

The emergence of geometric order in proliferating metazoan epithelia.

by Gibson *et al* - CBB seminar presentation

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April 25, 2020

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Introduction

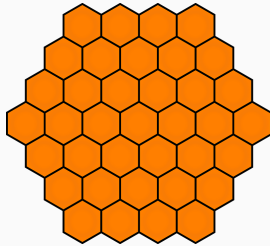
'The organisation of cells into epithelial sheets is an essential feature of animal design.'

Gibson *et al*

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

Examples

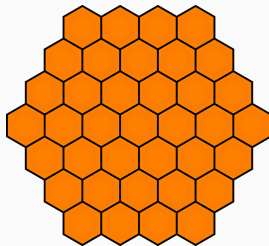
- *Drosophila Melanogaster*'s retinal cells
- Honeycombs
- Coins on tabletops



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

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.

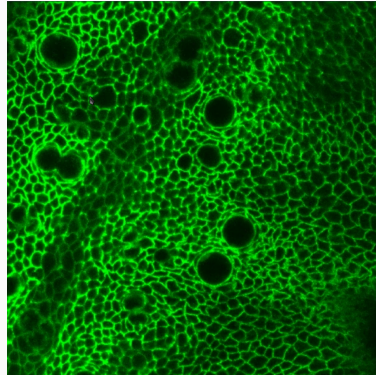


Figure 1: General hexagon-dominated topology of growing epithelial in *Drosophila Melanogaster*.

Post-mitotic relationship between two daughter cells

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

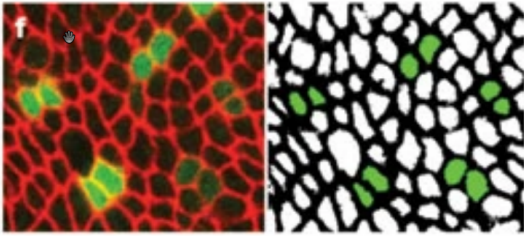


Figure 2: Daughter-cell relationship.

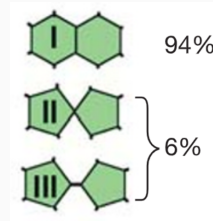


Figure 3: Observed post-mitotic relationship between daughter cells.

Recurrence system

Figure 4: Topology changes during cell division

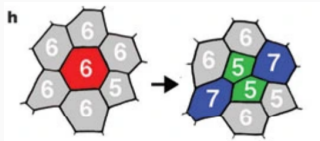


Table 1: Cell features, topological equivalence and evolution at division t

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:

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

A discrete Markov model to model stochastic cell proliferation

s	state of the cell as its number of sides, where $s > 3$.
p_s	the relative frequency of s -sided cells in the population
$\mathbf{p}^{(t)}$	the infinite row vector $\mathbf{p}^{(t)} = [p_4, p_5, p_6, \dots]$ state of the population at generation t .
$\mathbf{p}^{(t+1)} = \mathbf{p}^{(t)}PS$	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 j -sided.

Derivation of Markov state dynamics - transition matrix P

- P_{ij} : the probability 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.
- 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 $K_t - 2 \sim \mathcal{B}(n = s_{t-1}, p = \frac{1}{2})$.

- As a result

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

- Therefore, the unnormalized entries of P are the coefficients of Pascal's triangle:

$$P = \begin{matrix} & 4 & 5 & 6 & 7 & 8 & 9 & \dots \\ \begin{matrix} 4 \\ 5 \\ 6 \\ 7 \\ 8 \\ 9 \\ \vdots \end{matrix} & \left(\begin{array}{cccccc} 1 & & & & & \\ 1 & 1 & & & & \\ 1 & 2 & 1 & & & \\ 1 & 3 & 3 & 1 & & \\ 1 & 4 & 6 & 4 & 1 & \\ 1 & 5 & 10 & 10 & 5 & 1 \\ & & & & & \ddots \end{array} \right) \end{matrix}$$

Derivation of Markov state dynamics - transition matrix S

- S_{ij} : the probability of an i -sided cell will gain sides from dividing neighbour cells divisions
- On mitosis, a cell adds one side to each of two neighbouring cells.
- Assuming N cells in an epithelium, the number of cells after one round of divisions is $2N$
- 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.

$$S = \begin{matrix} & \begin{matrix} 4 & 5 & 6 & 7 & 8 & 9 & \dots \end{matrix} \\ \begin{matrix} 4 \\ 5 \\ 6 \\ 7 \\ 8 \\ 9 \\ \vdots \end{matrix} & \begin{pmatrix} 0 & 1 & & & & & \\ & 0 & 1 & & & & \\ & & 0 & 1 & & & \\ & & & 0 & 1 & & \\ & & & & 0 & 1 & \\ & & & & & 0 & 1 \\ & & & & & & 0 & 1 \\ & & & & & & & \ddots \end{pmatrix} \end{matrix}$$

Predicting the equilibrium E from $T = PS$

The Perron-Frobenius theorem guarantees that a Markov chain will converge to a unique equilibrium \mathbf{p}^* where \mathbf{p}^* is the principal eigenvector of the transition matrix T .

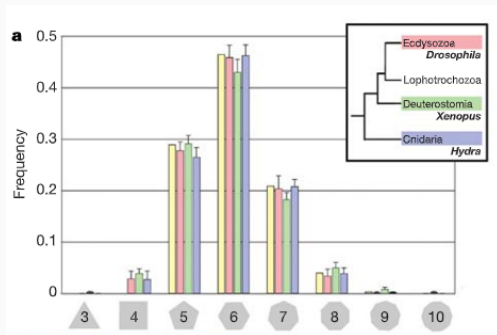


Figure 5: Distribution of the predicted and observed polygons

Questions?

Backup slides - Assumptions of the recurrence system

From the aforementioned 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.