Maintaining and escaping feedback control in hierarchically organised tissue: a case study of the intestinal epithelium

Matthias M. Fischer,[1, *] Hanspeter Herzel,[1, †] and Nils Blüthgen[1, ‡]

[1]*Institute for Theoretical Biology, Charité and*

Humboldt Universität zu Berlin, 10115 Berlin, Germany

The intestinal epithelium is one of the fastest renewing tissues in mammals. It shows a hierarchical organisation, where intestinal stem cells at the base of crypts give rise to rapidly dividing transit amplifying cells that in turn renew the pool of short-lived differentiated cells. Upon injury and stem-cell loss, cells can also dedifferentiate. Tissue homeostasis require a tightly regulated balance of differentiation and stem cell proliferation, and failure can lead to tissue extinction or to unbounded growth and cancerous lesions. Here, we present a two-compartment mathematical model of intestinal epithelium population dynamics that includes a known feedback inhibition of stem cell differentiation by differentiated cells. The model shows that feedback regulation stabilises the number of differentiated cells as these become invariant to changes in their apoptosis rate. Stability of the system is largely independent of feedback strength and shape, but specific thresholds exist after unbounded growth occurs. When dedifferentiation is added to the model, we find that the system can recover more gracefully after certain external perturbations. However, dedifferentiation makes the system more prone to loosing homoeostasis. Taken together, our mathematical model shows how a feedback-controlled hierarchical tissue can maintain homeostasis and can be robust to many external perturbations.

Keywords: Cancer, Colon, Dedifferentiation, Intestinal epithelium, Regeneration, Tissue homoeostasis

* matthias.fischer@charite.de

† h.herzel@biologie.hu-berlin.de

‡ **Corresponding author** − nils.bluethgen@charite.de

I. INTRODUCTION

In a hierarchically organised tissue, usually two classes of cells can be distinguished: Adult stem cells with unlimited capacity of self-renewal, which differentiate into cells with limited proliferative potential that perform the designated function of the tissue [1]. Additionally, tissue damage can lead to dedifferentiation of differentiated cells back into cycling stem cells Bonventre [2], Tata et al. [3], Liu and Chen [4], Liu et al. [5]. Homoeostasis of such a tissue in the face of external perturbations requires a tight regulation of the stem cell compartment. Upon tissue damage, stem cells need to increase proliferation; however proliferation of the stem cell compartment has to be tightly controlled to prevent unlimited growth of the tissue [6]. Such a control seems to be realised by specific feedback loops exerted by differentiated cells onto the stem cell compartment [7]. In contrast, control of the dedifferentiation of differentiated cells seems to be exerted by the stem cell compartment (Tata et al. [3], Beumer and Clevers [8] and the references therein).

The intestinal epithelium is a prime examples of hierarchically organised tissues. Despite its simple single-layered structure, it is able to withstand continuous mechanical, chemical and biological stresses due to its specific tissue architecture and a high rate of tissue turnover [9]: Stem cells at the bottom of the intestinal crypts divide continuously approximately once per day. Cells mature while migrating out of the crypt until they terminally differentiate and become part of the villi, eventually undergoing apoptosis and being shed into the intestinal lumen [10]. In the intestine and colon, the stem cell compartment is controlled via differentiated epithelial cells releasing Indian Hedgehog (Ihh), which stimulates mesenchymal cells to release Bone Morphogentic Proteins (BMPs). These interfere with intracellular effects of WNT signalling and thus stimulate stem cell differentiation [11–15].

Previous theoretical research on hierarchical tissues has often focussed on the case of arbitrary tissues: Rodriguez-Brenes et al. [16] considered a generic hierarchival tissue with a compartment of cycling and differentiating stem cells, and another compartment of non-cycling differentiated cells undergoing apoptosis, assuming the differentiated cell compartment may exert feedbacks onto the stem cells by decreasing their rate of proliferation and reducing the probability of symmetrical stem cell division. They then studied the order in which

mutations in feedbacks need to arise to yield a selective advantage. The same model was
used in Rodriguez-Brenes et al. [17] to reveal that during recovery from an injury dampened
oscillations may occur, which is more pronounced when the stem cells are only a small
fraction of the cell population. The same model has also been studied by Sun and Komarova
[18] to obtain analytical solutions for the mean and variance of the cell compartment sizes.
Recently, Wodarz [19] has extended the model by including dedifferentiation. Assuming
a specific sigmoidal feedback onto stem cell cycling rate and self-renewal probability, he
studied the effect of a linear and a sigmoidal dedifferentiation term, showing how unbounded
cancerous tissue growth may arise as a consequence of escaping this feedback. He also
demonstrated how dedifferentiation may allow for speedier regeneration after perturbations
using numerical simulations.

Other tissue-specific theoretical studies include models of the haematopoietic system [20–24],
the mammalian olfactory epithelium [25, 26], and breast cancer [27]. Models of the intestinal
epithelium have been reviewed in Carulli et al. [28]. These include Paulus et al. [29], a model
for its post-irradiation recovery that showed that damage-control response may reside solely
in the stem cell compartment without requiring a feedback from the villus, challenging earlier
views. Another important model of the colon epithelium [30] showed that a "bang-bang"
control of proliferation is optimal to minimise the time for crypt construction. However,
this model assumes continuous exponential growth, thus it is not suitable to address tissue
homoeostasis and stability.

Theoretical examinations that address mechanisms of homoeostasis and dynamics of healthy
intestinal epithelium have been rare. A noteworthy exception is Johnston et al. [31], who
derived a three-compartment ODE model distinguishing stem, transit-amplifying and termi-
nally differentiated cells, assuming that the first two compartments limit their own growth
via a negative saturating or a negative quadratic term, reminiscent of classical single-species
population dynamic models [32]. However, no mechanistic justification of these assumptions
has been provided, possibly owing to the fact that our mechanistic understanding of the
biology of intestinal stem cells has been limited until very recently [9]. Therefore, in this
work, we set out to close this gap in the literature and derive a m¸odel of intestinal epithelial
population dynamics based on our current mechanistic understanding of the involved under-

lying biological processes. This model allowed us to study specifically the role of feedbacks
and dedifferentiation in tissue homoeostasis and regeneration for the specific case of the colon
epithelium.

II.   MATERIALS AND METHODS

A.   Colon epithelium model

Our main model consists of two cell compartments S and D, denoting stem cells, and
differentiated cells, respectively, following earlier approaches Rodriguez-Brenes et al. [16].
Stem cells cycle at a constant rate β, differentiated cells are cell-cycle arrested and die with
an apoptosis rate ω. Finally, stem cells differentiate with a rate $\delta(D)$ that is a function of
the size of the differentiated cell compartment. Following earlier models in the literature
such as Rodriguez-Brenes et al. [16], we consider transit-amplifying cells to be a part of the
differentiated cell compartment D, since their limited proliferative potential distinguishes
them from stem cells [9].
85
86 Overall, the model is described by two coupled ordinary differential equations:

$$
\begin{aligned}
\frac{\mathrm{d}S(t)}{\mathrm{d}t} &= \beta S(t) - \delta(D)S(t) \\
\frac{\mathrm{d}D(t)}{\mathrm{d}t} &= \delta(D)S(t) - \omega D(t),
\end{aligned}
\tag{1}
$$

87 where $\beta, \omega \in \mathbb{R}^+$ and δ is a continuously differentiable, monotonically increasing, invertible
88 function $\mathbb{R}^+ \to \mathbb{R}^+$. This function δ models the results of experimental studies on intestinal
89 tissue that demonstrated the stimulation of stem cell differentiation by differentiated cells
90 via Indian Hedgehog (*Ihh*), and Bone Morphogenic Protein (*BMP*) [11, 12]. A sketch of the
91 model is shown in Figure 1a..
92

B.   Colon epithelium model with dedifferentiation

94 We extend the model by a dedifferentiation process of differentiated cells into stem cells. In
95 this model, the rate of dedifferentiation is a continuously differentiable and monotonically

96 decreasing function $\varrho : \mathbb{R}^+ \to \mathbb{R}^+$ of the stem cell compartment, where a higher number of
97 stem cells reduces dedifferentiation. Thus, the model reads:

$$\begin{aligned}
\frac{\mathrm{d}S(t)}{\mathrm{d}t} &= \beta S(t) - \delta(D)S(t) + \varrho(S)D \\
\frac{\mathrm{d}D(t)}{\mathrm{d}t} &= \delta(D)S(t) - \varrho(S)D(t) - \omega D(t).
\end{aligned} \tag{2}$$

98 A sketch of the model is shown in Figure 4a.

C.   Numerical simulations

100 Numerical simulations have been implemented in the Python programming language, version
101 3.7.3. We provide the complete commented source code under: `https://github.com/`
102 `Matthias-M-Fischer/Epithelium`.

III.   RESULTS

A.   A population dynamics model of the colon epithelium

105 We developed a model of the intestinal epithelial population dynamics (Fig. 1a, 'Materials
106 and Methods'), distinguishing between two compartments: Stem cells cycling at a rate β
107 and differentiating at a rate $\delta(D)$; and differentiated cells undergoing apoptosis and leaving
108 the system at a rate ω. We model the differentiation rate as a positive function of the size of
109 the differentiated cell compartment D, reflecting earlier experimental results [11, 12].
110

1.   Stability can be lost via two routes

112 First we study the steady-states of system (1). For now, we will not use any specific
113 function δ and only demand that δ is a positive, monotonic, continuously differentiable and
114 invertible function. We also assume $\mathrm{d}\delta/\mathrm{d}D \geq 0$, since differentiated cells stimulate stem cell
115 differentiation.
116

By solving $\mathrm{d}S(t)/\mathrm{d}t = \mathrm{d}D(t)/\mathrm{d}t = 0$ we find two steady-states, which in the following we

will mark with an over-bar. We find a trivial steady-state $(\bar{S} = 0, \bar{D} = 0)$ that exists for all

parameter values. Under some conditions, also a non-trivial steady-state exists at $\bar{S} = \omega\bar{D}/\beta$

and $\bar{D} = \delta^{-1}(\beta)$. Here, δ^{-1} denotes the inverse function of δ. This non-trivial steady-state

exists if δ can reach the value of β. Thus, this second steady-state is only present if the

maximum differentiation rate is higher than the proliferation rate. One may see that the

steady-state does not depend on the apoptosis rate ω. This is a biologically interesting

property which we will return to later.

To address the stability of the steady-states, we investigate the Jacobian [33] of the system:

$$\mathbf{J} = \begin{pmatrix} \beta - \delta(D) & -S\delta'(D) \\ \delta(D) & S\delta'(D) - \omega \end{pmatrix},$$

127 which at the trivial steady-state has eigenvalues $\lambda_1 = \beta - \delta(0), \lambda_2 = -\omega < 0$. Hence, the

128 trivial steady-state is only stable if β is smaller than the basal differentiation rate $\delta(0)$.

130 At any non-trivial steady-state, the Jacobian is given by

$$\mathbf{J}_{s-s} = \begin{pmatrix} 0 & -S\delta'(\bar{D}) \\ \beta & S\delta'(\bar{D}) - \omega \end{pmatrix}.$$

The steady-state is stable if $\det \mathbf{J}_{s-s} > 0$ and $\operatorname{tr} \mathbf{J}_{s-s} < 0$ (Routh–Hurwitz criterion, see

Strogatz [33]), which leads to the condition that $0 < \delta'(\bar{D}) < \delta(\bar{D})/\bar{D}$.

These results reveal two routes of how the intestinal epithelium can lose homeostasis

(Fig. 1b): First, the tissue might show unbounded growth, which could be interpreted

as the emergence of a cancerous lesion (Fig. 1b, second column). This occurs once the

proliferation rate β exceeds the maximum differentiation rate δ_{max} (either by increased pro-

liferation or decreased maximal differentiation). In this case, only the (unstable) steady-state

$(0, 0)$ remains, and any positive perturbation of S away from it will lead to unbounded growth.

Second, the steady-state $\bar{D} = \delta^{-1}(\beta)$ can get unstable if $\delta'(\bar{D}) < \delta(\bar{D})/\bar{D}$. In other words,

at the steady-state the slope of the feedback function might exceed $\delta(\bar{D})/\bar{D}$. Then, the

behaviour of the system depends on the feedback function. This can be illustrated with the following two examples:

First, consider a linear function of the form $\delta(D) = \delta_0 + \delta_{slope}D$. The intercept δ_0 denotes the basal differentiation rate of stem cells in the absence of external stimuli, whereas δ_{slope} quantifies the increase in differentiation as a function of $D(t)$. For instability, we require $\delta'(\bar{D}) > \beta/\bar{D}$, hence the system will only be stable for a strictly positive δ_0. This is biologically plausible, since the basal stem cell differentiation rate can be expected to exceed zero. If, however, $\delta_0 = 0$, then the fix point will be unstable. This case is equivalent to the classical Lotka-Volterra model of predator-prey population dynamics, exhibiting undampened oscillations. As a second example, consider a sigmoidal function of the form $\delta(D) = \delta_0 + \delta_{max}D^p/(D_{mid}^p + D^p)$. Here, δ_{max} denotes the maximum increase in differentiation rate the stem cell compartment may experience, D_{mid} is the position of the mid-point of the sigmoidal function, and p quantifies the steepness of the feedback function around it. For instability, we require $\delta'(\bar{D}) > \beta/\bar{D}$. Hence, at $p > 4\beta/\delta_{max}$ the steady-state loses its stability, and sustained oscillations start to occur.

2. Exponential growth and convergence to a stable cell type ratio after escaping control

Next, we analyse the behaviour of system (1) during unbounded growth. This case corresponds to a tissue that has escaped homeostatic control and has degenerated into a cancerous lesion. In our model, this occurs when β exceeds the maximum of δ, implying that δ has a maximum value $\delta_{max} < \beta$. Such saturation of the differentiation rate could arise from saturated signalling or thermodynamical constraints.

Interestingly, during such unbounded growth the system will converge to a stable cell type ratio ($S(t)/D(t) = \text{const}$ for $t \to \infty$): After feedback saturation, the dynamics of the system is governed by the following differential equations:

$$\frac{\mathrm{d}S(t)}{\mathrm{d}t} = (\beta - \delta_{max})S(t)$$
$$\frac{\mathrm{d}D(t)}{\mathrm{d}t} = \delta_{max}S(t) - \omega D(t).$$

169 Directly solving for $S(t)$ yields a simple exponential function, and subsequently we may also
170 solve $D(t)$. Overall, we get:

$$S(t) = S_0 e^{(\beta - \delta_{max})t}$$

$$D(t) = D_0 e^{-\omega t} + S_0 \delta_{max} e^{-\omega t} \int_{s=0}^{s=t} e^{(\beta - \delta_{max} + \omega)s} \, ds \,,$$

171 where $S_0 = S(0), D_0 = D(0)$ denote the initial condition of the system. This yields

$$D(t) = D_0 e^{-\omega t} + \frac{S_0 \delta_{max}}{\beta - \delta_{max} + \omega} e^{(\beta - \delta_{max})t} \,.$$

172 This allows us to take the following limit

$$\lim_{t \to \infty} \frac{S(t)}{D(t)} = \frac{\beta + \omega}{\delta_{max}} - 1 \,,$$

173 which denotes the ratio of stem to differentiated cells the system will converge to during
174 explosive growth.

176 Additionally, one may see that after the transient period the overall growth of the system
177 amounts to

$$S(t) + D(t) = S_0 \frac{\beta + \omega}{\beta + \omega - \delta_{max}} e^{(\beta - \delta_{max})t},$$

178 which implies an exponential growth at a constant rate $\beta - \delta_{max}$. Figure 1c provides an
179 exemplary numerical simulation of the system with a piecewise linear feedback function
180 $\delta(D) = \min\{0.9 + 10^{-4}D, 1.0\}$, and parameters $\beta = 1.1$ and $\omega = 0.1$. Observe the convergence
181 of the ratio $S(t)/D(t)$ to a stable ratio of $-1 + (\beta + \omega)/\delta_{max} \approx 0.2$, which is in agreement
182 with our theoretical analysis.

183   *3.   Bifurcation analysis for the case of piecewise linear δ*

184 Now, we further analyse the influence of system parameters on the behaviour of system 1.
185 To this end, we now choose a concrete differentiation rate function δ. For simplicity, we
186 choose a piecewise linear function with a positive intercept $\delta_0 > 0$, denoting the basal stem

187 cell differentiation rate. We assume that δ grows linearly in D with a slope of $\delta_{slope} > 0$,
188 until it reaches an upper bound δ_{max}, where it stops increasing. Hence, we have

$$\delta(D) = \min\{\delta_0 + \delta_{slope}D, \delta_{max}\}, \tag{3}$$

189 where $\delta_0, \delta_{slope}, \delta_{max} > 0$.

191 The non-trivial steady-state resides at $\bar{S} = \omega\bar{D}/\beta, \bar{D} = (\beta - \delta_0)/\delta_{slope}$ and only exists if
192 $\beta < \delta_{max}$. The steady-state is only biologically feasible $(\bar{S}, \bar{D} > 0)$ if $\delta_0 < \beta$. Then, the
193 Jacobian at the steady-state is given by

$$\mathbf{J}_{s-s} = \begin{pmatrix} 0 & -\frac{(\beta - \delta_0)\omega}{\beta} \\ \beta & \frac{(\beta - \delta_0)\omega}{\beta} - \omega \end{pmatrix},$$

194 with eigenvalues

$$\lambda_{1,2} = \frac{-\delta_0\omega \pm \sqrt{-4\beta^3\omega + 4\beta^2\delta_0\omega + \delta_0^2\omega^2}}{2\beta}.$$

195 If

$$\beta - \delta_0 < \frac{\delta_0^2\omega}{4\beta^2}$$

the radicand is positive. Then, due to $\delta_0 < \beta$ both eigenvalues will be negative and real,
hence the steady-state is a stable node. If, however, the radicand is negative, oscillations
occur and the real part of the eigenvalues is always negative due to $\delta_0, \omega > 0$, making the
steady-state a stable focus. In any case, any biologically feasible steady-state of this system
will always be stable.

Biologically, it is interesting that no other constraints on δ_0 and δ_{slope} are required except
for $\delta_0 < \beta$, which we require for a feasible steady-state, and $\beta < \delta_{max}$. In particular,
$\delta_0 > 0, \delta_{slope} > 0$ can be arbitrarily small, yet the system remains stable. Similarly, changes
in $\omega \geq 0$ will not affect the stability of the system. Figure 1d illustrates these findings and
shows bifurcation diagrams of the system for varying parameters $\beta, \omega, \delta_0, \delta_{slope}$.

4.   The feedback loop can cause oscillatory behaviour in cell numbers

Next, we investigate the case of the system showing dampened oscillations around the
steady-state. For any given steady-state $\bar{S}, \bar{D}$ it holds that $\delta_0 = \beta - \delta_{slope}\bar{D}$. This permits
to express $\lambda_{1,2}$, and thus the amplitude decay and the angular frequency of the dampened
oscillations in terms of δ_{slope}. We find that the amplitude of the oscillations will decay with

$$\sim e^{\dfrac{\delta_{slope}\bar{D}\omega - \beta\omega}{2\beta}t}.$$

213 Note that at any equilibrium the exponent is negative because from $\bar{D} = (\beta - \delta_0)/\delta_{slope}$ it
214 follows that $\delta_{slope}\bar{D} - \beta = -\delta_0 < 0$. The angular frequency of oscillations reads

$$\omega_0 = \sqrt{\delta_{slope}\bar{D}(4\beta^2\omega + \omega^2) - \beta\omega^2}/2\beta.$$

215 Hence, a greater slope of the feedback function δ will result in a slower decay of the amplitude,
216 as well a higher cycling frequency. For illustration, we provide a set of numerical simulations
217 of the system with a piecewise linear function δ as defined before and varying values of δ_{slope}
218 in Figure 1e.
219

The relationship between function slope and oscillatory behaviour of the system is a biologi-
cally interesting result. Observe how large values of δ_{slope} cause strong oscillations which are
biologically unlikely. This suggests that to minimise oscillations in cell numbers, a smaller
slope of the feedback function is desirable. For very small slopes, the difference between β
and δ_0 needs to become sufficiently small as well, if the position of the steady-state $(\bar{S}, \bar{D})$
should remain constant. Then, the oscillations will vanish, and the steady-state becomes a
stable node, as shown previously.
227

228   **B.   Comparison of different model topologies**

229 In the previous section, we have shown that the colon epithelium model can cause homoeosta-
230 sis through a feedback from differentiated cells onto the stem cell differentiation rate. Now,
231 we explore how alternative feedback topologies in our two-compartment colon model might

FIG. 1. The basic colon epithelium model. **(a)** Schematic sketch of the model. **(b)** The feedback function δ (upper row) determines the qualitative behaviour of the model (lower row). First column: Stable steady-state. Second column: unbounded growth. Third column: sustained oscillations. **(c)** In case of explosive growth, the system converges to a stable ratio of stem to differentiated cells. Exemplary numerical integration. **(d)** Bifurcation diagram of the system with linear feedback function δ. Standard parametrisation: $\beta = 1.0, \omega = 0.1, \delta_0 = 0.9, \delta_{slope} = 10^{-4}, \delta_{max} = 2.0$. Note the switch from a stable steady-state to unbounded growth for $\beta > \delta_{max}$ (upper-left panel). Also note the invariance of $\bar{D}$ to changes in ω (upper-right panel). Both properties hold regardless of how we chose δ. Finally, notice that $\delta_0 > 0, \delta_{slope} > 0$ can be made arbitrarily small without switching to unbounded growth; however, if $\delta_0 > \beta$, the system will go extinct (lower two panels). **(e)** A steeper feedback functions causes stronger and longer oscillations. Exemplary numerical simulations with linear function δ for varying δ_{slope}; steady-state fixed at $\bar{S} = 100, \bar{D} = 1000$.

232 alter tissue homeostasis, and compare the properties of these models. We therefore generate

233 the family of all six one-looped model topologies.

1. Controlling stem cells is required for stability

First, we consider those two model topologies where the apoptosis rate ω is regulated by either S or D, respectively. These feedback loops are biologically unlikely, but we mention them for the sake of completeness. Both topologies are not able to show homoeostasis: If both β and δ are unregulated and hence constant, we have $\mathrm{d}S(t)/\mathrm{d}t = \beta S - \delta S$, which for any non-trivial steady-state requires $\beta = \delta$. This, in turn, implies $\forall t : \mathrm{d}S(t)/\mathrm{d}t = 0$, hence after any perturbation of S, S will not return to its pre-perturbation state. Accordingly, these models are biologically not realistic, and we will not consider them in the rest of this paper.

2. Indirect regulation of the stem cell compartment decouples the steady-state number of differentiated cells from their apoptosis rate

Next, we show that all models in which the stem cell compartment is exclusively regulated by the differentiated cell compartment D following arbitrary continuously differentiable functions $\beta(D), \delta(D)$ enjoy the property of $\bar{D}$ being independent of ω: Let $\mathrm{d}S(t)/\mathrm{d}t = \beta(D)S - \delta(D)S$. At any non-trivial steady-state $(\bar{S}, \bar{D})$, from $\mathrm{d}S(t)/\mathrm{d}t = 0$ we get that $\beta(\bar{D}) = \delta(\bar{D})$. Define $\alpha(D) := \beta(D) - \delta(D)$, then $\bar{D} = \alpha^{-1}(0)$, which does indeed not depend on our choice of ω.

3. Saturating feedback functions β and δ cause convergence to a stable cell type ratio during unbounded growth

Finally, it may also be briefly remarked that all systems in which proliferation and differentiation rates of the stem cell compartment are functions saturating to β_{min} and δ_{max} will in case of unbounded growth converge to a stable ratio $S(t)/D(t) = \mathrm{const}$ for $t \to \infty$, given by:

$$\lim_{t\to\infty} \frac{S(t)}{D(t)} = \frac{\beta_{min} + \omega}{\delta_{max}} - 1.$$

4. *Differences in relaxation dynamics of different model topologies after perturbations*

In this section, we compare the relaxation dynamics of the four remaining model topologies after perturbations from steady-state, again using simple piecewise linear feedback functions to keep analyses tractable. We examine three different kinds of perturbations: first, removing differentiated cells; second, removing stem cells; and third, both of these perturbations combined. The first of these perturbations corresponds to a loss of differentiated cells which e.g. may arise as a consequence of physical stresses, inflammation or interactions of intestinal cells with microbes or parasites [9]. The second perturbation describes the loss of intestinal stem cells, which for instance arises as a consequence of radiation-induced cell damage [34]. Alternatively, stem cells numbers also might be reduced due to temporarily increased stem cell differentiation due to stochastic effects which may arise in such small populations.

Because of bilinear terms in the equations, we cannot in all cases obtain exact analytical solutions of the occurring dynamics, but use approximations of the dynamics based on a linearisation of the systems around their respective non-trivial steady-state (see Supplement for details).

The Jacobians and their Eigensystems for the remaining four models are shown in Figure 2, along with schematic model sketches and exemplary numerical simulations, illustrating typical solutions.

Physiologically, it is important that the number of differentiated cells recovers quickly, as these are responsible for carrying out the function of a tissue. We hence compare the models based on their 'defect' χ of differentiated cells after perturbation – see Panel (a) of Figure 3 for an illustration. We define the defect as the total area between $D(t)$a nd $\bar{D}$ after a perturbation at $t = 0$, i.e.

$$\chi := \int\limits_{\substack{t=0 \\ D(t)<\bar{D}}}^{\infty} \bar{D} - D(t) \, \mathrm{d}t \, .$$

The defect integrates only those time intervals in which $D(t) < \bar{D}$, as this defines the lack of tissue functionality.

FIG. 2. All four one-looped model topologies that can show homoeostasis. First column: schematic sketches; second column: exemplary numerical simulations; third column: model equations, as well as Jacobian at the non-trivial steady-state and its eigensystem for the case of a linear feedback function $\beta(x) = \beta_0 + \beta_{slope}x$ or $\delta(x) = \delta_0 + \delta_{slope}x$, respectively.

For all four models and the three types of perturbations, we derived either an analytical

expression for the defect, or – if these are too complicated for a meaningful interpretation –

an approximate solution (see Supplement). Three relations simplify their analysis: First, for

all models we obtain $\omega = \beta\bar{S}/\bar{D}$. Second, the slope of the feedback function (β_{slope} or δ_{slope},

respectively) is not required for describing the dynamics relative to the steady-state, as the

slope is not part of the Eigensystems of the steady-state Jacobians of the models. Third,

the displacements of all models from steady-state of the first and second perturbation are

linear in the initial displacement ($\Delta D(0)$ or $\Delta S(0)$, respectively). This indicates that in both

cases the choice of initial displacement will not affect the comparison of the model defects,

and we can express all defects as multiples of initial displacement. In case of the third perturbation, we remove the same fraction of stem and differentiated cells, and we express the defect as multiples of the intial displacement of differentiated cells. Overall, the complete parameter space we need to consider thus consists only of the ratio of stem to differentiated cells at steady-state and two additional free parameters (depending on the model β_0, δ or β, δ_0).

a.   Colon model vs. direct stimulation of stem cell differentiation   We start by comparing our basic colon epithelium model (model 1), with model 2, where stem cell differentiation is stimulated by the stem cell compartment itself (Figure 3b). Figure 3c shows the difference in model defects throughout the parameter space (see Supplement for the raw values), where areas shaded in red indicate regions where the colon epithelium model has a bigger defect than the alternative model. In case of removing differentiated cells (first row) or removing both differentiated and stem cells at the same time (third row), there exist ample regions (shaded in blue) where the colon model recovers more efficiently – namely, whenever δ_0 is close to β. This makes sense, because in the colon epithelium model removing differentiated cells will cause the stem cells to differentiate more slowly and thus grow in numbers quickly, thus being able to replenish the differentiated cell compartment more quickly. However, in case of removing only stem cells (second row), the colon epithelium model always performs worse than the alternative model, except for cases where the fraction of stem cells at the steady-state becomes sufficiently big (middle and right column). However, even then the difference between δ_0 and β still needs to be small for the colon model to recover more efficiently.

b.   Colon model vs. indirect inhibition of stem cell cycling rate   Next, we compare the recovery dynamics of the colon epithelium model with alternative model 3, where the differentiated cell compartment inhibits stem cell cycling. Because the two models have different system parameters (β, δ_0 vs. β_0, δ) we cannot compare them pointwise in parameter space. However, we can still compare the ranges of model defects for the three different perturbations and steady-state stem cell fractions (1%, 10%, and 25%) if we systematically vary the two other free parameters within biologically plausible intervals. In all cases, the defects of the two models fall into similar ranges (see Supplement). Hence, there does not seem to be any relevant difference in recovery behaviour between indirectly regulating stem cell differentiation vs. indirectly regulating stem cell cycling rate.

FIG. 3. **(a)** Illustration of our model comparison: After a perturbation at $t = 0$, the model relaxes to its steady-state (continuous blue and orange lines, depicting differentiated and stem cells, respectively). We use a first-order approximation (continuous red line) to not have to rely on costly numerical solutions of the system, and compute the 'defects' χ of the models we want to compare (shaded red area). **(b)** The two models we compare in this figure. Top: model 1, where stem cell differentiation is stimulated by the differentiated cell compartment; bottom: model 2, where the stem cell compartment stimulates its own differentiation. Note that model 1 is equivalent to our basic colon epithelium model derived earlier. **(c)** Difference in defects of model 2 and 1 throughout the parameter space. Areas shaded in red depict regions where model 1 shows a bigger defect, i.e. where the colon model recovers less gracefully than the alternative model; blue areas indicate the opposite. Columns represent different cases of stem cell fraction at steady-state (1, 10, and 25% respectively), rows represent the three different kinds of perturbations (removing differentiated cells, removing stem cells, and removing both, respectively).

c. Colon model vs. self-inhibition of stem cell cycling rate   Finally, we compare the recovery

dynamics of the colon epithelium model with the behaviour of model 4, where the stem cell

compartment inhibits its own cycling rate. We again face the problem of the two models

having some different system parameters (β, δ_0 vs. β_0, δ), thus we again compare the ranges

of defects occurring in different scenarios. For the first and third perturbation, model 4

recovers slower by several orders of magnitude if its stem cell differentiation rate δ is small

(see Supplement), since a small differentiation rate will cause the system to take longer to

replenish the pool of differentiated cells. The only exception is when the steady-state stem cell

fraction is very large. In contrast, if differentiated cells are removed from the colon epithelium

model, the stem cell differentiation rate will decrease, causing the stem cell compartment to

temporarily overshoot, until the growing differentiated stem cell compartment stimulates

differentiation again. This way, the colon epithelium model is able to recover more quickly.

For the remaining case of removing stem cells, the model defects fall into similar ranges.

However the colon epithelium model shows its largest defect in case of a small δ_0, whereas

the alternative model performs worst in case of a very small difference between β_0 and δ.

C.   Dedifferentiation improves recovery from perturbations, however offers an

additional route of losing homoeostasis

Previously, we have seen that upon loss of stem cells, the colon epithelium model often

does not recover as gracefully as model 2 (Figure 3), as removing stem cells will not alter

the rate of stem cell differentiation. Hence, removing stem cells will lead to a significant

loss of differentiated cells first, before differentiation rate drops enough for the stem cell

compartment to replenish itself. This way, the transient behaviour is characterised by large

oscillations in the number of differentiated cells, causing a large model defect.

We now study how allowing for the dedifferentiation of differentiated cells affects these

recovery dynamics (Figure 4a). We again assume for simplicity that the rate of stem cell

differentiation is given by a linear function with intercept δ_0 and slope δ_{slope}. We also assume

that the function saturates to a maximum value δ_{max} after some value of D. Hence, we have

$\delta(D) = \min\{\delta_0 + \delta_{slope}D, \delta_{max}\}$, where $\delta_0, \delta_{slope}, \delta_{max} > 0$. Next, we also for now assume the

same, but horizontally mirrored shape for ϱ, giving $\varrho(S) = \max\{\varrho_0 + \varrho_{slope}S, \varrho_{min}\}$, where $\varrho_0 > 0, \varrho_{slope} < 0, \varrho_{min} \geq 0$.

For brevity purposes, the exact calculations are presented in the Supplement. Briefly, adding a linear dedifferentiation function causes a faster decay of the oscillations after perturbations. Additionally, we can find a critical value ϱ_0^*, which, when exceeded by ϱ_0 will also reduce the frequency of oscillations after perturbations. It is given by

$$\varrho_0^* = (\beta - \delta_0/2)4\omega^2/\beta^2.$$

By means of a Taylor expansion around the steady-state, we can also generalise this finding to arbitrary decreasing differentiable functions ϱ. Figure 4b shows some exemplary numerical simulations of this behaviour. Observe, how the transient period after removing stem cells is characterised by smaller oscillation amplitudes and frequencies when dedifferentiation is allowed.

Next, we the influence of dedifferentiation on the stability of the non-trivial steady-state of the system. Regardless of the concrete functions δ, ϱ, we find that the non-trivial steady-state needs to satisfy $\beta - \delta(\bar{D}) + (\beta/\omega)\varrho(\bar{S}) = 0$ and $\bar{D} = \beta\bar{S}/\omega$. Hence, a non-trivial steady-state exists, if and only if we can solve

$$\beta - \delta((\beta/\omega)\bar{S}) + (\beta/\omega)\varrho(\bar{S}) = 0, \bar{S} > 0. \tag{4}$$

Importantly, this shows that dedifferentiation enables the system to lose its non-trivial steady-state through a different route, namely via a sufficient increase of the values of ϱ (see Panel (c) of Figure 4 for an illustration).

Then, the system will again converge to a stable cell type composition

$$\lim_{t\to\infty} \frac{S(t)}{D(t)} = \frac{a + b + w}{2\delta_{max}},$$

where $a := \sqrt{4\delta_{max}\varrho_{min} + (b+w)^2}, b := \beta - \delta_{max}, w := \omega + \varrho_{min}$ (see Supplement for details).

FIG. 4. The colon epithelium model with dedifferentiation and its most important properties. **(a)**: Schematic sketch of the model. **(b)**: Exemplary numerical simulations of the system with piecewise linear differentiation rate function δ and different dedifferentiation rate functions ϱ: no dedifferentiation (first column), linear dedifferentiation with $\varrho_0 = 0.5, \varrho_{slope} = -0.01$ (second column) and a faster linear dedifferentiation $\varrho_0 = 0.9, \varrho_{slope} = -0.01$ (third column). System parameters are $\beta = 1, \delta_0 = 0.9, \delta_{slope} = 10^{-4}, \omega = 0.1$. Note how adding dedifferentiation, as well as increasing the higher maximum dedifferentiation rate ϱ_0 makes the system recover more gracefully after removing stem cells. **(c)**: Adding dedifferentiation opens up a second way the system can lose homeostatic stability. First column shows the case of homoeostasis. Equilibrium stem cell pool size $\bar{S}$ is given by solving $\beta \varrho(\bar{S})/\omega - \delta(\beta \bar{S}/\omega) = \beta$. Second column: Sufficient decrease of differentiation rate destroys the non-trivial steady-state and unbounded growth occurs. Third column: Sufficient increase of dedifferentiation rate has the same consequence.

IV. DISCUSSION

In this work, we derived and analysed a simple population dynamics model of the colon epithelium, taking into account the stimulating effect of differentiated cells onto the differentiation of stem cells [11–15]. Despite its simplicity, the model recapitulated known experimental results, such as the necessity to control the stem cell compartment for tissue homoeostasis [6], the balance between stem cell proliferation and differentiation required for constant tissue size [35, 36], and the emergence of stable intratumoural heterogeneity in a growing cancer (see below).

We showed that at any stable steady-state the number of differentiated cells is not affected by changes in their apoptosis rate, which is a biologically useful property, since differentiated cells are responsible for carrying out the primary function of a tissue [1, 10] and changes in apoptosis rate may regularly happen locally as a consequence of infections or mechanical wounding [9]. Additionally, we have seen that strong alterations in system parameters are required for the steady-state to be destroyed and unbounded growth to occur. In contrast, other single alterations will only change the position of the stable steady-state. This is equivalent to the case of a pre-cancerous lesion, where only a subset of 'canonical' colorectal cancer mutations is yet present and which shows a higher number of cells, but requires further alterations to switch to unbounded growth [37]. Hence, despite its simplicity, our model recapitulates the observation of colorectal tumorigenesis being a characteristic multi-step process of subsequent system alterations in vivo [38], showcasing the intrinsic resilience of the system towards isolated mutations affecting system properties.

However, the indirect feedback loop in the model may cause oscillations. This is reminiscent of e.g. the dampened oscillations observed in the healthy haematopoietic system [39]. In case of very steep feedback functions, these oscillations are more pronounced or can even become undampened; thus, we may expect rather flat feedbacks in nature. Oscillations in cell numbers do not serve any immediately obvious biological purpose and indeed may not be desirable for upholding tissue homoeostasis. This led us to the question whether the colon model topology may offer additional beneficial properties compared to other topologies, to which we compared our model.

We revealed pronounced differences of the plausible model topologies regarding their recovery after perturbations. We have shown the existence of ample regions in parameter space where the colon epithelium model recovers more gracefully than the other examined model or at least equally well. If, however, a perturbation removes stem cells, recovery behaviour was poor.

This prompted us to study the behaviour of the model if we allow for dedifferentiation of differentiated cells. We revealed that this lets the model recover more gracefully; however, dedifferentiation may also cause the destruction of the non-trivial steady-state. This is biologically interesting because it suggests that a sufficient degree of dedifferentiation may offer an alternative route towards unbounded growth. Experimental data seems to confirm the possibility of tumours arising from the differentiated intestinal epithelium [40]. Interestingly, the mutations required for this indeed occur rather often in colorectal cancers [41, 42], suggesting that increased dedifferentiation might regularly contribute to tumorigenesis. Experimental data [43] point into the same direction.

Obviously, dedifferentiation interferes with the invariance of the steady-state number of differentiated cells to changes in apoptosis rate. However, if the dedifferentiation rate around the steady-state is small and only becomes noticeable if a sizeable fraction of stem cells is removed (which seems to be suggested by the literature, see Tata et al. [3] and Beumer and Clevers [8]), then around the steady-state dedifferentiation becomes negligible and the biologically desirable invariance still holds.

We also revealed that a tissue can be expected to converge to a stable cell-type ratio during unbounded growth. From a biological point of view, this relates to the topic of intra-tumoural heterogeneity [44–46], and suggests that in case of cancerous growth the colon epithelium can still be expected to recapitulate known cellular hierarchies and differentiation gradients. In case of breast cancer [47] and cancers of the haematopoietic system [48] this has already been observed experimentally.

An interesting avenue for future research lies in comparing healthy with tumoural intestinal tissue to clarify, whether one indeed observes similar cellular subpopulations and differen-

tiation gradients. Recent research indeed suggests this [49, 50]. Tracking the numbers of different cell types over time during development and after perturbations could therefore be expected to yield data for validating and parametrising the presented model. Additionally, comparing the quantitative behaviour of tissues in different stages of tumorigenesis could elucidate at which stage which system properties change in which way, and thus increase our mechanistic insight into the dynamics of tumour development. This, in turn, might ultimately help to contribute to the development of novel approaches to limiting tumoural growth *in vivo*.

ACKNOWLEDGMENTS

We thank Christine Sers and Markus Morkel for fruitful discussions and helpful comments during an early stage of this project and DFG for funding (RTG2424, CompCancer). MMF would like to thank J. F. W. Maas for help during the preparation of the schematic model sketches.

DECLARATIONS

A. Author contributions

All authors conceived of the presented ideas. MMF carried out model derivations and analyses, and wrote the initial version of the manuscript, both with help from NB and HH. HH and NB contributed to the final version of the manuscript. NB supervised the project. All authors have read and approve of the final version of this manuscript.

---

[1] Irving L Weissman. Stem cells: units of development, units of regeneration, and units in evolution. *Cell*, 100(1):157–168, 2000.

[2] Joseph V Bonventre. Dedifferentiation and proliferation of surviving epithelial cells in acute renal failure. *Journal of the American Society of Nephrology*, 14(suppl 1):S55–S61, 2003.

[3] Purushothama Rao Tata, Hongmei Mou, Ana Pardo-Saganta, Rui Zhao, Mythili Prabhu, Brandon M Law, Vladimir Vinarsky, Josalyn L Cho, Sylvie Breton, Amar Sahay, et al.

Dedifferentiation of committed epithelial cells into stem cells in vivo. *Nature*, 503(7475): 218–223, 2013.

[4] Yuan Liu and Ye-Guang Chen. Intestinal epithelial plasticity and regeneration via cell dedifferentiation. *Cell Regeneration*, 9(1):1–11, 2020.

[5] Yuan Liu, Xiaochen Xiong, and Ye-Guang Chen. Dedifferentiation: the return road to repair the intestinal epithelium. *Cell Regeneration*, 9:1–3, 2020.

[6] Benoit Biteau, Christine E Hochmuth, and Heinrich Jasper. Maintaining tissue homeostasis: dynamic control of somatic stem cell activity. *Cell stem cell*, 9(5):402–411, 2011.

[7] Fiona M Watt and Brigid LM Hogan. Out of eden: stem cells and their niches. *Science*, 287 (5457):1427–1430, 2000.

[8] Joep Beumer and Hans Clevers. Regulation and plasticity of intestinal stem cells during homeostasis and regeneration. *Development*, 143(20):3639–3649, 2016.

[9] Helmuth Gehart and Hans Clevers. Tales from the crypt: new insights into intestinal stem cells. *Nature Reviews Gastroenterology & Hepatology*, 16(1):19–34, 2019.

[10] Laurens G Van Der Flier and Hans Clevers. Stem cells, self-renewal, and differentiation in the intestinal epithelium. *Annual review of physiology*, 71:241–260, 2009.

[11] Gijs R Van Den Brink, Sylvia A Bleuming, James CH Hardwick, Berber L Schepman, G Johan Offerhaus, Josbert J Keller, Corinne Nielsen, William Gaffield, Sander JH Van Deventer, Drucilla J Roberts, et al. Indian hedgehog is an antagonist of wnt signaling in colonic epithelial cell differentiation. *Nature genetics*, 36(3):277–282, 2004.

[12] Blair B Madison, Katherine Braunstein, Erlene Kuizon, Kathleen Portman, Xiaotan T Qiao, and Deborah L Gumucio. Epithelial hedgehog signals pattern the intestinal crypt-villus axis. *Development*, 132(2):279–289, 2005.

[13] Lorene E Batts, D Brent Polk, Raymond N Dubois, and Holger Kulessa. Bmp signaling is required for intestinal growth and morphogenesis. *Developmental dynamics: an official publication of the American Association of Anatomists*, 235(6):1563–1570, 2006.

[14] Willemijn A Van Dop, Jarom Heijmans, Nikè VJA Büller, Susanne A Snoek, Sanne L Rosekrans, Elisabeth A Wassenberg, Marius A van den Bergh Weerman, Beate Lanske, Alan R Clarke, Douglas J Winton, et al. Loss of indian hedgehog activates multiple aspects of a wound healing response in the mouse intestine. *Gastroenterology*, 139(5):1665–1676, 2010.

[15] Nikè VJA Büller, Sanne L Rosekrans, Jessica Westerlund, and Gijs R van den Brink. Hedgehog signaling and maintenance of homeostasis in the intestinal epithelium. *Physiology*, 27(3): 148–155, 2012.

[16] Ignacio A Rodriguez-Brenes, Natalia L Komarova, and Dominik Wodarz. Evolutionary dynamics of feedback escape and the development of stem-cell–driven cancers. *Proceedings of the National Academy of Sciences*, 108(47):18983–18988, 2011.

[17] Ignacio A Rodriguez-Brenes, Dominik Wodarz, and Natalia L Komarova. Stem cell control, oscillations, and tissue regeneration in spatial and non-spatial models. *Frontiers in oncology*, 3: 82, 2013.

[18] Zheng Sun and Natalia L Komarova. Stochastic control of proliferation and differentiation in stem cell dynamics. *Journal of Mathematical Biology*, 71(4):883–901, 2015.

[19] Dominik Wodarz. Effect of cellular de-differentiation on the dynamics and evolution of tissue and tumor cells in mathematical models with feedback regulation. *Journal of theoretical biology*, 448:86–93, 2018.

[20] Ingo Roeder and Markus Loeffler. A novel dynamic model of hematopoietic stem cell organization based on the concept of within-tissue plasticity. *Experimental hematology*, 30(8):853–861, 2002.

[21] Ingmar Glauche, Michael Cross, Markus Loeffler, and Ingo Roeder. Lineage specification of hematopoietic stem cells: mathematical modeling and biological implications. *Stem cells*, 25 (7):1791–1799, 2007.

[22] I Glauche, M Horn, and I Roeder. Leukaemia stem cells: hit or miss? *British journal of cancer*, 96(4):677–678, 2007.

[23] Franziska Michor et al. Mathematical models of cancer stem cells. *Journal of Clinical Oncology*, 26(17):2854–2861, 2008.

[24] Anna Marciniak-Czochra, Thomas Stiehl, Anthony D Ho, Willi Jäger, and Wolfgang Wagner. Modeling of asymmetric cell division in hematopoietic stem cells—regulation of self-renewal is essential for efficient repopulation. *Stem cells and development*, 18(3):377–386, 2009.

[25] Arthur D Lander, Kimberly K Gokoffski, Frederic YM Wan, Qing Nie, and Anne L Calof. Cell lineages and the logic of proliferative control. *PLoS Biol*, 7(1):e1000015, 2009.

[26] Wing-Cheong Lo, Ching-Shan Chou, Kimberly K Gokoffski, Frederic Y-M Wan, Arthur D Lander, Anne L Calof, and Qing Nie. Feedback regulation in multistage cell lineages. *Mathematical*

biosciences and engineering: MBE, 6(1):59, 2009.

[27] Heiko Enderling, Mark AJ Chaplain, Alexander RA Anderson, and Jayant S Vaidya. A

mathematical model of breast cancer development, local treatment and recurrence. *Journal of*

Theoretical Biology, 246(2):245–259, 2007.

[28] Alexis J Carulli, Linda C Samuelson, and Santiago Schnell. Unraveling intestinal stem cell

behavior with models of crypt dynamics. *Integrative Biology*, 6(3):243–257, 2014.

[29] U Paulus, Christopher S Potten, and M Loeffler. A model of the control of cellular regeneration

in the intestinal crypt after perturbation based solely on local stem cell regulation. *Cell*

proliferation, 25(6):559–578, 1992.

[30] Shalev Itzkovitz, Irene C Blat, Tyler Jacks, Hans Clevers, and Alexander van Oudenaarden.

Optimality in the development of intestinal crypts. *Cell*, 148(3):608–619, 2012.

[31] Matthew D Johnston, Carina M Edwards, Walter F Bodmer, Philip K Maini, and S Jonathan

Chapman. Mathematical modeling of cell population dynamics in the colonic crypt and in

colorectal cancer. *Proceedings of the National Academy of Sciences*, 104(10):4008–4013, 2007.

[32] Pierre-François Verhulst. Notice sur la loi que la population suit dans son accroissement.

Corresp. Math. Phys., 10:113–126, 1838.

[33] Steven H Strogatz. *Nonlinear dynamics and chaos with student solutions manual: With*

applications to physics, biology, chemistry, and engineering. CRC press, 2018.

[34] Christopher S Potten. Radiation, the ideal cytotoxic agent for studying the cell biology of

tissues such as the small intestine. *Radiation research*, 161(2):123–136, 2004.

[35] Benjamin D Simons and Hans Clevers. Strategies for homeostatic stem cell self-renewal in

adult tissues. *Cell*, 145(6):851–862, 2011.

[36] Bo Yang, Lin Cao, Bin Liu, Colin D McCaig, and Jin Pu. The transition from proliferation to

differentiation in colorectal cancer is regulated by the calcium activated chloride channel a1.

PloS one, 8(4):e60861, 2013.

[37] T Muto, HJR Bussey, and BC Morson. The evolution of cancer of the colon and rectum.

Cancer, 36(6):2251–2270, 1975.

[38] Eric R Fearon and Bert Vogelstein. A genetic model for colorectal tumorigenesis. *cell*, 61(5):

759–767, 1990.

[39] Anna Marciniak-Czochra and Thomas Stiehl. Mathematical models of hematopoietic reconsti-

tution after stem cell transplantation. In *Model Based Parameter Estimation*, pages 191–206.

Springer, 2013.

[40] Ansu O Perekatt, Pooja P Shah, Shannon Cheung, Nidhi Jariwala, Alex Wu, Vishal Gandhi,
Namit Kumar, Qiang Feng, Neeket Patel, Lei Chen, et al. Smad4 suppresses wnt-driven
dedifferentiation and oncogenesis in the differentiated gut epithelium. *Cancer research*, 78(17):
4878–4890, 2018.

[41] Michiko Miyaki, Takeru Iijima, Motoko Konishi, Kimiyo Sakai, Aki Ishii, Masamichi Yasuno,
Tsunekazu Hishima, Morio Koike, Nobuyuki Shitara, Takeo Iwama, et al. Higher frequency of
smad4 gene mutation in human colorectal cancer with distant metastasis. *Oncogene*, 18(20):
3098–3103, 1999.

[42] Nicholas I Fleming, Robert N Jorissen, Dmitri Mouradov, Michael Christie, Anuratha Sakthi-
anandeswaren, Michelle Palmieri, Fiona Day, Shan Li, Cary Tsui, Lara Lipton, et al. Smad2,
smad3 and smad4 mutations in colorectal cancer. *Cancer research*, 73(2):725–735, 2013.

[43] Michitaka Nakano, Yoshikane Kikushige, Kohta Miyawaki, Yuya Kunisaki, Shinichi Mizuno,
Katsuto Takenaka, Shingo Tamura, Yuta Okumura, Mamoru Ito, Hiroshi Ariyama, et al.
Dedifferentiation process driven by tgf-beta signaling enhances stem cell properties in human
colorectal cancer. *Oncogene*, 38(6):780–793, 2019.

[44] MG Brattain, AE Levine, S Chakrabarty, LC Yeoman, JKV Willson, and B Long. Heterogeneity
of human colon carcinoma. *Cancer and Metastasis Reviews*, 3(3):177–191, 1984.

[45] David Buob, Harold Fauvel, Marie-Pierre Buisine, Stéphanie Truant, Christophe Mariette,
Nicole Porchet, Agnès Wacrenier, Marie-Christine Copin, and Emmanuelle Leteurtre. The
complex intratumoral heterogeneity of colon cancer highlighted by laser microdissection.
Digestive diseases and sciences, 57(5):1271–1280, 2012.

[46] Chelsie K Sievers, Alyssa A Leystra, Linda Clipson, William F Dove, and Richard B Halberg.
Understanding intratumoral heterogeneity: lessons from the analysis of at-risk tissue and
premalignant lesions in the colon. *Cancer prevention research*, 9(8):638–641, 2016.

[47] Muhammad Al-Hajj, Max S Wicha, Adalberto Benito-Hernandez, Sean J Morrison, and
Michael F Clarke. Prospective identification of tumorigenic breast cancer cells. *Proceedings of
the National Academy of Sciences*, 100(7):3983–3988, 2003.

[48] Dominique Bonnet and John E Dick. Human acute myeloid leukemia is organized as a hierarchy
that originates from a primitive hematopoietic cell. *Nature medicine*, 3(7):730–737, 1997.

[49] Winston R Becker, Stephanie A Nevins, Derek C Chen, Roxanne Chiu, Aaron Horning, Rozelle Laquindanum, Meredith Mills, Hassan Chaib, Uri Ladabaum, Teri Longacre, et al. Single-cell analyses reveal a continuum of cell state and composition changes in the malignant transformation of polyps to colorectal cancer. *bioRxiv*, 2021.

[50] Florian Uhlitz, Philip Bischoff, Stefan Peidli, Anja Sieber, Benedikt Obermayer, Eric Blanc, Alexandra Trinks, Mareen Luethen, Yana Ruchiy, Thomas Sell, et al. Mitogen-activated protein kinase activity drives cell trajectories in colorectal cancer. *bioRxiv*, pages 2020–01, 2021.