# A Model for Cell Proliferation in a Developing Organism 

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ACEMS Research Group Meeting

19 September 2016

ACEM. $\int$

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Disclaimer. Work in progress!

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Starting point. Talks given by Kerry Landman at recent UQ Mathematics Colloquia (August 2015 and July 2016).

The model in question is described in ...

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Specific instance. Neural crest cells, present in the gut tissue of the developing human embryo, form neurons that construct the enteric nervous system (ENS). Failure of these cells to invade the gut tissue completely can cause imperfect formation of the ENS, and lead to Hirschsprung's disease, which can result in potentially life-threating complications.

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The state $\boldsymbol{X}(t)$ of the system at time $t$ is a binary vector of length $N(t)$, whose $i$-th entry is 1 or 0 according to whether site $i$ is occupied by a marked cell. It takes values in the (countable) subset of $\{0,1\}^{\mathbb{N}}$ whose elements have only finitely many 1 s .

## A model for cell proliferation



## Q \& A. What can we say about the model?

- The process $(\boldsymbol{X}(t), t \geq 0)$ is a
- Consider a particular marked cell. Is its position $I(t)$ at time $t$ Markovian?
- If that cell is in position $i$, it moves to the right at rate
- The position of any particular marked cell evolves as a
- Therefore, the distance travelled by any particular marked cell up to time $t$ (its position relative to its starting site $j$ ) has a distribution.
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- If that cell is in position $i$, it moves to the right at rate $\lambda i$.
- The position of any particular marked cell evolves as a Yule Process, that is, a pure-birth process with birth rates $\lambda_{i}=\lambda i$ (in the usual notation).
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- Finally note that if we start with all marked cells in adjacent sites, then, at any future time, the gaps between them will be independent, and thus the positions of the marked cells at any fixed time $t$ will follow a discrete renewal process with negative binomial lifetimes. (We have not exploited this fact in our analysis so far, but we have plans!)


## The approach of Hywood, Hackett-Jones, and Landman

They focussed attention on the expected occupancy (occupancy probability) $C_{i}(t)$, the chance that site $i$ is occupied by a marked cell at time $t$, and derived a continuum model for occupation density $C(x, t)$ at position $x$ (taking $\Delta$ to its limit 0 ).

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## The approach of Hywood, Hackett-Jones, and Landman

(a)


(b)

Solutions, in red, to $\frac{\partial}{\partial t} C(x, t)=-\lambda \frac{\partial}{\partial x}[x C(x, t)]+\frac{\lambda \Delta}{2} \frac{\partial^{2}}{\partial x^{2}}[x C(x, t)]$, and expected occupancy estimates (1000 runs), in black, for $t=1,2,3,4$, with $L(0)=24, \lambda=0.69$, and marked cells initially in [12, 18]: (a) $\Delta=1$, (b) $\Delta=1 / 2$.

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If the marked cells are initially located at adjacent sites $j=r+1, \ldots s$, the expected occupancy for site $k$ at time $t$ is

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C_{k}(t)=\sum_{j=r+1}^{\min \{s, k\}}\binom{k-1}{k-j} e_{t}^{j}\left(1-e_{t}\right)^{k-j}, \quad \text { where } e_{t}=\exp (-\lambda t)
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To see this, generate a Yule process starting $j$ using an ensemble of independent rate- $\lambda$ Poisson processes $\left\{N_{i}^{(j)}(t), i=1,2, \ldots\right\}$ (one for each site $i$ ), and superimpose these processes ( $s-r$ of them).

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If the $\left\{N^{(j)}\right\}, j=r+1, \ldots s$, are independent ensembles of Poisson processes, we get trajectories the approximating model. However, if they are the same ensemble, we get trajectories of the original model. The above formula is a simple consequence of noticing that the approximating model is the empirical process, which counts the numbers of "particles" in each state.

## The expected occupancy




Estimates (blue), based on 10,000 runs, of the expected occupancy of the proliferation process (left) and the corresponding ensemble of Yule processes (right) over 600 sites, with $\Delta=1, t=4.0$, proliferation rate $\lambda=0.69$, and initially 7 marked cells located at sites 12 up to 18 . Also plotted (solid red) is $C_{i}(t)$ for $i=1, \ldots, 600$.

## The expected occupancy




Estimates (blue), based on 10, 000 runs, of the expected occupancy of the proliferation process (left) and the corresponding ensemble of Yule processes (right) over 2,500 sites, with $\Delta=1, t=4.0$, proliferation rate $\lambda=0.69$, and initially 107 marked cells located at sites 12 up to 118 . Also plotted (solid red) is $C_{i}(t)$ for $i=1, \ldots, 2,500$.

## The occupancy density

Since the PDE model for the occupation density $C(x, t)$ obtained by Hywood, Hackett-Jones, Landman was not really a continuum model, we tried to derive an explicit expression for occupation density, as follows:

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Suppose cells are situated at points in the interval $[a, b]$ with equal spacing $\Delta(>0)$ :

$$
a+\Delta, a+2 \Delta, \ldots, a+N \Delta(=b)
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where $N=(b-a) / \Delta$ is the number of points. The idea is that the initial "cell mass" $b-a$ is distributed evenly among these $N$ points. We are now interested in the expected occupancy at position $x$ at time $t$, namely

$$
C_{\Delta}(x, t)=\sum_{j=a / \Delta+1}^{\min \{b / \Delta, x / \Delta\}}\binom{x / \Delta-1}{x / \Delta-j} e_{t}^{j}\left(1-e_{t}\right)^{x / \Delta-j}
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$$

What happens to $C_{\Delta}(x, t)$ as $\Delta \rightarrow 0$ ?

## The occupancy density




The left-hand pane shows $C_{\Delta}(x, t)$ at $t=4.0$ for values of $\Delta$ from 1 down to $1 / n$, where $n=10$. The initial cell mass is on the interval [12, 18], and $\lambda=0.69$. The right-hand pane shows $C_{\Delta}(x, t)$ for $x=228$ (approximately where the peaks occur) for values of $\Delta=1 / n$ up to $n=100$. The code uses nbinpdf ( $k-j, j, e$ ).

## The occupancy density


$C_{\Delta}(x, t)$ at $t=4.0$ for values of $\Delta$ from 1 down to $1 / n$, where $n=5$. The initial cell mass is on the interval $[12,118]$, and $\lambda=0.69$. Decreasing $\Delta$ corresponds to an increasing amount of flatness in the curves and decreasing tail mass.

## The occupancy density - an approximation

Recall that

$$
C_{\Delta}(x, t)=\sum_{j=a / \Delta+1}^{\min \{b / \Delta, x / \Delta\}}\binom{x / \Delta-1}{x / \Delta-j} e_{t}^{j}\left(1-e_{t}\right)^{x / \Delta-j}, \quad \text { where } e_{t}=\exp (-\lambda t) .
$$

Suppose that all quantities are chosen so that $i:=x / \Delta-1, l:=a / \Delta-1, m:=b / \Delta-1$, and $n:=(b-a) / \Delta=m-l$ are integers, and in particular when $\Delta$ becomes small.

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Then, we may write

$$
C_{\Delta}(x, t)=\sum_{j=1}^{\min \{n, i-/\}}\binom{i}{i-I-j} \theta^{j+l+1}(1-\theta)^{i-I-j}, \quad \text { where } \theta=e_{t}
$$

noting that $i, I, m$, and $n$, increase at the same rate when $\Delta \rightarrow 0$, and in particular, $I / i \rightarrow a / x$ and $m / i \rightarrow b / x$.

## The occupancy density - an approximation

Next observe that

$$
C_{\Delta}(x, t)=\theta \sum_{j=l+1}^{\min \{n+l, i\}}\binom{i}{j} \theta^{j}(1-\theta)^{i-j}=\theta \operatorname{Pr}\left(I+1 \leq S_{i} \leq \min \{n+I, i\}\right)
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where $S_{i}$ has a binomial $B(i, \theta)$ distribution $(\theta=\exp (-\lambda t)$ ).
So, we may employ the normal approximation to the binomial distribution to approximate $C_{\Delta}(x, t)$ where $\Delta$ is small (and hence $i$ is large). We get $C_{\Delta}(x, t) \simeq C_{\text {approx }}(x, t)$, where

$$
C_{\text {approx }}(x, t)=\theta \operatorname{Pr}\left(\frac{a / x-\theta}{\sqrt{\theta(1-\theta)}} \sqrt{i} \leq Z \leq \frac{\min \{b / x, 1\}-\theta}{\sqrt{\theta(1-\theta)}} \sqrt{i}\right)
$$

where $Z$ is a standard normal random variable.

## The occupancy density - normal approximation



Evaluation of $C_{\Delta}(x, t)$ at $t=4.0$ with $\lambda=0.69$, and with initial cell mass on the interval $[12,18]$ (left pane) and on the interval [12, 118] (right pane). The corresponding normal approximation is shown in bold red. The code uses normcdf.

## The occupancy density - normal approximation



The normal approximation with $\Delta$ quite small ( $\Delta=0.001$ ). A clearer picture is emerging of the shape of occupation density curve $C(x, t)$.

## The occupancy density

Theorem. If, initially, the marked cells lie in the interval $[a, b]$, the occupation density at time $t$ is given by

$$
C(x, t):=\lim _{\Delta \rightarrow 0} C_{\Delta}(x, t)= \begin{cases}0 & \text { if } 0<x<a e^{\lambda t} \\ \frac{1}{2} e^{-\lambda t} & \text { if } x=a e^{\lambda t} \\ e^{-\lambda t} & \text { if } a e^{\lambda t}<x<b e^{\lambda t} \\ \frac{1}{2} e^{-\lambda t} & \text { if } x=b e^{\lambda t} \\ 0 & \text { if } x>b e^{\lambda t} .\end{cases}
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Proof. Recall that $C_{\Delta}(x, t)=\theta \operatorname{Pr}\left(I+1 \leq S_{i} \leq \min \{m, i\}\right)$, where $S_{i}$ has a binomial $B(i, \theta)$ distribution $(\theta=\exp (-\lambda t)$ ), and $I / i \rightarrow a / x$ and $m / i \rightarrow b / x$ as $i \rightarrow \infty$.

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We use Theorem 2 of ...
Arratia, R. and Gordon, L. (1989) Tutorial on large deviations for the binomial distribution. Bulletin of Mathematical Biology 51, 125-131.
$\ldots$ which provides an approximation for $\operatorname{Pr}\left(S_{i} \geq a i\right)$, $a>0$, when $i$ is large.

## The result of Arratia and Gordon

Let $H(\epsilon, \theta)$ be Kullback-Leibler divergence between an $\epsilon$-coin and a $\theta$-coin:

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H(\epsilon, \theta)=\epsilon \log \left(\frac{\epsilon}{\theta}\right)+(1-\epsilon) \log \left(\frac{1-\epsilon}{1-\theta}\right) .
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r=r(\epsilon, \theta)=\left(\frac{\theta}{1-\theta}\right) /\left(\frac{\epsilon}{1-\epsilon}\right)=\frac{\theta(1-\epsilon)}{\epsilon(1-\theta)}
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This satisfies $0<r<1$ whenever $\theta<\epsilon<1$.
Suppose $S_{i}$ has a binomial $B(i, \theta)$ distribution. If $\theta<\epsilon<1$,

$$
\operatorname{Pr}\left(S_{i} \geq i \epsilon\right) \sim \frac{1}{(1-r) \sqrt{2 \pi \epsilon(1-\epsilon) i}} e^{-i H(\epsilon, \theta)}, \quad \text { as } i \rightarrow \infty
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Evaluation of the occupation density $C(x, t)$ at times $t=0,0.5,1.0, \ldots, 5.0$, with $\lambda=0.69$, and with initial cell mass on the interval $[12,18]$.

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## The occupancy density



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Evaluation of the occupation density $C(x, t)$ at times $t=0,0.5,1.0, \ldots, 5.0$, with $\lambda=0.69$, and with initial cell mass on the interval $[0,18]$. The green bars indicate relative cell mass.

