleura*, the regular, spontaneous Ca²⁺ signals exhibit the same waveform as first seen pre-fertilization, with a small “mini-peak” followed by a larger, main peak (Mikhaleva et al., 2019). Pharmacological blockade of Ca²⁺ channels (using divalent cations or the more specific T-type Ca²⁺ channel blocker mibefradil) has various effects on the amplitude and shape of the signals, and in the case of the divalent cations Cd²⁺ and Ni²⁺ disrupts the

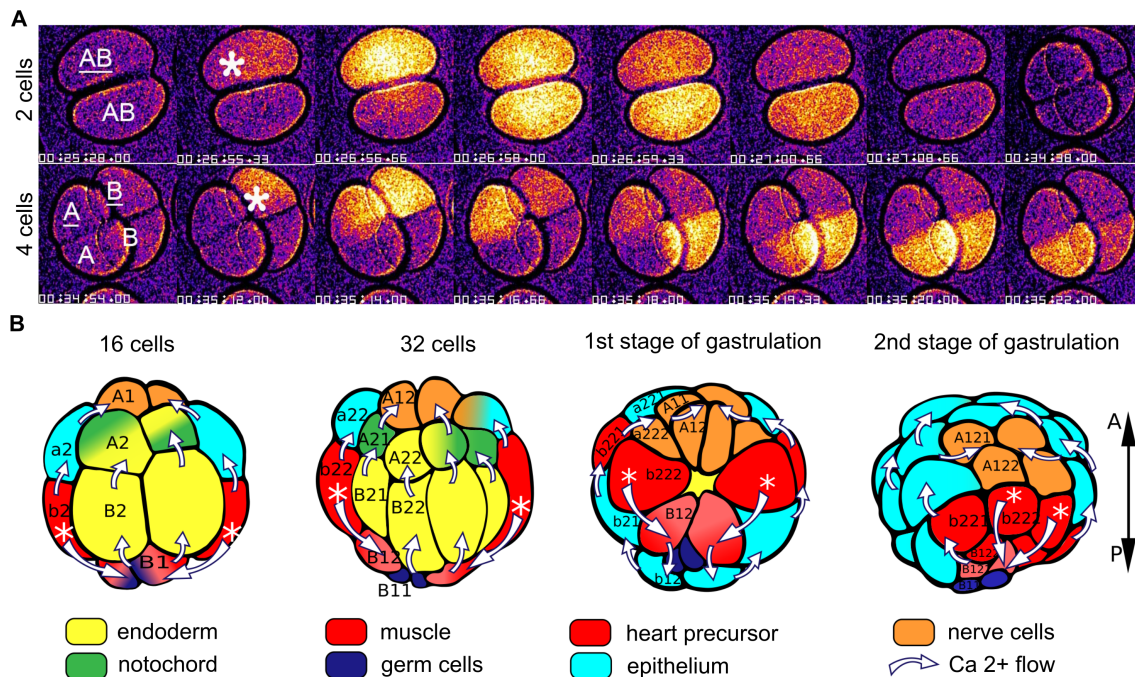


FIGURE 4

Illustration of intercellular Ca²⁺ signals in *Oikopleura dioica*. (A) Video recordings of Ca²⁺ signals in 2- and 4-cell embryos, showing the spread from the single blastomere of origin (indicated by asterisk) to the other blastomeres. (B) Diagrammatic representation of the intercellular spread of Ca²⁺ signals at later stages, oriented according to the anteroposterior (A-P) axis at the right. The white asterisks indicate the origin of the signals (in cells that give rise to the muscle cell lineage) and the white arrows indicate the trajectory of the signals among the cells. Tissue types are color coded as shown. Modified from Mikhaleva et al. (2019), Figures 2, 3.

coordinated progression of the Ca²⁺ signals among the blastomeres. Ca²⁺ channel blockade also leads to perturbed and arrested development, underscoring the importance of Ca²⁺ signaling in regulating early developmental processes. A similar effect of mibefradil has been reported in *Styela plicata*, in which it diminishes voltage gated Ca²⁺ currents recorded electrophysiologically, slows the cleavage rate and disrupts larval development (Gallo et al., 2013).

The stereotyped pattern of Ca²⁺ signal progression, always starting in cells of the muscle lineage and spreading from these throughout the embryo in a choreographed sequence, suggests a selective intercellular communication, and prompted Mikhaleva et al. (2019) to test whether gap junctional signaling is involved. This they did by knocking down the expression of the two (of 29 total) connexin genes that are expressed at early developmental stages in *Oikopleura*, Od-CxA and Od-CxB, through injection of inhibitory dsRNA into oocytes prior to fertilization. No effects were seen through the 32-cell stage, as prior to this the embryo is still supplied by maternally derived transcripts. But by gastrulation, Od-CxA and Od-CxB transcripts were diminished to 20-50% of normal levels. The effect was a severe disruption of Ca²⁺ signal progression from cell to cell, which was slowed by a factor of 10, and perturbed development (failure of ingression and epiboly at gastrulation, and later tail structure defects in embryos that succeeded to gastrulate). Thus, the patterned intercellular spread of Ca²⁺ signals depends on the presence of gap junctions, and its disruption is paralleled by abnormal development.

Calcium signaling post-gastrulation

The characteristics and functions of Ca²⁺ signaling at post-gastrulation stages are less well studied and understood than at earlier stages, particularly in a comparative context (reviewed in Slusarski and Pelegri, 2007; Webb et al., 2005). In vertebrates, Ca²⁺ transients have been shown to be associated with, and in some cases mechanically linked to, convergent extension of the body axis (Créton et al., 2000; Prudent et al., 2013; Wallingford et al., 2001; Whitaker, 2006), the establishment of left-right asymmetry (Chuang et al., 2007; Kreiling et al., 2008; Onuma et al., 2020), neural induction and neurulation (Abdul-Wajid et al., 2015; Christodoulou and Skourides, 2015; Créton et al., 1998; Gilland et al., 1999; Leclerc et al., 2000; Moreau et al., 1994; Smedley and Stanisstreet, 1985), somitogenesis (Cheung et al., 2011; Webb et al., 2012; Webb and Miller, 2011) and muscle cell differentiation (Ferrari and Spitzer, 1999; Ohtsuka and Okamura, 2007; Webb et al., 2012).

Calcium signaling during neurulation

The role of Ca²⁺ signaling during the process of neurulation has been studied in *Ciona* (Abdul-Wajid et al., 2015; Hackley et al., 2013). Abdul-Wajid et al. (2015) have demonstrated a requirement for Ca²⁺ currents mediated by a T-type calcium channel (coded by the CAV3 gene) for proper closure of the anterior neural tube.

When expression of CAV3 is prevented by the *bugeye* mutation, the normal downregulation of the *ephrinA-d* gene at the end of neurulation is prevented. This disrupts EphrinA-EphA-mediated cell-cell interactions, which are pivotal for neural tube closure. Inhibition of EphrinA activity rescues the *bugeye* phenotype. Treatment of embryos with mibefradil phenocopies the *bugeye* mutation, supporting the importance of T-type Ca²⁺ channels in mediating *ephrinA-d* downregulation. Using morpholino-based knockdown of CAV3 homologues in *Xenopus laevis* embryos, Abdul-Wajid et al. (2015) showed that the dependence of anterior neural tube closure on T-type Ca²⁺ channel activity is conserved from tunicates to vertebrates.

Another mutation in *Ciona intestinalis*, *frimousse*, leads to abnormal specification of anterior neural plate cells, which adopt an epidermal rather than a neural fate. The *frimousse* mutation lies in a connexin gene (*cx11*) and abolishes Ca²⁺ signals in the anterior neural plate (Hackley et al., 2013). Pharmacological blockade of gap junctions and exposure to low-Ca²⁺ sea water phenocopy the mutation, underscoring the essential role of gap junction-mediated intercellular Ca²⁺ signals in core developmental processes.

Calcium signaling during general organogenesis

A complete assessment of the roles of Ca²⁺ during post-gastrulation development requires the recording of Ca²⁺ signals throughout the organism. Only a few studies have characterized whole-body Ca²⁺ signaling spanning extended periods of post-gastrulation development, and in only a few species (zebrafish: Créton et al., 1998; Gilland et al., 1999; Cheung et al., 2011; Webb et al., 2012; Tsuruwaka et al., 2017; *Xenopus*: Leclerc et al., 2000; *Ciona*: Abdul-Wajid et al., 2015; Akahoshi et al., 2017; *Oikopleura*: Tolstenkov et al., 2022).

Whole-embryo Ca²⁺ signals have been characterized in various tissues including the epidermis, palps, central nervous system, peripheral nervous system, muscle, heart, and notochord of *Ciona*, through expression of the genetically encoded Ca²⁺ indicator GCaMP (Abdul-Wajid et al., 2015; Akahoshi et al., 2017). At the early tailbud stage, oscillating Ca²⁺ signals have been observed at the midbrain-hindbrain boundary, in differentiating neuronal precursors in the neural plate and nerve cord, and in a rhythmic pattern in a small population of neurons in the visceral ganglion, an activity suggestive of the central pattern generator for swimming (Abdul-Wajid et al., 2015; Akahoshi et al., 2017). At the mid-tailbud stage, Ca²⁺ signals have been observed in epidermal cells and in posterior notochord cells. Starting at the late tailbud stage, frequent Ca²⁺ signals have been observed in the tail musculature during maturation of the muscle cells, corroborating an earlier study that showed that Ca²⁺ influx through voltage-gated Ca²⁺ channels was associated with muscle cell development (Ohtsuka and Okamura, 2007).

In *Oikopleura*, whole-embryo signals have been characterized using GCaMP6 expressed throughout the body (Tolstenkov et al., 2022). Initially, Ca²⁺ signals originate in the muscle cells and

spread from these into other tissues (Figures 5, 6), continuing the situation through gastrulation in which Ca²⁺ signals originate from blastomeres of the muscle lineage and spread from these to other blastomeres. The similar spread of Ca²⁺ signals from differentiated muscle cells to other parts of the body is likely the result of gap junction-mediated coupling across tissue types, although this has not been tested directly by connexin gene knockdown as it has at pre-gastrulation and gastrulation stages (Tolstenkov et al., 2022; Mikhaleva et al., 2019). However, as organogenesis proceeds, Ca²⁺ signals begin to originate instead in the central nervous system and then spread from there to the muscle cells (Figures 5, 6). At the same time, Ca²⁺ signals in the epithelium are no longer temporally coupled with the signals in the CNS-muscle axis. Tolstenkov et al. (2022) have proposed that this transition from a muscle origin to separate origins in the CNS and epithelium involves the selective loss of general gap junction-mediated coupling across different tissue types, supplanted by more tissue-specific coupling, perhaps mediated by tissue-specific connexin expression. This hypothesis will require extensive investigation to test, since the number of connexin genes expressed increases as different tissues develop.

With respect to the CNS-muscle axis, the reversal of direction evidently relates to the development of neuromuscular synapse function. The transition is temporally linked to the developmental appearance of neuromuscular junctions (Søviknes et al., 2007), and the spread from CNS to muscle involves a much shorter temporal delay, consistent with neuromuscular synaptic transmission, and is prevented by pharmacological agents that block that transmission (Tolstenkov et al., 2022).

Calcium signals in the nervous system and relationship to early behavior patterns

In the neural plate of *Ciona*, Ca²⁺ signals appear in specific neural progenitor/neuronal precursor cells that evidently correspond to future motoneurons (MNs) and inhibitory interneurons (Akahoshi et al., 2017). Once the late tailbud stage is reached, repetitive Ca²⁺ signals appear in the visceral ganglion where MNs reside (Akahoshi et al., 2017), and Ca²⁺ signals also reappear (after a period of quiescence at early tailbud stages) in the tail musculature. In *Oikopleura dioica*, the repetitive, neurally driven Ca²⁺ signals in the tail musculature are correlated with tail movements, and the signal in the muscle cells is protracted relative to the CNS signal, suggesting a link to excitation-contraction coupling, which lasts substantially longer than nerve impulse conduction and synaptic transmission. Tolstenkov et al. (2022) observed an interesting correlation between a given tail muscle contraction and the next initiating Ca²⁺ signal in the CNS: a new CNS signal arises shortly after and never before the cessation of the muscle signal and associated tail movement. A similar relationship has been observed in *Ciona* between Ca²⁺ signals recorded in a subpopulation of central neurons (GnRH+ neurons) and tail movements: a new signal in the neurons does

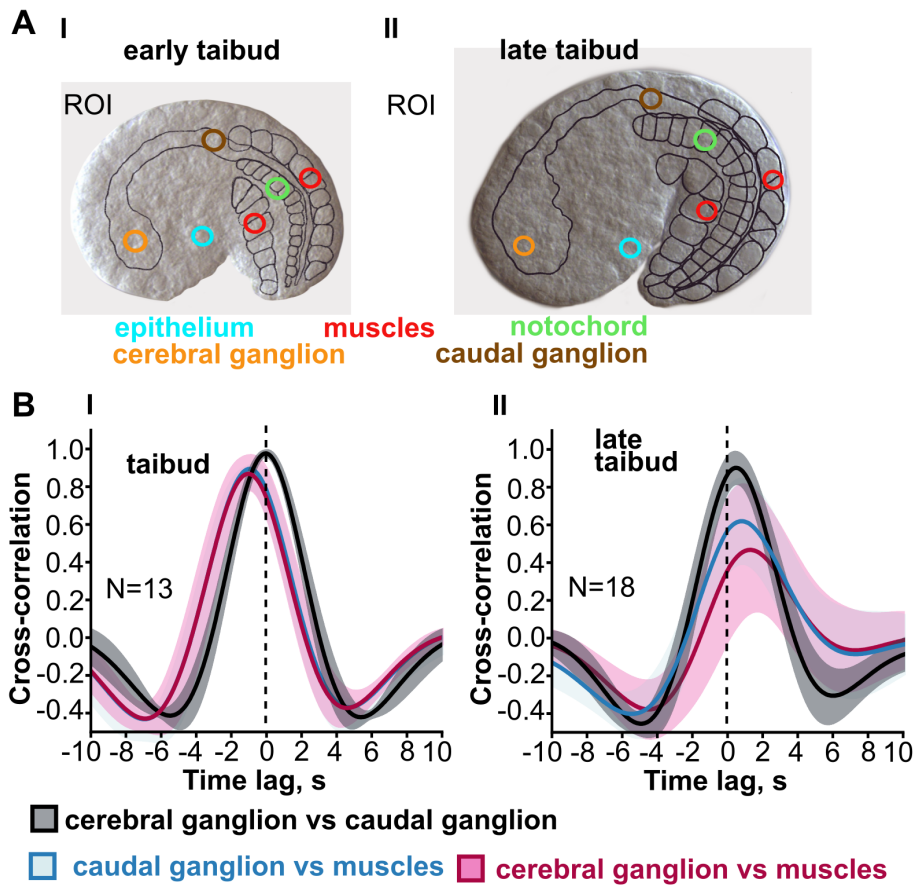


FIGURE 5 Transition from muscle lineage driven intercellular Ca²⁺ signals to neurally driven signals in *Oikopleura dioica*. (A) Signals were recorded from tissue-specific regions of interest (color coded in early and late tailbud embryos that express GCaMP6 throughout the body). (B) Cross-correlograms include (among others) the temporal relationship between signals in the cerebral ganglion and in muscle (maroon trace) and between signals in the caudal ganglion and muscle (blue trace). The time lag shift in these traces (in seconds, s) indicates that at the early tailbud stage Ca²⁺ signals in the tail muscle cells precedes that in either ganglion, whereas at the late tailbud stage the sequence is reversed. Modified from Tolstenkov et al. (2022), Figure 2.

not arise until the end of a tail movement (Okawa et al., 2020). This led Okawa et al. (2020) to postulate a sensory feedback loop that activates the central neurons once a tail movement is completed. An alternative explanation is that the cyclic nature of these phenomena is driven by an intrinsic clock that is independent of feedback

signals. Of interest here is the observation by Tolstenkov et al. (2022) that the neurally driven Ca²⁺ signals have the same regularity as the earlier muscle-driven signals. One might expect that signals driven by sensory input would lose regularity and rather relate more to the vagaries of environmental conditions.

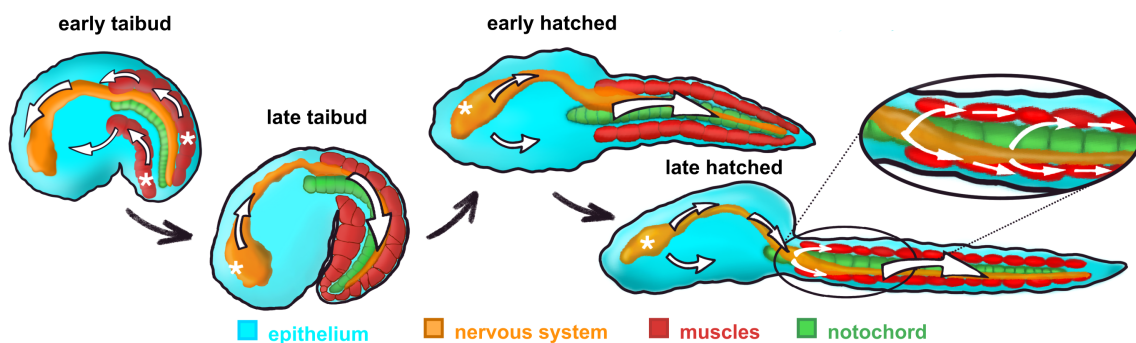


FIGURE 6 Diagrammatic illustration of the change in intercellular Ca²⁺ signal origin (white asterisks) and trajectory (white arrows) during post-gastrulation development in *Oikopleura dioica*. From Tolstenkov et al. (2022).

Calcium signals in the epidermis/epithelium

In both *Ciona* and *Oikopleura*, Ca²⁺ signals arise in the epidermis and epithelium, respectively, once these are established as definitive tissues (Akahoshi et al., 2017; Tolstenkov et al., 2022). Initially, signals in presumptive epidermis/epithelium are driven by the muscle lineage origin, but at about the time muscle signals begin to be driven by the CNS, the epidermal/epithelial signals begin to follow their own separate temporal pattern. At first haphazard, in separated individual cells, the signals eventually organize into a wave that typically starts in the head/trunk region and then travels caudally, often spreading into the notochord after reaching the tip of the tail. The function of these signals is unclear.

Conclusion

Calcium signaling in tunicates, as in other animal groups, plays a key role in a variety of developmental processes. From oocyte maturation to polyspermy block at fertilization, to the differentiation of specific cell types, in particular muscle cells, tunicates share calcium-dependent mechanisms that appear to be conserved throughout much of the chordate radiation. The evolutionary significance of other aspects of calcium signaling that have been characterized in tunicates deserves a more comprehensive assessment – both between different tunicate classes and between tunicates and vertebrates. Of note is the muscle lineage origin of whole embryo Ca²⁺ signals, clearly demonstrated in the appendicularian *Oikopleura dioica*, and its transition to neurally driven signals mediated by neuromuscular synaptic transmission. The role of gap junctions in coordinating the muscle lineage-driven whole embryo signals should be investigated in other tunicate species and compared to vertebrates. The same holds for the way this transitions to more tissue-specific signals, presumably through the regulation of tissue-specific connexin expression. The nature and underlying mechanisms of the pacemakers that drive Ca²⁺ signals, both in the single oocyte and in the multicellular embryo, need to be elucidated in more detail, again in a comparative context. As the embryo matures, the fate of the clock(s) that drive(s) intercellular Ca²⁺ signals needs to be

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determined, as the nervous system gains primacy in directing muscle activity, and other tissues and organs attain their specific functions. Of particular interest is the function of the primitive movements triggered by intrinsic waves of intercellular Ca²⁺ signals, as the forerunner of more complex, goal-directed movement repertoires that are engaged by sensory inputs. In all of these questions, tunicates – with their small larval size, low cell number, and rapid development – are well positioned to contribute insight into the developmental and evolutionary significance of calcium signaling in the chordate lineage.

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