The emergence of geometric order in proliferating metazoan epithelia.

by Gibson *et al. Nature*, 2006 Computational Biology and Bioinformatics Seminar

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Introduction

Motivation

'The organisation of cells into epithelial sheets is an essential feature of animal design.'

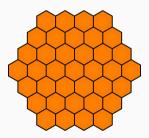
Gibson et al [1]

Context

Predictable geometries arise when *minimising surface energy* or *maximising space filling* in biological and non-biological structures:

Examples

- Drosophila Melanogaster's retinal cells
- Honeycombs
- Coins on tabletops

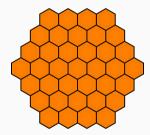


Context

Predictable geometries arise when *minimising surface energy* or *maximising space filling* in biological and non-biological structures:

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How is the geometry of growing epithelial cells organized and how can it be explained?

Emergent behaviour

 $Understanding\ emergent\ behaviours\ is\ required\ to\ model\ geometries\ of\ growing\ epithelial\ cells.$

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Emergent behaviour

It is the behaviour of a system that can *only* be explained by examination of a system's **parts** and their **relationships**.

Model Assumptions

Post-mitotic relationship between two daughter cells

Stochastically marked cell lineages with GFP revealed stable post-mitotic patterns.

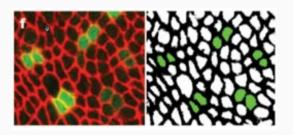


Figure 1: Daugher-cell relationship.

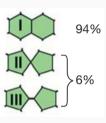


Figure 2: Observed post-mitotic relationship between daughter cells.

Post-mitotic relationship between two daughter cells

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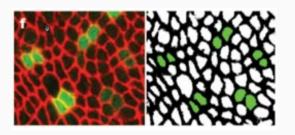


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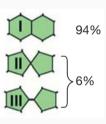
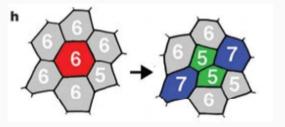


Figure 2: Observed post-mitotic relationship between daughter cells.

Typical topology changes of a dividing cell

Figure 3: Topology changes during cell division



A Discrete Markov Model for

Proliferating Epithelia

Markov models

A Markov chain is a discrete stochastic process with the Markov property:

$$P(X_t|X_{t-1},...,X_1) = P(X_t|X_{t-1})$$

where X_t is the time-dependent r.v. describing the state of the system at time t.

It is determined by a probability transition matrix P and an initial probability distribution.

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S

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$\mathbf{p}^{(t+1)} = \mathbf{p}^{(t)} PS$	the state dynamics, where P and S are the probabilistic transition matrices.
P_{ij}	the probability that an i sided cell divides to produce a j -sided daughter cell.
S_{ij}	the probability of an i-sided cell will gain sides from dividing neighbour cells divisions
-	to become <i>j</i> -sided.

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Transition matrices

$$P = \begin{bmatrix} 4 & 5 & 6 & 7 & 8 & 9 & \dots \\ 4 & 1 & & & & & \\ 5 & 1 & 1 & & & & & \\ 6 & 1 & 2 & 1 & & & & \\ 1 & 3 & 3 & 1 & & & & \\ 8 & 1 & 4 & 6 & 4 & 1 & & \\ 9 & 1 & 5 & 10 & 10 & 5 & 1 & & \\ \vdots & & & & \ddots & \end{bmatrix}$$

 P_{ij} : the probability that an i sided cell divides to produce a j-sided daughter cell.

 S_{ij} the probability of an *i*-sided cell will gain sides from dividing neighbour cells divisions

Predicting the equilibrium E from T = PS

The Perron-Frobenius theorem guarantees that a Markov chain will converge to a unique equilibrium E which is the principal eigenvector of the transition matrix T = PS.

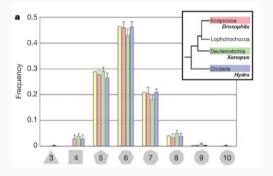


Figure 4: Distribution of the predicted and observed polygons (yellow: predicted equilibirum)

Robust equilibrium topology

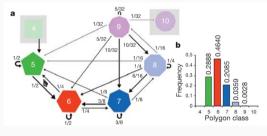


Figure 5: Discrete state dynamics

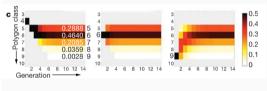


Figure 6: Figure showing the same attained equilibirum, no matter the initial conditions

Additional insights on mitotic cells

As seen in *S*, a mitotic cell gains on average one side, and daughter cells have one less side. Experimental evidence confirms this:

Non-mitotic cells	(5.94 ± 0.06)
Mitotic cells	(6.99 ± 0.07)

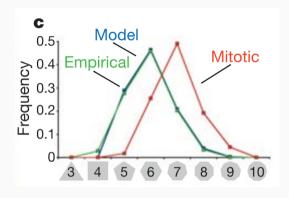


Figure 7: Polygon distributions of mitotic cells vs. non-mitotic cells

^{ightarrow} cells accumulate sides until division.

Polygon distribution of mitotic cells in non-proliferating tissues

- The model does not apply to non-proliferating tissue
- Forcing mitosis using string (stg) in tissues does not replicate the proliferating tissue distribution.

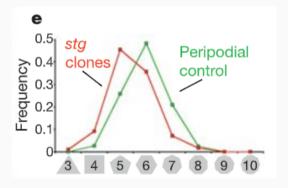


Figure 8: Distribution of polygons in non-proliferating tissue

ightarrow Differential proliferation influences cell shape and morphogenesis.

Subsequent works

Subsequent work

Work on 3D structures

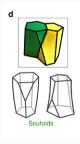


Figure 9: The scutoid has been discovered as a solution to 3D cell packing.[2]

Work on dynamics

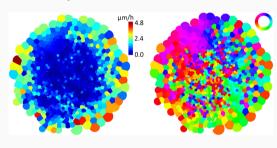


Figure 10: Modelling and prediction of speed and direction of proliferating epithelia. [3]

Conclusions

Take home messages

The proposed discrete Markov chain:

1. shows how epithelial topology can be irregular, but **not random**.

Take home messages

The proposed **discrete Markov chain**:

- 1. shows how epithelial topology can be irregular, but **not random**.
- 2. **explains the predominantly hexagonal topology** of growing epithelia as well as the distribution of other polygons.

Take home messages

The proposed **discrete Markov chain**:

- 1. shows how epithelial topology can be irregular, but **not random**.
- explains the predominantly hexagonal topology of growing epithelia as well as the distribution of other polygons.
- 3. shows an **emergent mechanism** by which epithelia accommodate **rapid proliferation** while maintaining **structural integrity**.

Questions?

https://github.com/pjhartout/CBB_Seminar

Geometry of growing epithelial cells

Time-lapse movies collected revealed the following findings:

- Initially polygonal prophase cells rounded up and divided
- Cell-neighbour relationships were stably maintained throughout the cell cycle
- The topology of the divided cells is invariant over time.

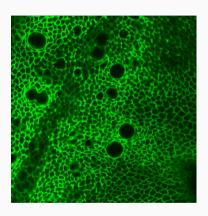


Figure 11: General hexagon-dominated topology of growing epithelial in *Drosophila Melanogaster*.

- P_{ij}: the probabilty that an i sided cell divides to produce a j-sided daughter cell.
- K_t: the number of junctions distributed to one daughter cell on division at generation
 t. s_{t-1} - K_t are left for the other.

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- Because no triangular cells are observed, each daughter receives at least two junctions, leaving s_t - 4 junctions to be distributed among the daughters with equal probability.
- So, the number of junctions received by the first daughter is $K_t 2 \sim \mathcal{B}(n = s_{t-1}, p = \frac{1}{2}).$

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As a result

$$P_{ij} = P(K_t + 2 = j | s_{t-1} = i) = {i-4 \choose i-4} \cdot \frac{1}{2^{i-4}}$$

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- As a result $P_{ij} = P(K_t + 2 = j | s_{t-1} = i) = \binom{i-4}{i-4} \cdot \frac{1}{2^{i-4}}$
- Therefore, the unnormalized entries of P are the coefficients of Pascal's triangle:

Derivation of Markov state dynamics - transition matrix ${\mathcal S}$

 S_{ij}: the probability of an *i*-sided cell will gain sides from dividing neighbour cells divisions

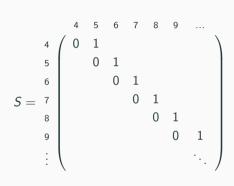
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- On average the number of sides gained per cell is $\frac{2N}{2N} = 1$. That is, a cell will gain 1 side every cycle on average.
- Thus, $S_{ij} = 1$ if j = i + 1 and 0 otherwise.



Backup slides - Assumptions of the recurrence system

From the aforementioned observations observations we make the following assumptions about the mathematical modelling of cell epithelia:

- Cells are polygons with a minimum of four sides
- Cells do not resort
- Mitotic siblings retain a common common junctional interface
- Cells have asynchronous but uniform cell cycle times
- Cleavage places cut a side rather than a vertex of the mother polygon
- Mitotic cleavage orientation randomly distributes existing tricellular junctions to both daughter cells.

Recurrence system

Table 1: Cell features, topological equivalence and evolution at division t

Cell feature	Graph equivalent	Evolution at division t
Tricellular junction	Vertex, v_t	$v_t = v_{t-1} + 2f_t$
Cell side	Edge, e_t	$e_t = e_{t-1} + 3f_{t-1}$
Apical cell surface	Face, f_t	$e_t = e_{t-1} + 3f_{t-1}$

So we can construct the following system:

$$s_t = \frac{2(e_t + 3f_{t-1})}{2f_{t-1}} = \frac{s_{t-1}}{2} + 3$$

which exponentially converges to 6 provided the initial condition:

$$s_t = 6 + 2^{-t}(s_0 - 6)$$

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