

RESEARCH ARTICLE

Symbiotic Cell Differentiation and Cooperative Growth in Multicellular Aggregates

Jumpei F Yamagishi¹, Nen Saito^{2*}, Kunihiro Kaneko^{2*}

¹ College of Arts and Sciences, The University of Tokyo, Meguro-ku, Tokyo, Japan, ² Graduate School of Arts and Sciences, The University of Tokyo, Meguro-ku, Tokyo, Japan

* saito@complex.c.u-tokyo.ac.jp (NS); kaneko@complex.c.u-tokyo.ac.jp (KK)



OPEN ACCESS

Citation: Yamagishi JF, Saito N, Kaneko K (2016) Symbiotic Cell Differentiation and Cooperative Growth in Multicellular Aggregates. *PLoS Comput Biol* 12(10): e1005042. doi:10.1371/journal.pcbi.1005042

Editor: Richard (Rick) Michod, The University of Arizona, UNITED STATES

Received: February 10, 2016

Accepted: June 29, 2016

Published: October 17, 2016

Copyright: © 2016 Yamagishi et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

Data Availability Statement: All relevant data are within the paper and its Supporting Information files.

Funding: The Platform Project for Supporting in Drug Discovery and Life Science Research (Platform for Dynamic Approaches to Living System) from Japan Agency for Medical Research and Development (AMED) supported this work. The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

Abstract

As cells grow and divide under a given environment, they become crowded and resources are limited, as seen in bacterial biofilms and multicellular aggregates. These cells often show strong interactions through exchanging chemicals, as evident in quorum sensing, to achieve mutualism and division of labor. Here, to achieve stable division of labor, three characteristics are required. First, isogenous cells differentiate into several types. Second, this aggregate of distinct cell types shows better growth than that of isolated cells without interaction and differentiation, by achieving division of labor. Third, this cell aggregate is robust with respect to the number distribution of differentiated cell types. Indeed, theoretical studies have thus far considered how such cooperation is achieved when the ability of cell differentiation is presumed. Here, we address how cells acquire the ability of cell differentiation and division of labor simultaneously, which is also connected with the robustness of a cell society. For this purpose, we developed a dynamical-systems model of cells consisting of chemical components with intracellular catalytic reaction dynamics. The reactions convert external nutrients into internal components for cellular growth, and the divided cells interact through chemical diffusion. We found that cells sharing an identical catalytic network spontaneously differentiate via induction from cell-cell interactions, and then achieve division of labor, enabling a higher growth rate than that in the unicellular case. This symbiotic differentiation emerged for a class of reaction networks under the condition of nutrient limitation and strong cell-cell interactions. Then, robustness in the cell type distribution was achieved, while instability of collective growth could emerge even among the cooperative cells when the internal reserves of products were dominant. The present mechanism is simple and general as a natural consequence of interacting cells with limited resources, and is consistent with the observed behaviors and forms of several aggregates of unicellular organisms.

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

Author Summary

Unicellular organisms, when aggregated under limited resources, often exhibit behaviors akin to multicellular organisms, possibly without advanced regulation mechanisms, as observed in biofilms and bacterial colonies. Cells in an aggregate have to differentiate into several types that are specialized for different tasks, so that the growth rate should be enhanced by the division of labor among these cell types. To consider how a cell aggregate can acquire these properties, most theoretical studies have thus far assumed the fitness of an aggregate of cells and the ability of cell differentiation *a priori*. In contrast, we developed a dynamical-systems model consisting of cells without assuming predefined fitness. The model consists of catalytic-reaction networks for cellular growth. By extensive simulations and theoretical analysis of the model, we showed that cells growing under the condition of nutrient limitation and strong cell-cell interactions can differentiate with distinct chemical compositions. They achieve cooperative division of labor by exchanging the produced chemicals to attain a higher growth rate. The conditions for spontaneous cell differentiation and collective growth of cells are presented. The uncovered symbiotic differentiation and collective growth are akin to economic theory on division of labor and comparative advantage.

Introduction

As unicellular organisms grow and divide, they often form a crowded aggregate. As exemplified by bacterial biofilms [1–3] and slime molds [4, 5], these aggregates are not merely crowded passively, but sometimes form a functional cell aggregate, in which cells strongly interact with each other by exchanging chemicals, as demonstrated with quorum sensing [6]. Such a “multicellular aggregate” is often observed to form under a limited resource condition, which may indicate that formation of an aggregate is a universal strategy for a unicellular organism to survive in a severe environment and for cells to grow collectively and cooperatively. Interestingly, mutualistic behaviors, cell differentiation, and division of labor are ubiquitously observed in such aggregates with isoclonal cells [1–3, 7–9] as well as with heterogeneous cells (e.g., bacterial ecosystem) [1–3, 10–12]. This raises the questions of how aggregates of identical cells achieve division of labor for cooperative growth, and what are the necessary conditions? These questions are important to be addressed in order to understand the formation of multicellular aggregates, including the formation of biofilms, which has attracted much attention recently [1–3].

From this point of view, there are at least three characteristics required to achieve stably growing aggregates with division of labor.

- (i) Cell differentiation: starting from a single cell, cell states become differentiated as their numbers increase and they coexist and grow.
- (ii) Cooperative growth: these different cell types mutually cooperate for their growth, possibly by division of labor, to avoid being driven to extinction by competition with the isolated, undifferentiated cells.
- (iii) Robustness in the distribution of the number of differentiated cells: balance in the population distribution of cell types is maintained so that cells of different types can coexist stably and grow together.

To achieve stable task differentiation, (i) cell differentiation through cell-cell interaction would be necessary, whereas (ii) cooperative growth is also required, since otherwise community formation through cell-cell communication would not be advantageous (or might even be deleterious) compared with the case of isolated cells without any interaction. However, simply achieving this cooperative growth does not necessarily imply that this state is robust, since if one cell type reproduces faster than any other type, the fastest type would dominate the population and the appropriate cell type ratio for division of labor would be easily lost. Therefore, (iii) coexistence of diverse cell types is also an important issue to be addressed for the stability and survival of a cell colony.

Indeed, such characteristics have also been studied as a primitive form of multicellularity. In experimental evolution, aggregation of unicellular organisms with division of labor has been recently investigated with the use of yeast [13] and algae [14]. With respect to theoretical approaches, a related issue of the survival of an aggregate of cells has been conventionally discussed in multi-level evolution theory, by introducing a fitness parameter at the cellular and multi-cellular levels, and investigating how these two fitness values are aligned [15–21]. In most of the previous studies based on the prescribed fitness, however, the existence of differentiated cell types is presumed, and thus the capacity of cell differentiation and the fitness alignment are separated [15–23]. Related criticisms of these previous approaches are discussed in [24, 25]. Specifically in [24], intracellular dynamics is introduced as the optimization of resource allocation to different tasks under a given artificial fitness function, and it is shown that division of labor emerges when it increases the fitness. However, in nature, in general, cell differentiation does not result from optimization of a given fitness but rather results from intra-cellular metabolic reaction dynamics, and thus the division of labor is not guaranteed even when it increases fitness. The fitness, i.e., the rate of cellular growth, is also obtained through the reaction dynamics. Therefore, it will be important to take cell differentiation and growth rate into account simultaneously, as a result of intra-cellular reaction dynamics, where the growth rate of each cell type is not predetermined, but rather changes according to the cellular states. Furthermore, this growth state also depends on the states of surrounding cells, which may alter the abundances of available resources and the strength of cell-cell interactions. To consider these issues that have not been addressed in the previous studies, we here present a dynamical-systems model of cells with intracellular reactions, cell-cell interactions, and uptake of resources, by which the fitness is determined as the cellular growth rate, rather than being prescribed in advance.

In fact, such models of interacting and growing cells with intracellular reaction dynamics have been introduced previously, where the concept of isologous diversification [26] has been proposed, to address differentiation from a single cell type (property (i)). A previous mathematical model [27] demonstrated that an ensemble of cells sharing a common genotype could achieve niche differentiation through cell differentiation, and thereby relax the strength of resource competition. Although this indirect cooperation through niche differentiation would be beneficial for cell aggregates, we here address cooperative growth in a stronger sense, where differentiated cells help each other so that interacting cells in an aggregate grow faster than the isolated undifferentiated cells (unicellular organisms) under the condition of limited resources (property (ii)). For this purpose, we here consider an environment in which only a single resource exists, and in such situation, property (ii) is considered as the property of the cell ensemble to help each other achieve a higher growth than the isolated cells, rather than specializing to each resource.

In the present paper, by using a simple model of cells that contain diverse components and interact with each other through the exchange of chemicals, we address the question of whether the above three characteristic behaviors are a necessary outcome of an ensemble of interacting

cells. Specifically, we show that a cell ensemble under strong cell-cell interactions with limited resources fulfills cell differentiation, cooperative growth, and robustness in the cell type distribution.

Models

We consider a mathematical model proposed in [26–29], which describes a simple, primitive cell that consists of k chemical components $\{X_0, \dots, X_{k-1}\}$. As illustrated in Fig 1, we assume that n cells globally interact with each other in a well-mixed medium, and each of them grows by uptake of the nutrient chemical X_0 , which is supplied into the medium from the external environment. The internal state of each cell is characterized by a set of variables $(x_0^{(m)}, \dots, x_{k-1}^{(m)}, v^{(m)})$, where $x_i^{(m)}$ is the concentration of the i -th chemical X_i , and $v^{(m)}$ is the volume of the m -th cell ($m = 1, \dots, n$). As a simple model, we consider a situation with only catalysts and resources, where these k components are mutually catalyzed for their synthesis, thus forming a catalytic reaction network. A catalytic reaction from a substrate X_i to a product X_j by a catalyst X_l as $X_i + \alpha X_l \rightarrow X_j + \alpha X_l$ occurs at a rate $\epsilon x_i^{(m)} x_l^{(m)\alpha}$, where α refers to the order of the catalytic reaction and is mostly set as $\alpha = 2$. Here, ϵ is the rate constant for this reaction, and, for simplicity, all the rate constants are equally fixed at $\epsilon = 1$. The parameters and variables in this model are listed in Table 1.

Cell states change through intracellular biochemical reaction dynamics and the in- and out-flow of chemicals, leading to cell-cell interactions via the medium. The medium's state is characterized by concentrations $(x_0^{(med)}, \dots, x_{k-1}^{(med)})$, and a constant volume V_{med} . Then, the dynamics of the concentration of X_i in the m -th cell are represented as:

$$\frac{dx_i^{(m)}}{dt} = \sum_{j,l=0}^{k-1} \epsilon P(j,i,l) x_j^{(m)} x_l^{(m)\alpha} - \sum_{j,l=0}^{k-1} \epsilon P(i,j,l) x_i^{(m)} x_l^{(m)\alpha} + D \sigma_i (x_i^{(med)} - x_i^{(m)}) - x_i^{(m)} \mu^{(m)}, \quad (1)$$

where $P(i,j,l)$ takes the value 1 if there is a reaction $X_i + \alpha X_l \rightarrow X_j + \alpha X_l$, and is 0 otherwise. In Eq (1), the third term describes the influx of X_i from the medium, and the fourth term gives the dilution owing to the volume growth of the cell, and $\mu^{(m)}$ denotes the cellular growth rate. Here, only a subset of chemical species is diffusible across the cell membranes with the rate of diffusion D . X_i is transported from the medium into the m -th cell at a rate $D \sigma_i (x_i^{(med)} - x_i^{(m)})$, where σ_i is 1 if X_i is diffusible, and is 0 otherwise. Therefore, the m -th cell grows in volume according

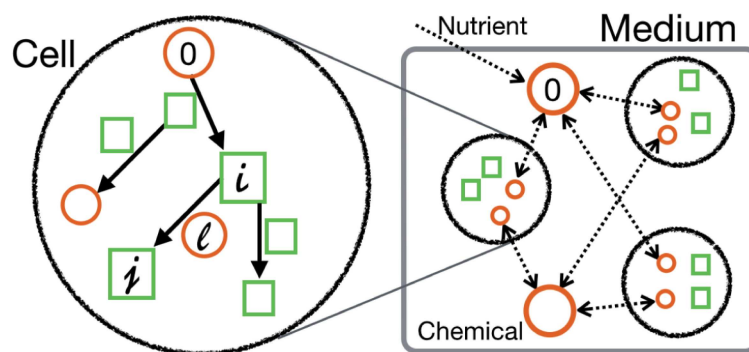


Fig 1. Schematic illustration of the N -cell model. Each cell has a common catalytic network, while the nutrient X_0 transported from the medium is transformed to catalytic components for cellular growth. The nutrient X_0 is supplied to the medium from the exterior. The reaction from X_i to X_j is catalyzed by another component, X_l . Some components (orange circles) are diffusible and exchanged via the medium, while others (green squares) are not. In the medium, n cells coexist and interact with each other ($1 \leq n \leq N$).

doi:10.1371/journal.pcbi.1005042.g001

Table 1. The parameters and variables of the N-cell model.

Symbol	Value	Interpretation
N	100, 1	The maximum number of cells coexisting in a medium
k	20	The number of chemicals
α	2	The order of catalytic reactions
ϵ	1	The rate constant for catalytic reactions
C	0.05, 0.15	The concentration of the nutrient X_0 in the medium's exterior
V_{med}	100	The volume of the medium
D	0.1, 1	The diffusion coefficient between cells and the medium
D_{med}	0.1, 1	The diffusion coefficient between the medium and its exterior
$P(i, j, l)$	0, 1	Whether or not the reaction $X_i + \alpha X_j \rightarrow X_j + \alpha X_i$ exists
$x_i^{(m)}$	variable	The concentration of the i -th chemical X_i in the m -th cell
$x_i^{(med)}$	variable	The concentration of the i -th chemical X_i in the medium
$v^{(m)}$	variable	The volume of the m -th cell
$\mu^{(m)}$	variable	The growth rate of the m -th cell

doi:10.1371/journal.pcbi.1005042.t001

to the rate $\mu^{(m)} \equiv \sum_{i=0}^{k-1} D\sigma_i(x_i^{(med)} - x_i^{(m)})$ by assuming that the cellular volume is in proportion to the total amount of chemicals. The volume dynamics are given by $dv^{(m)}/dt = \mu^{(m)}v^{(m)}$. As the abundances of chemicals are conserved by the intracellular reactions, with this form of volume growth, $\sum_{i=0}^{k-1} x_i^{(m)} = 1$ is time-invariant [28]. The nutrient chemical X_0 , which is necessary for cellular growth, is supplied into the medium from the external environment according to the rate $D_{med}(C - x_0^{(med)})$, where D_{med} denotes the diffusion coefficient of the nutrient across the medium's boundary, whereas C is the constant external concentration of the nutrient X_0 (for simplicity, the flow of the other diffusible chemicals to the outside of the medium is not included, although its inclusion does not alter the result below as long as their D_{med} values are not large).

Therefore, the temporal change of $x_i^{(med)}$ is given by

$$\frac{dx_i^{(med)}}{dt} = D_{med}\sigma'_0(C - x_i^{(med)}) - \sum_{m=1}^n \frac{D\sigma_i(x_i^{(med)} - x_i^{(m)})v^{(m)}}{V_{med}}, \quad (2)$$

where σ'_0 takes unity only if $i = 0$, i.e., if X_i is the nutrient. For simplicity, D_{med} was set as $D_{med} = D$, though the results reported here do not greatly depend on the value of D_{med} .

According to these processes, each cell grows by converting nutrient chemicals into non-diffusible chemicals and storing them within the cell until its volume doubles, and then divides into two cells with almost the same chemical compositions. Here, the catalytic network in daughter cells is identical to that in their mother cell. As the initial condition, only a single cell exists with a randomly determined chemical composition. In addition, we set the carrying capacity of a medium N , which is an upper limit to the number of cells that can coexist in the medium. When the cell number exceeds its upper limit N due to cell division, the surplus cells are randomly eliminated. Hereafter, this model is referred to as the N -cell model.

Results

Cell differentiation in the N-cell model: Brief summary

We simulated the N -cell model over hundreds of randomly generated reaction networks. Each catalytic network is generated in the following manner. Each chemical is set to be diffusible

with probability $q = 0.15$ and has $\rho = 4$ outward reaction paths to other chemicals; i.e., each chemical works as a substrate in ρ reactions. Each reaction $X_i + \alpha X_l \rightarrow X_j + \alpha X_l$ ($i \neq j$, and X_j and X_l are not nutrients) is randomly determined so that $j \neq l$ is fulfilled. We did not allow for autocatalytic reactions ($j = l$) as they are rare in nature. However, the described results were also obtained without these restrictions.

We are particularly interested in if and how the cells differentiate, and whether the growth rate would increase as a result of differentiation. For this purpose, cell differentiation is defined as the emergence of cells with different chemical compositions within the population that share an identical catalytic network. For the case where the concentrations asynchronously oscillate in time, we evaluated whether cells have different compositions even after taking the temporal average over a sufficiently longer time scale than the oscillation period. To evaluate the growth enhancement, we compared two different situations, “interacting” ($N = 100$) and “isolated” ($N = 1$) cases, and then we computed R_μ , the ratio of the growth rate of interacting cells to that of isolated cells. Then the growth enhancement is defined as $R_\mu > 1$.

The behavior of the N -cell model is classified into four categories. In category (a), interacting cells differentiate into two or more types and grow faster than isolated cells, i.e., $R_\mu > 1$ (Fig 2; see also Figure A in S1 Text). In category (b), interacting cells differentiate but their growth is slower than that of isolated cells ($R_\mu < 1$); in this category, as far as we have examined, cells of a certain type gain chemicals diffused from another type, which are used as catalysts for conversion to non-diffusible chemicals. The latter cell type has a composition similar to that of the isolated cell, and its growth is decreased by this cell-cell interaction (see Figure B in S1 Text). Hence, the former cell type is considered to exploit the latter as it receives the unidirectional chemical inflow. In category (c), cells do not differentiate with respect to the average composition, but chemical concentrations asynchronously oscillate in time. In category (d), the behavior of each cell is identical, regardless of the presence or absence of cell-cell interactions, and therefore $R_\mu = 1$.

Here, we are mainly concerned with category (a), as this case enables both cell differentiation and cooperative growth. We found four common properties in this category. (1) A state with homogeneity among cells becomes unstable as the cell number increases, and is replaced by two (or more) distinct cellular states. (2) In differentiated cells, the compositions are concentrated for only a few chemicals, whereas the concentrations of the other chemicals are nearly zero; i.e., each cell type uses only a sub-network of the total reaction network. (3) Different cell types share only a few common components, and each of the other components mostly exists in one cell type. (4) The components that predominate in one cell type diffuse to the other cell type, where they function as catalysts, and vice versa. Thus, the two cell types help each other to achieve higher cooperative growth.

Reduction of the N -cell model

After examining a number of networks in category (a), we extracted a common core structure in the reaction network topology, designated as networks 1-3 (Fig 3A and 3B; see also Figure C in S1 Text). In these networks, cells in the N -cell model differentiate into two types, type-1 and type-2, as exemplified in Fig 3C. In type-1, x_1 is high while x_2 is close to zero, and in type-2, x_2 is high and x_1 is close to zero. Accordingly, X_3 (X_4) can be produced only in the former (latter) type, and the two types of cells complement each other by exchanging X_3 and X_4 . Consequently, the differentiated cells grow faster than the isolated cells (Fig 3D).

To analyze the mechanism of this cooperative differentiation, we reduced the N -cell dynamics to two effective groups of cells represented by $(x_0^{(i)}, \dots, x_{k-1}^{(i)}, v^{(i)})$, where $v^{(i)}$ denotes the total volume of each cell group ($i = 1, 2$). Considering that the total cell number is sustained at

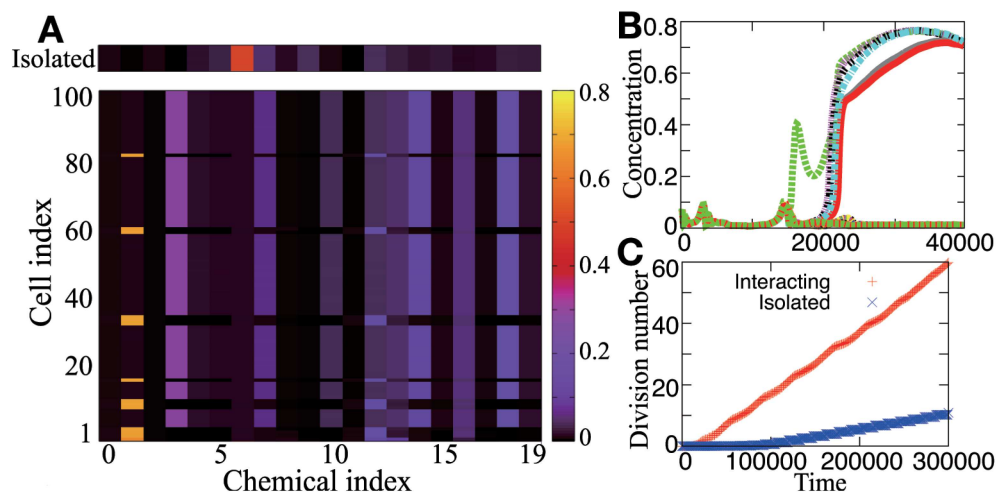


Fig 2. Typical behavior of the N -cell model in category (a). (A) Chemical compositions of N cells surviving at time $t = 5 \times 10^5$. The concentration $x_i^{(m)}$ is plotted with a color code, with the vertical axis indicating the cell index m , and the horizontal axis indicating the chemical index i , while the top band designates the composition of an “isolated” cell. Cells differentiate into two types with distinct compositions. (B) The time series of $x_1^{(m)}$ in interacting N cells surviving at time $t = 4 \times 10^4$, overlaid for all cells shown as different colors. Each line represents the concentration of the chemical X_1 in a different cell, plotted with different colors over 100 cells. (C) The time series of the number of cell divisions per cell for interacting (red) and isolated (blue) cells. Interacting cells grow faster than isolated cells; i.e., $R_\mu > 1$.

doi:10.1371/journal.pcbi.1005042.g002

its maximum N , the total cellular volume is also bounded. Therefore, $v = v^{(1)} + v^{(2)}$ is regarded as a constant in the reduced version of interacting cells, termed the reduced-2cell (r2cell) model. This model obeys Eqs (1) and (2) with $\mu^{(m)} \equiv \sum_{i=0}^{k-1} D\sigma_i(x_i^{(med)} - x_i^{(m)})$, and the effect of random cell elimination associated with cell division is implicitly incorporated into dilution due to volume growth. Besides, by considering the symmetry in networks 1-3, we can assume $v^{(1)} = v^{(2)} = v/2$ for symmetric differentiation with the same number of cells of the two types, while the case with $v^{(1)} \neq v^{(2)}$ will be discussed later.

Likewise, we also consider the reduced-1cell (r1cell) model corresponding to the “isolated” case of the N -cell model, by ignoring cell division and assuming that the cellular volume is constant at $v^{(iso)} = v$.

Cell differentiation

The behavior of the r1cell and r2cell models (i.e., isolated and interacting cells) can be classified into several phases, depending on parameters (C, V, D) , where $V \equiv V_{med}/v$ is the volume ratio between the cells and the medium.

The phase diagram with network 1 for $D = 1$ is shown in Fig 4A, and Figure E in S1 Text shows phase diagrams of networks 1-3 for various D values. The blue area in Fig 4A represents phase (I), in which the cells cannot differentiate, and always reach a single fixed point attractor in both the r1cell and r2cell models. In phase (II), differentiation into two fixed points occurs in the r2cell model from a stable fixed point in the r1cell model, as shown in Fig 4B. In phase (III), the r1cell model exhibits oscillation, while two cells in the r2cell model reach two distinct fixed points (Fig 4C). In terms of dynamical systems theory, this loss of oscillation is referred to as oscillation death [30, 31]. In phase (IV), both “oscillation-death” differentiation and synchronous oscillation (i.e., non-differentiation) can occur depending on the initial condition, whereas the r1cell model always exhibits oscillation.

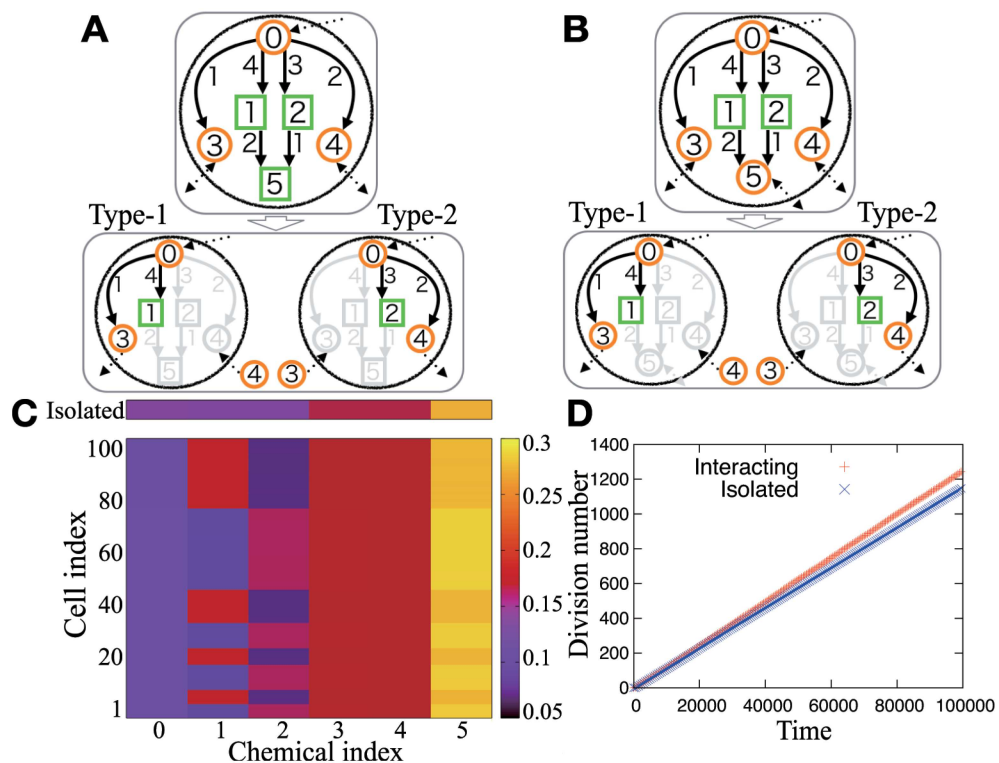


Fig 3. Behavior of simplified networks. (A) Network 1. (B) Network 2. In (A) and (B), each number represents the chemical index, with orange-circle and green-square nodes representing diffusible and non-diffusible chemicals, respectively. The difference between networks 1 and 2 lies only in the diffusibility of X_5 . The dashed arrows denote the diffusive fluxes of chemicals, and the thick arrows indicate catalytic reactions. The chemical at the arrowtail is transformed to the chemical at the arrowhead, catalyzed by the chemical labeled on the edge; e.g., the left arrow from X_0 in (A) denotes the catalytic reaction $X_0 + \alpha X_1 \rightarrow X_3 + \alpha X_1$. When the cells differentiate, type-1 (type-2) cells mainly produce the chemicals X_1 and X_3 (X_2 and X_4), and receive X_4 (X_3) from type-2 (type-1) cells, which are illustrated in the lower panel with color. (C) and (D) are examples of the behavior of network 1 for $(C, V_{med}, D) = (0.15, 100, 1)$. (C) Chemical concentrations $X_i^{(m)}$ in N cells surviving at time $t = 10^5$ are plotted according to the color code shown in the sidebar, with the vertical axis as the cell index m and the horizontal axis as the chemical index i . The top band designates an “isolated” cell. (D) The time series of the number of cell divisions per cell for interacting (red) and isolated (blue) cells.

doi:10.1371/journal.pcbi.1005042.g003

Thus, differentiation occurs in phases (II)-(IV) (i.e., at the left of the green line in Fig 4A), while stable differentiation without falling into synchronized oscillation is achieved only in phases (II)-(III), i.e., with small C and V values, representing a limited resource and strong cell-cell interaction condition. In Fig 4A, phases (II)-(III) are divided by the red line, and the red and green lines are determined according to linear stability analysis (see S1 Text for details).

With respect to the network structure, the catalytic reactions $X_1 + \alpha X_2 \rightarrow X_5 + \alpha X_2$ and $X_2 + \alpha X_1 \rightarrow X_5 + \alpha X_1$ function as two mutually repressive reactions, i.e., forming a double-negative feedback loop. Further, the product X_5 consumes X_1 and X_2 , and is maintained within the cell, which enhances the dilution of X_1 and X_2 . Thus, each of these reactions works as a composite negative feedback loop, leading to instability of the homogeneous cell state. Since nonlinearity is a necessary condition for multi-stability, a high order of catalytic reactions α tends to facilitate cell differentiation.

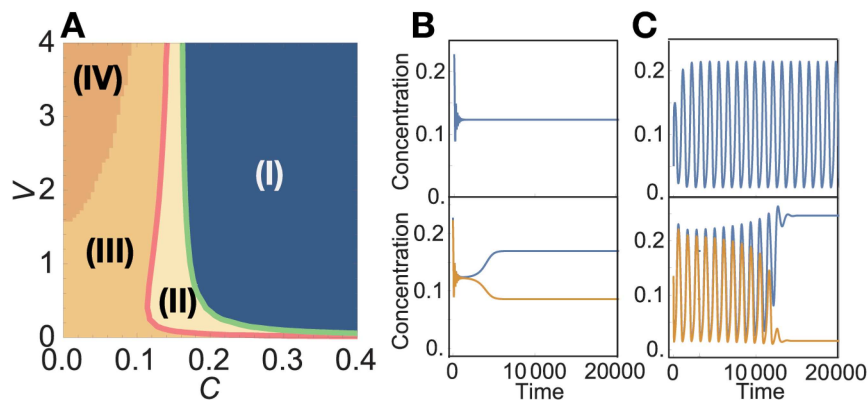


Fig 4. The behavior of the r1cell and r2cell models with network 1. (A) A phase diagram ($D = 1$). The blue area designates phase (I) in which cells cannot exhibit differentiation, and reach a single fixed point in both the r1cell and r2cell models. In phases (II) and (III), cells always differentiate. In phase (II), shown in cream color, cells exhibit pitch-fork bifurcation from a homogeneous state. In phase (III), “oscillation-death” occurs in the dynamics of the chemical compositions of the two cells. In phase (IV), depicted by orange, if the initial difference between the compositions of the two cells is large enough, “oscillation-death” occurs; otherwise, the r2cell model exhibits non-differentiated, synchronized oscillation. (B) The time series of concentrations of X_1 in phase (II) for $(C, V, D) = (0.27, 0.1, 1)$. (C) The time series in phase (III) for $(C, V, D) = (0.1, 0.8, 0.1)$. For both (B) and (C), the time series for the r1cell (r2cell) model is displayed on the upper (lower) column.

doi:10.1371/journal.pcbi.1005042.g004

Cooperative growth by differentiation

Fig 5A shows the dependence of R_μ on parameters in network 1, exemplifying that differentiation increases the growth rate. Surprisingly, this differentiation-induced growth enhancement was always observed for any set of parameters in networks 1-3 (network 3 is shown in Figure C in S1 Text).

We next sought to determine the mechanism contributing to the faster growth of differentiated cells. An intuitive explanation is as follows. On one hand, an isolated cell must contain all chemical components required for self-reproduction (e.g., X_0 - X_5 in the upper panel of Fig 3A), leading to lower concentrations of each chemical on average. On the other hand, differentiated cells can achieve division of labor; each type of differentiated cell exclusively produces a portion of the required chemical species, and cells exchange these chemicals with each other. Since

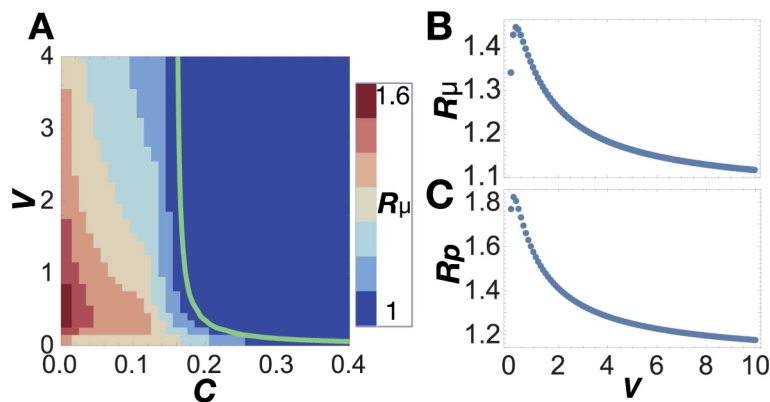


Fig 5. Growth enhancement with network 1 ($D = 1$). (A) Dependence of R_μ on parameters C and V . Cells can differentiate given the parameters to the left of the green line. (B) Dependence of R_μ on V ($C = 0.1$). R_μ is the ratio of the growth rate of differentiated cells to that of isolated cells. (C) Dependence of R_p on V ($C = 0.1$). R_p is the degree of increase in the production of X_3 - X_4 conferred by differentiation.

doi:10.1371/journal.pcbi.1005042.g005

catalytic reactions occur only in a sub-network of the original network (e.g., a network in the lower panel of Fig 3A), the chemicals are concentrated on fewer components, which increases the efficiency of chemical reactions and promotes cellular growth.

This suggests that stronger cell-cell interactions support higher growth. Indeed, Fig 5B shows that a smaller V , i.e., stronger cell-cell interaction, causes larger R_μ . A smaller V also increases R_p , the ratio of the total production of X_3 - X_4 in the r2cell model to that in the r1cell model (Fig 5C); that is, the production of exchanged chemicals is enhanced. To conclude, stronger cell-cell interactions reinforce the division of labor, whereby differentiated cells can grow more efficiently.

The rate of growth enhancement through cell differentiation R_μ can be roughly estimated by recalling that the growth rate of a cell is given by the average influx of the nutrient chemical it receives. We compared the growth rate of an isolated cell $\mu^{(iso)}$ to that of a differentiated cell $\mu^{(dif)}$ by assuming that the concentration of each chemical species is equally distributed, except for the nutrient chemical. Considering that an isolated cell has a catalytic network with k chemical components and q reaction paths from the nutrient X_0 , each concentration of X_1 - X_{k-1} is calculated as $x^{(iso)} = (1 - x_0^{(iso)})/(k - 1)$. Hence, for the steady state, the growth rate is estimated by $\mu^{(iso)} = qx_0^{(iso)}x^{(iso)\alpha}/(1 + x_0^{(iso)})$. On the other hand, the sub-network in a differentiated cell is considered to have k' chemicals and q' reaction paths from the nutrient ($k' < k$, $q' < q$). Then, each chemical concentration and the growth rate are given by $x^{(dif)} = (1 - x_0^{(dif)})/(k' - 1)$ and $\mu^{(dif)} = q'x_0^{(dif)}x^{(dif)\alpha}/(1 + x_0^{(dif)})$, respectively.

Here, we also assume that $x_0^{(iso)} \simeq x_0^{(dif)}$, because these concentrations mostly depend on the supplied nutrient concentration C rather than on the internal dynamics of individual cells. From these assumptions, the growth ratio $R_\mu \equiv \mu^{(dif)}/\mu^{(iso)}$ is calculated as $R_\mu = (q'/q)[(k - 1)/(k' - 1)]^\alpha$. For example, with network 1 or 2 (Fig 3A and 3B), $k = 6$, $q = 4$, $k' = 4$, $q' = 2$, and thus $R_\mu = (1/2)(5/3)^\alpha$, which is greater than unity, at least when $\alpha \geq 2$. Although this estimate is not strictly accurate, it nevertheless demonstrates how cell differentiation can enhance cellular growth, which is facilitated by greater α . Even when the chemical concentrations were non-uniform, division of labor could accelerate growth when α was sufficiently large.

Robustness in the number distribution of differentiated cells

The cells in our models achieved (i) cell differentiation and (ii) cooperative growth. However, if one cell type grows faster than the other type, the cooperation between the differentiated cells collapses. Thus, the third condition is necessary: the growth rate of each cell type needs to be in conformity, through mutual regulation by cell-cell interactions.

Thus far, we have considered the case with equal populations of the two cell types by imposing the condition $v^{(1)} = v^{(2)}$. Here, we examine the case with $v^{(1)} \neq v^{(2)}$ for fixed $v^{(1)}$ and $v^{(2)}$, to evaluate whether the increases in cell volume (or number) are identical between the two types to meet the requirement (iii). Therefore, Fig 6A and 6B show plots of the growth rate versus $r^{(1)}$, where $r^{(i)} \equiv v^{(i)}/(v^{(1)} + v^{(2)})$ is the volume proportion between type-1 and type-2 cells. Now, let us denote the dependence of $\mu^{(1)}$ on $r^{(1)}$ by a function $F(r^{(1)})$. Then, the growth rate of the type-2 cell $\mu^{(2)}$ is given by $G(r^{(2)}) = G(1 - r^{(1)})$, which is equal to $F(1 - r^{(1)})$ due to symmetry in the catalytic network.

Since differentiated cells help each other, balanced growth would be expected; if the volume or relative number of one cell type is larger than the other, a larger (smaller) amount of chemicals would be supplied from the majority (minority) type to the minority (majority) type, so that the growth rate of the minority type is enhanced compared to that of the other type. This is the case for network 2, where $F(r^{(1)}) < F(1 - r^{(1)})$ for $r^{(1)} > 1/2$, and the difference in volume (or number) decreases over time, leading to a balanced cell distribution (Fig 6B).

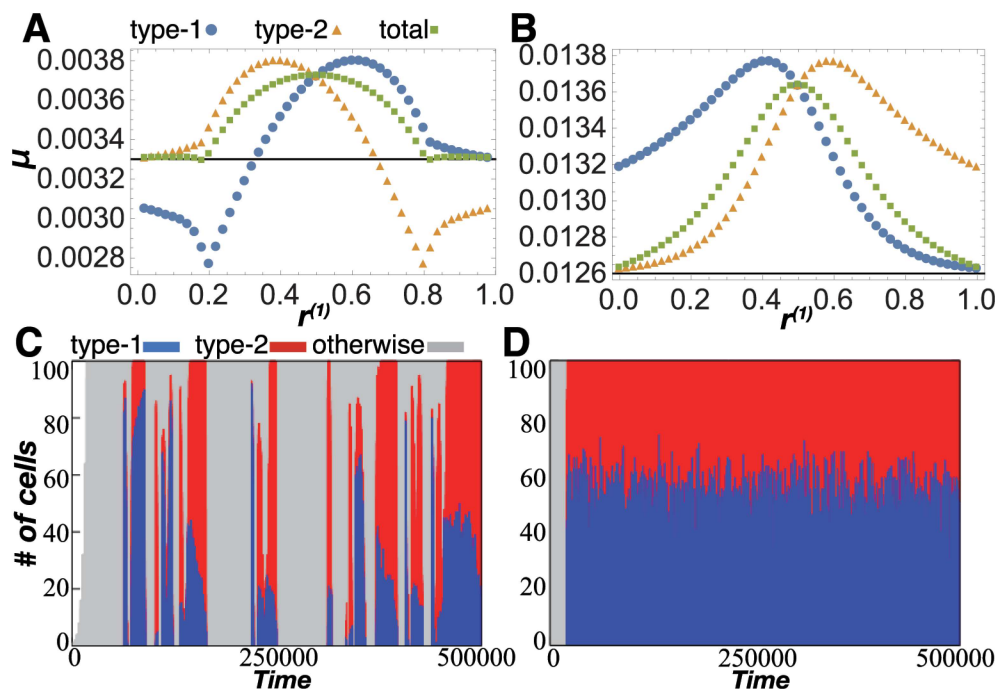


Fig 6. The stability of collective growth. (A) Network 1 in the r2cell model for $(C, V, D) = (0.1, 1, 0.1)$, representing unstable collective growth. (B) Network 2 in the r2cell model for $(C, V, D) = (0.15, 3, 1)$, representing balanced collective growth. In (A) and (B), the horizontal axis denotes the volume ratio of the two cells, $r^{(1)} \equiv v^{(1)}/(v^{(1)} + v^{(2)})$, and each black line displays the growth rate in the r1cell model $\mu^{(iso)}$. Blue circles and orange triangles denote the growth rate of type-1 $\mu^{(1)}$ and type-2 $\mu^{(2)}$, respectively. Green rectangles indicate the total growth rate of the cell aggregate $\mu^{(tot)} \equiv r^{(1)}\mu^{(1)} + r^{(2)}\mu^{(2)}$. (C) and (D) show the cell number distribution in the N -cell model ($N = 100$). Blue, red, and gray bars designate the numbers of type-1, type-2, and non-differentiated cells, respectively. (C) Case for network 1 with $(C, V_{med}, D) = (0.01, 100, 0.1)$. (D) Case for network 2 with $(C, V_{med}, D) = (0.3, 50, 0.1)$.

doi:10.1371/journal.pcbi.1005042.g006

Nonetheless, this is not always the case. Fig 6A shows that the growth rate of the majority type is larger than that of the minority type in network 1; that is, $F(r^{(1)}) > F(1-r^{(1)})$ for $r^{(1)} > 1/2$. Accordingly, the difference in volume increases over time, and thus the two different cell types cannot stably coexist. This instability of collective growth differs from a scenario of parasitic behavior, because $\mu^{(tot)} \equiv r^{(1)}\mu^{(1)} + r^{(2)}\mu^{(2)}$ is higher than $\mu^{(iso)}$ for almost the entire range of $r^{(1)}$.

The condition for the stability is analytically expressed as follows. First, the temporal change of $r^{(1)}$ is represented by $dr^{(1)}/dt = r^{(1)}(1-r^{(1)})[F(r^{(1)}) - F(1-r^{(1)})]$, which has a trivial fixed point solution $r^{(1)} = 1/2$. This fixed point, where the two cell types coexist, is unstable if $F'(1/2) > 0$, and is stable if $F'(1/2) < 0$. To estimate $F'(1/2)$, recall that $dx_i^{(m)}/dt = 0$ is fulfilled in a cell with steady growth, and $D_{med} V \gg D$. Then, from the definition of the growth rate μ , we get

$$\frac{F'(1/2)}{D} \simeq -\frac{\partial x_0^{(1)}}{\partial r^{(1)}} - \frac{1}{2} \sum_{i=1}^{k-1} \sigma_i \frac{\partial (x_i^{(1)} - x_i^{(2)})}{\partial r^{(1)}} \bigg|_{r^{(1)}=\frac{1}{2}}. \quad (3)$$

Since the first term is always negative as described in S1 Text, Eq (3) shows that if the difference in exchanged chemicals between the majority and minority cells $[\sum_{i=1}^{k-1} \sigma_i (x_i^{(1)} - x_i^{(2)})]$ increases in proportion to the increase in volume ratio of the majority type, then $F'(1/2)$ is negative and thus the collective growth is balanced.

Now, let us consider how the difference in volume alters the states of two interacting cells in networks 1 and 2. When $r^{(1)} > 1/2$, the population ratio of type-1 cells is increased, and the amount of X_4 supplied from the minority type-2 cells is not sufficient for the majority type-1 cells to maintain their differentiated chemical composition. In contrast, the minority type-2 cells receive sufficient amounts of X_3 from the majority type-1 cells, and maintain their differentiated composition. Consequently, the chemical composition of the type-1 cells approaches that of the isolated case, which contains X_1 and X_2 equally. This indicates that the majority of type-1 cell produces more X_5 than the minority of type-2 cell does, and thus $\partial(x_5^{(1)} - x_5^{(2)})/\partial r^{(1)} > 0$ holds around $r^{(1)} = 1/2$. Therefore, the diffusibility of X_5 contributes to the stability of network 2, and the non-diffusibility of X_5 contributes to the instability of network 1. Intuitively, this mechanism can also be explained as follows: with network 1, the majority cell type can produce a greater fraction of a non-diffusible chemical X_5 for itself and a smaller fraction of a diffusible chemical X_3 or X_4 for the other cell type, and thus the majority cell type grows faster than the minority cell type.

The stability and instability of collective growth are also observed in the original N -cell model. With an “unstable” network 1, the N -cell model repeats the following dynamic behavior, as shown in Fig 6C: the medium is dominated by cells of one type, and cells of the minority type become extinct. Then, their differentiated compositions cannot be maintained with a single cell type, leading to de-differentiation. Thus, the coexistence of differentiated cells is temporally regained.

In contrast, in a “stable” network 2, the two differentiated cell types stably coexist and their growth is balanced (Fig 6D), in which a perturbation to increase the population of one cell type leads to a decrease in the growth rate of that type.

Discussion

In this paper, we have shown that an aggregate of identical cells achieves metabolic division of labor, with strong cell-cell interactions under limited resources. We have revealed how (i) cell differentiation, (ii) growth enhancement, and (iii) robustness in a cell population can be simultaneously self-organized without assuming the ability of differentiation *a priori*, in a simple system consisting only of intracellular reaction dynamics and cell-cell interactions through chemical diffusion.

First, cells sharing a common genotype (i.e., an identical reaction network and identical parameters for reaction and diffusion) differentiate into several types with different chemical compositions as a result of the instability of a homogeneous cell state induced by cell-cell interactions. This differentiation is facilitated under a condition of limited resources and strong cell-cell interactions, given a high order of catalytic reactions. This dynamical-systems mechanism has also been proposed in previous studies using models of metabolic networks [27] and gene regulation networks [32].

Second, the differentiated cells can achieve cooperative division of labor. The explanation of the division of labor given in the Results section can be simply sketched as follows. Let us consider two reactions in the r2cell model, $X_0 + \alpha X_1 \rightarrow X_3 + \alpha X_1$ and $X_0 + \alpha X_2 \rightarrow X_4 + \alpha X_2$, with $x_1 + x_2 = c$. If $x_1 = x_2 = c/2$, the total production rate of X_3 and X_4 is $2x_0(c/2)^\alpha$. In contrast, if the concentrations are biased either to X_1 or X_2 due to differentiation, $x_1 \sim c$ (or $x_2 \sim c$), the total production rate of X_3 and X_4 per cell is $x_0 c^\alpha$, which is $2^{\alpha-1}$ times greater than that of the previous isolated, generalist cell. Thus the higher the order of the reaction α , the greater the advantage of division of labor. A more precise argument is given in the Results (ii). This growth enhancement due to division of labor is clearly distinguishable from a scenario of relaxation of the competition for resources through niche differentiation reported previously [27], in which the growth rate is not increased relative to that of an isolated cell. In our model, the growth rate

can be enhanced by concentrating chemicals on one of the modules in the network, while the other module is necessary for catalyzing the reaction. A related mechanism for division of labor was proposed by Michod and colleagues (see also [21, 24, 33, 34] for discussion on the trade-off for division of labor). In the framework of Michod et al., the convexity of the trade-off function is important for division of labor, and the condition of $(q'/q)[(k-1)/(k'-1)]^\alpha > 1$ in our model may be related to the convexity of the trade-off function.

Although the proposed model describes the division of chemical production among a cell aggregate, the above mechanism can be seen as analogous to the theory for division of labor in economics: the theory of comparative advantage proposed by Ricardo [35] states that the mutual use of surplus from a different country is more advantageous than producing all necessary resources in a single country, unless the transport cost is too high. In this sense, Ricardo's theory parallels the present mechanism, because two cell types help each other by exchanging the products that are necessary to the other cell type. Indeed, our mechanism works best when cell density is high, so that chemicals are easily exchanged without much loss within the medium. Note, however, that in Ricardo's theory, trade is assumed to occur between countries that differ in their relative ability of producing multiple goods, in contrast to the current model in which cells share an identical chemical network. With regard to this point, a better comparison would be with Taylorism [36], which refers to increases in a group's productivity by each member specializing to each task without assuming individuals with different abilities [26].

Remarkably, this cooperative differentiation is not sufficient to satisfy condition (iii), robustness in the number distribution of differentiated cells. If one cell type begins to dominate the population, production of the chemicals needed by the minority type will increase. Thus, a feedback mechanism to reduce the majority population is expected. However, if the fraction of non-diffusible chemicals is increased for the majority cell type, this storage of chemicals within a cell would suppress the supply of chemicals for the other cell types. Consequently, the majority cell type would further increase its population. This suggests that to achieve a balanced population state, the mutual transport of necessary chemicals must work efficiently beyond any possible increase in internal reserves. From this perspective, it may be interesting to consider a possible economic analogy: reducing internal reserves and sharing a higher degree of wealth will be relevant to the stabilization of groups with division of labor. We here stress that this instability of cooperation can emerge only when cells simultaneously achieve differentiation and division of labor.

In theoretical studies for multi-level evolution, a game theory approach has been sometimes adopted to address the evolution and dynamics of conflict between individuals and society. Although our approach differs from game theory, it might be worth discussing our result in light of this perspective. From a game theory perspective, the cellular growth rate is regarded as a measure of fitness or score. Hence, when two cell types stably coexist in network 2, stable Nash equilibrium is achieved at $r^{(1)} = 1/2$. In contrast, in network 1, no stable equilibrium exists for $0 < r^{(1)} < 1$, and only unstable Nash equilibrium exists at $r^{(1)} = 1/2$, and thus one type dominates the population. Interestingly, after extinction of one type, re-differentiation of the remaining cells into two types increases the fitness (i.e., growth rate) for both types, as shown in Fig 6A. This dominance of one cell type and re-differentiation are repeated as a result of symbiotic growth and differentiation due to the instability of a homogeneous cell society. We expect that such dynamic behavior will be observed in an artificial symbiosis experiment with *Escherichia coli* and diffusible amino acids [37, 38].

Considering the difference between networks 1 and 2, the stability and instability of the system can be switched by even a slight change in the diffusibility of a single chemical species. This implies that slight epigenetic changes and transcriptional errors occurring during the lifetime of an organism can lead to instability in the cell distribution, which may relate to the phenomena of metamorphosis during development and carcinogenesis.

Our results demonstrate that an aggregate of simple cells consisting only of catalytic reactions and the diffusive transport of chemicals can fulfill differentiation with division of labor, collective growth with symbiotic relationship, and stability. Note that these basic characteristics in our model emerge without a fine-tuned mechanism, and are facilitated by the conditions of strong cell-cell interactions, limited resources, and a high order of catalytic reactions.

From this point of view, it is interesting to compare the present results to some characteristics of multicellular aggregates. First, filaments of the cyanobacterium *Anabaena* are known to differentiate, with each cell metabolically specializing in photosynthesis or nitrogen fixation, enabling more efficient growth [39]. Second, such cell differentiation with metabolic division of labor in some cyanobacteria occurs in response to combined nitrogen limitation [40, 41]. Third, the biofilm of *Bacillus subtilis* exhibits metabolic co-dependence between interior and peripheral cells by chemical oscillation [9], suggesting the relevance of nonlinear dynamics and cell-cell interactions for differentiation.

In contrast to the present model of symbiotic growth, however, it has been pointed out that most multicellular aggregates and organisms have achieved division of labor between reproductive and non-reproductive cells throughout evolution [42]. Nevertheless, at some developmental stage of multicellular aggregates, symbiotic growth of different cell types may be expected to exist, by achieving differentiation and functional division of labor for biofilm formation [43, 44]. Furthermore, in our model, whether or not both types of differentiated cells reproduce strongly depends on the conditions. For example, in network 1, the growth rate is different between the major and minor cell types. Depending on the parameters, there are also cases in which one cell type would cease growing. In addition, we considered the symmetric differentiation case for clarity, but if the reaction rates are different by different chemicals (which are natural), the growth rates of differentiated cell types could generally be different. Further, if we assume that cellular growth is determined by a certain chemical (e.g., X_1 in networks 1-3) rather than by the total amount of chemicals, after differentiation, one cell type will be reproductive, and the other non-reproductive, while maintaining functional division of labor.

Interestingly, characteristics (i)-(iii) can be part of the requirements for multicellularity. Thus, such characteristics may provide a primitive step to the evolution of multicellular organisms, which has been a topic of much attention from both theorists and experimentalists over the last few decades [15, 27, 45–50]. In this context, our results are also related to the experimental emergence of multicellularity from unicellular organisms [13, 14]. However, the three characteristics may not be sufficient for the emergence of multicellular organisms. For example, besides the metabolic division of labor, multicellular organisms ubiquitously show germ-soma differentiation and apoptosis. Therefore, determining how the cell aggregates with metabolic division of labor considered here might achieve this universal property of multicellularity remains an important issue to be addressed.

Supporting Information

S1 Text. Supplementary information about the description and analysis of the *N*-cell, *r1*cell, and *r2*cell models. Figures A and B Typical behavior of categories (a) and (b); Figure C Network 3; Figure D Dependence of the concentration of the nutrient X_0 on $r^{(1)}$; Figure E Phase diagrams of the *r1*cell and *r2*cell models. (PDF)

Acknowledgments

We would like to acknowledge helpful discussions with Chikara Furusawa.

Author Contributions

Conceived and designed the experiments: JFY NS KK.

Performed the experiments: JFY.

Analyzed the data: JFY NS KK.

Wrote the paper: JFY NS KK.

References

- Shapiro JA. Thinking about bacterial populations as multicellular organisms. *Annu Rev Microbiol.* 1998; 52: 81–104. doi: [10.1146/annurev.micro.52.1.81](https://doi.org/10.1146/annurev.micro.52.1.81) PMID: [9891794](https://pubmed.ncbi.nlm.nih.gov/9891794/)
- Jefferson KK. What drives bacteria to produce a biofilm? *FEMS Microbiol Lett.* 2004; 236: 163–173. doi: [10.1111/j.1574-6968.2004.tb09643.x](https://doi.org/10.1111/j.1574-6968.2004.tb09643.x) PMID: [15251193](https://pubmed.ncbi.nlm.nih.gov/15251193/)
- Claessen D, Rozen DE, Kuipers OP, Søgaard-Andersen L, van Wezel GP. Bacterial solutions to multicellularity: a tale of biofilms, filaments and fruiting bodies. *Nat Rev Microbiol.* 2008; 6: 199–210. doi: [10.1038/nrmicro3178](https://doi.org/10.1038/nrmicro3178) PMID: [24384602](https://pubmed.ncbi.nlm.nih.gov/24384602/)
- Bonner JT. *The cellular slime molds.* 2nd ed. Princeton, NJ: Princeton University Press; 1967. doi: [10.1515/9781400876884](https://doi.org/10.1515/9781400876884)
- Bonner JT. Evolution of development in the cellular slime molds. *Evol Dev.* 2003; 5: 305–313. doi: [10.1046/j.1525-142X.2003.03037.x](https://doi.org/10.1046/j.1525-142X.2003.03037.x) PMID: [12752769](https://pubmed.ncbi.nlm.nih.gov/12752769/)
- Parsek MR, Greenberg EP. Sociomicrobiology: the connections between quorum sensing and biofilms. *Trends Microbiol.* 2005; 13: 27–33. doi: [10.1016/j.tim.2004.11.007](https://doi.org/10.1016/j.tim.2004.11.007) PMID: [15639629](https://pubmed.ncbi.nlm.nih.gov/15639629/)
- Cole SP, Harwood J, Lee R, She R, Guiney DG. Characterization of monospecies biofilm formation by *Helicobacter pylori*. *J Bacteriol.* 2004; 186: 3124–3132. doi: [10.1128/JB.186.10.3124-3132.2004](https://doi.org/10.1128/JB.186.10.3124-3132.2004) PMID: [15126474](https://pubmed.ncbi.nlm.nih.gov/15126474/)
- George S, Kishen A, Song P. The role of environmental changes on monospecies biofilm formation on root canal wall by *Enterococcus faecalis*. *J Endod.* 2005; 31: 867–872. doi: [10.1097/01.don.0000164855.98346.fc](https://doi.org/10.1097/01.don.0000164855.98346.fc) PMID: [16306820](https://pubmed.ncbi.nlm.nih.gov/16306820/)
- Liu J, Prindle A, Humphries J, Gabalda-Sagarra M, Asally M, Lee DD, et al. Metabolic co-dependence gives rise to collective oscillations within biofilms. *Nature.* 2015; 523: 550–554. doi: [10.1038/nature14660](https://doi.org/10.1038/nature14660) PMID: [26200335](https://pubmed.ncbi.nlm.nih.gov/26200335/)
- James GA, Beaudette L, Costerton JW. Interspecies bacterial interactions in biofilms. *J Ind Microbiol.* 1995; 15: 257–262. doi: [10.1007/BF01569978](https://doi.org/10.1007/BF01569978)
- Rickard AH, Gilbert P, High NJ, Kolenbrander PE, Handley PS. Bacterial coaggregation: an integral process in the development of multi-species biofilms. *Trends Microbiol.* 2003; 11: 94–100. doi: [10.1016/S0966-842X\(02\)00034-3](https://doi.org/10.1016/S0966-842X(02)00034-3) PMID: [12598132](https://pubmed.ncbi.nlm.nih.gov/12598132/)
- Stewart PS, Franklin MJ. Physiological heterogeneity in biofilms. *Nat Rev Microbiol.* 2014; 12: 115–124. doi: [10.1038/nrmicro1838](https://doi.org/10.1038/nrmicro1838)
- Ratcliff WC, Denison RF, Borrello M, Travisano M. Experimental evolution of multicellularity. *Proc Natl Acad Sci USA.* 2012; 109(5): 1595–1600. doi: [10.1073/pnas.1115323109](https://doi.org/10.1073/pnas.1115323109) PMID: [22307617](https://pubmed.ncbi.nlm.nih.gov/22307617/)
- Ratcliff WC, Herron MD, Howell K, Pentz JT, Rosenzweig F, Travisano M. Experimental evolution of an alternating uni- and multicellular life cycle in *Chlamydomonas reinhardtii*. *Nat Commun.* 2013; 4: 2742. doi: [10.1038/ncomms3742](https://doi.org/10.1038/ncomms3742) PMID: [24193369](https://pubmed.ncbi.nlm.nih.gov/24193369/)
- Michod RE, Roze D. Cooperation and conflict in the evolution of multicellularity. *Heredity.* 2001; 86: 1–7. doi: [10.1046/j.1365-2540.2001.00808.x](https://doi.org/10.1046/j.1365-2540.2001.00808.x) PMID: [11298810](https://pubmed.ncbi.nlm.nih.gov/11298810/)
- Roze D, Michod RE. Mutation, multilevel selection, and the evolution of propagule size during the origin of multicellularity. *Am Nat.* 2001; 158: 638–654. doi: [10.1086/323590](https://doi.org/10.1086/323590) PMID: [18707357](https://pubmed.ncbi.nlm.nih.gov/18707357/)
- Nowak MA. Five rules for the evolution of cooperation. *Science.* 2006; 314: 1560–1563. doi: [10.1126/science.1133755](https://doi.org/10.1126/science.1133755) PMID: [17158317](https://pubmed.ncbi.nlm.nih.gov/17158317/)
- Willensdorfer M. On the evolution of differentiated multicellularity. *Evolution* 2009; 63(2): 306–323. doi: [10.1111/j.1558-5646.2008.00541.x](https://doi.org/10.1111/j.1558-5646.2008.00541.x) PMID: [19154376](https://pubmed.ncbi.nlm.nih.gov/19154376/)
- Gavrilets S. Rapid transition towards the division of labor via evolution of developmental plasticity. *PLoS Comput Biol.* 2010; 6(6): e1000805. doi: [10.1371/journal.pcbi.1000805](https://doi.org/10.1371/journal.pcbi.1000805) PMID: [20548941](https://pubmed.ncbi.nlm.nih.gov/20548941/)
- Michod RE, Viossat Y, Solari CA, Hurand M, Nedelcu AM. Life-history evolution and the origin of multicellularity. *J Theor Biol.* 2006; 239: 257–272. doi: [10.1016/j.jtbi.2005.08.043](https://doi.org/10.1016/j.jtbi.2005.08.043) PMID: [16288782](https://pubmed.ncbi.nlm.nih.gov/16288782/)

21. Vászárhelyi Z, Meszéna G, Scheuring I. Evolution of heritable behavioural differences in a model of social division of labour. *PeerJ*. 2015; 3:e977. doi: [10.7717/peerj.977](https://doi.org/10.7717/peerj.977) PMID: [26038732](https://pubmed.ncbi.nlm.nih.gov/26038732/)
22. Traulsen A, Nowak MA. Evolution of cooperation by multilevel selection. *Proc Natl Acad Sci USA*. 2006; 103(29): 10952–10955. doi: [10.1073/pnas.0602530103](https://doi.org/10.1073/pnas.0602530103) PMID: [16829575](https://pubmed.ncbi.nlm.nih.gov/16829575/)
23. Vural DC, Isakov A, Mahadevan L. The organization and control of an evolving interdependent population. *J R Soc Interface*. 2015; 12: 20150044. doi: [10.1098/rsif.2015.0044](https://doi.org/10.1098/rsif.2015.0044) PMID: [26040593](https://pubmed.ncbi.nlm.nih.gov/26040593/)
24. Ispolatov I, Ackermann M, Doebeli M. Division of labour and the evolution of multicellularity. *Proc R Soc B*. 2011; 279: 1768–1776. doi: [10.1098/rspb.2011.1999](https://doi.org/10.1098/rspb.2011.1999) PMID: [22158952](https://pubmed.ncbi.nlm.nih.gov/22158952/)
25. Ferrante E, Turgut AE, Duéñez-Guzmán E, Dorigo M, Wenseleers T. Evolution of self-organized task specialization in robot swarms. *PLoS Comput Biol*. 2–15; 11(8): e1004273. doi: [10.1371/journal.pcbi.1004273](https://doi.org/10.1371/journal.pcbi.1004273) PMID: [26247819](https://pubmed.ncbi.nlm.nih.gov/26247819/)
26. Kaneko K, Yomo T. Isologous diversification for robust development of cell society. *J Theor Biol*. 1999; 199: 243–256. doi: [10.1006/jtbi.1999.0952](https://doi.org/10.1006/jtbi.1999.0952) PMID: [10433890](https://pubmed.ncbi.nlm.nih.gov/10433890/)
27. Furusawa C, Kaneko K. Origin of complexity in multicellular organisms. *Phys Rev Lett*. 2000; 84(26): 6130–6133. doi: [10.1103/PhysRevLett.84.6130](https://doi.org/10.1103/PhysRevLett.84.6130) PMID: [10991141](https://pubmed.ncbi.nlm.nih.gov/10991141/)
28. Furusawa C, Kaneko K. Emergence of rules in cell society: differentiation, hierarchy, and stability. *Bull Math Biol*. 1998; 60:659–687. doi: [10.1006/bulm.1997.0034](https://doi.org/10.1006/bulm.1997.0034) PMID: [9659010](https://pubmed.ncbi.nlm.nih.gov/9659010/)
29. Furusawa C, Kaneko K. Theory of robustness of irreversible differentiation in a stem cell system: chaos hypothesis. *J Theor Biol*. 2001; 209: 395–416. doi: [10.1006/jtbi.2001.2264](https://doi.org/10.1006/jtbi.2001.2264) PMID: [11319890](https://pubmed.ncbi.nlm.nih.gov/11319890/)
30. Aronson DG, Ermentrout GB, Kopell N. Amplitude response of coupled oscillators. *Physica D*. 1990; 41(3): 403–449. doi: [10.1016/0167-2789\(90\)90007-C](https://doi.org/10.1016/0167-2789(90)90007-C)
31. Koseska A, Volkov E, Kurths J. Parameter mismatches and oscillation death in coupled oscillators. *Chaos*. 2010; 20: 023132. doi: [10.1063/1.3456937](https://doi.org/10.1063/1.3456937) PMID: [20590328](https://pubmed.ncbi.nlm.nih.gov/20590328/)
32. Goto Y, Kaneko K. Minimal model for stem-cell differentiation. *Phys Rev E*. 2013; 88: 032718. doi: [10.1103/PhysRevE.88.032718](https://doi.org/10.1103/PhysRevE.88.032718) PMID: [24125305](https://pubmed.ncbi.nlm.nih.gov/24125305/)
33. Rueffler C, Hermisson J, and Wagner GP. Evolution of functional specialization and division of labor. *Proc Natl Acad Sci USA*. 2012; 109(6): E326–335. doi: [10.1073/pnas.1110521109](https://doi.org/10.1073/pnas.1110521109) PMID: [22308336](https://pubmed.ncbi.nlm.nih.gov/22308336/)
34. Goldsby HJ, Knoester DB, Ofria C, Kerr B. The evolutionary origin of somatic cells under the dirty work hypothesis. *PLoS Biol*. 2014; 12(5): e1001858. doi: [10.1371/journal.pbio.1001858](https://doi.org/10.1371/journal.pbio.1001858) PMID: [24823361](https://pubmed.ncbi.nlm.nih.gov/24823361/)
35. Ricardo D. Principles of political economy and taxation. London, UK: John Murray; 1817. doi: [10.1017/CBO9781107589421](https://doi.org/10.1017/CBO9781107589421)
36. Taylor FW. The Principles of scientific management, New York, USA: Harper & Brothers; 1911.
37. Hosoda K, Yomo T. Designing symbiosis. *Bioeng Bugs*. 2011; 2(6): 338–341. doi: [10.4161/bbug.2.6.16801](https://doi.org/10.4161/bbug.2.6.16801) PMID: [22008942](https://pubmed.ncbi.nlm.nih.gov/22008942/)
38. Hosoda K, Suzuki S, Yamauchi Y, Shiroguchi Y, Kashiwagi A, Ono N, et al. Cooperative adaptation to establishment of a synthetic bacterial mutualism. *PLoS ONE*. 2011; 6(2): e17105. doi: [10.1371/journal.pone.0017105](https://doi.org/10.1371/journal.pone.0017105) PMID: [21359225](https://pubmed.ncbi.nlm.nih.gov/21359225/)
39. Flores E, Herrero A. Compartmentalized function through cell differentiation in filamentous cyanobacteria. *Nat Rev Microbiol*. 2010; 8: 39–50. doi: [10.1038/nrmicro2242](https://doi.org/10.1038/nrmicro2242) PMID: [19966815](https://pubmed.ncbi.nlm.nih.gov/19966815/)
40. Adams DG, Duggan PS. Heterocyst and akinete differentiation in cyanobacteria. *New Phytol*. 1999; 144: 3–33. doi: [10.1046/j.1469-8137.1999.00505.x](https://doi.org/10.1046/j.1469-8137.1999.00505.x)
41. Laurent S, Chen H, Bédu S, Ziarelli F, Peng L, Zhang C-C. Nonmetabolizable analogue of 2-oxoglutarate elicits heterocyst differentiation under repressive conditions in *Anabaena* sp. strain PCC 7120. *Proc Natl Acad Sci USA*. 2005; 102: 9907–9912. doi: [10.1073/pnas.0502337102](https://doi.org/10.1073/pnas.0502337102) PMID: [15985552](https://pubmed.ncbi.nlm.nih.gov/15985552/)
42. Simpson C. The evolutionary history of division of labour. *Proc R Soc B*. 2012; 279: 116–121. doi: [10.1098/rspb.2011.0766](https://doi.org/10.1098/rspb.2011.0766) PMID: [21561969](https://pubmed.ncbi.nlm.nih.gov/21561969/)
43. Vlamakis H, Aguilar C, Losick R, Kolter R. Control of cell fate by the formation of an architecturally complex bacterial community. *Genes Dev*. 2008; 22: 945–953. doi: [10.1101/gad.1645008](https://doi.org/10.1101/gad.1645008) PMID: [18381896](https://pubmed.ncbi.nlm.nih.gov/18381896/)
44. van Gestel J, Vlamakis H, Kolter R. From cell differentiation to cell collectives: *Bacillus subtilis* uses division of labor to migrate. *PLOS Biol*. 2015; 13(4): e1002141. doi: [10.1371/journal.pbio.1002141](https://doi.org/10.1371/journal.pbio.1002141) PMID: [25894589](https://pubmed.ncbi.nlm.nih.gov/25894589/)
45. Szathmáry E, Maynard-Smith J. *The Major Transitions in Evolution*. Oxford: Oxford University Press; 1997.
46. Shapiro JA, Dworkin M, editors. *Bacteria as Multicellular Organisms*. Oxford: Oxford University Press; 1997. doi: [10.1038/scientificamerican0688-82](https://doi.org/10.1038/scientificamerican0688-82)

47. Niklas KJ, Newman SA. The origins of multicellular organisms. *Evol Dev.* 2013; 15: 41–52. doi: [10.1111/ede.12013](https://doi.org/10.1111/ede.12013) PMID: [23331916](https://pubmed.ncbi.nlm.nih.gov/23331916/)
48. Lyons NA, Kolter R. On the evolution of bacterial multicellularity. *Curr Opin Microbiol.* 2015; 24: 21–28. doi: [10.1016/j.mib.2014.12.007](https://doi.org/10.1016/j.mib.2014.12.007) PMID: [25597443](https://pubmed.ncbi.nlm.nih.gov/25597443/)
49. Solé RV, Duran-Nebreda S. 2015 In Silico Transitions to Multicellularity. In: Ruiz-Trillo I, Nedelcu AM, editors. *Evolutionary Transitions to Multicellular Life*. Heidelberg: Springer; 2015. pp. 245–266.
50. Niklas KJ, Newman SA, editors. *Multicellularity: Origins and Evolution*. Cambridge: MIT Press; 2016.