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Citation: Soundarrajan DK, Huizar FJ, Paravitorghabeh R, Robinett T, Zartman JJ (2021) From spikes to intercellular waves: Tuning intercellular calcium signaling dynamics modulates organ size control. PLoS Comput Biol 17(11): e1009543. https://doi.org/10.1371/journal. pcbi.1009543

Editor: Jason M. Haugh, North Carolina State University, UNITED STATES

Received: April 16, 2021

Accepted: October 7, 2021

Published: November 1, 2021

Peer Review History: PLOS recognizes the benefits of transparency in the peer review process; therefore, we enable the publication of all of the content of peer review and author responses alongside final, published articles. The editorial history of this article is available here: https://doi.org/10.1371/journal.pcbi.1009543

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Data Availability Statement: All the data and simulation codes are available on our lab GitHub repository. On the linked webpage below are

RESEARCH ARTICLE

From spikes to intercellular waves: Tuning intercellular calcium signaling dynamics modulates organ size control

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Abstract

Information flow within and between cells depends significantly on calcium (Ca²⁺) signaling dynamics. However, the biophysical mechanisms that govern emergent patterns of Ca²⁺ signaling dynamics at the organ level remain elusive. Recent experimental studies in developing Drosophila wing imaginal discs demonstrate the emergence of four distinct patterns of Ca²⁺ activity: Ca²⁺ spikes, intercellular Ca²⁺ transients, tissue-level Ca²⁺ waves, and a global "fluttering" state. Here, we used a combination of computational modeling and experimental approaches to identify two different populations of cells within tissues that are connected by gap junction proteins. We term these two subpopulations "initiator cells," defined by elevated levels of Phospholipase C (PLC) activity, and "standby cells," which exhibit baseline activity. We found that the type and strength of hormonal stimulation and extent of gap junctional communication jointly determine the predominate class of Ca²⁺ signaling activity. Further, single-cell Ca²⁺ spikes are stimulated by insulin, while intercellular Ca²⁺ waves depend on Gag activity. Our computational model successfully reproduces how the dynamics of Ca²⁺ transients varies during organ growth. Phenotypic analysis of perturbations to Gag and insulin signaling support an integrated model of cytoplasmic Ca²⁺ as a dynamic reporter of overall tissue growth. Further, we show that perturbations to Ca²⁺ signaling tune the final size of organs. This work provides a platform to further study how organ size regulation emerges from the crosstalk between biochemical growth signals and heterogeneous cell signaling states.

Author summary

Calcium (Ca²⁺) is a universal second messenger that regulates a myriad of cellular processes such as cell division, cell proliferation and apoptosis. Multiple patterns of Ca²⁺ signaling including single-cell spikes, multicellular Ca²⁺ transients, large-scale Ca²⁺ waves, and global "fluttering" have been observed in epithelial systems during organ instructions on how to run a sample demonstration of the code on Google Colaboratory and how to run the code on a personal computer. Simulations corresponding to the main text figure conclusions were repeated five separate times with five different random number generator seeds (S13 Fig). This was done to ensure reproducibility of the drawn conclusions, and the corresponding outputs are documented in the GitHub repository and the linked GitHub webpage: https://

multicellularsystemslab.github.io/MSELab_ Calcium_Cartography_2021/.

Funding: The work in this paper was supported by NIH Grant R35GM124935 (JJZ) and NSF Award CBET-1553826 (JJZ). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Competing interests: The authors have declared that no competing interests exist.

development. Key molecular players and biophysical mechanisms involved in formation of these patterns during organ development are not well understood. In this work, we developed a generalized multicellular model of Ca^{2+} that captures all the key categories of Ca^{2+} activity as a function of key hormonal signals. Integration of model predictions and experiments reveals two subclasses of cell populations and demonstrates that Ca^{2+} signaling activity at the organ scale is defined by a general decrease in gap junction communication as an organ grows. Our experiments also reveal that a "goldilocks zone" of optimal Ca^{2+} activity is required to achieve optimal growth at the organ level.

Introduction

Mechanisms of intercellular communication are critical during epithelial morphogenesis when cells communicate and coordinate their activities to generate functioning organs [1,2]. One modality of intercellular communication occurs through gap junctions (GJ), intercellular channels that permit direct cell-cell transfer of ions and other small molecules [3]. Calcium ions (Ca^{2+}) act as second messengers that regulate a myriad of cellular processes such as proliferation, differentiation, transcription, metabolism, cellular motility, fertilization, and neuronal communication [4–13]. Ca^{2+} signaling also regulates developmental processes at the multicellular level. For instance, Ca^{2+} signaling has been shown to regulate scale development in butterfly wings [14]. It also mediates autophagic and apoptotic processes required for hearing acquisition in the developing cochlea [15–17]. However, a systems-level description of Ca^{2+} signaling during organ development is lacking.

A major challenge in reverse engineering Ca^{2+} signaling during organ development is the lack of an in vivo model system to identify how cells interpret and integrate information across the broad range of input molecules that dynamically vary concentrations of cytosolic Ca^{2+} ions. In particular, it remains unclear how single-cell Ca^{2+} dynamics are coordinated to regulate tissue-level Ca^{2+} patterns. To overcome these challenges, we developed a computational model based on a realistic geometry of epithelial cells to model Ca^{2+} signaling in the *Drosophila* wing imaginal disc. The *Drosophila* wing imaginal disc is an experimentally amenable system for investigating systems-level regulation of cell signaling (Fig 1A) [18–20]. Overall, *Drosophila* wing imaginal discs are a premier system to gain insights into several organ-intrinsic and organ-extrinsic mechanisms that control organ growth [21–26].

Previous experimental investigations revealed four classes of Ca^{2+} signaling activity in the developing wing disc: single-cell Ca^{2+} spikes, multicellular transients, intercellular waves, and global fluttering (Fig 1B and 1C). We have shown recently that the patterns depend on the strength of agonist stimulation [19]. We and others have previously reported that Ca^{2+} patterns observed in the wing disc are dependent on phospholipase C (PLC) and the inositol trisphosphate receptor (IP₃R) pathway mediated by gap junctional communication [18–20]. In non-excitable cells, stimulation of receptors in the cell surface results in activation of PLCs to generate IP₃, which binds to and activates IP₃R (Fig 1D) [6]. Upon binding, IP₃Rs channel Ca^{2+} from the endoplasmic reticulum (ER) to the cytosolic space [6,27,28]. However, the specific receptors involved in stimulation of PLCs in the *Drosophila* wing imaginal disc remain to be more fully defined. The key physical/chemical parameters and their interactions that define multicellular Ca^{2+} dynamics in response to agonist stimulation is not fully characterized. How these different spatiotemporal modes of signaling encode information from upstream signals that impact downstream cellular processes during organ development is poorly understood. We have previously shown that inhibition of Ca^{2+} regulators of IP₃, including PLC21C, Gαq



Fig 1. Multicellular Ca^{2+} signaling in a developing organ. (A) (*Left panel*) Image of third instar *Drosophila* wing imaginal disc. The larval wing disc includes main four regions: pouch (blue), hinge (red), notum (green) and peripodium (yellow). The pouch cells are the region of interest for this study. (*Right panel*) A schematic of the side view of the wing disc showing the peripodial membrane composed of squamous epithelial cells. (**B**) Kymographs illustrate two-dimensional slices of three-dimensional (X, Y and t planes respectively) spatiotemporal signaling. Ca^{2+} signaling activity is related to the fluorescence intensity of the Ca^{2+} sensor, GCaMP6f. A view of the X-Y plane (top left), the X-t plane (top right), and Y-t plane (bottom left) are combined to illustrate a 3D view of the signaling activity (bottom right). Yellow dashed lines indicate the pouch region. X coordinates roughly correspond to the A/P direction of the wing disc pouch whereas the Y coordinate principally describes the D/V axis of the pouch. (**C**) Four classes of Ca^{2+} signaling patterns are observed both in vivo and ex vivo: single-cell spikes, intercellular transient, intercellular waves and fluttering [19]. In ex vivo cultures, the occurrence of these patterns depend on the concentration of fly extract added to the culture media. Red arrows highlight subtle cellular Ca^{2+} activity. (**D**) Major components of Ca^{2+} toolkit: G protein–coupled receptors (GPCRs), receptor tyrosine kinase (RTKs), gap junctions (GJ), Inositol trisphosphate (IP₃), diacylglycerol (DAG), Phospholipase C (PLC), Phosphatidylinositol 4,5-bisphosphate (PIP₂), sarco/endoplasmic reticulum Ca^{2+} -ATPase (SERCA) and IP₃ receptors (IP₃R). Parameters for our computational model are denoted in purple. K_{PLC} and V_{PLC} values are lumped activity parameters that are describe the stimulus-dependent activity of specific PLC isoforms, specifically PLCγ downstream of the insulin receptor and PLCβ downstream of GPCR signaling.

https://doi.org/10.1371/journal.pcbi.1009543.g001

and the gap junction protein Innexin 2 (Inx2) in the wing disc results in reduction in size of adult wing blade [19]. Whether changes to multiscale Ca^{2+} signaling patterns in wing disc alters overall adult blade wing size remains unknown.

Overall, the computational models of calcium signaling in developing epithelial systems have received sparse attention to date. Here, we report the necessary conditions to generate the full spectrum of experimentally observed spatiotemporal patterns by employing a computational modeling approach. To do so, we built a geometrically accurate 2D-model of a wing disc based on experimental data. We discovered that in silico replication of wing disc Ca²⁺ patterns requires two distinct classes of cells, which we term as "initiator cells" and "standby cells." Here, we show how the standby cells organize themselves with respect to a Hopf

bifurcation threshold of the model's V_{PLC} parameter, and how the range of standby cell V_{PLC} values determine the final patterns of Ca²⁺ signaling. Next, computational simulations and experiments demonstrate that gap junction communication alters Ca²⁺ signaling response resulting in more Ca²⁺ spikes in the absence of external stimuli. Finally, we provide computational and experimental evidence for the role of Ca²⁺ signaling in imaginal disc morphogenesis. Our findings suggest a "goldilocks zone" of integrated Ca²⁺ signaling where lower levels of Ca²⁺ is correlated with reduced organ growth and higher levels of Ca²⁺ signal. Overall, we identify crucial crosstalk between biochemical growth signals, such as insulin and Gaq, and heterogeneous cell signaling states during the growth of an organ.

Methods

Computational model

Several mathematical models have been proposed to describe intra- and intercellular Ca^{2+} wave propagation [29]. Here we extended a previously formulated model reported by Politi and colleagues that described single-cell Ca^{2+} oscillations observed in Chinese hamster ovarian (CHO) cells [30]. This model serves as a baseline model in cells for IP₃R-mediated Ca^{2+} signaling in the *Drosophila* wing disc. This model accounts for the formation and degradation of IP₃, Ca^{2+} flux across the endoplasmic reticulum (ER) through IP₃R, and sarco/endoplasmic reticulum Ca^{2+} -ATPase (SERCA), IP₃R, and ER Ca^{2+} dynamics. The model consists of four state variables: cytosolic IP₃ (*p*), cytosolic Ca^{2+} (*c*), the ER Ca^{2+} concentration (*s*), and the fraction of IP₃ receptors that have not been inactivated by Ca^{2+} (*r*).

IP₃ dynamics

IP₃ is generated in the cytosol by phospholipases (PLC) [31]. The *Drosophila* genome consists of three PLC genes. They include *PLC21C* and *norpA*, which are related to the PLC β 1–4 sub-family of *Homo sapien* homologs, and a single PLC γ (*sl*) [32]. While different classes of PLC can hydrolyze PI(4,5)P₂ to generate IP₃ and DAG, they are activated by different receptors on the cell surface. For instance, PLC β homolog PLC21C is activated by the heterotrimeric G-protein α q subunit in response to G-protein receptor signaling [33]. On the other hand, PLC γ is recruited via its SH2 domain to activated receptor tyrosine kinase, such as the insulin receptor, at the plasma membrane [32]. In our model, we describe all the combined production of IP₃ as dependent on the total PLC activity of the cell:

$$V_{PLC_i} = V_{PLC} \frac{c_i^2}{K_{PLC}^2 + c_i^2},$$
 (1)

where V_{PLC} describes the maximal production rate of IP₃, and K_{PLC} describes the sensitivity of PLC to Ca²⁺. The parameter V_{PLC} depends on agonist concentration, we assume that V_{PLC} describes a summed activity of PLCs activated by upstream receptors.

Our model also considers degradation of IP_3 by other factors such as IP_3 kinases, which converts IP_3 to IP_4 . We generalize the degradation of IP_3 using first order kinetics. Collectively, the equation describing the dynamics of IP_3 in a cell is:

$$\frac{dp_i}{dt} = J_{p_i} + v_{PLC_i} - k_{5,P} p_i,$$
(2)

where J_{p_i} is the flux of cell *i* IP₃ through gap junction communication and $k_{5,P}$ is the IP₃ dephosphorylation rate constant. We assume that IP₃ diffuses from one cell to the adjacent

cells through gap junctional coupling. We model the flux through gap junctions (GJs) using the following equation:

$$J_{p_i} \approx F_p \Big[\sum_{j \in N_i} p_j l_{ij} - p_j \Big(\sum_{j \in N_j} l_{ij} \Big) \Big], \tag{3}$$

where F_p refers to the IP₃ permeability of the gap junctions, p_j refers to the IP₃ concentration in neighboring cell *j* and l_{ij} refers to the length of cell boundary shared by cells *i* and *j* respectively. We assume that the intracellular diffusion of IP₃ is fast relative to the diffusion of IP₃ between cells through GJs. Consequently, we have neglected terms that describe intracellular diffusion.

Ca²⁺ dynamics

 Ca^{2+} is released through IP₃R from the ER. Similarly, cytosolic Ca^{2+} is pumped into the ER using SERCA pumps. In many cell types, Ca^{2+} is also pumped out from the cytosol to the extracellular space through the plasma membrane. In our model, we ignore the flux of Ca^{2+} through the cell's plasma membrane, and we only consider the transport of Ca^{2+} from the ER to cytosol. This assumption is based on our previous experimental studies showing that the observed Ca^{2+} dynamics in the wing disc are due to the IP₃ mediated Ca^{2+} release through IP₃R from ER [18,19]. To describe the IP₃R dynamics, we followed Politi et al.'s derivation [30,33,34]. Thus, the dynamics of cytosolic Ca^{2+} in a cell is given by:

$$\frac{dc_i}{dt} = J_{c_i} + \left[k_1 \left(r_i \cdot \frac{c_i}{K_a + c_i} \frac{p_i}{K_p + p_i} \right)^3 + k_2 \right] (s_i - c_i) - V_{SERCA} \frac{c_i^2}{c_i^2 + K_{SERCA}^2},$$
(4)

where k_1 refers to maximal rate of Ca^{2+} release, K_a is the rate constant characterizing Ca^{2+} binding to activating site in IP₃R, K_p is the rate constant characterizing IP₃ binding to IP₃R, k_2 refers to Ca^{2+} leak out of ER, V_{SERCA} is the maximum rate of SERCA pump and K_{SERCA} is the half activation constant. We assume that Ca^{2+} acts as both a positive and negative regulator of IP₃R which is consistent with experimental observations of the single channel properties of the wild-type *Drosophila* receptor that has been studied using a lipid bilayer reconstitution technique [35]. Similar to IP₃, we also model diffusion of Ca^{2+} through GJs by the following equation:

$$J_{c_i} \approx F_c \Big[\sum_{j \in N_i} c_j l_{ij} - c_j \Big(\sum_{j \in N_j} l_{ij} \Big) \Big],$$
(5)

where F_c refers to the permeability of Ca^{2+} through GJs, c_j refers to the concentration of Ca^{2+} in neighboring cell *j* and l_{ij} refers to the length of cell boundary shared by cells *i* and *j* respectively.

Similarly, we describe the dynamics of Ca^{2+} concentration in the ER of a cell as:

$$s_i(t) = \frac{c_{tot} - c_i(t)}{\beta},\tag{6}$$

where s_i is the Ca²⁺ concentration in the ER of the cell and β is the ratio of effective cytoplasmic and effective ER volume, c_i refers to the cytosolic Ca²⁺ concentration in the cell and c_{tot} refers to the total Ca²⁺ concentration in the cell which includes both ER and the cytosol. We modified the rate of IP₃R inactivation term from the Politi model, *r*, to replicate our experimental Ca²⁺ dynamics. The modified equation is described below:

$$\frac{dr_{i}}{dt} = \frac{1}{\tau_{max}} \frac{k_{\tau}^{4} + c_{i}^{4}}{k_{\tau}^{4}} \left(1 - r_{i} \frac{K_{r} + c_{i}}{K_{r}}\right)$$

A similar modification to the rate of inactivation term, *r*, has been proposed previously [36].

IP₃R dynamics

We assume that the Ca^{2+} binding to the inactivating site on the IP₃R is a slow process [36]. Consequently, we consider the dynamics of IP₃R inactivation by Ca^{2+} in a cell as a separate differential equation given below:

$$t_r \frac{dr_i}{dt} = \left[1 - r_i \frac{(K_r + c_i)}{K_r}\right],\tag{7}$$

where r_i refers to the fraction of IP₃Rs of cell *i* that are not inactivated by Ca²⁺, K_r refers to the binding coefficient characterizing Ca²⁺ binding to the inactive site on the IP₃R and τ_r refers to the characteristic time of IP₃R inactivation.

Results

The relative rate of IP₃ production governs transitions between classes of spatiotemporal Ca²⁺ patterns at the tissue level

Multiple spatiotemporal classes of Ca^{2+} activity are observed in vivo and ex vivo in the wing disc. However, an understanding of how this activity is regulated requires developing a systems-level description. To summarize, these include: (i) single-cell Ca^{2+} spikes, (ii) intercellular Ca^{2+} transients (ICTs), (iii) intercellular Ca^{2+} waves (ICWs), and a (iv) global fluttering phenomenon (Fig 1C and Table 1 and S1–S8 Movies) [18–20]. The frequencies of these observed classes are dependent on the age of the larvae in both in vivo and ex vivo experiments. Younger larva with smaller discs (4–5 days after egg laying) exhibit higher occurrences of ICWs and fluttering states while older, larger larval discs (6–8 days after egg laying) predominantly show ICTs and spikes [19]. For ex vivo cultures, the transition from limited to tissue-wide Ca^{2+} signaling activity depends on the amount of fly extract (FEX) added to the culture. Low concentrations of FEX stimulated Ca^{2+} spikes. Progressively increasing levels of FEX resulted in ICTs, ICWs, and eventually fluttering. Further, FEX-stimulated Ca^{2+} dynamics is based on IP₃R-based release of Ca^{2+} from the ER to the cytosol as shown in Fig 1D [19].

These experimental findings motivated us to ask what specific cellular properties of the wing disc result in the emergence of these distinct patterns. To systematically investigate the underlying principles governing the emergence of these patterns, we formulated a two-dimensional image-based, geometrically realistic computational model of Ca^{2+} signaling in the wing disc pouch where columnar epithelial cells are connected by gap junctions (S1 Fig), [37–40]. Image-based modelling enables the holistic characterization of molecular mechanisms and tissue dynamics during organogenesis [41]. Given the near universal conservation of the Ca^{2+} signaling pathway across model systems [27,28], the baseline single-cell Ca^{2+} model in our study was adapted from Politi and colleagues [30]. The model equations, biological relevance and descriptions of the parameters are shown in Fig 1D and Table 1.

To reproduce the four distinct patterns in silico, we varied the V_{PLC} parameter progressively in individual cells across a range of values (Table 1). Given that our patterns were dependent on the concentration of FEX, we varied V_{PLC} as a lumped-parameter representing the

Parameter	Description	Value
$k_{5,P}$	IP ₃ dephosphorylation rate constant	0.66 s ⁻¹
K_{PLC}	Half activation of PLC	0.2 μΜ
V_{PLC}	Maximum production rate of PLC	0.1–1.5 μM s ⁻¹
β	Ratio of effective volumes ER/cytosol	0.185
V _{SERCA}	Maximum SERCA pump rate	$0.9 \ \mu M \ s^{-1}$
K _{SERCA}	Half activation constant for SERCA pump	0.1 μΜ
k_1	Maximum rate of Ca ²⁺ release from IP ₃ R	1.11 s ⁻¹
k_2	Ca ²⁺ leak from ER	0.0203 s ⁻¹
K _a	Ca ²⁺ binding to activating site of IP ₃ R	0.08 μM
K _r	Ca ²⁺ binding to inactivating site of IP ₃ R	0.4 μΜ
K _p	IP ₃ binding to IP ₃ R	0.13 μΜ
τ _{max}	Maximum time constant of IP ₃ R inactivation	800 s ⁻¹
$k_{ au}$	Ca ²⁺ dependent rate of IP ₃ R inactivation	1.5 μΜ
F _p	GJ permeability for IP ₃	$0.005 \mu M^2 s^{-1}$
	GJ permeability for Ca ²⁺	$0.0005 \mu\text{M}^2\text{s}^{-1}$
C _{tot}	Total Ca ²⁺ concentration in ER and cytosol	2 μΜ

Table 1. Baseline parameters used in the model.

Most baseline parameters were adopted from Politi and colleagues [30]. V_{PLC} was varied in this report in order to investigate the effects of IP₃ production on spatiotemporal calcium patterns.

https://doi.org/10.1371/journal.pcbi.1009543.t001

level of agonist stimulation (Fig 1D). We and others have demonstrated by FEX, which contains a mixture of agonists, stimulates PLC activity through GPCR and RTK signaling [18,19]. From these results, it can be inferred that the stimulation of PLC activity from FEX would subsequently increase the maximal rate of production of IP₃ (higher V_{PLC}). Thus, V_{PLC} is not *directly* a parameter representing the concentration of FEX, but is a parameter that describes the net activity of PLC through FEX stimulation of upstream receptors. The computational model successfully reproduced the four different spatiotemporal classes of Ca²⁺ signaling dynamics observed in vivo and ex vivo (Fig 2A-2D and S9-S12 Movies). Interestingly, we discovered that the formation of these patterns is dependent on the number of cells in the simulated tissue having a V_{PLC} value below, above, or equal to the Hopf bifurcation threshold for single-cells ($V_{PLC} = 0.774$) (Fig 2E). The Hopf threshold was identified from a single-cell version of the model wherein Ca²⁺ oscillations occur in the cell when V_{PLC} is at or above the value of 0.774 (Figs 2F and S2A and S3A). Simulated cells that have a V_{PLC} value above the Hopf threshold, in the absence of agonist stimulation, are termed "initiator cells" and are posed to exhibit high levels of IP3 production. Neighboring simulated cells with VPLC values below the Hopf threshold are termed "standby cells" that receive a signal from initiator cells to propagate a signal. For instance, if a majority of standby cells have V_{PLC} values significantly below the critical Hopf bifurcation threshold (standby cell VPLC randomly uniformly distributed between 0.1–0.5), single-cell Ca²⁺ spikes occur only where initiator cells oscillate (Fig 2A and 2E). When we increased standby cell V_{PLC} values close to the lower end of the Hopf bifurcation point (Fig 2B and 2E), we noticed the formation of ICTs (standby cell V_{PLC} randomly uniformly distributed between 0.25–0.60). Finally, we observed the formation of ICWs (standby cell V_{PLC} randomly uniformly distributed between 0.4–0.8) and fluttering phenotypes (standby cell V_{PLC} randomly uniformly distributed between 1.4–1.5) (Fig 2C–2E) for cases when the majority of cells in the system were assigned a V_{PLC} close to or above the bifurcation threshold, thereby placing more cells in an initiator state. In the absence of initiator cells, Ca²⁺ activity is



Fig 2. The level of hormonal stimulation governs the spatial extent of intercellular Ca^{2+} communication. (A-D) Computer simulations recapitulating the key classes of multicellular Ca^{2+} activity observed in vivo and ex vivo. (A) When the majority of cells have V_{PLC} values below the Hopf bifurcation threshold (*left*), single-cell Ca^{2+} spikes are seen (*right*). Initiator cells (red arrows) are cells with V_{PLC} values set between 1.4 and 1.5 in the simulation. A line through the A/P direction (green) demonstrates where the kymograph line is drawn that produces the simulated tissue's corresponding kymograph. (B) Intercellular Ca^{2+} transients are observed (*right*) as the distribution of V_{PLC} in cells is increased (*left*). (C) A further increase in V_{PLC} (*left*) results in the emergence of periodic intercellular Ca^{2+} waves (*right*). (D) "Fluttering" occurs (*right*) when V_{PLC} levels in all of the cells in the disc is above Hopf bifurcation (*left*). (E) Quantification of V_{PLC} distribution in the initiator and the standby cells for each of the classes of Ca^{2+} signaling activity. The first three Ca^{2+} signaling classes have a small distribution of initiator cells (red box) that are necessary for signal initiation. The dashed line indicates the threshold of the V_{PLC} parameter that permits Ca^{2+} oscillations. (F-G) The single-cell version of our model predicts that the frequency and width at half maximum (WHM) of Ca^{2+} oscillations is altered by varying V_{PLC} and k_r . This prediction matches the WHM of Ca^{2+} activity observed in ex vivo discs cultured with variable concentrations of fly extract. Error bars are reported as standard error of the means (SEM).

https://doi.org/10.1371/journal.pcbi.1009543.g002

not observed. This suggests that initiator cells are necessary for the formation of Ca²⁺ transients in developing tissues.

The "single-cell" version of the model predicts that differences in Ca²⁺ signal amplitude and frequency are tunable by varying the V_{PLC} and k_{τ} parameters (Figs 2F and 2G and S2). An output Ca²⁺ signal was observably tunable with V_{PLC} variations in replication of the four distinct patterns (Fig 2A–2E). Because the single-cell model predicted Ca²⁺ signal perturbations by tuning k_{τ} , we sought to investigate how the characteristic time associated with inactivation of IP₃R would influence tissue-scale signaling. To do this, we performed a sensitivity analysis of the two model parameters that influence characteristic time associated with inactivation of IP₃R: k_{τ} and τ_{max} . Two-dimensional computational model simulations of the tissue were performed by varying parameter values as a percentage of their baseline values (Table 1) while holding other parameters constant. The baseline parameter set was selected such that the simulation generated intercellular Ca²⁺ waves (S4 Fig, red box). This was done by each baseline simulation having the exact same V_{PLC} profile for standby and initiator cells with standby cell V_{PLC} values being uniformly random distributed between 0.7 and 1.0 and parameter values being set to those detailed in Table 1. Reducing k_{τ} leads to a narrower width at half maximum (WHM) of Ca²⁺ transients (S4A Fig). A similar result is observed where a reduction of τ_{max} results in decreased WHM, whereas increased τ_{max} increased WHM (S4B Fig). Further, a decrease in the τ_{max} parameter increased the frequency of signals in the simulated tissue, while an increase reduced the frequency of Ca²⁺ transients. This suggests that the system is more sensitive to τ_{max} , and variations in τ_{max} have much greater impact in how quickly the system responds to stimulus.

These results indicate that tissue-level Ca^{2+} patterns depend on the spatial distribution of cell states defined by their maximal IP₃ production rates, in relation to the effective tissue-level Hopf bifurcation threshold. Further, other key parameters in the model can be modified in a manner to allow tunable Ca^{2+} signaling patterns. A systematic sensitivity analysis of other parameters in the model demonstrated perturbations to Ca^{2+} signal strength, frequency, duration, and propagation (S4–S6 Figs). Further exploration and quantitative analysis of all parameters could allow an avenue to explore how tunable Ca^{2+} signaling can induce a desired physiological outcome.

GJ communication limits Ca²⁺ spikes in the absence of hormonal stimulation

To elucidate how gap junction proteins alter Ca^{2+} signals, we simulated a scenario where the initiator cells chosen at random had their V_{PLC} values set to the Hopf bifurcation threshold value of 0.774 and standby cells had a VPLC values that were randomly uniformly distributed between 0.1 and 0.5 (Fig 3A). Under these conditions, no Ca^{2+} activity was observed in the presence of normal functioning GJ communication (Fig 3A'). Next, we compared the effect of blocking gap junction communication in silico. To do so, we set the permeability terms for $Ca^{2+}(F_c)$ and for IP₃ (F_p) to zero. Strikingly, we observed Ca^{2+} spikes in simulated wing disc cells in the absence of GJ communication (Fig 3A"). We explored this phenomenon computationally by considering a single stimulated cell connected to neighboring cells by GJ communication by performing bifurcation analysis on our modified model for a single cell. We observed the emergence of a Hopf bifurcation as expected (S3A Fig). Next, the effect of the initial Hopf bifurcation point (HB₁) on gap junctional (GJ) permeability of IP₃, F_p, was analyzed. Setting F_c to zero and progressively increasing F_p increased the critical maximum rate of IP₃ activation threshold V_{PIC}^* where HB₁ occurred (<u>S3B Fig</u>). Similar trends were observed when F_c was increased to 0.05. Thus, our model suggests that inhibition of GJ communication lowers the Hopf threshold necessary to generate Ca²⁺ activity in wing disc epithelial cells.

A closer look into the importance of GJ permeability on the formation of Ca^{2+} signals was taken by varying F_c and F_p in computational simulations. Similar to the sensitivity analysis performed on k_τ and τ_{max} , GJ permeability was varied by fixed percentages (S4C Fig). We discovered that decreased GJ permeability decreased the propagation of the Ca^{2+} signal across the simulated tissue. However, the fixed percentage values ranging from 50% to 150% of the baseline parameter values (Table 1) did not produce notable changes of the ICW Ca^{2+} pattern. To further investigate this, a wider range of fixed percentages were tested between 1% and 1000% of the baseline parameter values (S7A Fig). Starting from a baseline ICW, a 90% decrease in GJ permeability resulted in a transition from ICWs to ICTs, and eventually to single-cell Ca^{2+} spikes, while a 100% increase in GJ permeability increased Ca^{2+} signal propagation and decreased the frequency (S7A Fig). These findings show that GJ permeability alters the cytosolic residence time for critical messengers such as IP₃ and Ca^{2+} whose cytosolic



Fig 3. Gap junction (GJ) communication decreases the proportion of cells exhibiting Ca²⁺ spikes. (A) Simulation of Ca^{2+} signaling in wing disc where the V_{PLC} values of initiator cells were set to the Hopf bifurcation threshold 0.774, and standby cell V_{PLC} values were randomly distributed between 0.1 and 0.5. (A') Allowing GJ communication by letting permeability of IP₃ (F_p) and Ca²⁺ (F_c) be 0.005 μ M² s⁻¹ and 0.0005 μ M² s⁻¹ respectively, resulted in no Ca²⁺ activity (GJ Enabled). (A") F_p and F_c were set to zero to simulate inhibition of GJ communication (GJ Blocked). Inhibition of GJ communication resulted in Ca²⁺ spike activity. A.U. indicates arbitrary units that correspond to the intensity of the signal. X coordinates roughly correspond to the A/P direction of the wing disc pouch whereas the Y coordinate principally describes the D/V axis of the pouch. (B) Ex vivo time lapses of nub>GCaMP6f x UAS-RyR^{RNAi} (control) wing discs in Grace's low ecdysone media were generated by imaging for 1 h at 10 sec intervals. Under this condition, we observed no Ca²⁺ activity in the wing disc cells. (B') Time-projection of the time lapse stack. The wing disc boundary is indicated with the yellow dashed line. (B") A kymograph generated further demonstrates no instances of Ca²⁻ activity. (C) GJ communication was blocked by culturing wing discs in Grace's low ecdysone media with 100 mM of Carbenoxolone (CBX). Instances of spike activity are denoted by red arrows. The intensity of a region of interest (yellow dashed circle) is overlaid to demonstrate a spike in local Ca^{2+} activity. Ca^{2+} spike is observed when the intensity normalized to basal intensity is plotted (yellow line, F/F₀). (C') Time-projection of the time lapse movies. We observed a significant number of Ca²⁺ spikes in the 1 h time interval when GJs were inhibited. Yellow dashed lines indicated disc boundary. (C") A kymograph generated demonstrates instances of Ca²⁺ spike activity.

https://doi.org/10.1371/journal.pcbi.1009543.g003

concentrations affects Ca^{2+} release from ER in both initiator cells and standby cells. This is consistent with bifurcation analysis demonstrating that GJ permeability influences stimulation threshold required for Ca^{2+} oscillations in cells (S3B Fig).

We tested these computational modeling predictions experimentally. To do so, we pharmacologically inhibited the gap junctional protein Inx2 using Carbenoxolone (CBX) and observed the emergence of Ca^{2+} spikes in the absence of FEX in the culture media (Fig <u>3B-3B</u>" and <u>3C-3C</u>", and <u>S13</u> and <u>S14</u> Movies). This further demonstrates that gap junction communication regulates Ca^{2+} dynamics in the wing disc pouch.

GJ permeability modulates Ca²⁺ signaling during development

Of note, the integrated intensity of Ca^{2+} signaling throughout the wing disc pouch decreases during development suggesting an inverse relationship between Ca^{2+} signaling activity and organ growth rates [19]. Therefore, we investigated the role of tissue size in altering Ca^{2+} signaling dynamics to propose an explanation for this finding. To do so, we simulated Ca^{2+} signaling in different sized wing discs. We hypothesized two different possible scenarios that could explain a decrease in integrated Ca^{2+} signaling activity. In the first scenario, the total number initiator cells was allowed to decrease in a power law fashion as development proceeds while GJ permeability remained constant. This resulted in a decay in total Ca^{2+} signaling activity with a transition from intercellular waves to predominantly single-cell spiking activity (Fig 4A). The fraction of initiator cells was varied according to the following equation:

$$N_{\text{initiators}} = 8,000 \cdot N_{\text{cells}}^{-0.8},\tag{8}$$

where N_{cells} is the total number of cells in the simulated tissue. The power-law relationship exponential value of -0.8 was used from the discovery that integrated Ca²⁺ intensity scales in a similar power-law fashion detailed in one of our previous publications [19]. The constant in front of the equation, 8,000, was chosen to output physiologically realistic fractions of initiator cells with the simulated tissue sizes. In the second scenario, GJ permeability was set to decrease with increasing organ size while the total number of initiator cells was held constant at N_{initia $tors} = 65$ (Fig 4B). GJ permeability was varied according to the following equation:

$$F_{p} = 800 \cdot N_{cells}^{-1.8},\tag{9}$$

where F_c is directly proportional to F_p such that $F_c = 0.1 \cdot F_p$. Similar to the scenario where initiator cell count was scaled, we investigated a scenario of scaling gap junction communication. We selected a power-law relationship value of -1.8 as an analogy to the relationship between integrated Ca²⁺ signaling activity we reported previously [19]. To ensure consistency across the two simulation setups, all other parameter values aside from initiator cell count and GJ permeability are those listed in Table 1 with V_{PLC} values of standby cell equal to 0.40. Simulations corresponding to each scenario show a decrease in progression of Ca²⁺ signaling activity starting from ICWs and intercellular transients in smaller simulated discs and decaying to intermittent single-cell spikes in larger simulated discs (Fig 4A and 4B). This suggests that both scenarios provide a possible explanation for the decrease in integrated Ca²⁺ signaling activity observed in wing discs as development progresses.

To distinguish between these scenarios, we cultured wing discs from multiple developmental stages from days 5–7 after egg laying (AEL) without any agonist stimulation and observed the resulting Ca^{2+} dynamics. We reasoned that the lack of agonist stimulation would reveal cells with phospholipase activity sufficient to create spikes, and this could further be considered initiator cells. We observed single-cell Ca^{2+} spikes that we interpret as characteristic of "initiator" cells in all samples independent of sizes. We did not observe Ca^{2+} transients or waves in unstimulated smaller wing discs obtained day 5 AEL. This is consistent with agonist stimulation increase in PLC activity in standby cells. We next characterized the total number of spikes in the discs of all sizes. We found a positive correlation between the total number of spikes and the size of the disc pouch (Fig 4C). The difference in spiking activity between discs of varying ages was not significant when scaled for pouch size (Fig 4D). This is consistent with



Fig 4. GJ permeability determines total tissue-level signaling activity as development progresses. (A-B) Simulations of Ca^{2+} signaling for wing discs of increasing size. (A) (Left column) The total number of initiator cells was varied with tissue size according to the following equation: $N_{initiators} = 8,000 \cdot N_{cells}^{-0.8}$ while GJ permeability was held constant. The V_{PLC} of all standby cells was restricted to values of 0.40, with initiator cells being denoted by dark blue cells with V_{PLC} values between 1.3 and 1.5. (Right column) Associated 2D kymographs of the simulated pouches shown in A. (B) (Left column) The GJ permeability is varied while holding the total number of initiator cells constant. GJ permeability was varied according to the following equation: $F_p = 800 \cdot N_{cells}^{-1.8}$ and $F_c = 0.1 \cdot F_p$. The total number of initiator cells were held constant in this scenario (N_{initiator} = 65) and standby cells were restricted to V_{PLC} values of 0.40. (Right column) Associated kymographs of the simulated pouches shown in **B**. Both scaling models demonstrate high Ca²⁺ signaling activity in small discs and gradually regress to low Ca²⁺ signaling activity in large discs. (C-D) Experimental validation of the computational predictions in which the discs were cultured ex vivo in Grace's low 20E media (basal media) for 1 h in the absence of any stimulus triggering Ca^{2+} . (C) Quantification of total number of Ca^{2+} spikes in different sized wing disc pouch during 1 h culture. A linear regression line was fit to the data set and the p-value for the slope of the fitted line is shown. Since the *p*-value is less than 0.05 (level of significance), a positive correlation between size and the number of spikes could be inferred. Grey region corresponds to 95% confidence bands of the trend line. (D) Quantification of total number of spikes in disc pouch scaled with respect to pouch size during various stages of larval development. p-values were obtained by Mann Whitney U test.

https://doi.org/10.1371/journal.pcbi.1009543.g004

a scenario of a relatively constant number of initiator cells in the system with overall GJ permeability decreasing as the organ reaches its terminal size. This scenario is further supported by findings from previous reports showing a decrease in GJ permeability as larval development proceeds [39,40]. Furthermore, a decrease in GJ permeability increases the net cytosolic residence time and effective concentrations of IP₃ and Ca²⁺ within the cytosol leading to an increased instance of cytosolic Ca²⁺ spikes.

Gaq overexpression induces intercellular Ca²⁺ waves and reduces wing size

Next, using the GAL4/UAS system (S8 Fig), we overexpressed the wild type splice 3 variant of Drosophila $G\alpha q$ in the wing disc to characterize how different classes of upstream signals impact the spatiotemporal patterns of Ca^{2+} signaling [42]. G protein-coupled receptor (GPCR) activation stimulates Goq-driven PLC β activity to generate IP₃ and Ca²⁺ [43]. Strikingly, ectopic Gaq expression was sufficient to generate robust formation of intercellular waves independent of the presence of FEX in the media (Fig 5D and 5D'). The waves were periodic in nature and were similar to FEX-induced waves (Fig 1C). This experimental finding most likely resembles our previous simulation of ICWs with a small fraction of randomly located initiator cells surrounded by standby cells (Fig 2C). Additionally, the wing disc size (day 6 AEL) and adult wing size were significantly reduced when G α q was overexpressed (Fig 5E and 5F). To understand whether the reduction in the wing size was due to changes in proliferation or cell growth, we quantified the total number of cells in the region bounded by the LIII, LIV, ACV wing veins and the wing margin. We observed a reduction in the total number of trichomes, where each individual trichome corresponds to a cell [44]. Furthermore, we found that cell size was reduced when Goq was overexpressed (S9F Fig). However, the wing shape is not significantly affected when $G\alpha q$ is overexpressed (S9G Fig). In sum, increasing the concentration of Goq in the pouch is sufficient to generate periodic Ca^{2+} waves. Further, these periodic Ca^{2+} waves are correlated with reduction in wing and disc size suggesting that tissue-wide Ca²⁺ wave activity may play a role in determining final organ size via growth inhibition.

Insulin signaling increases wing size but only generates localized Ca²⁺ signals

Because FEX is an undefined cocktail of biochemical factors, we tested whether specific ligands added to the organ culture affects Ca²⁺ signaling activity and growth. In addition to FEX, insulin is often added to organ culture media to stimulate cell proliferation [45,46]. Hence, we tested whether insulin signaling regulates Ca²⁺ signaling activity independent of FEX. Similar to other experiments, we upregulated and downregulated insulin signaling in the wing disc using the GAL4/UAS expression system (S8 Fig). As expected, wing disc size and adult wing size were decreased when insulin signaling is downregulated (Fig 5C", 5E, and 5F). Strikingly, we observed that activation of insulin stimulated pathways results in localized Ca^{2+} spikes and ICTs (Fig 5B and 5B'). Titrated concentration of insulin in the culture media demonstrated that a higher concentration of insulin increased the number of spikes (Fig 5G). Quantification of the Ca²⁺ spikes showed a positive correlation between spikes normalized to area of the pouch with log of insulin concentration (Fig 5G). However, even high concentrations of insulin were not sufficient to generate periodic ICWs. In contrast, expressing a dominant negative form of the insulin receptor resulted in minimal Ca^{2+} spiking activity (Fig 5C and 5C') [47]. These results demonstrate that insulin signaling stimulates Ca^{2+} activity in the wing disc. Overall, these results indicate that the agonist class encodes different spatiotemporal dynamics of Ca^{2+} signaling at the tissue scale.



Fig 5. GPCR and insulin signaling regulate wing size and differentially regulate Ca²⁺ signaling. (A-D) Montages of time-lapse movies of wing discs cultured ex vivo. (**A'-D'**) Kymographs of the corresponding time-lapse movies. (**A"-D"**) Adult wings from the indicated genetic perturbation. (**E**) Quantification of the wing disc sizes for the indicated genetic perturbations. (**F**) Quantification of the adult wing sizes for the indicated genetic perturbations.

perturbations. (G) Quantification of Ca^{2+} spikes when insulin dose is progressively increased in ex vivo cultures. A linear regression trend line was fit to the data and the *p*-value of the slope is shown. Since the *p*-value is less than 0.05, a positive correlation between spikes and the log (concentration) can be inferred. Grey region illustrates 95% confidence bands of the linear regression (H) Summary of key findings based on the proposed model for tissue-level regulation of dynamics in epithelial tissues. Different cell surface receptor stimulation produces varying V_{PLC} profiles of developing tissues causing Ca^{2+} signaling and varied tissue responses. Scale bars in (A-D) and (A"-D") represent 100 µm, while yellow dotted lines indicate pouch boundaries, and red lines indicate x-y positions in the kymograph. Horizontal scale bars in (A'-D') represent 50 µm. Vertical scale bars in (A'-D') represent 6 min. Student t-test was performed. Labels in (E) represent crosses of UAS-transgene to parental nub>GCaMP6f in the case of InsR^{CA} and InsR^{DN} or nub>GCaMP6f; mcherry in the case of RyR^{RNAi} or Gaq. The UAS lines used are UAS-RyR^{RNAi} (BL#31540), UAS InsR^{CA} (BL#8248), UAS-InsR^{DN} (BL#8252) and UAS-Gq (BL#30734) respectively.

https://doi.org/10.1371/journal.pcbi.1009543.g005

In silico simulations with parameter variations in half activation of the PLC parameter, K_{PLC} , and GJ permeability were performed to test scenarios of how insulin signaling can only generate Ca^{2+} spiking activity even under saturating conditions (S7B and S10 Figs). These two parameters were chosen because K_{PLC} can potentially be influenced by downstream of insulin receptor activity, a receptor tyrosine kinase (Fig 1D) or by the biochemical activity of phospholipase C γ . Additionally, single-cell spikes were only observed when gap junction communication was inhibited either experimentally (Fig 3B and 3C) or in silico (Figs 3A and S7A). Although large K_{PLC} variations influenced signal frequency (S7B Fig), there was no instance in which varying K_{PLC} resulted in exclusive production of Ca^{2+} spikes. To further investigate this, both GJ permeability and K_{PLC} were varied simultaneously (S10 Fig). From a baseline intercellular wave (S10A Fig), K_{PLC} was increased from its baseline value while GJ permeability was simultaneously decreased from its baseline value. Only when GJ permeability was decreased was there observance of single-cell Ca^{2+} spikes (S10 Fig). Thus, the computational model suggests that insulin signaling must not only stimulate IP₃ production, but also inhibit GJ permeability to account for the limitations of spatial spread of Ca^{2+} signaling.

As a first step to assess whether Ca²⁺ levels in the cell directly control organ sizes, we exploited the Ca^{2+} buffering effects induced by high levels of GCaMP6f sensor expression. The GCaMP6f sensor consists of a M13 fragment of myosin light chain kinase, GFP and Calmodulin (CaM) to which Ca^{2+} binds [48]. To elucidate the role of Ca^{2+} signaling in insulin mediated growth, we compared the effects of co-expressing the GCaMP6f sensor ($K_d = 375 \pm 14 \text{ nM}$) to decrease the amount of free cytosolic Ca^{2+} [49]. We found that co-expressing GCaMP6f sensor increased wing size when insulin signaling was also stimulated (S11 Fig). This increase in size was also observed in control wing disc without insulin upregulation (S11A and S11B Fig). This analysis provides additional evidence that buffering cytosolic Ca²⁺ signaling influences the final tissue size. In contrast, when we expressed Gaq and GCaMP6f, we did not observe a significant decrease in size when compared to just expressed Gaq without GCaMP6f (S11C and S11D Fig). This may be due to the inability of GCaMP6f expression to buffer the high levels of Ca^{2+} in the cytosol when Goq is overexpressed. We also observed a severe reduction in the wing size, along with vein defects, in flies that were homozygous for the GCaMP6f sensor, consistent with a buffering role for high concentrations of GCaMP6f expression (S12 Fig). To validate this finding, we compared the adult wing sizes from flies expressed one and two copies of GCaMP6f. Strikingly, increasing the dose of the relative GCaM6f expression decreased the size (S12C and S12D Fig). We observed a severe reduction in size as expression of the transgene was further increased with the presence of two copies of the GAL4 driver. To further elucidate the sponging effects of Ca^{2+} , we expressed CaMKII, which binds to Ca^{2+}/CaM complex [50]. Similar to overexpressing GCaM6f, we observe a significant increase in the wing area (S12E and S12F Fig). Collectively, these results support a role of cytosolic Ca²⁺ concentration as either a growth enhancer or suppressor and that an optimal amount of Ca²⁺ signaling is required for robust size control.

Discussion

The main finding of this work is the discovery of a parsimonious mechanistic model that links global hormonal stimulation of Ca^{2+} signaling to emergent spatiotemporal classes of signaling dynamics. Further, we identify downstream correlations to the final organ size that suggest these signaling dynamics may mediate growth-related information used by the system to tune organ size. To do so, we developed a geometrically accurate computational model of Ca^{2+} signaling in the *Drosophila* wing imaginal disc, a premier model for studying conserved cell signaling mechanisms [23,51,52]. Previously, we discovered a correlation between Ca^{2+} signaling during larval growth and final organ size. Four distinct patterns of Ca^{2+} signaling activity occur in the wing imaginal disc pouch as observed in vivo and ex vivo experiments [19]. Through systems-level computational analysis, we established the essential conditions required for generating the different patterns.

The model predicts two cell types with different levels of IP₃ production: "initiator cells" with high IP₃ production and "standby cells" with baseline IP₃ production levels. The presence of initiator cells is necessary, but not sufficient, to induce multicellular Ca²⁺ signaling. Additionally, the distribution of the maximal rate of IP₃ production, V_{PLC} , in the standby cells determines the spatial range of Ca²⁺ signaling. As V_{PLC} values of standby cells approaches the Hopf bifurcation threshold, Ca²⁺ activity transitions from single-cell signals toward global signals.

What are the possible functional implications of a tissue consisting of initiator and standby cells? A recent study on electrical signal transmission in bacterial communities suggests that the transition from localized short-range signaling to global community-level communication is associated with a cost-benefit balance [53]. In that context, long-range signaling increases the overall fitness of the community against chemical attacks, while a reduction in growth rate is the cost to individual cells. In the wing disc, a similar analogy can be drawn where significant generation of long-range Ca²⁺ signals due to overexpression of G α q results in reduced wing disc growth. Thus, the proposed model in this work can also be characterized as a cost-benefit tradeoff within the context of tissue-level signaling. For instance, it has been suggested that the fast Ca²⁺ waves facilitate migration and proliferation of the healing cells by inhibiting excessive apoptotic response during wound healing in epithelia [54].

Our model also predicts that the inhibition of GJs lowers the Hopf threshold necessary for generating Ca^{2+} spikes. We have validated this prediction experimentally where inhibition of GJs results in the formation of Ca^{2+} spikes in the absence of external agonist. Further, computational simulations demonstrate that as GJ permeability decreases, there is a transition of activity from synchronous global to asynchronous local Ca^{2+} activity. How gap junctional mediated Ca^{2+} signaling is connected to the regulation of cell mechanics is not currently understood and warrants further investigation. One possibility is that tension impacts the level of gap junction communication between cell and may also influence the activity of mechanosensitive ion channels [55]. Feedback between Ca^{2+} signaling and cell mechanics may play important roles in ensuring tissue growth and morphogenesis. This is evident from our previous experimental work and other experimental studies in the literature where knockdown of gap junctional proteins such as Inx2 leads to a reduction in wing and eye size in *Drosophila* [19,56].

Upregulated insulin signaling increases the formation of Ca^{2+} spikes. One possible implication is that insulin signaling inhibits GJs in addition to increasing PLC γ activity. This implication is consistent with the role of insulin signaling in inhibiting gap junction proteins Connexin43 in vertebrates [57–59]. Thus, the increase of adult wing and developing wing disc size from higher insulin activity correlates with higher levels of localized Ca^{2+} spiking activity, which has a limited total integrated calcium signal at that tissue level. In contrast, Ca^{2+} waves induced by Gaq overexpression are correlated with smaller adult wings and wing discs. Higher Gaq expression, which activates PLC β activity results in robust production of tissue-scale intercellular Ca^{2+} waves. We show experimentally that insulin signaling controls Ca^{2+} spike activity in the wing disc, potentially through GJ inhibition or PLC γ activation, whereas GPCRbased Gaq signaling is sufficient to generate global Ca^{2+} waves. These results are consistent with previous reports of Ca^{2+} spikes being observed in discs that have reached their final size, and Ca^{2+} waves being observed in smaller developing discs [19]. A recent study found that Ca^{2+} signals are initiated in response to wounding by the G-protein coupled receptor Methuselah like 10 (Mthl10) [60]. Mthl10 is activated by Growth-blocking peptides (Gbps) [60]. Whether Mthl10 also is involved in developmental growth requires further investigation, but may be consistent with our findings that intercellular calcium wave activity inhibits organ growth. These findings suggest that Ca^{2+} acts as both a growth enhancing and growth inhibiting signal dependent upon the tissue-level scale of the activity and level of gap junction coupling.

The spatial range of tissue-level signaling is determined by how the IP₃ production is organized with respect to a Hopf bifurcation threshold throughout the tissue. Localized transients are correlated with larger wings induced by insulin-stimulated growth whereas global signaling is correlated with smaller wings that are stimulated by upstream GPCRs and G α q upregulation (Fig 5H). This resembles a paradoxical network motif [61] where Ca²⁺ signaling has two opposite effects on the same downstream target, dependent upon the tissue-level magnitude of the Ca²⁺ signaling. Within the context of the hypothesized IP₃/shunt model proposed in our previous study, the strong induction of Ca²⁺ waves will reduce the level of phosphatidylinositol 4,5-bisphosphate (PIP₂), a key substrate for growth [19]. This may occur as high levels of G α q/ PLC activity are proposed to deplete PIP₂ levels [62]. This would occur due to substrate depletion of PIP₂ through promotion of IP₃ generation and downstream activity by stimulation of PLC activity. In turn, this would lead to reduced availability of PIP₂ for conversion of PIP₂ to phosphatidylinositol-trisphosphate (PIP₃), a key second messenger for stimulating protein kinase AKT and downstream growth promotion [63].

Similar to the reduced wing size observed in this study due to overexpression of G α q, we have also reported that knockdown of G α q gene decreases wing size in our previous study (S2 Table) [19]. Comparing the reduction in wing size due to perturbation of Ca²⁺ signaling components with known size control genes such as morphogens (Dpp, Wg, Hippo) and mechanical transducers (RoK) indicate that the reduction in size is comparable to when Ca²⁺ signaling is perturbed (S2 Table). Taken together, these experimental findings imply that G α q signaling is paradoxical in nature. In the context of conserved network motif observed in biological systems, paradoxical components have the ability to activate and inhibit the downstream target despite a single source of stimulus [61]. G α q could possibly be a growth promoter and growth inhibitor. This correlation motivates further studies that map out the exact molecular players that are downstream of G α q signaling. This will require careful quantification of PIP₂ and PIP₃ under genetic perturbations of GPCR signaling. Additionally, future work is needed to quantify the metabolic benefits and costs of Ca²⁺ signaling during tissue growth to observe if abundant use of metabolic resources to consistently propagate global activity is explanatory for the reduction in size in G α q overexpression wings.

Materials and methods

Drosophila genetics

We used the GAL4/UAS system to express modulators of the Ca²⁺ signaling pathway in the wing disc [64,65]. A nub-GAL4, UAS-GCaMP6f reporter tester line was created by

recombining nub-GAL4 and UAS-GCaMP6f lines [66]. Additionally, a second tester line was used that also includes UAS-mcherry. Gene perturbations were generated by crossing the tester line to either RNAi-based transgenic lines (UAS-Gene X^{RNAi}) or gene overexpression (UAS-Gene X). The following UAS transgenic lines were used: UAS-RyR^{RNAi} (BL#31540) [67], UAS-Gq (BL#30734) [67], UAS-InsR^{CA} (BL#8248) [68], UAS-InsR^{DN} (BL#8252) [69]. Progeny wing phenotypes are from F1 male progeny emerging form the nub-Gal4, UAS-G-CaMP6f/CyO x UAS-X cross or nub-Gal4, UAS-GCaMP6f/CyO; UAS-mcherry x UAS-X cross. Flies were raised at 25°C and on a 12-hour light cycle.

Live imaging

Wandering third instar larva approximately 6 days after egg laying were dissected in ZB media with 15% fly extract to obtain wing discs [19]. ZB media + 15% fly extract contains 79.4% (v/v) ZB media, 0.6% (v/v) of 1 mg/ml of insulin (Sigma aldrich), 15% ZB-based fly extract and 5% penicillin/streptomycin (Gibco). Wing discs were loaded into the previously described REM-Chip [70] and imaged using Nikon Eclipse Ti confocal microscope with a Yokogawa spinning disc and MicroPoint laser ablation system. Image data were collected on an IXonEM +colled CCD camera (Andor technology, South Windsor, CT) using MetaMorph v7.7.9 software (Molecular devices, Sunnyvale, CA). Discs were imaged at three z-planes with a step size of 10 µm, 20x magnification and 10-seconds intervals for a total period of one hour, with 200 ms exposure time, and 50 nW, 488 nm laser exposure at 44% laser intensity. We blocked GJ by inhibiting innexin-2 using Carbenoxolone (Cbx, Sigma Aldrich) drug [66]. Wing discs were incubated in ZB + 15% FEX with 30 μ M Cbx for one hour before imaging. To induce Ca²⁺ transients, we imaged wing discs in ZB media + 2.5% FEX [71]. Ca^{2+} waves were induced by imaging the wing disc in ZB media + 15% FEX. Ca²⁺ fluttering was observed when discs were imaged in ZB media + 40% FEX respectively. For experiments reported in Figs 3, 4, and 5, wing imaginal discs were cultured in Grace's media with low 20E (Dye et al., 2017). Briefly, basal Grace's media was prepared by addition of Grace's medium (Sigma, G9771) with 5 mM BisTris, 5% fetal bovine serium (FBS; ThermoFisher/Invitrogen, 10370098) and Pennicillin-Streptomyocin (Sigma P4333, 100x stock solution) along with 20 nM 20E (Sigma, H5142). For the Cbx experiment reported in Fig 3, we cultured wing discs in Grace's cocktail with 30 uM Cbx (Sigma Aldrich). For insulin dose response experiments reported in Fig 5, we added appropriate volume of Bovine insulin (Sigma, I5500) to the Grace's cocktail respectively.

Image processing

All the images were processing in FIJI. Volume viewer plugin was used to generate 3D Kymographs. Briefly, TIFF stacks had background subtracted using the rolling ball background subtraction algorithm with a rolling ball radius of 15. The TIFF stack was then processed by the volume viewer plugin. Stacks were adjusted using the base Fire LUT setting to portray the signaling intensities. A similar approach was followed for the simulation outputs.

Model formulation

Fig 1 summarizes the experimental system and data. Different classes of patterns emerge at the tissue-level as the level of global stimulation increases: spikes, intercellular Ca^{2+} transients (ICTs), intercellular Ca^{2+} waves (ICWs) and global fluttering [19]. However, a mechanistic understanding linking hormonal stimulation levels to transitions in these qualitative classes of organ-level signaling is lacking. We therefore formulated a computational model to test mechanistic hypotheses that could explain the observed Ca^{2+} signaling dynamics.

Intracellular model

A modified model of Ca^{2+} signaling toolkit is based on adaptation of previous single-cell model of calcium signaling [30]. The model is summarized in the computational model section of the main text. To recapitulate the same time resolution as the experiments, the simulation time is 1 hour and for generating movies, samples are obtained every 10 s.

Tissue model

For constructing a realistic model of the tissue, we used experimental images of a wing pouch to build an accurate model of the tissue structure. S1 Fig depicts the structure of the tissue used for simulations and the statistics of the corresponding network. A pouch was constructed computationally using EpiTools. We first segmented the apical cell boundaries using images from Ecad::GFP line. Then, the centroids of segmented cells were used to define cellular positions in the simulated wing disc. A Voronoi tessellation followed by multiple rounds of Lloyd's relaxation was used to define a template wing disc that matches the experimentally observed network topology.

Quantification and statistical analysis

Quantification of adult wings and statistics. Total wing area was measured using ImageJ. We traced the wing margin by following veins L1 and L5 and the wing hinge region was excluded from the size analysis. All statistical analyses were performed using MATLAB or R. For comparisons, we used student t-tests to assess the statistical significance. p-value, standard deviation and sample size (n) are as indicated in each figure and legend.

Supporting information

S1 Fig. Computational framework. (A) Experimental *Drosophila* imaginal disc showing cell boundaries marked with Ecad::GFP. The developing wing pouch has been segmented using ImageJ. The genotype of the *Drosophila* used is *yw;;dECad::GFP* (BL# 46556) (**B**) A pouch constructed computationally using EpiTools that served as a basis for Ca²⁺ signaling simulations. In brief, cells were segmented from a wing disc. Centroids of segmented cells were used to define cellular positions in the simulated wing disc. A Voronoi tessellation followed by multiple rounds of Lloyd's relaxation [72] was used to define a template wing disc that matches the experimentally observed network topology. (TIF)

S2 Fig. Single-cell Ca²⁺ dynamics. The model was calibrated to match experimental single-cell frequency and amplitude. Perturbations to stimulation strength V_{PLC} (**A**) alters the frequency and amplitude of Ca²⁺ oscillations whereas perturbations to k_{τ} (**B**) only alters the frequency.

(TIF)

S3 Fig. Bifurcation analysis of the modified model. (A) Bifurcation diagram for the modified model used in this study; shown are the maxima and minima of the Ca²⁺ oscillations (dots) and the Ca²⁺ steady states (solid and dashed lines) as a function of the stimulus (V_{PLC}). Solid and dashed lines in red indicate stable and unstable states, respectively. Red dots indicate the maxima and minima of unstable limit cycle and the black dots indicate maxima and the minima of the stable limit cycle. HB, Hopf bifurcation occurs when V_{PLC} is varied. Inset figure shows the period of Ca²⁺ oscillations as a function of V_{PLC} . (B) Blocking permeability of IP₃, F_p via gap junctions decreases V_{PLC} where the initial Hopf bifurcation point (HB₁) occurs in

the bifurcation diagram. Block dots indicate conditions where permeability of Ca^{2+} , F_c is set to 0. Red diamonds indicate conditions where F_c is set to 0.5. (TIF)

S4 Fig. Model sensitivity analysis to parameters: k_{τ} , τ_{max} , F_p , F_c , K_r and K_p . Five different parameters in the 2D model were varied from their baseline values (BV). V_{PLC} profiles of simulated tissues were selected to generate intercellular waves (red box) and are identical across all simulations to enable comparisons. Each row represents one parameter being varied in a scaling manner by fixed percentages listed in each column (*i.e.*, 50% of a BV of $1.5 \,\mu$ M would simulate a value of $0.75 \,\mu$ M). Simulations were performed varying only one parameter while holding all others constant at their BVs. Signal frequency is observed through number of bands in the kymograph, and signal duration is observed through thickness of the bands in the kymograph. (A) Decreased k_{τ} (BV of 1.5 μ M) increased frequency and decreased duration of the Ca²⁺ signal whereas increased k_{τ} did not influence the signal. (B) Decreased τ_{max} (BV of 800 s⁻¹) increased frequency and decreased duration of the Ca²⁺ signal whereas increased τ_{max} decreased frequency and increased duration. (C) Decreased gap junction (GJ) communication $F_{p/c}$ (BVs of 0.005 μ M² s⁻¹ for F_p ; 0.0005 μ M² s⁻¹ for F_c) decreased propagation of the Ca²⁺ signal whereas increased GJ communication increased propagation. Signal propagation is visualized by the uniformity of the signal across the tissue. (D) Decreased K_r (BV of 0.4 μ M) decreased frequency and duration of the Ca^{2+} signal whereas increased K_r decreased frequency but increased duration. (E) Decreased K_p (BV of 0.13 μ M) increased frequency of the Ca²⁺ signal whereas increased K_p decreased frequency.

(TIF)

S5 Fig. Model sensitivity analysis to parameters: $k_{5,P}$, K_a , K_{PLC} , V_{SERCA} and β . Five different parameters in the 2D model were varied from their baseline values (BV). V_{PLC} profiles of simulated tissues were selected to generate intercellular waves (red box) and are identical across all simulations to enable comparisons. Each row represents one parameter being varied in a scaling manner by fixed percentages listed in each column (*i.e.*, 50% of a BV of 0.66 s⁻¹ would simulate a value of 0.33 s⁻¹). Simulations were performed varying only one parameter while holding all others constant at their BVs. Signal frequency is observed through number of bands in the kymograph, and signal duration is observed through thickness of the bands in the kymograph. (A) Decreased $k_{5,P}$ (BV of 0.66 s⁻¹) increased frequency of the Ca²⁺ signal whereas increased $k_{5,P}$ decreased frequency. (B) Decreased K_a (BV of 0.08 μ M) increased frequency of the Ca²⁺ signal to the point of observing constant activity, whereas increased K_a decreased frequency to the point of loss of signal in a 60 minute simulation. (C) Decreased K_{PLC} (BV of 0.2 μ M) increased frequency of the Ca²⁺ signal whereas increased K_{PLC} decreased frequency to the point of loss of signal in a 60 minute simulation. (D) Decreased V_{SERCA} (BV of 0.9 μ M s⁻¹) increased frequency and duration of the Ca²⁺ signal to the point of observing constant activity, whereas increased V_{SERCA} decreased frequency to the point of loss of signal in a 60 minute simulation. (E) Decreased β (BV of 0.185) increased frequency and duration of the Ca²⁺ signal to the point of observing constant activity, whereas increased β decreased frequency to the point of loss of signal in a 60 minute simulation. (TIF)

S6 Fig. Model sensitivity analysis to parameters: K_{SERCA} , c_{tot} , k_1 and k_2 . Four different parameters in the 2D model were varied from their baseline values (BV). V_{PLC} profiles of simulated tissues were selected to generate intercellular waves (red box) and are identical across all simulations to enable comparisons. Each row represents one parameter being varied in a scaling manner by fixed percentages listed in each column (*i.e.*, 50% of a BV of 0.1 µM would simulate a value of 0.05 μ M). Simulations were performed varying only one parameter while holding all others constant at their BVs. Signal frequency is observed through number of bands in the kymograph, and signal duration is observed through thickness of the bands in the kymograph. (**A**) Decreased K_{SERCA} (BV of 0.1 μ M) decreased frequency of the Ca²⁺ signal whereas increased K_{SERCA} increased frequency. (**B**) Decreased c_{tot} (BV of 2 μ M) decreased frequency of the Ca²⁺ signal whereas increased frequency. (**C**) Decreased k_1 (BV of 1.11 s⁻¹) decreased frequency and duration of the Ca²⁺ signal whereas increased k_1 increased frequency and increased. (**D**) Decreased k_2 (BV of 0.0203 s⁻¹) decreased frequency of the Ca²⁺ signal whereas increased k_2 increased frequency.



S7 Fig. Model sensitivity analysis to extremes of the parameters: F_p , F_c and K_{PLC} . Two different parameters in the 2D model were varied from their baseline values (BV). VPLC profiles of simulated tissues were selected to generate intercellular waves (red box) and are identical across all simulations to enable comparisons. Each row represents one parameter being varied in a scaling manner by fixed percentages listed in each column (i.e., 50% of a BV of 0.1 µM would simulate a value of $0.05 \,\mu$ M). Simulations were performed varying only one parameter while holding all others constant at their BVs. Signal frequency is observed through number of bands in the kymograph, and signal duration is observed through thickness of the bands in the kymograph. (A) GJ permeability of IP₃ and Ca^{2+} influences synchronization of Ca^{2+} signaling among cells. Decreased gap junction (GJ) communication $F_{p/c}$ (BVs of 0.005 μ M² s⁻¹ for F_p ; $0.0005 \,\mu\text{M}^2 \,\text{s}^{-1}$ for F_c) results in a transition of intercellular waves to intercellular transients, and to single-cell spikes. Increased GJ communication increased Ca^{2+} signal propagation and decreased signal frequency. Signal propagation is visualized by the uniformity of the signal across the tissue. (B) Variations in the half-activation of V_{PLC} term, K_{PLC} (BV of 0.2 μ M), only changed the frequency of the ICWs. (TIF)

S8 Fig. Schematic of expression pattern using the Gal4/UAS system. (A) The GAL4/UAS system was used to express *GCaMP6f* transgene along with other transgenes. **(B)** *nubbin* is expressed in the wing disc pouch and the adult wing phenotype provide a readout of final phenotype after transgene expression in the wing disc pouch during larval stage. (TIF)

S9 Fig. Overexpression of Gaq decreases cell number and cell size. (A-B) Wings from adult males expressing RyR^{RNAi} and wild type Gaq splice 3 variant with *nubbin-Gal4, UAS GCaMP6f, UAS mcherry.* (A'-B') Region of interest (ROI) where the total number of setae was calculated. (C-F) Quantification of the wing size defined here as the area bounded by LIII, LIV, ACV and the wing margin, total cell number and cell area. Overexpression of Gaq in the pouch results in a decrease in total wing area, cell number and cell size. 10 samples were analyzed per condition. Error bars represent standard deviation. (G) Quantification of roundness of the adult wing. Gaq overexpression does not affect the roundness. Student t-test was used for statistical significance testing. (TIF)

S10 Fig. GJ inhibition is the key driver of single-cell Ca²⁺ spike activity. To replicate the ex vivo observations of insulin inducing single-cell Ca²⁺ spikes, GJ permeability and the half-activation of V_{PLC} were varied simultaneously in silico. (**A**) A baseline intercellular wave was used as the comparison for how parameter variations changed signal with the following parameter values: IP₃ gap junction permeability (F_p) of 0.005 μ M²s⁻¹, Ca²⁺ gap junction permeability (F_c) of 0.0005 μ M²s⁻¹, and K_{PLC} of 0.20 μ M. (**B-E**) K_{PLC} is increased left-to-right (red bar), and gap junction communication is

decreased left-to-right (blue bar). An increase in K_{PLC} results in a decrease in frequency, while decrease in gap junction communication results in single-cell Ca²⁺ spikes (red arrows). (TIF)

S11 Fig. Sponging cytosolic Ca²⁺ increases overall wing size. (A-F) Wings from adult males with the indicated crosses. (A) *nubbin-GAL4* x *UAS-RyR*^{RNAi} (i.e., nub4>RyR^{RNAi}), (B) *nub-bin-GAL4*, *UAS-GCaMP6f* x *UAS-RyR*^{RNAi} (i.e., nub4>GCaMP6f, RyR^{RNAi}), (C) *nubbin-GAL4*, *UAS-GGaq*^{OE} embryonic splice 3 variant of Gaq (i.e., nub4>Gaq^{OE}), (D) *nubbin-GAL4*, *UAS-GCaMP6f* x *UAS-Gaq*^{OE} (i.e., nub4>GCaMP6f, Gaq^{OE}), (E) *nubbin-GAL4* x *UAS-InsR*^{CA} (i.e., nub4>InsR^{CA}) gain of function mutant where the α subunit is partially deleted. (F) *nubbin-GAL4*, *UAS-GCaMP6f* x *InsR*^{CA} (i.e., nub4>GCaMP6f, InsR^{CA}). (G) Quantification of adult wings. The genetic encoded calcium sensor, GCaMP6f, binds to Ca²⁺ with high affinity, thus expression of the sensor will to some degree act as a sponge of cytosolic Ca²⁺. Interestingly, the presence of the GCaMP6f sponge with constitutively activated insulin signaling increases the adult wing size (E, F). Similar enhancement of wing size was observed in control wings when GCaMP6f sensor was expressed (A, B). No significant change in the adult wing size was observed when G α q was overexpressed, suggesting sponging effects are trivialized under G α q overexpression (C,D). Unpaired student t-test was used, and the p-values are indicated above.

(TIF)

S12 Fig. Increasing gene dosage of GCaMP6f Ca^{2+} sensor dramatically reduces wing size. (A-E) Wings from adult males of indicated genotypes (A) nubG4, UAS RyR^{RNAi}, (B) nubG4, UAS GCaMP6f, (C) nubG4, UAS GCaMP6f/UAS GCaMP6f, (D) nubG4, UAS GCaMP6f (Homozygous) (E) nubG4, UAS CaMKII (F) Quantification of adult wing sizes. As the gene dose of GCaMPf is increased in the wing disc, the overall wing area decreases in size (B, C, D). Overexpressing possible Ca^{2+} downstream target CaMKII increases the wing size consistent with B in which one copy of GCaMP6f was expressed. (TIF)

S13 Fig. Repeated simulations result in the same conclusions. Simulations corresponding to the main text figure conclusions (*i.e.*, Figs 2-4) were repeated five separate times with five different random number generator seeds. The random number generator seed value determines which cells in a simulated tissue are determined to be initiator cells. For the case of Fig 2, the V_{PLC} values of standby cells also rely on the random number generator seed as the values are sampled randomly from a uniform distribution with set boundaries. The simulations' resulting video and image outputs were randomized and their inputs were hidden to allow classification of the Ca^{2+} activity. (A) Using a graphical user interface (GUI) in MATLAB, the randomized video and kymograph simulations were drawn to show the output kymograph and play the video simulation. A user was tasked to classify the activity of the simulation output as having no activity, single-cell spikes, intercellular transient activity (ICT), intercellular wave activity (ICW), or global fluttering activity. For each main text figure, the five separate simulations had their Ca^{2+} activity classified in four independent runs. Each run corresponds to a brand new running of the classification GUI, each with a different randomization scheme to display the outputs of the simulations. (B) The proportions of Ca^{2+} activity are plotted for Fig 2's repeated simulations. Runs 1, 2, and 4 all had the same proportions, indicating reproducibility of the simulations' outputs. Run 3 had mismatched classifications between the ICT and spike class (red arrow), however, the difference in proportions was not significant using a proportions test without a continuity correction [73–75]. (C) The proportions of Ca^{2+} activity are plotted for Fig 3's repeated simulations. Because Fig 3 was designed to demonstrate either no activity

in the case of enabled gap junction communication, or spiking activity in the case of disabled gap junction communication, only two classes of activity appear. In each classification run, there were no differences in the user-recorded classifications. (**D**) The proportions of Ca^{2+} activity are plotted for Fig 4's repeated simulations. Runs 1, 2, and 4 all had the same proportions, indicating reproducibility of the simulations' outputs. Run 3 had mismatched classifications between the ICT and spike class (red arrow), however, the difference in proportions was not significant using a proportions test without a continuity correction. (TIF)

S1 Table. Extended data movies.

(DOCX)

S2 Table. Changes in wing area for known perturbations through Gal4/UAS system. Note that maximal deviations wing size for strong growth perturbations is in range of 20–50%. References for this table include this study, [19], [76], [77]. (DOCX)

S1 Movie. nub-Gal4>UAS-GCaMP6f, UAS-mcherry, ex vivo, spike. (AVI)

S2 Movie. nub-Gal4>UAS-GCaMP6f, UAS-mcherry, ex vivo, ICT. (AVI)

S3 Movie. nub-Gal4>UAS-GCaMP6f, ex vivo, ICW. (MP4)

S4 Movie. nub-Gal4>UAS-GCaMP6f, ex vivo, fluttering. (MP4)

S5 Movie. nub-Gal4>UAS-GCaMP6f, in vivo, spikes. (AVI)

S6 Movie. nub-Gal4>UAS-GCaMP6f, in vivo, ICT. (MP4)

S7 Movie. nub-Gal4>UAS-GCaMP6f, in vivo, ICW. (AVI)

88 Movie. nub-Gal4>UAS-GCaMP6f, in vivo, fluttering. (AVI)

S9 Movie. Spike, Simulation output. (MP4)

S10 Movie. ICT, Simulation output. (MP4)

S11 Movie. ICW, Simulation output. (MP4)

S12 Movie. Fluttering, Simulation output. (MP4)

S13 Movie. nub-Gal4>UAS-GCaMP6f, ex vivo in Grace's low 20E media, gap junctions not blocked (Control). (AVI) S14 Movie. nub-Gal4>UAS-GCaMP6f, ex vivo in Grace's low 20E media with Carbenoxolone, gap junctions blocked.

(AVI)

S15 Movie. nub-Gal4>UAS-GCaMP6f, UAS-RyRRNAi, ex vivo in Grace's low 20E media (Control).

(AVI)

S16 Movie. nub-Gal4>UAS-GCaMP6f, UAS-InsRCA, ex vivo in Grace's low 20E media. (AVI)

S17 Movie. nub-Gal4>UAS-GCaMP6f, UAS-InsRDN, ex vivo in Grace's low 20E media. (AVI)

S18 Movie. nub-Gal4>UAS-GCaMP6f, UAS-GaqOE, ex vivo in Grace's low 20E media. (AVI)

Acknowledgments

The authors gratefully acknowledge the Notre Dame Center for Research Computing (CRC) for providing computational facilities. The authors would also like to thank members of the Zartman lab for helpful discussions.

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References

- Leptin M. Drosophila Gastrulation: From Pattern Formation to Morphogenesis. Annual Review of Cell and Developmental Biology. 1995; 11: 189–212. https://doi.org/10.1146/annurev.cb.11.110195.001201 PMID: 8689556
- Schöck F, Perrimon N. Molecular Mechanisms of Epithelial Morphogenesis. Annual Review of Cell and Developmental Biology. 2002; 18: 463–493. https://doi.org/10.1146/annurev.cellbio.18.022602.131838 PMID: 12142280
- Levin M. Gap junctional communication in morphogenesis. Progress in Biophysics and Molecular Biology. 2007; 94: 186–206. https://doi.org/10.1016/j.pbiomolbio.2007.03.005 PMID: 17481700
- Berridge MJ. Unlocking the secrets of cell signaling. Annu Rev Physiol. 2005; 67: 1–21. https://doi.org/ 10.1146/annurev.physiol.67.040103.152647 PMID: 15709950
- Carafoli E. Calcium signaling: a tale for all seasons. Proceedings of the National Academy of Sciences. 2002; 99: 1115–1122. https://doi.org/10.1073/pnas.032427999 PMID: 11830654
- 6. Clapham DE. Calcium Signaling. Cell. 2007; 131: 1047–1058. https://doi.org/10.1016/j.cell.2007.11. 028 PMID: 18083096
- Cuthbertson KR, Cobbold P. Phorbol ester and sperm activate mouse oocytes by inducing sustained oscillations in cell Ca2+. Nature. 1985; 316: 541. https://doi.org/10.1038/316541a0 PMID: 4033751
- Giorgi C, Danese A, Missiroli S, Patergnani S, Pinton P. Calcium dynamics as a machine for decoding signals. Trends in cell biology. 2018; 28: 258–273. https://doi.org/10.1016/j.tcb.2018.01.002 PMID: 29409699
- 9. Humeau J, Bravo-San Pedro JM, Vitale I, Nuñez L, Villalobos C, Kroemer G, et al. Calcium signaling and cell cycle: progression or death. Cell calcium. 2017.
- La Rovere RM, Roest G, Bultynck G, Parys JB. Intracellular Ca2+ signaling and Ca2+ microdomains in the control of cell survival, apoptosis and autophagy. Cell calcium. 2016; 60: 74–87. <u>https://doi.org/10. 1016/j.ceca.2016.04.005</u> PMID: 27157108
- 11. Martin N, Bernard D. Calcium signaling and cellular senescence. Cell calcium. 2017. <u>https://doi.org/10.1016/j.ceca.2017.04.001</u> PMID: 28410770
- Orrenius S, Zhivotovsky B, Nicotera P. Calcium: Regulation of cell death: the calcium–apoptosis link. Nature reviews Molecular cell biology. 2003; 4: 552. https://doi.org/10.1038/nrm1150 PMID: 12838338
- 13. Wei C, Wang X, Chen M, Ouyang K, Song L-S, Cheng H. Calcium flickers steer cell migration. Nature. 2009; 457: 901. https://doi.org/10.1038/nature07577 PMID: 19118385
- Ohno Y, Otaki JM. Spontaneous long-range calcium waves in developing butterfly wings. BMC developmental biology. 2015; 15: 17. https://doi.org/10.1186/s12861-015-0067-8 PMID: 25888365
- Ceriani F, Pozzan T, Mammano F. Critical role of ATP-induced ATP release for Ca2+ signaling in nonsensory cell networks of the developing cochlea. Proceedings of the National Academy of Sciences. 2016; 113: E7194–E7201. https://doi.org/10.1073/pnas.1616061113 PMID: 27807138
- Mammano F, Bortolozzi M. Ca2+ signaling, apoptosis and autophagy in the developing cochlea: milestones to hearing acquisition. Cell calcium. 2018; 70: 117–126. https://doi.org/10.1016/j.ceca.2017.05. 006 PMID: 28578918
- Takeuchi Y, Narumi R, Akiyama R, Vitiello E, Shirai T, Tanimura N, et al. Calcium Wave Promotes Cell Extrusion. Current Biology. 2020; 30: 670–681.e6. https://doi.org/10.1016/j.cub.2019.11.089 PMID: 32004455
- Balaji R, Bielmeier C, Harz H, Bates J, Stadler C, Hildebrand A, et al. Calcium spikes, waves and oscillations in a large, patterned epithelial tissue. Scientific reports. 2017; 7: 42786. <u>https://doi.org/10.1038/</u> srep42786 PMID: 28218282
- Brodskiy PA, Wu Q, Soundarrajan DK, Huizar FJ, Chen J, Liang P, et al. Decoding calcium signaling dynamics during Drosophila wing disc development. Biophysical journal. 2019; 116: 725–740. https:// doi.org/10.1016/j.bpj.2019.01.007 PMID: 30704858
- Restrepo S, Basler K. Drosophila wing imaginal discs respond to mechanical injury via slow InsP 3 Rmediated intercellular calcium waves. Nature communications. 2016; 7: 12450. <u>https://doi.org/10.1038/ ncomms12450 PMID: 27503836</u>
- Buchmann A, Alber M, Zartman JJ. Sizing it up: The mechanical feedback hypothesis of organ growth regulation. Seminars in Cell & Developmental Biology. 2014; 35: 73–81. https://doi.org/10.1016/j. semcdb.2014.06.018 PMID: 25020200
- 22. Gou J, Stotsky JA, Othmer HG. Growth control in the Drosophila wing disk. WIREs Systems Biology and Medicine. 2020; 12: e1478. https://doi.org/10.1002/wsbm.1478 PMID: 31917525
- 23. Hariharan IK. Organ Size Control: Lessons from Drosophila. Developmental Cell. 2015; 34: 255–265. https://doi.org/10.1016/j.devcel.2015.07.012 PMID: 26267393

- Restrepo S, Zartman JJ, Basler K. Coordination of Patterning and Growth by the Morphogen DPP. Current Biology. 2014; 24: R245–R255. https://doi.org/10.1016/j.cub.2014.01.055 PMID: 24650915
- Vollmer J, Iber D. An Unbiased Analysis of Candidate Mechanisms for the Regulation of Drosophila Wing Disc Growth. Scientific Reports. 2016; 6: 39228. <u>https://doi.org/10.1038/srep39228</u> PMID: 27995964
- Vollmer J, Casares F, Iber D. Growth and size control during development. Open biology. 2017; 7: 170190. https://doi.org/10.1098/rsob.170190 PMID: 29142108
- Berridge MJ. Inositol Trisphosphate and Diacylglycerol: Two Interacting Second Messengers. Annual Review of Biochemistry. 1987; 56: 159–193. https://doi.org/10.1146/annurev.bi.56.070187.001111 PMID: 3304132
- 28. Berridge MJ, Lipp P, Bootman MD. The versatility and universality of calcium signalling. Nature reviews Molecular cell biology. 2000; 1: 11. https://doi.org/10.1038/35036035 PMID: 11413485
- Sera T, Kudo S. Mathematical models for intra- and inter-cellular Ca²⁺ wave propagations. J Biorheol. 2020; 34: 9–17. https://doi.org/10.17106/jbr.34.9
- Politi A, Gaspers LD, Thomas AP, Höfer T. Models of IP3 and Ca2+ oscillations: frequency encoding and identification of underlying feedbacks. Biophysical journal. 2006; 90: 3120–3133. https://doi.org/10. 1529/biophysj.105.072249 PMID: 16500959
- Berridge MJ, Irvine RF. Inositol phosphates and cell signalling. Nature. 1989; 341: 197–205. <u>https://doi.org/10.1038/341197a0 PMID: 2550825</u>
- Balakrishnan SS, Basu U, Raghu P. Phosphoinositide signalling in Drosophila. Biochimica et Biophysica Acta (BBA)—Molecular and Cell Biology of Lipids. 2015; 1851: 770–784. https://doi.org/10.1016/j. bbalip.2014.10.010 PMID: 25449646
- Lyon AM, Tesmer VM, Dhamsania VD, Thal DM, Gutierrez J, Chowdhury S, et al. An autoinhibitory helix in the C-terminal region of phospholipase C-β mediates Gα q activation. Nature structural & molecular biology. 2011; 18: 999–1005.
- Li Y-X, Rinzel J. Equations for InsP3 Receptor-mediated [Ca2+]i Oscillations Derived from a Detailed Kinetic Model: A Hodgkin-Huxley Like Formalism. Journal of Theoretical Biology. 1994; 166: 461–473. https://doi.org/10.1006/itbi.1994.1041 PMID: 8176949
- Srikanth S, Wang Z, Tu H, Nair S, Mathew MK, Hasan G, et al. Functional Properties of the Drosophila melanogaster Inositol 1,4,5-Trisphosphate Receptor Mutants. Biophysical Journal. 2004; 86: 3634– 3646. https://doi.org/10.1529/biophysj.104.040121 PMID: 15189860
- Sneyd J, Han JM, Wang L, Chen J, Yang X, Tanimura A, et al. On the dynamical structure of calcium oscillations. Proceedings of the National Academy of Sciences. 2017; 201614613. <u>https://doi.org/10. 1073/pnas.1614613114</u> PMID: 28154146
- Jursnich VA, Fraser SE, Held LI, Ryerse J, Bryant PJ. Defective gap-junctional communication associated with imaginal disc overgrowth and degeneration caused by mutations of the dco gene in Drosophila. Developmental Biology. 1990; 140: 413–429. <u>https://doi.org/10.1016/0012-1606(90)90090-6</u> PMID: 2373260
- Ryerse JS. Gap junctions are non-randomly distributed inDrosophila wing discs. Wilhelm Roux' Archiv. 1982; 191: 335–339. https://doi.org/10.1007/BF00848494 PMID: 28305287
- Weir MP, Lo CW. Gap junctional communication compartments in the Drosophila wing disk. PNAS. 1982; 79: 3232–3235. https://doi.org/10.1073/pnas.79.10.3232 PMID: 6954475
- 40. Weir MP, Lo CW. An anterior/posterior communication compartment border in engrailed wing discs: Possible implications for Drosophila pattern formation. Developmental Biology. 1985; 110: 84–90. https://doi.org/10.1016/0012-1606(85)90066-1 PMID: 4007269
- Gómez HF, Georgieva L, Michos O, Iber D. Image-Based In silico Models of Organogenesis. Systems Biology. John Wiley & Sons, Ltd; 2017. pp. 319–340. https://doi.org/10.1002/9783527696130.ch12
- Ratnaparkhi A, Banerjee S, Hasan G. Altered Levels of Gq Activity Modulate Axonal Pathfinding inDrosophila. J Neurosci. 2002; 22: 4499–4508. https://doi.org/20026385 PMID: 12040057
- Hanlon CD, Andrew DJ. Outside-in signaling-a brief review of GPCR signaling with a focus on the Drosophila GPCR family. J Cell Sci. 2015; 128: 3533–3542. <u>https://doi.org/10.1242/jcs.175158</u> PMID: 26345366
- García-Bellido A, Cortés F, Milán M. Cell interactions in the control of size in Drosophila wings. Proc Natl Acad Sci U S A. 1994; 91: 10222–10226. <u>https://doi.org/10.1073/pnas.91.21.10222</u> PMID: 7937866
- Wyss C., Ecdysterone, insulin and fly extract needed for the proliferation of normal Drosophila cells in defined medium. Experimental cell research. 1982; 139: 297–307. <u>https://doi.org/10.1016/0014-4827</u> (82)90254-3 PMID: 6806111

- Zartman J, Restrepo S, Basler K. A high-throughput template for optimizing Drosophila organ culture with response-surface methods. Development. 2013; 140: 667–674. <u>https://doi.org/10.1242/dev.</u> 088872 PMID: 23293298
- 47. Hevia CF, López-Varea A, Esteban N, de Celis JF. A Search for Genes Mediating the Growth-Promoting Function of TGFβ in the Drosophila melanogaster Wing Disc. Genetics. 2017; 206: 231–249. https://doi.org/10.1534/genetics.116.197228 PMID: 28315837
- Nakai J, Ohkura M, Imoto K. A high signal-to-noise Ca 2+ probe composed of a single green fluorescent protein. Nature Biotechnology. 2001; 19: 137–141. https://doi.org/10.1038/84397 PMID: 11175727
- Chen T-W, Wardill TJ, Sun Y, Pulver SR, Renninger SL, Baohan A, et al. Ultrasensitive fluorescent proteins for imaging neuronal activity. Nature. 2013; 499: 295–300. https://doi.org/10.1038/nature12354 PMID: 23868258
- Yamauchi T. Neuronal Ca²⁺/Calmodulin-Dependent Protein Kinase II—Discovery, Progress in a Quarter of a Century, and Perspective: Implication for Learning and Memory. Biological and Pharmaceutical Bulletin. 2005; 28: 1342–1354. https://doi.org/10.1248/bpb.28.1342 PMID: 16079472
- Johnston LA, Gallant P. Control of growth and organ size in Drosophila. BioEssays. 2002; 24: 54–64. https://doi.org/10.1002/bies.10021 PMID: 11782950
- 52. Potter C. Mechanisms of size control. Current Opinion in Genetics & Development. 2001; 11: 279–286. https://doi.org/10.1080/14728222.2018.1464141 PMID: 29634426
- Larkin JW, Zhai X, Kikuchi K, Redford SE, Prindle A, Liu J, et al. Signal Percolation within a Bacterial Community. Cell systems. 2018; 7: 137–145. https://doi.org/10.1016/j.cels.2018.06.005 PMID: 30056004
- Justet C, Hernández JA, Torriglia A, Chifflet S. Fast calcium wave inhibits excessive apoptosis during epithelial wound healing. Cell and tissue research. 2016; 365: 343–356. https://doi.org/10.1007/ s00441-016-2388-8 PMID: 26987821
- 55. Kaouri K, Maini PK, Skourides PA, Christodoulou N, Chapman SJ. A simple mechanochemical model for calcium signalling in embryonic epithelial cells. Journal of Mathematical Biology. 2019. <u>https://doi.org/10.1007/s00285-019-01333-8 PMID: 30826846</u>
- Richard M, Bauer R, Tavosanis G, Hoch M. The gap junction protein Innexin3 is required for eye disc growth in Drosophila. Developmental Biology. 2017; 425: 191–207. https://doi.org/10.1016/j.ydbio. 2017.04.001 PMID: 28390801
- Nobuo Homma, Alvarado José Luis, Coombs Wanda, Kathleen Stergiopoulos, Taffet Steven M., Lau Alan F., et al. A Particle-Receptor Model for the Insulin-Induced Closure of Connexin43 Channels. Circulation Research. 1998; 83: 27–32. https://doi.org/10.1161/01.res.83.1.27 PMID: 9670915
- Lin D, Boyle DL, Takemoto DJ. IGF-I-induced phosphorylation of connexin 43 by PKCgamma: regulation of gap junctions in rabbit lens epithelial cells. Invest Ophthalmol Vis Sci. 2003; 44: 1160–1168. https://doi.org/10.1167/iovs.02-0737 PMID: 12601045
- Warn-Cramer BJ, Lau AF. Regulation of gap junctions by tyrosine protein kinases. Biochim Biophys Acta. 2004; 1662: 81–95. https://doi.org/10.1016/j.bbamem.2003.10.018 PMID: 15033580
- O'Connor JT, Stevens AC, Shannon EK, Akbar FB, LaFever KS, Narayanan N, et al. A protease-initiated model of wound detection. bioRxiv. 2020; 2020.12.08.415554. https://doi.org/10.1101/2020.12.08. 415554
- 61. Hart Y, Alon U. The Utility of Paradoxical Components in Biological Circuits. Molecular Cell. 2013; 49: 213–221. https://doi.org/10.1016/j.molcel.2013.01.004 PMID: 23352242
- Loew LM. Where does all the PIP2 come from? The Journal of physiology. 2007; 582: 945–951. https://doi.org/10.1113/jphysiol.2007.132860 PMID: 17395623
- Czech MP. PIP2 and PIP3: complex roles at the cell surface. Cell. 2000; 100: 603–606. <u>https://doi.org/10.1016/s0092-8674(00)80696-0 PMID</u>: 10761925
- Brand AH, Perrimon N. Targeted gene expression as a means of altering cell fates and generating dominant phenotypes. Development. 1993; 118: 401–415. PMID: 8223268
- **65.** Duffy JB. GAL4 system in Drosophila: a fly geneticist's Swiss army knife. genesis. 2002; 34: 1–15. https://doi.org/10.1002/gene.10150 PMID: 12324939
- 66. Narciso C, Wu Q, Brodskiy P, Garston G, Baker R, Fletcher A, et al. Patterning of wound-induced intercellular Ca2+ flashes in a developing epithelium. Physical biology. 2015; 12: 056005. https://doi.org/10. 1088/1478-3975/12/5/056005 PMID: 26331891
- Perkins LA, Holderbaum L, Tao R, Hu Y, Sopko R, McCall K, et al. The transgenic RNAi project at Harvard Medical School: resources and validation. Genetics. 2015; 201: 843–852. https://doi.org/10.1534/genetics.115.180208 PMID: 26320097

- Werz C, Köhler K, Hafen E, Stocker H. The Drosophila SH2B family adaptor Lnk acts in parallel to chico in the insulin signaling pathway. PLoS genetics. 2009; 5: e1000596. <u>https://doi.org/10.1371/journal.pgen.1000596</u> PMID: 19680438
- Kakanj P, Moussian B, Grönke S, Bustos V, Eming SA, Partridge L, et al. Insulin and TOR signal in parallel through FOXO and S6K to promote epithelial wound healing. Nature communications. 2016; 7: 12972. https://doi.org/10.1038/ncomms12972 PMID: 27713427
- Narciso CE, Contento NM, Storey TJ, Hoelzle DJ, Zartman JJ. Release of applied mechanical loading stimulates intercellular calcium waves in Drosophila wing discs. Biophysical journal. 2017; 113: 491– 501. https://doi.org/10.1016/j.bpj.2017.05.051 PMID: 28746859
- Burnette M, Brito-Robinson T, Li J, Zartman J. An inverse small molecule screen to design a chemically defined medium supporting long-term growth of Drosophila cell lines. Molecular BioSystems. 2014; 10: 2713–2723. https://doi.org/10.1039/c4mb00155a PMID: 25096480
- Lloyd S. Least squares quantization in PCM. IEEE transactions on information theory. 1982; 28: 129– 137.
- Wilson EB. Probable Inference, the Law of Succession, and Statistical Inference. Journal of the American Statistical Association. 1927; 22: 209–212. https://doi.org/10.2307/2276774
- 74. Newcombe RG. Interval estimation for the difference between independent proportions: comparison of eleven methods. Stat Med. 1998; 17: 873–890. https://doi.org/10.1002/(sici)1097-0258(19980430) 17:8<873::aid-sim779>3.0.co;2-i PMID: 9595617
- 75. Newcombe RG. Two-sided confidence intervals for the single proportion: comparison of seven methods. Stat Med. 1998; 17: 857–872. https://doi.org/10.1002/(sici)1097-0258(19980430)17:8<857::aid-sim777>3.0.co;2-e PMID: 9595616
- 76. Su T, Ludwig MZ, Xu J, Fehon RG. Kibra and Merlin Activate the Hippo Pathway Spatially Distinct from and Independent of Expanded. Developmental Cell. 2017; 40: 478–490.e3. <u>https://doi.org/10.1016/j. devcel.2017.02.004 PMID: 28292426</u>
- Rauskolb C, Sun S, Sun G, Pan Y, Irvine KD. Cytoskeletal Tension Inhibits Hippo Signaling through an Ajuba-Warts Complex. Cell. 2014; 158: 143–156. https://doi.org/10.1016/j.cell.2014.05.035 PMID: 24995985