

A New Perspective on Tumor progression



Frédéric THOMAS



COMMENTARY

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EVOLUTION,
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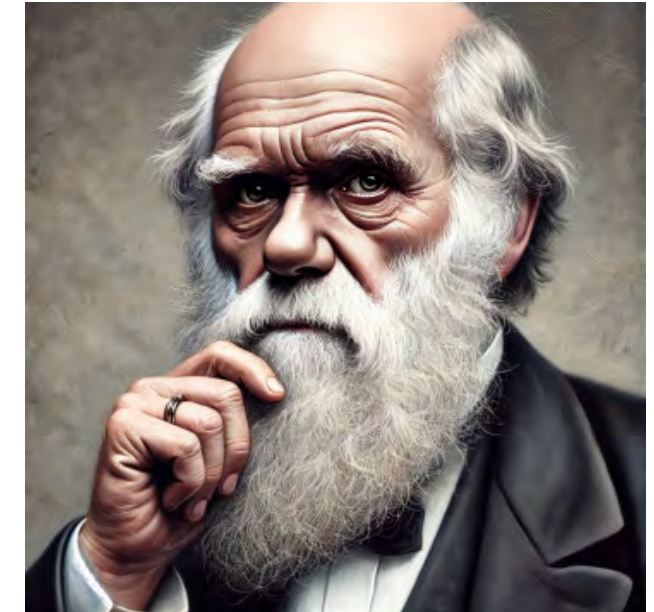
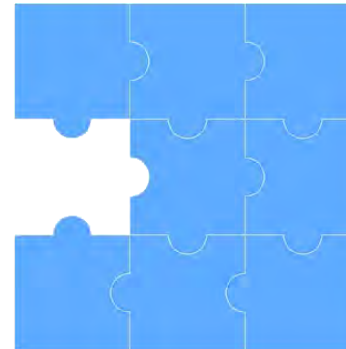
A new perspective on tumor progression

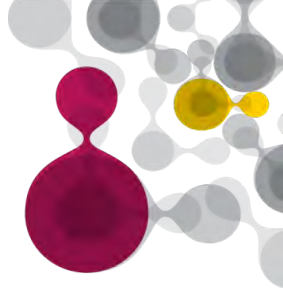
Evolution via selection for function

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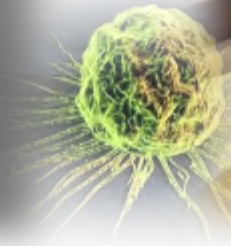
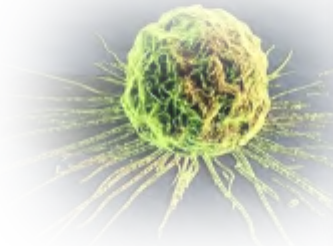
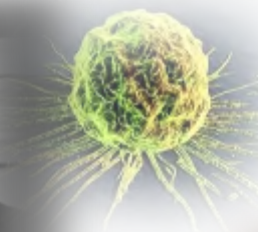
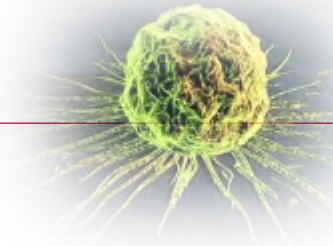
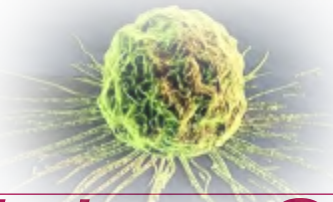
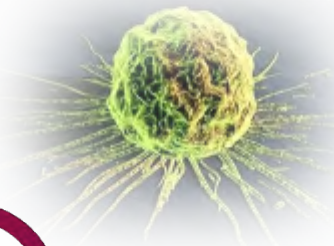


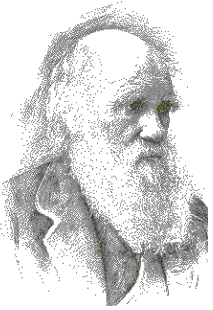


*“Nothing in Biology Makes Sense
Except in the Light of Evolution”*

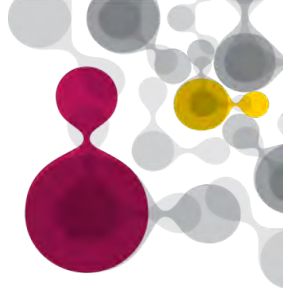
- Theodosius Dobzhansky (1973)

 **CANCER**





Cancer is a Disease of Clonal Evolution Within the Body



Nature Vol. 255 May 15 1975

review article

Mutation selection and the natural history of cancer

John Cairns*

Survival of the rapidly renewing tissues of long-lived animals like man is dependent on the ability of the cells to resist the natural selection of fitter variant cells (that is, the cells that are better adapted to the environment). This article discusses three possible protective mechanisms and some features of the natural history of certain common cancers of man.

We are accustomed to thinking of the combination of natural variation and natural selection as a force for the good, that creates and maintains the fittest in a species and discards the unfit. This is the fundamental theorem of biology. But when we turn from the competition between the individuals of a species to the competition between the individual cells within a single animal, we see that natural selection has now become a liability. The dangerous mutations are now those that confer on a cell an increased survival advantage. We may therefore expect to find, especially in animals which undergo continual cell multiplication during their adult life, the evolution of mechanisms that protect the animal from being taken over by any "fitter"

tumours of such sarcomas and leukaemias. This paper discusses the risk of cancer arising from somatic mutations and the features of natural carcinogenesis as a programme.

Nature 1975

1
Between
cells



Science 1976

The Clonal Evolution of Tumor Cell Populations

ability permits stepwise selection and underlies tumor progression.

Peter C. Nowell

Despite this wide recognition that most neoplasms have a unicellular origin and clonal growth pattern, relatively little emphasis has been placed on the developmental evolution of tumor cell populations, and the apparent genetic instability underlying the sequential acquisition of biological characteristics that we associate with tumor progression. This article suggests a model for the evolution of tumor cell populations in terms of stepwise genetic variation, and considers some of the evidence that this model is a valid one for most mammalian neoplasms. Some of the theoretical and practical implications of this concept of

variants are eliminated, because of metabolic disadvantage or immunologic destruction (for example, T₁), but occasionally one has an additional selective advantage with respect to the original tumor cells as well as normal cells, and this mutant becomes the precursor of a new predominant subpopulation.

Over time, there is sequential selection by an evolutionary process of sublines which are increasingly abnormal, both genetically and biologically. Because this sequence is not completely random, certain similarities are acquired by different tumors as they progress; but divergence also occurs as local conditions in each neoplasm differently effect the emergence of variant sublines. Ultimately, the fully developed malignancy as it appears histologically has a unique, aneuploid karyotype associated with aberrant morphology, and specific antigens that are also associated with the capability of metastasis. The relative positions in the evolutionary sequence of solid tumors, both in terms of their origin, as well as certain histological characteristics, are indicated in Fig. 1, along with the biological characteristics associated with the various stages of neoplastic development.

In the following sections, particular aspects of the model are considered in more detail, with major emphasis on evi-

interest. That neoplasms frequently develop as a clone from a single cell of origin is a concept gaining increased acceptance, and various investigators, beginning with chromosome studies on trans-



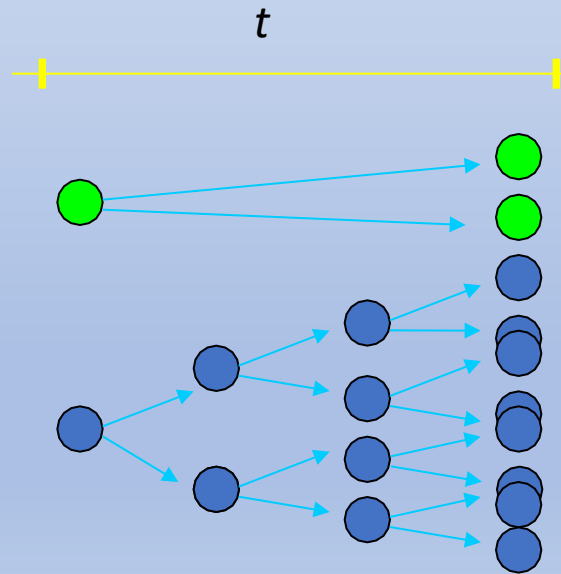
Natural selection

- Natural selection occurs in neoplasms because (epi) genetic mutations generate heritable variation, and some mutations confer a selective advantage or disadvantage on the cell

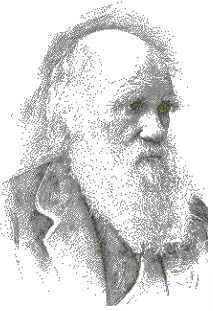
- Resources are limited



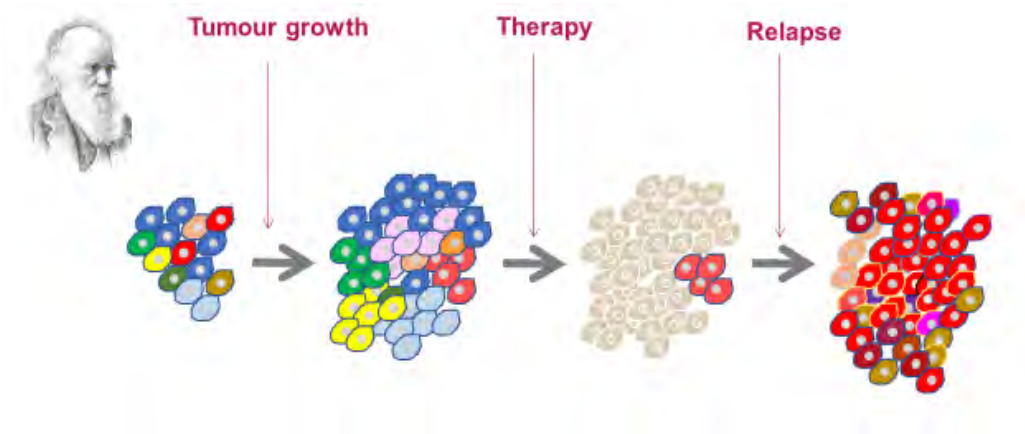
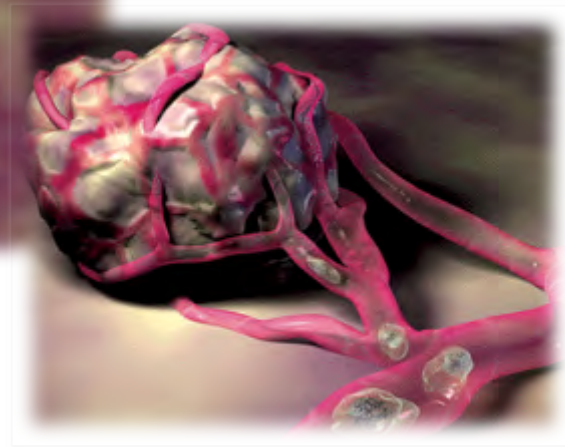
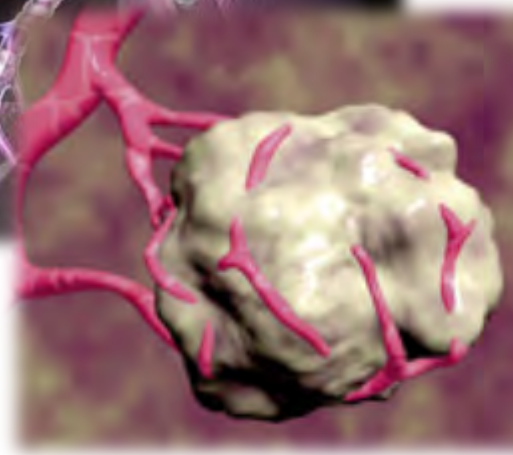
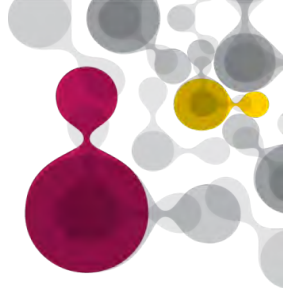
Differential reproductive success between clones

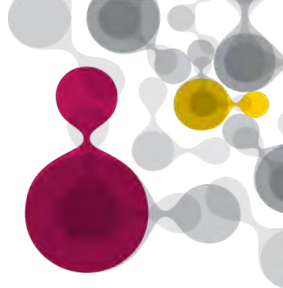


Best adapted survivors emerge...



Natural Selection / Oncogenic Selection



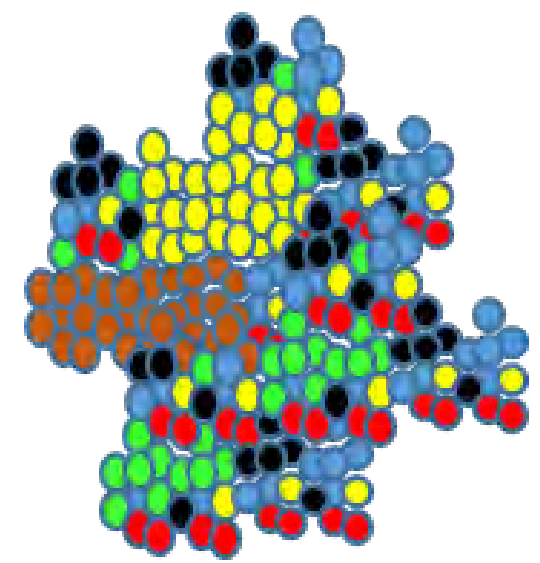
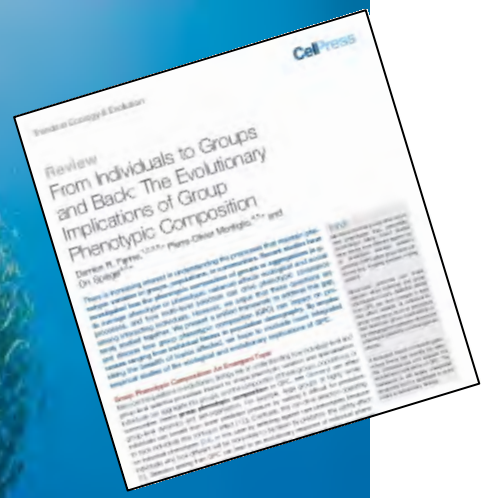
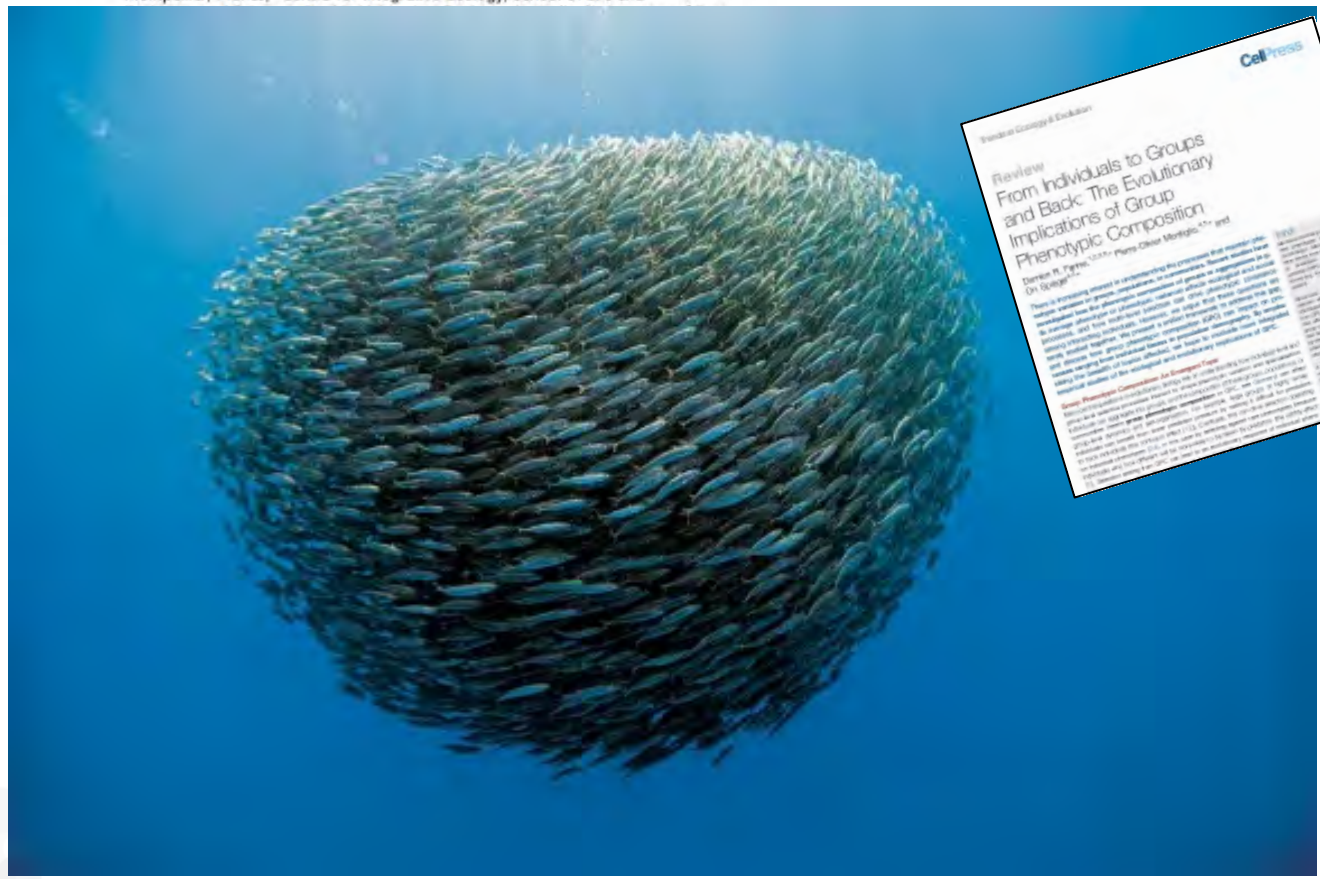


Group phenotypic composition in cancer

Jean-Pascal Capp^{1†}, James DeGregori^{2†}, Aurora M Nedelcu^{2†}, Antoine M Dujon^{4,5}, Justine Boutry⁴, Pascal Pujol⁴, Catherine Alix-Panabières^{4,*}, Rodrigo Hamede⁷, Benjamin Roche⁴, Beata Ujvari^{1,3†}, Andriy Marusyk⁸, Robert Gatenby^{9†}, Frédéric Thomas^{4†*}

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Cells but **clusters of cells** too...



Group Phenotypic Composition
GPC

An appropriate spatio-functional distribution of intra-tumoral heterogeneity conducive to tumor progression

Tumors capable of progressing are those that possess a suitable group phenotypic composition (GPC)

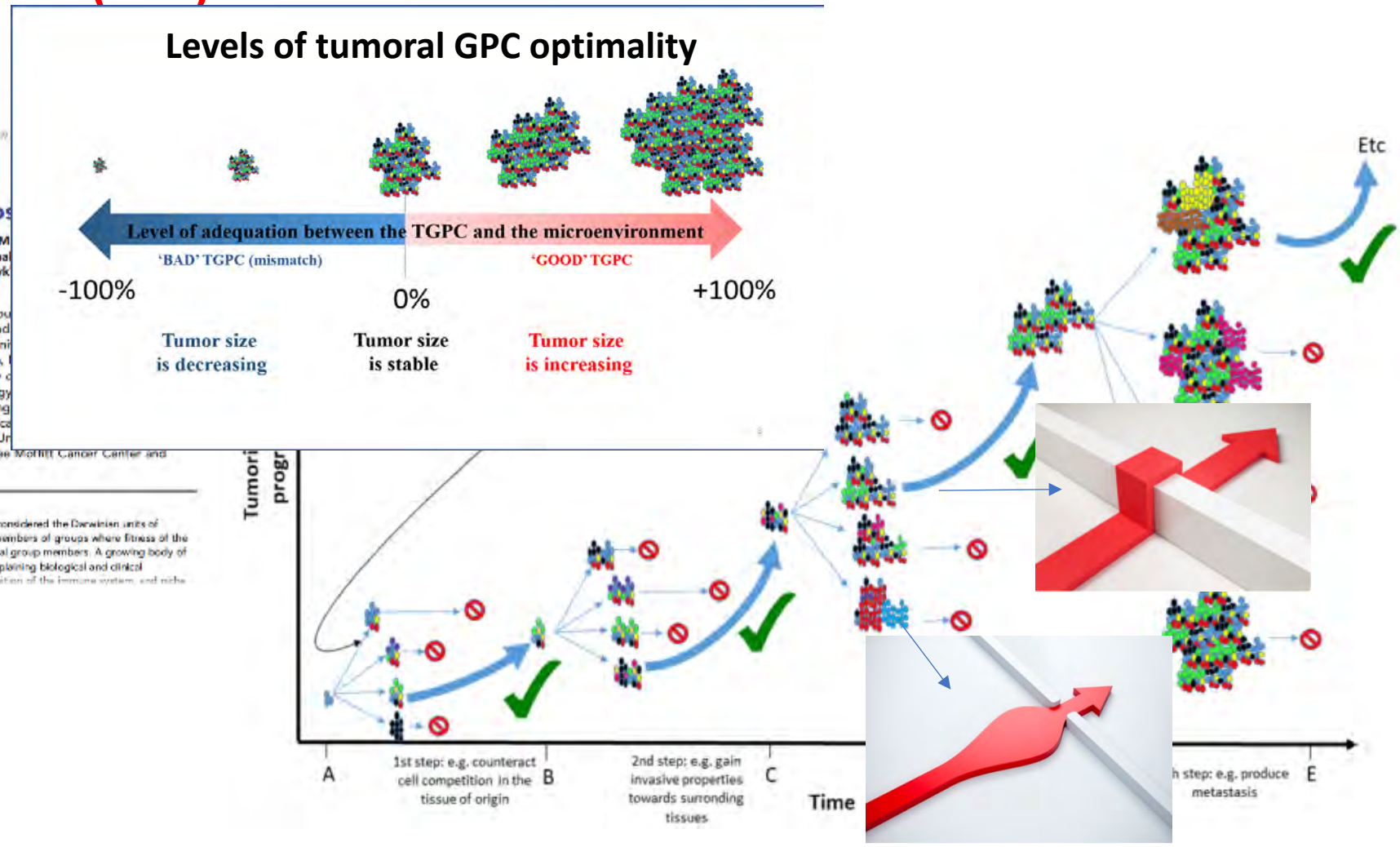


Group phenotypic composition

Jean-Pascal Capp^{1*}, James DeGregori^{2†}, Aurora M. Justine Boutry³, Pascal Pujol⁴, Catherine Alix-Panati⁵, Benjamin Roche⁶, Beata Ujvari^{6,7*}, Andriy Marusyk⁸, Frédéric Thomas^{1†*}

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Abstract Although individual cancer cells are generally considered the Darwinian units of selection in malignant populations, they frequently act as members of groups where fitness of the group cannot be reduced to the average fitness of individual group members. A growing body of studies revisits limitations of reductionist approaches to explaining biological and clinical observations. For example, induction of angiogenesis, inhibition of the immune system, and niche

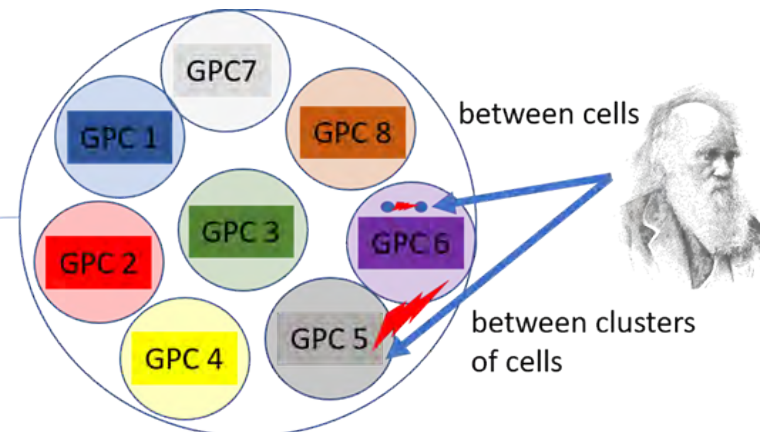


So it is widely accepted that tumorigenesis is a process that can be explained in its entirety by Darwinian processes **essentially based on interactions between cells and groups of cells.**



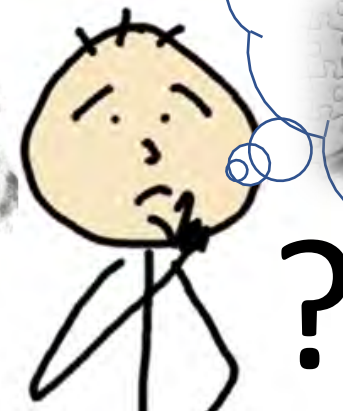
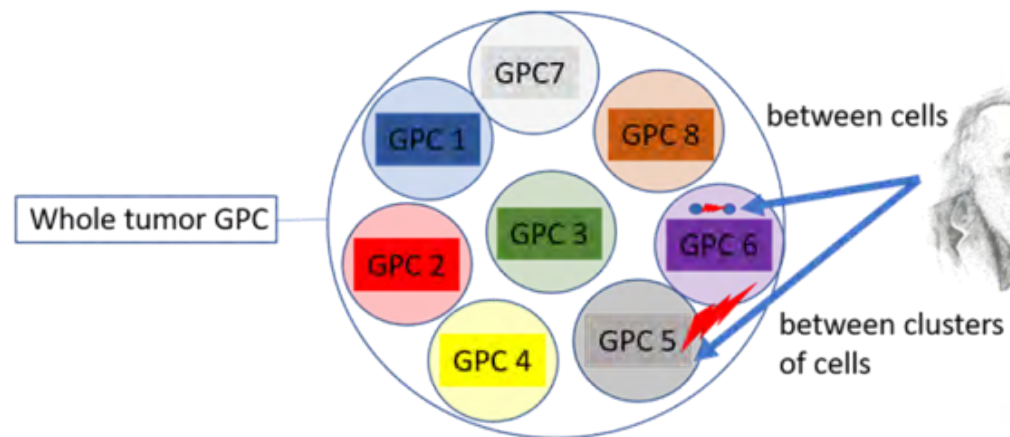
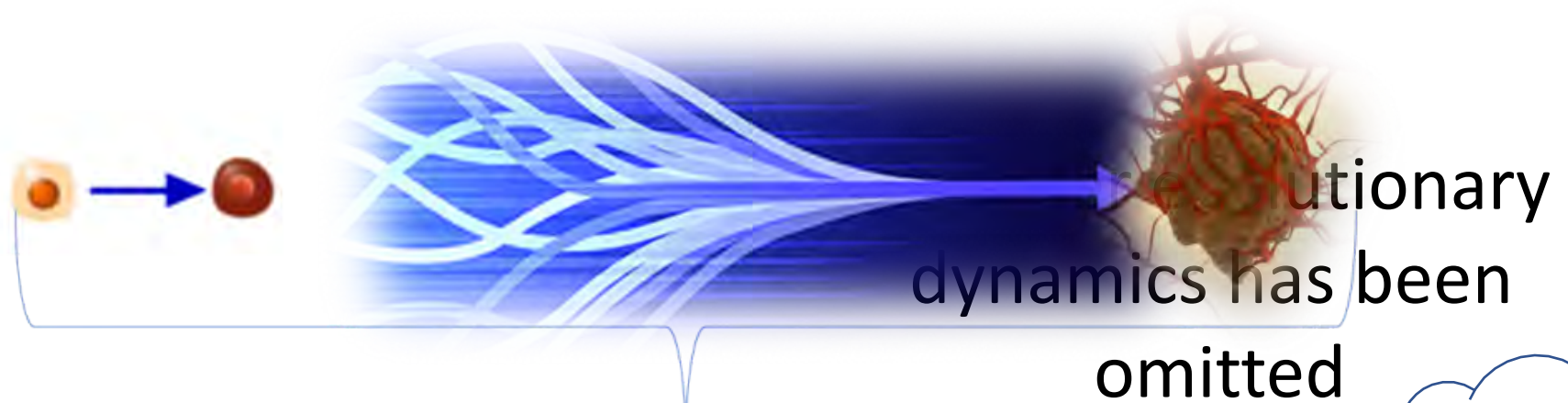
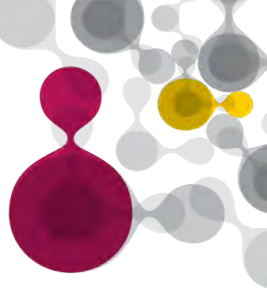
1
Between cells

Whole tumor GPC

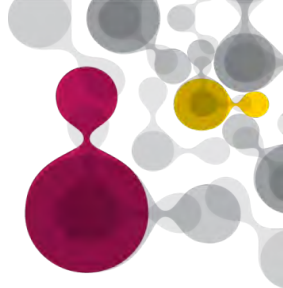


2
Between cells + Clusters of cells

The aim today is to challenge this point of view, by stating that...



Law of increasing functional information



La loi de l'augmentation de l'information fonctionnelle

PNAS RESEARCH ARTICLE BIOPHYSICS AND COMPUTATIONAL BIOLOGY EVOLUTION OPEN ACCESS Check for updates

On the roles of function and selection in evolving systems

Michael L. Wong^{1,2}, Carol E. Cleland¹, Daniel Arend Jr.³, Stuart Bartlett⁴, H. James Cleaves II^{5,6*}, Heather Demarest⁷, Anirudh Prabhu⁸, Jonathan I. Lunine^{2,1}, and Robert M. Hazen⁴

Contributed by Jonathan I. Lunine, received July 8, 2023; accepted September 10, 2023; reviewed by David Deamer, Andrea Roli, and Corday Seldon

Physical laws—such as the laws of motion, gravity, electromagnetism, and thermodynamics—codify the general behavior of varied macroscopic natural systems across space and time. We propose that an additional, hitherto-unarticulated law is required to characterize familiar macroscopic phenomena of our complex, evolving universe. An important feature of the classical laws of physics is the conceptual equivalence of specific characteristics shared by an extensive, seemingly diverse body of natural phenomena. Identifying potential equivalencies among disparate phenomena—for example, falling apples and orbiting moons or hot objects and compressed springs—has been instrumental in advancing the scientific understanding of our world through the articulation of laws of nature. A pervasive wonder of the natural world is the evolution of varied systems, including stars, minerals, atmospheres, and life. These evolving systems appear to be conceptually equivalent in that they display three notable attributes: 1) They form from numerous components that have the potential to adopt combinatorially vast numbers of different configurations; 2) processes exist that generate numerous different configurations; and 3) configurations are preferentially selected based on function. We identify universal concepts of selection—static persistence, dynamic persistence, and novelty generation—that underpin function and drive systems to evolve through the exchange of information between the environment and the system. Accordingly, we propose a “law of increasing functional information”: The functional information of a system will increase (i.e., the system will evolve) if many different configurations of the system undergo selection for one or more functions.

Significance
The universe is replete with complex evolving systems, but the existing macroscopic physical laws do not seem to adequately describe these systems. Recognizing that the identification of conceptual equivalencies among disparate phenomena were foundational to developing previous laws of nature, we approach a potential “missing law” by looking for equivalencies among evolving systems. We suggest that all evolving systems—including but not limited to life—are composed of diverse components that can combine into configurational states that are then selected for

selection | natural laws | evolving systems | functional information | Titan

Three types of functions according To Wong et al. 2024:


Static persistence

Dynamic persistence

Novelty generation



This law of increasing functional information, expanded Charles Darwin's theory of evolution by natural selection to include non-living systems.

Minerals consistently become more complex over time when subject to selection pressures

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nexus

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<https://doi.org/10.1093/pnasnexus/pgae248>
Advance access publication 25 June 2024
Research Report

Open-ended versus bounded evolution: Mineral evolution as a case study

Robert M. Hazen * and Michael L. Wong b

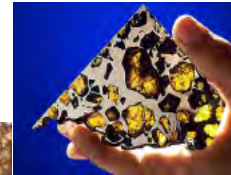
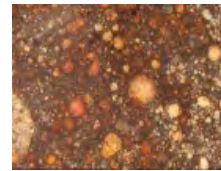
*Earth and Planets Laboratory, Carnegie Institution for Science, 5251 Broad Branch Road NW, Washington, DC 20015, USA
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*To whom correspondence should be addressed; Email: rhazen@carnegiescience.edu
Edited by: Harry McSweeney

Abstract

To what extent are naturally evolving systems limited in their potential diversity (i.e. “bounded”) versus unrestricted (“open-ended”)? Minerals provide a quantitative model, evolving system, with well-documented increases in mineral diversity through multiple stages of planetary evolution over billions of years. A recent framework that unifies behaviors of both biotic and abiotic evolving systems posits that all such systems are characterized by combinatorial richness subject to selection. In the case of minerals, combinatorial richness derives from the possible combinations of chemical elements coupled with permutations of their formulas’ coefficients. Observed mineral species, which are selected for persistence through deep time, represent a minuscule fraction of all possible element configurations. Furthermore, this model predicts that as planetary systems evolve, stable minerals become an ever-smaller fraction of the “possibility space.” A postulate is that “functional information,” defined as the negative log₂ of that fraction, must increase as a system evolves. We have tested this hypothesis for minerals by estimating the fraction of all possible chemical formulas observed from one stage of mineral evolution to the next, based on numbers of different essential elements and the maximum chemical formula complexity at each of nine chronological stages of mineral evolution. We find a monotonic increase in mineral functional information through these nine stages—a result consistent with the hypothesis. Furthermore, analysis of the chemical formulas of minerals demonstrates that the modern Earth may be approaching the maximum limit of functional information for natural mineral systems—a result demonstrating that mineral evolution is not open-ended.

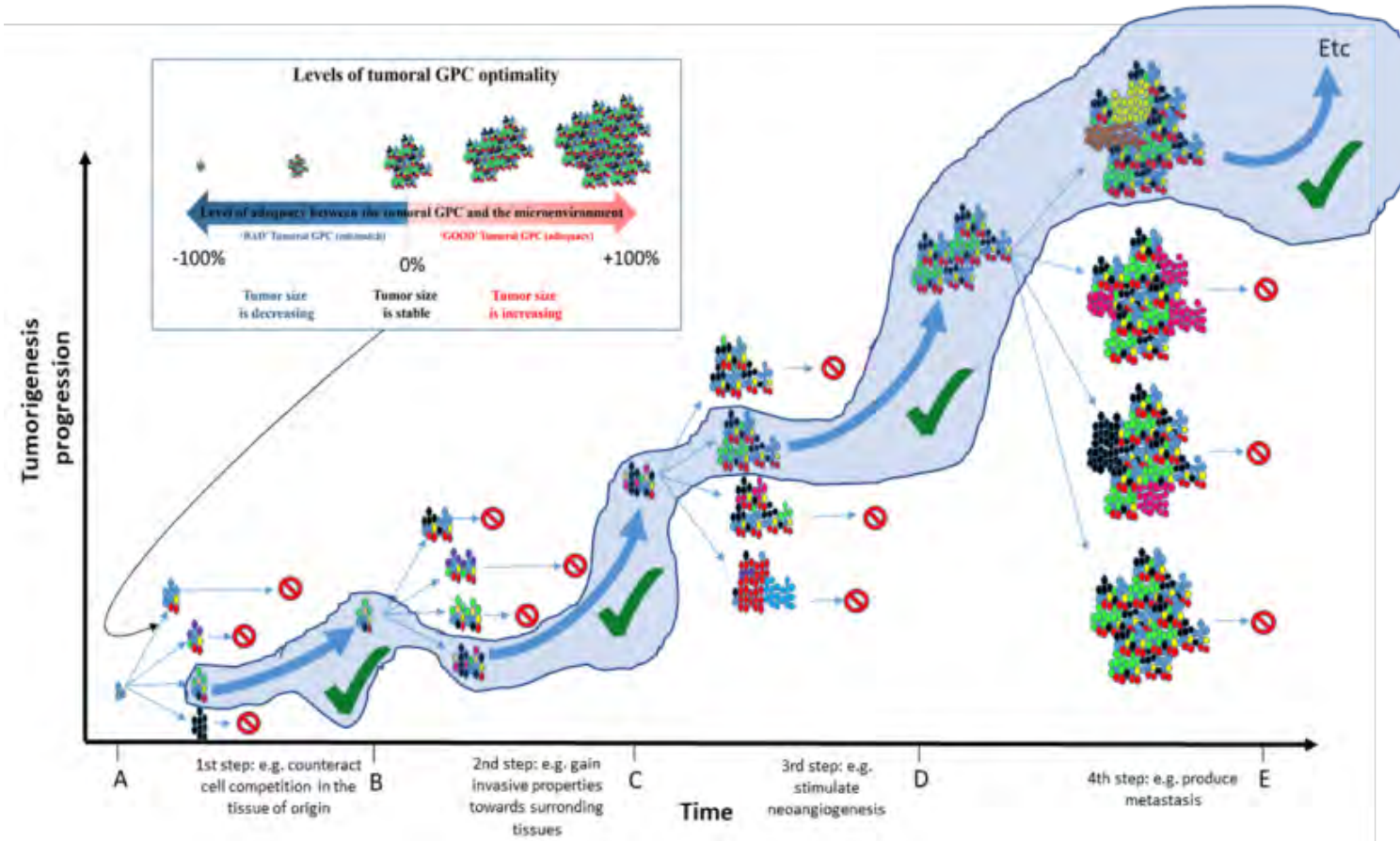
Keywords: evolution, open-ended evolution, complex systems, functional information, mineral evolution

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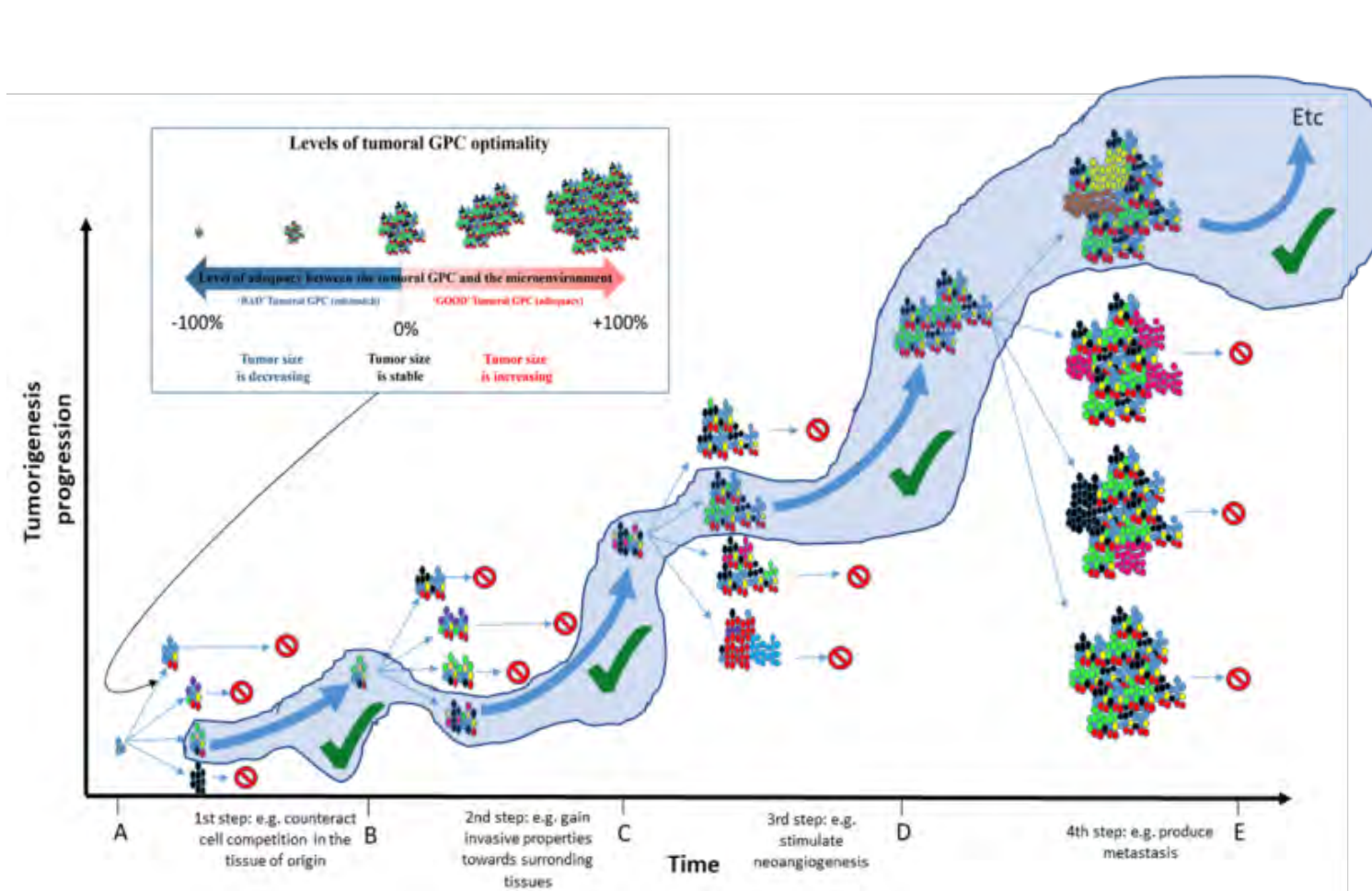


Over a period of more than 4.6 billion years, starting with the earliest known minerals from before Earth formed around 4.54 billion years ago and ending with all of the minerals on our planet today, **the number of mineral types increased from 27 to around 9,000.** This increase in Earth's mineral complexity also occurred at each mineral evolution stage — the first stage being the formation of the earliest minerals and the final stage being modern-day Earth, where mineral creation is facilitated by life.

The ability to develop a favorable GPC is thus subject to selection

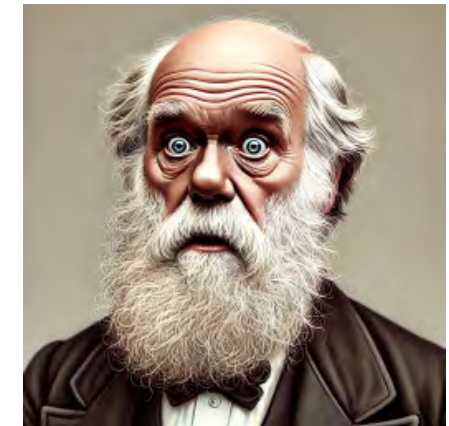


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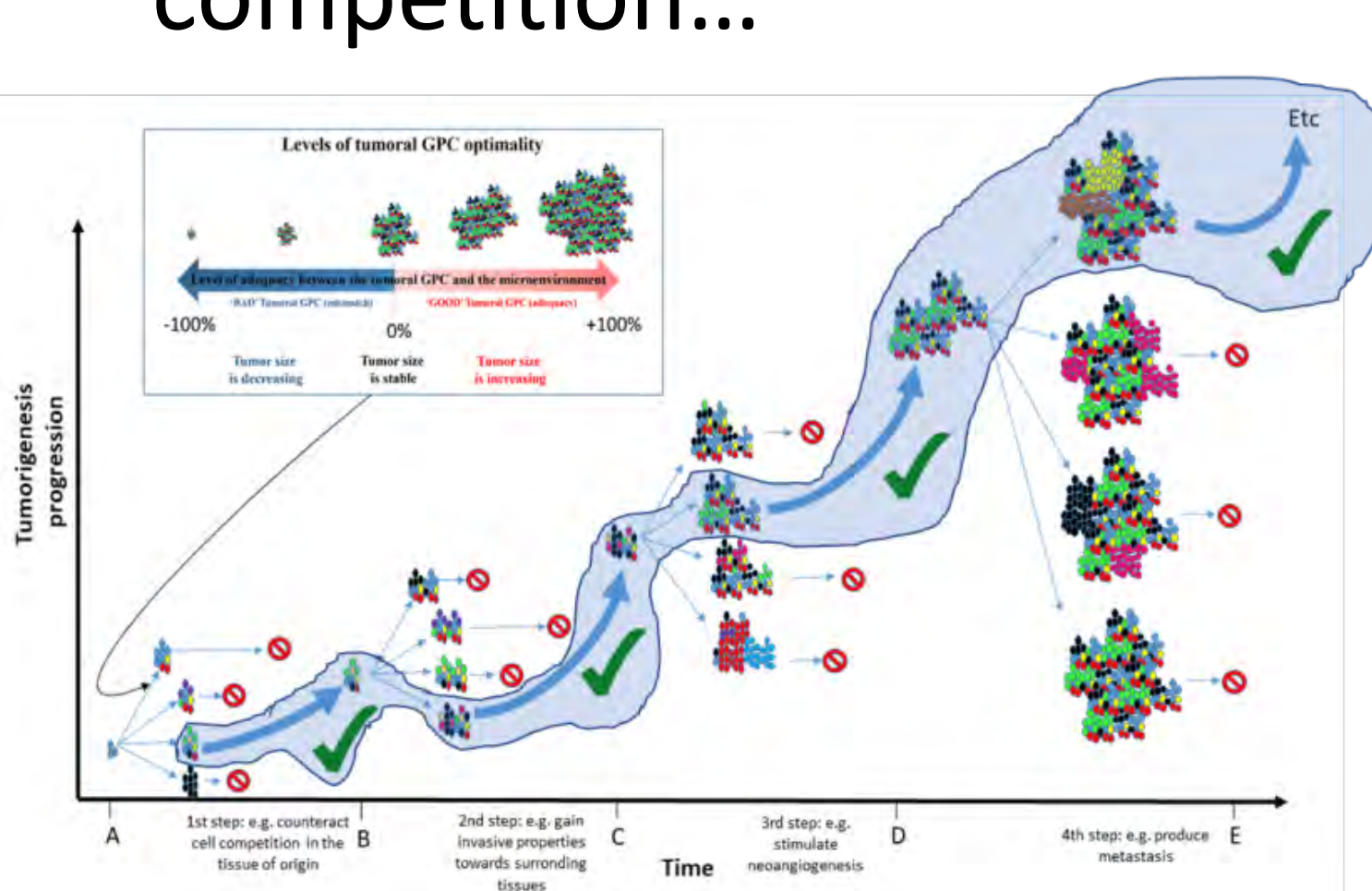


Selective process

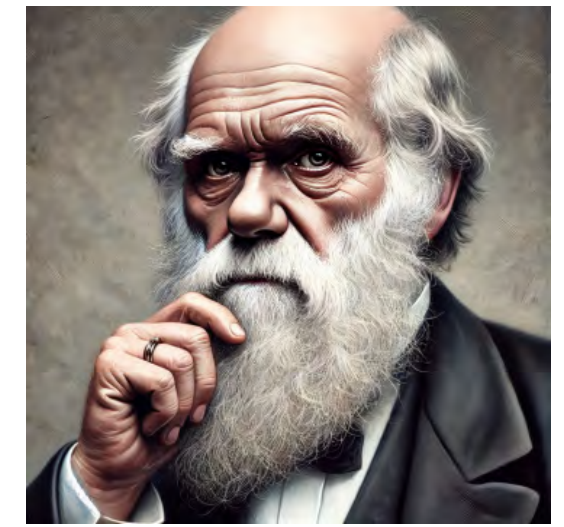
But NOT a classical Darwinian one !



There is no notion of heritability or reproductive success, etc. Tumors are not in competition...



Yet, they evolve...



Tumor functionality can include either of the three types of functions envisioned by Wong et al.: **static persistence**, **dynamic persistence**, and **novelty generation**.

PNAS RESEARCH ARTICLE BIOPHYSICS AND COMPUTATIONAL BIOLOGY EVOLUTION OPEN ACCESS

On the roles of function and selection in evolving systems

Michael L. Wong^{1,2}, Carol E. Oland^{1,2}, Daniel Arend Jr.¹, Stuart Bartlett¹, H. James Cleaves III^{1,3*}, Heather Demarest¹, Anirudh Prathu¹, Jonathan I. Lunine^{2,4}, and Robert M. Hazen¹

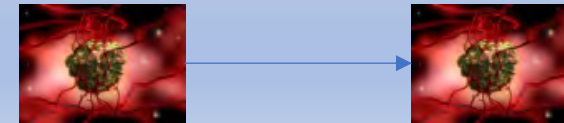
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Physical laws—such as the laws of motion, gravity, electromagnetism, and thermodynamics—codify the general behavior of varied macroscopic natural systems across space and time. We propose that an additional, hitherto-unarticulated law is required to characterize familiar macroscopic phenomena of our complex, evolving universe. An important feature of the classical laws of physics is the conceptual equivalence of specific characteristics shared by an extensive, seemingly diverse body of natural phenomena. Identifying potential equivalencies among disparate phenomena—for example, falling apples and orbiting moons or hot objects and compressed springs—has been instrumental in advancing the scientific understanding of our world through the articulation of laws of nature. A pervasive wonder of the natural world is the evolution of varied systems, including stars, minerals, atmospheres, and life. These evolving systems appear to be conceptually equivalent in that they display three notable attributes: 1) They form from numerous components that have the potential to adopt combinatorially vast numbers of different configurations; 2) processes exist that generate numerous different configurations; and 3) configurations are preferentially selected based on function. We identify universal concepts of selection—static persistence, dynamic persistence, and novelty generation—that underpin function and drive systems to evolve through the exchange of information between the environment and the system. Accordingly, we propose a “law of increasing functional information”: The functional information of a system will increase (i.e., the system will evolve) if many different configurations of the system undergo selection for one or more functions.

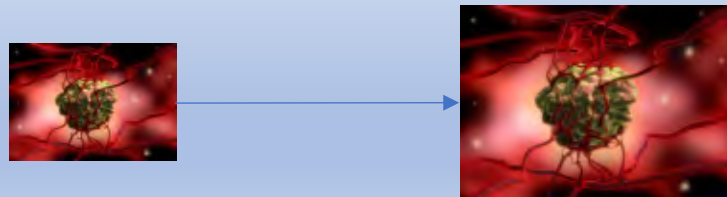
Significance
The universe is replete with complex evolving systems, but the existing macroscopic physical laws do not seem to adequately describe these systems. Recognizing that the identification of conceptual equivalencies among disparate phenomena were foundational to developing previous laws of nature, we approach a potential “missing law” by looking for equivalencies among evolving systems. We suggest that all evolving systems—including but not limited to life—are composed of diverse components that can combine into configurational states that are then selected for

selection | natural laws | evolving systems | functional information | Titan

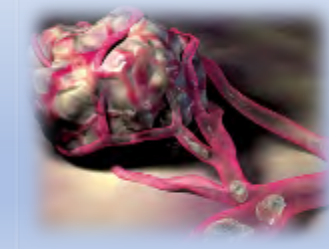
For instance, the ability of a tumor to maintain its size (through adaptively changing its GPC) in response to immune attacks and microenvironmental changes can be viewed as selection for “**static persistence**”



whereas its growth in response to various intra-tumour and microenvironmental changes (e.g., through increased plasticity, inducing angiogenesis) might involve selection for “**dynamic persistence**”



Similarly, the acquisition of the ability to invade and migrate (i.e., a novel capability) can be considered the result of selection for “**novelty generation**”.



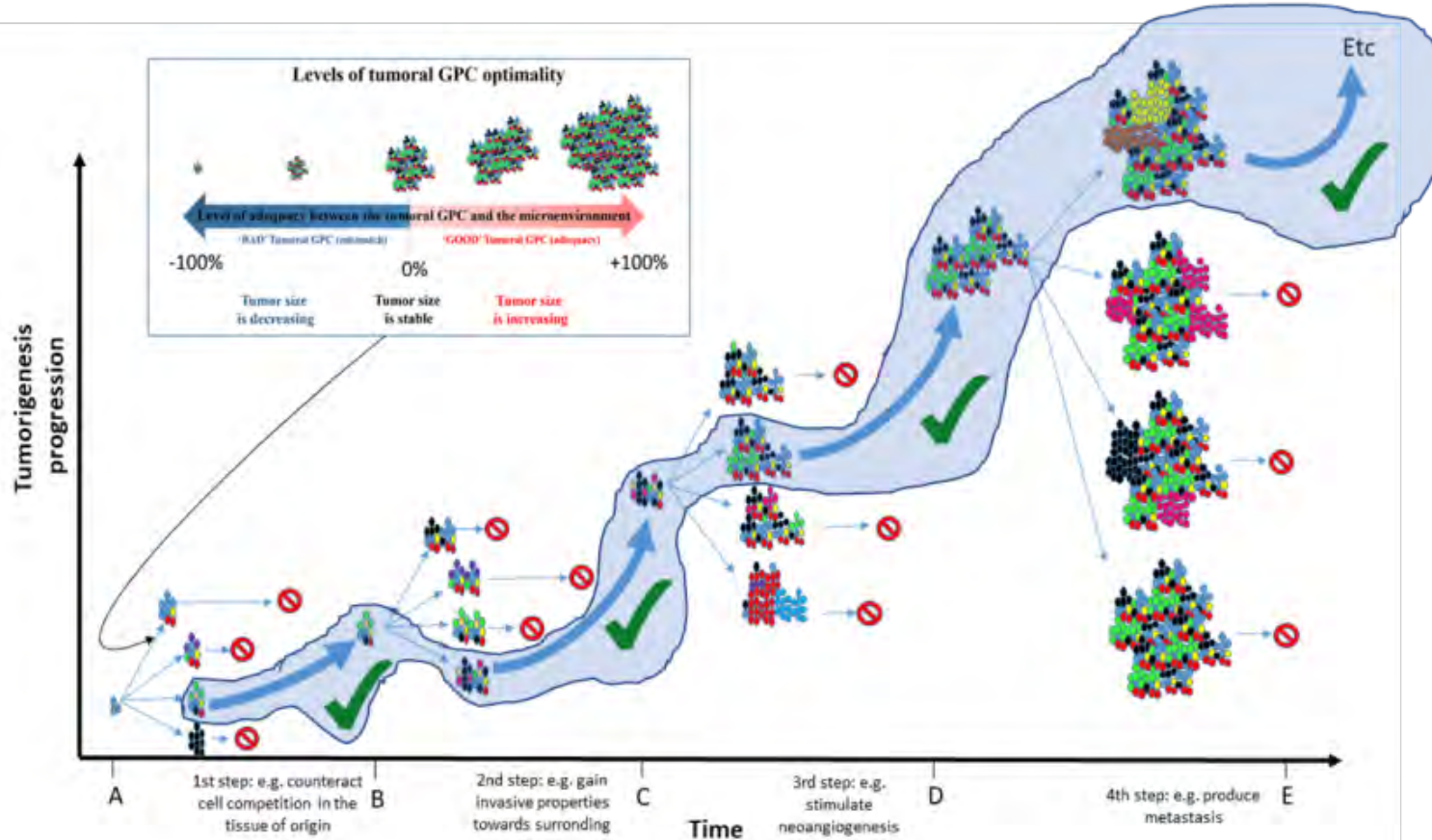
A new perspective on tumor progression

Evolution via selection for function

Frédéric Thomas^{1,2}, James DeGregori³, Andriy Marusyk⁴, Antoine M. Dujon^{1,4}, Beata Ujvari⁵, Jean-Pascal Cappi¹, Robert Gatenby⁶ and Aurora M. Nedelcu¹

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Tumors progress through selection based on function



PNAS RESEARCH ARTICLE PHYSICS AND COMPUTATIONAL BIOLOGY EVOLUTION

On the roles of function and selection in evolving systems

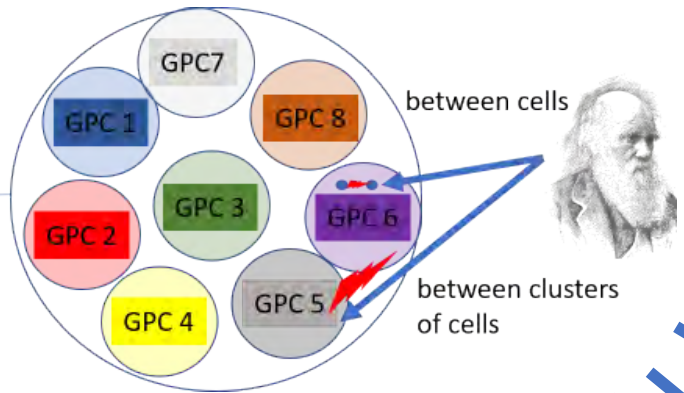
Mohamed Wong¹, Lily E. Baker², David A. Clark³, David R. Nelson⁴, James D. Watson⁵, Robert D. Levine⁶, and Robert M. May⁷

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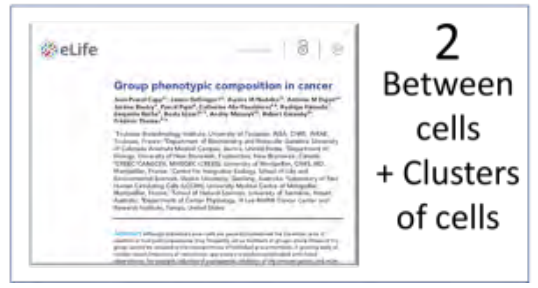
Physical laws—such as the laws of motion, gravity, electromagnetism, and thermodynamics—govern the general behavior of varied macroscopic natural systems across space and time. We propose that an additional, higher-order, articulated law is required to characterize familiar macroscopic phenomena of our complex, evolving universe. An important feature of the classical laws of physics is the conceptual equivalence of specific characteristics shared by an extensive, seemingly diverse body of natural phenomena. Identifying potential equivalencies among disparate phenomena—for example, falling apples and orbiting moons or ion objects and compressed springs—has been instrumental in advancing the scientific understanding of our world through the articulation of laws of nature. A pervasive wonder of the natural world is the evolution of varied systems, including stars, minerals, atmospheres, and life. These evolving systems appear to be conceptually equivalent in that they display three notable attributes: 1) They form from numerous configurations that have the potential to adopt combinatorially vast numbers of different configurations; 2) processes exist that generate numerous different configurations; and 3) configurations are preferentially selected based on function. We identify universal concepts of selection—static persistence, dynamic persistence, and novelty generation—that underpin function and drive systems to evolve through the exchange of information between the environment and the system. Accordingly, we propose a “law of increasing functional informativity”. The functional informativity of a system will increase (i.e., the system will evolve) if many different configurations of the system undergo selection for use or more function.

Significance
 The universe is replete with complex evolving systems, but the existing macroscopic physical laws do not seem to adequately describe these systems. Recognizing that the identification of conceptual equivalencies among disparate phenomena were foundational to developing previous laws of nature, we approach a potential “missing law” by looking for equivalencies among evolving systems. We suggest that all evolving systems—including but not limited to life—are composed of diverse components that can combine into configurational states that are then selected for.

Whole tumor GPC



1
Between cells

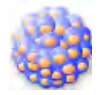
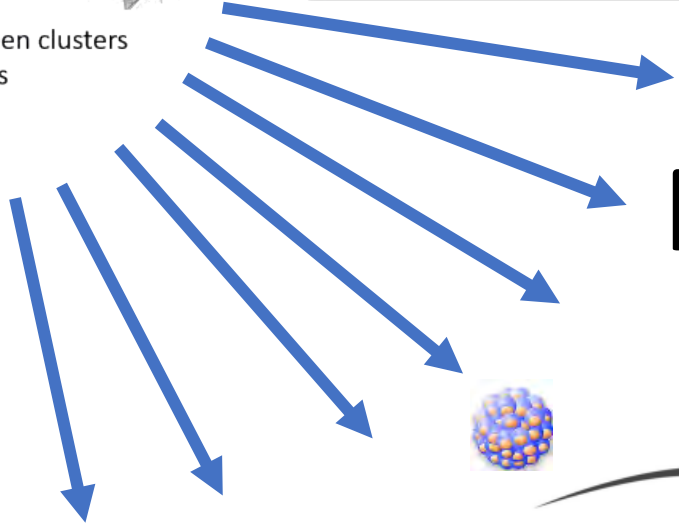


2
Between cells + Clusters of cells

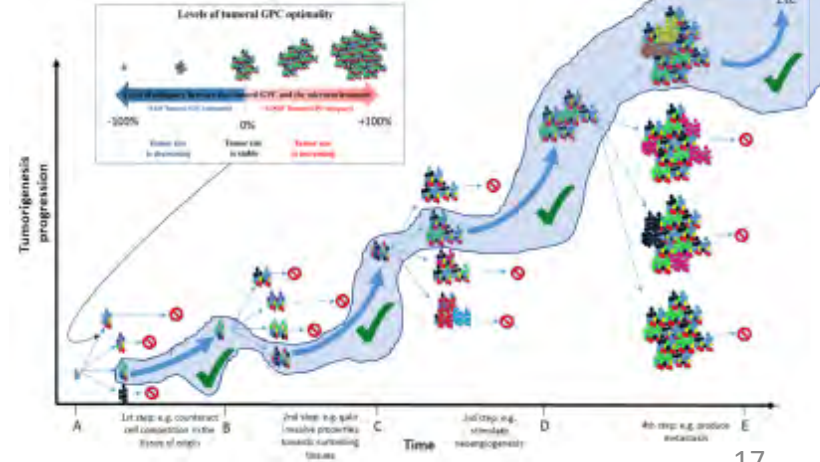
Myriads of GPCs



Most of them ...



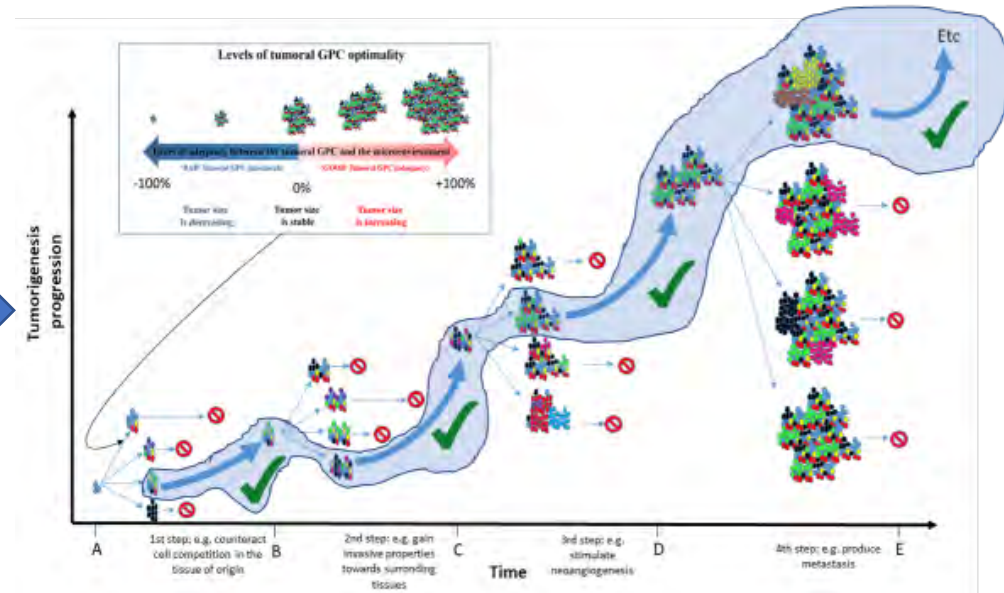
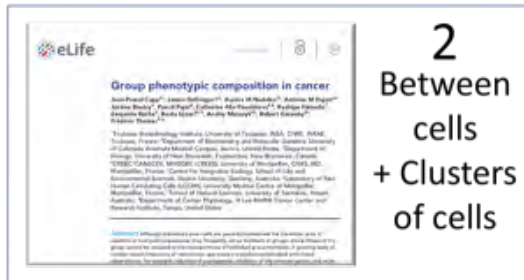
Selection on function



Darwinian processes between cells and groups of cells do not alone explain tumorigenesis; they merely feed into another form of selection, non-Darwinian, at the tumor level (selection based on function).



+



Selection on function

In other words, tumorigenesis is the result of **three distinct nested selective processes**

Life evolves. So do minerals. How about everything else?
 Proposed "natural law" broadening evolution finds support
 1 NOV 2024 · 11:20 AM ET · BY PAUL VOISEN

Tumorigenesis progression

1st s... e.g. co... cell cor... in the tissi...

Darwinian between cells and groups of cells

Between cells
 Between clusters of cells

Level of adequacy between the tumoral GPC and the microenvironment

-100% 'BAD' Tumoral GPC (mismatch) 0% 'GOOD' Tumoral GPC (adequacy) +100%

COMMENTARY

Evolution, Medicine, and Public Health [2024] pp.172-177
<https://doi.org/10.1093/emph/eeae021>
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EVOLUTION, MEDICINE, & PUBLIC HEALTH

A new perspective on tumor progression

Evolution via selection for function

Frédéric Thomas^{1,*,†}, James DeGregori², Andriy Marusyk³, Antoine M. Dujon^{4,†}, Beata Ujvari⁵, Jean-Pascal Capp⁶, Robert Gatenby⁷ and Aurora M. Nedelcu⁸

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ABSTRACT
 Tumorigenesis is commonly attributed to Darwinian processes involving natural selection among cells and groups of cells. However, progressing tumors are those that also achieve an appropriate group phenotypic composition (GPC). Yet, the selective processes acting on tumor GPCs are distinct from that associated with classical Darwinian evolution (i.e. natural selection based on differential reproductive success) as tumors are not genuine evolutionary individuals and do not exhibit heritable variation in fitness. This complex evolutionary scenario is analogous to the recently proposed concept of 'selection for function' invoked for the evolution of both living and non-living systems. Therefore, we argue that it is inaccurate to assert that Darwinian processes alone account for all the aspects characterizing tumorigenesis and cancer progression; rather, by producing the genetic and phenotypic diversity required for creating novel GPCs, these processes fuel the evolutionary success of tumors that is dependent on selection for function at the tumor level.

KEYWORDS: perspective; tumors; progression; evolution; selection; function; group phenotypic composition

THE PREMISE
 Following the pioneering work of Cairns [1] and Nowell [2], tumorigenesis has been generally viewed as underpinned by a classical Darwinian evolutionary process (i.e. somatic evolution) primarily governed by natural selection among mutant clones differing in fitness (i.e. survival and reproduction), starting from the emergence

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Exploiting the role of ‘selection for function’ in tumor progression to develop new therapeutic strategies against cancer

Frédéric THOMAS^{1*}, Jean-Pascal CAPP², Antoine M. DUJON^{1,3}, Andriy MARUSYK⁴, Mario CAMPONE⁵, Pascal PUJOL^{1,6}, Catherine ALIX-PANABIERES^{1,7,8}, Benjamin ROCHE¹, Beata UJVARI³, Robert GATENBY⁴ & Aurora M. NEDELCO^{9*}, *En preparation*

Targeting functional networks within a tumor's



Targeting functional networks involves identifying and disrupting specific pathways or cellular interactions that are essential for tumor growth and survival. For example, blocking molecular signals that allow cancer cells to communicate, cooperate with stromal cells, or adapt to environmental changes can inhibit tumor progression.



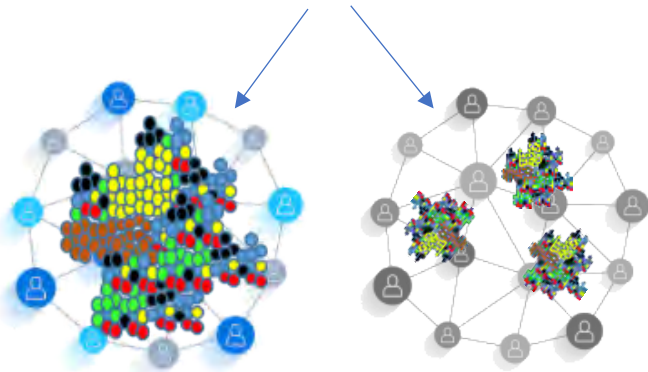
Traditional cancer treatments often aim to eliminate as many tumor cells as possible, focusing on targeting the physical form of individual cells.

GPC-based therapies



Therapy that does not aim to eradicate the maximum number of cells but rather focuses on destroying the interacting components of the functional network

The tumor does not disappear, but it takes time to regenerate the interacting components of a functional network, which may **(on the left)** or may not **(on the right)** yield an oncogenic GPC.



Tumor with an oncogenic GPC, including cellular and non-cellular components as well as their interaction network.



The tumor dies and disappears because the functional network has been too severely altered.

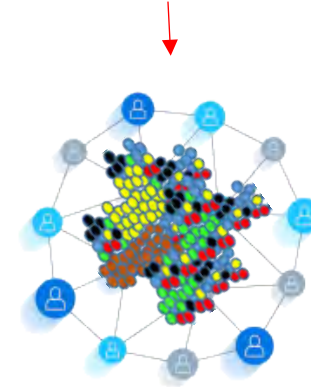


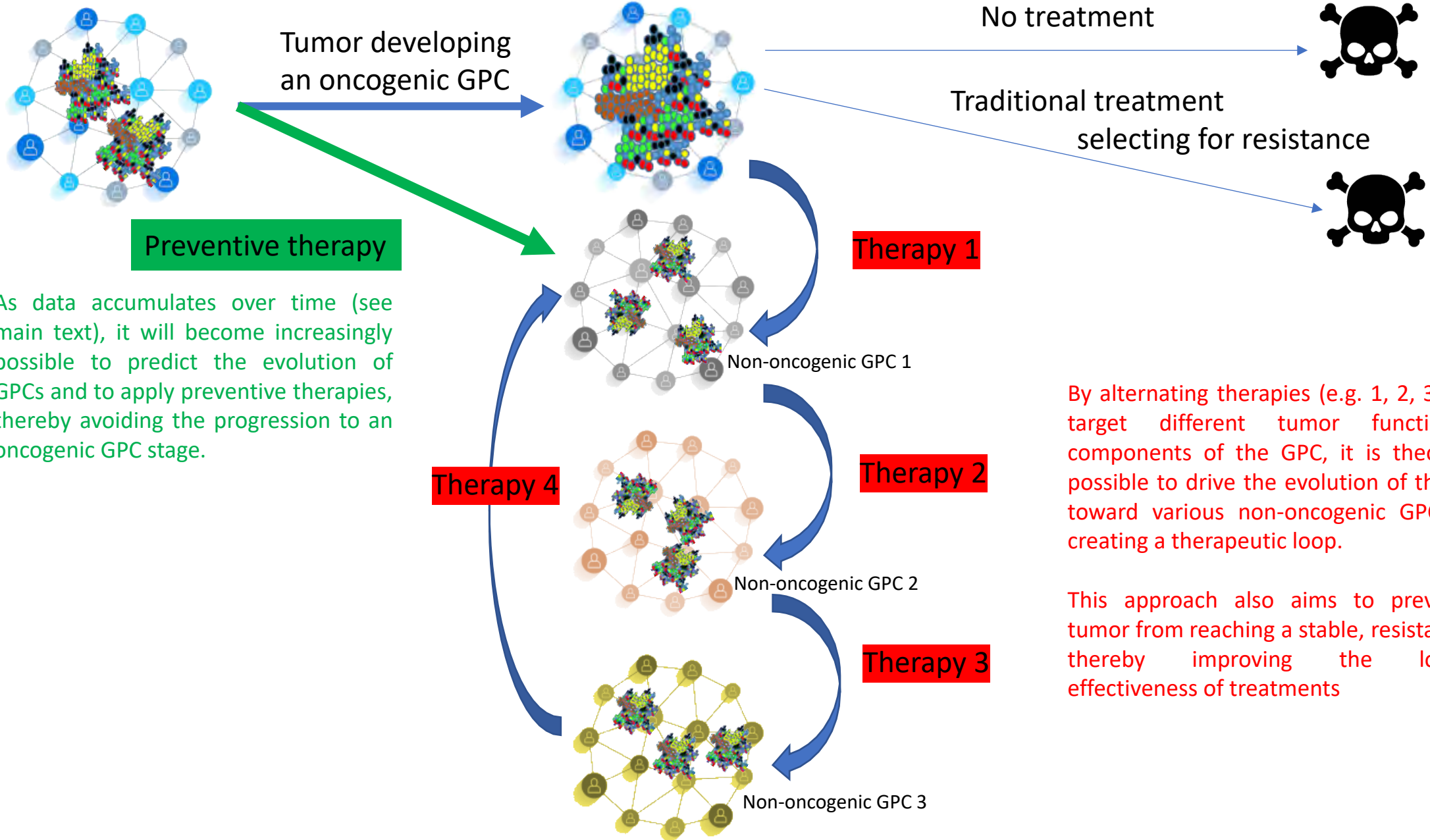
Traditional cancer treatments



Therapy that kills the maximum number of cells without necessarily targeting the interacting components of the functional network.

The tumor can restore an oncogenic GPC because the previous functional network remains active.





Tumor developing an oncogenic GPC

No treatment



Traditional treatment selecting for resistance



Preventive therapy

Therapy 1

Non-oncogenic GPC 1

Therapy 2

Non-oncogenic GPC 2

Therapy 3

Non-oncogenic GPC 3

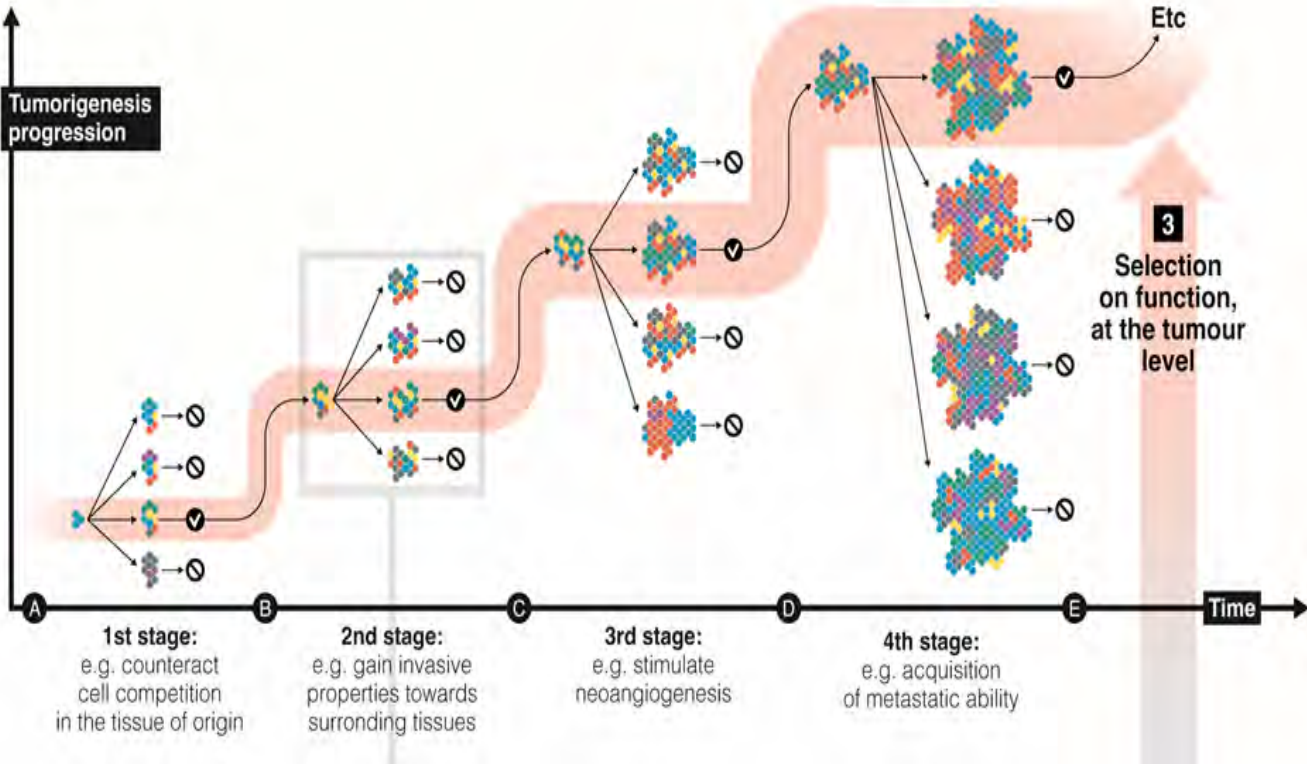
Therapy 4

As data accumulates over time (see main text), it will become increasingly possible to predict the evolution of GPCs and to apply preventive therapies, thereby avoiding the progression to an oncogenic GPC stage.

By alternating therapies (e.g. 1, 2, 3, 4) that target different tumor functions or components of the GPC, it is theoretically possible to drive the evolution of the tumor toward various non-oncogenic GPCs, even creating a therapeutic loop.

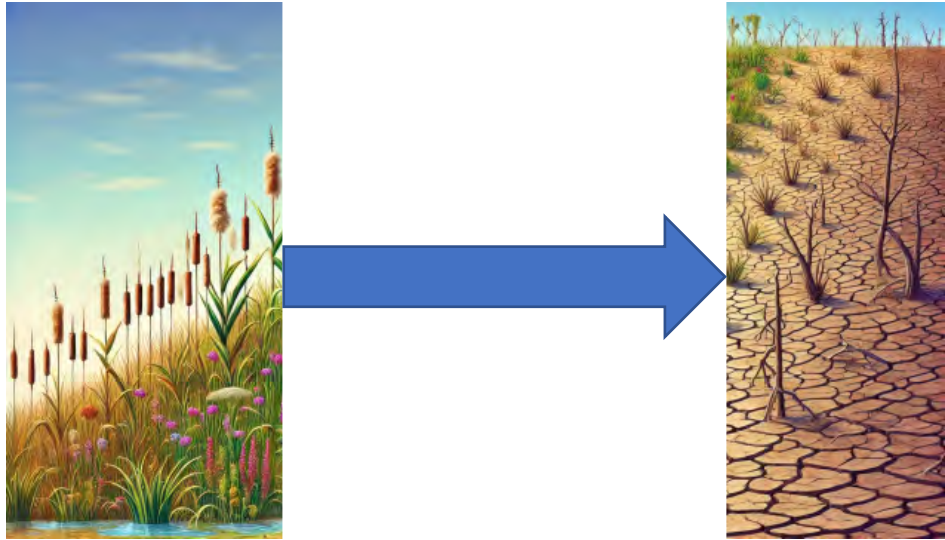
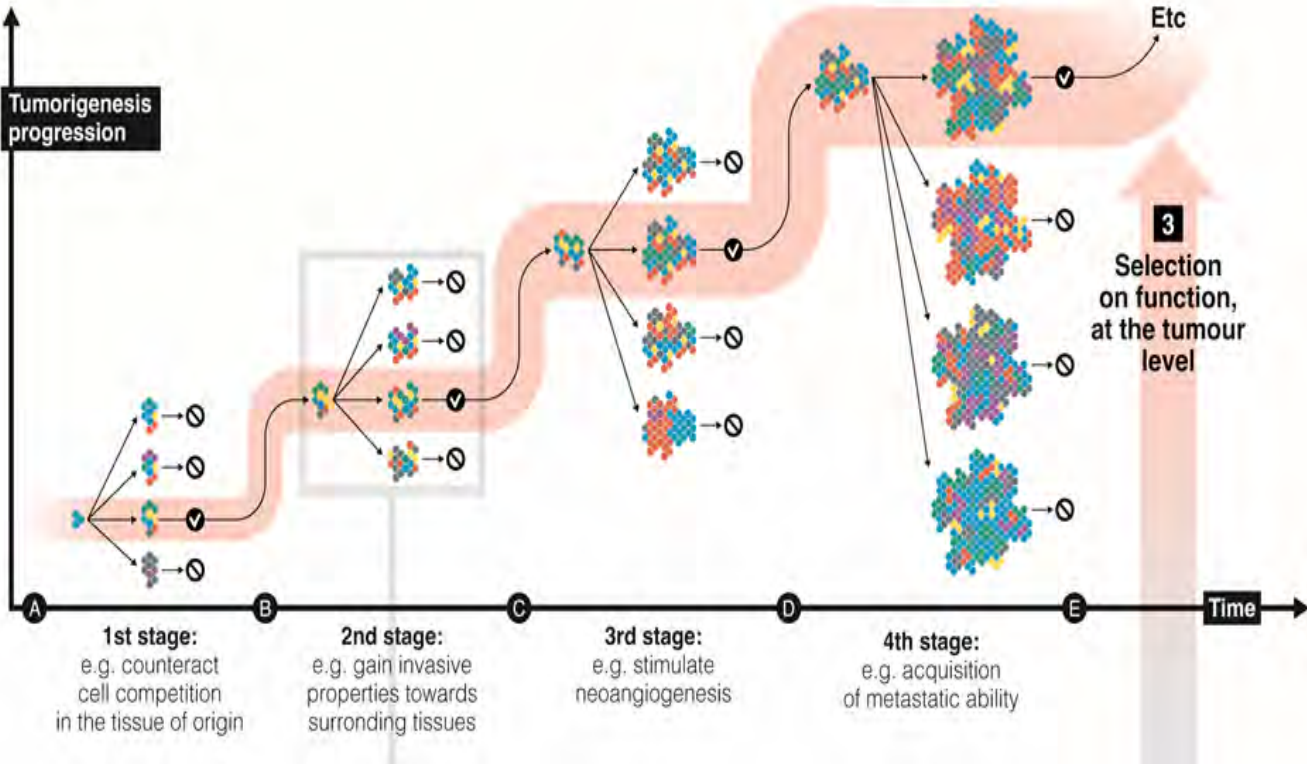
This approach also aims to prevent the tumor from reaching a stable, resistant state, thereby improving the long-term effectiveness of treatments

Modifying the microenvironment can also destabilize tumors by disrupting the GPC/microenvironment coupling responsible for tumor growth. In other words, therapies can be developed to **create a mismatch between the tumor GPC and its surrounding microenvironment**.



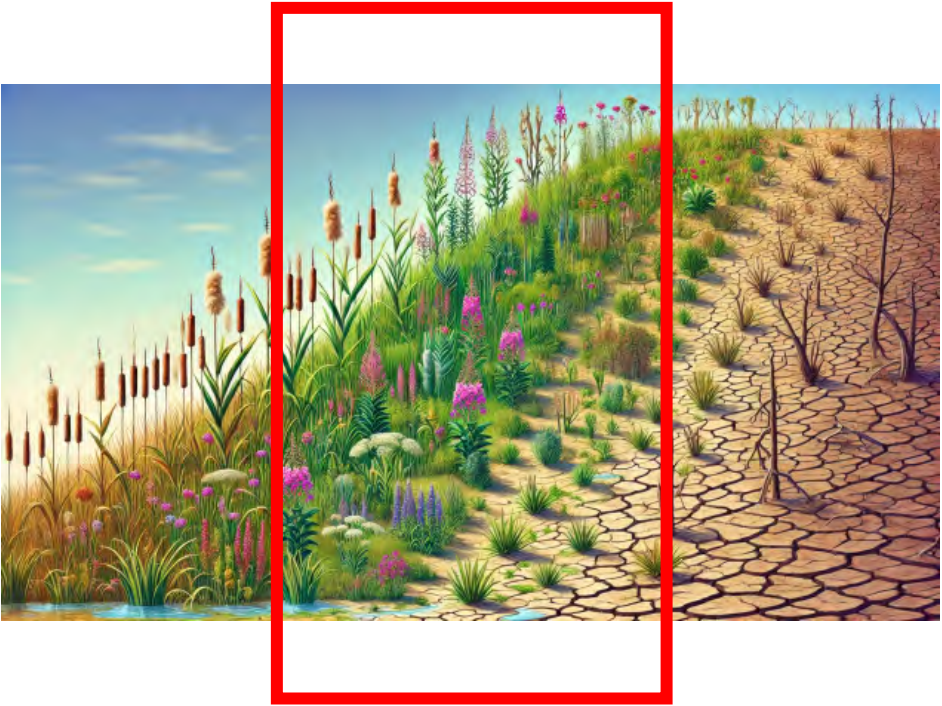
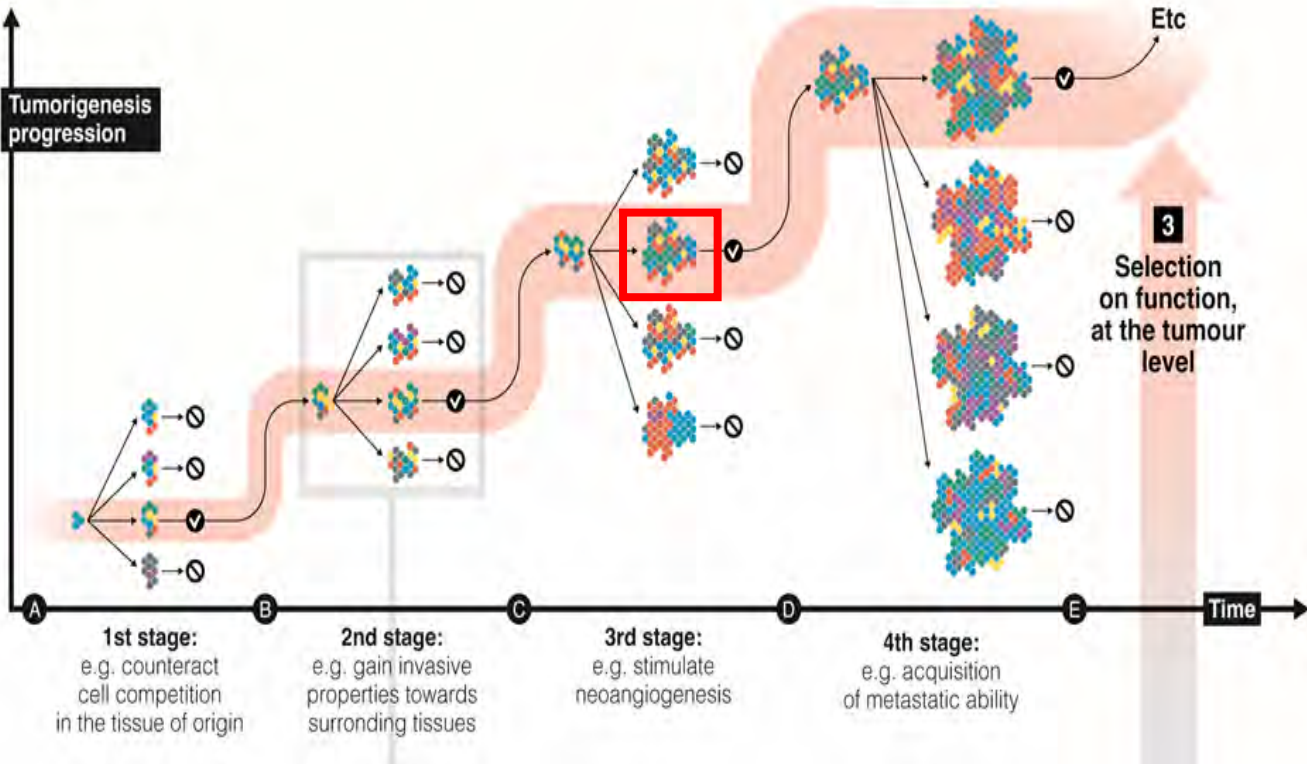
The tumor microenvironment, which includes immune cells, fibroblasts, blood vessels, and the extracellular matrix, plays a crucial role in forming and maintaining a tumor's GPC. Interactions between tumor cells and the microenvironment can either support or hinder tumor progression. Therefore, modifying the tumor microenvironment, such as inhibiting angiogenesis or enhancing the anti-tumor immune response, can potentially disrupt the oncogenic GPC and inhibit tumor growth.

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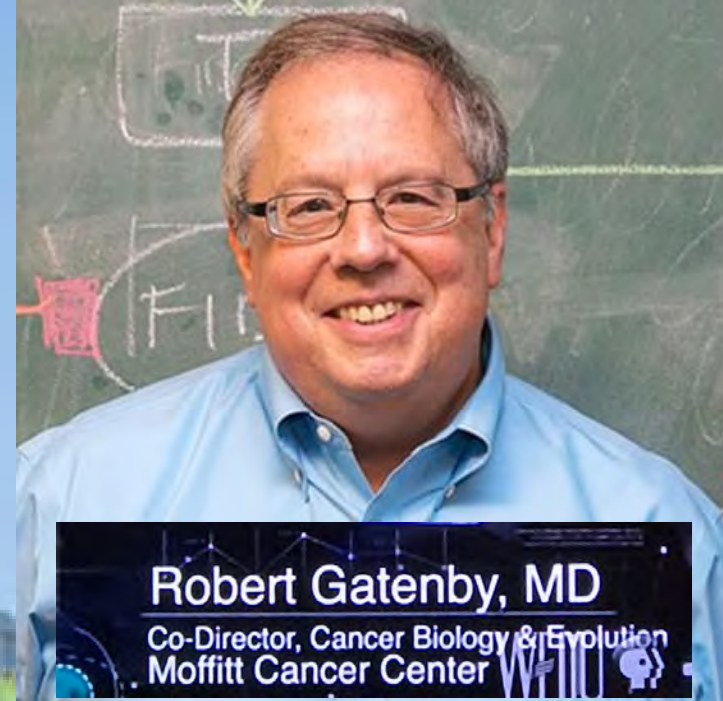
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Adaptive Therapy

Modifier la compétition entre cellules malignes agressives et cellules bénignes



A National Cancer Institute
Comprehensive Cancer Center



Cancer Res. 2009 June 1; 69(11): 4894–4903. doi:10.1158/0008-5472.CAN-08-3658.

Adaptive Therapy

Robert A. Gatenby¹, Ariosto S. Silva¹, Robert J. Gillies¹, and B. Roy Frieden²

¹Department of Integrative Mathematical Oncology, Moffitt Cancer Center, Tampa, Florida

²School of Optical Sciences, University of Arizona, Tucson, Arizona

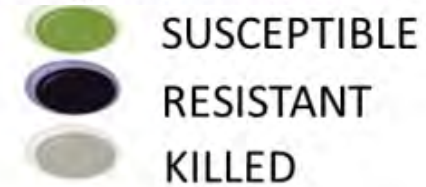
Abstract

A number of successful systemic therapies are available for treatment of disseminated cancers. However, tumor response is often transient, and therapy frequently fails due to emergence of resistant populations. The latter reflects the temporal and spatial heterogeneity of the tumor microenvironment as well as the evolutionary capacity of cancer phenotypes to adapt to therapeutic perturbations. Although cancers are highly dynamic systems, cancer therapy is typically administered according to a fixed, linear protocol. Here we examine an adaptive

Adaptive Therapy



A change of strategy in the war on cancer



nature COMMUNICATIONS

ARTICLE

DOI: 10.1038/n41467-017-01968-5 OPEN

Integrating evolutionary dynamics into treatment of metastatic castrate-resistant prostate cancer

Jingsong Zhang¹, Jessica J. Cunningham², Joel S. Brown^{2,3} & Robert A. Gatenby^{2,4}

Abiraterone treats metastatic castrate-resistant prostate cancer by inhibiting CYP17A, an enzyme for testosterone auto-production. With standard dosing, evolution of resistance with treatment failure (radiographic progression) occurs at a median of ~16.5 months. We hypothesize time to progression (TTP) could be increased by integrating evolutionary dynamics into therapy. We developed an evolutionary game theory model using



DOUBLE BIND

Owls facilitate the hunting success of snakes and vice versa



Published in final edited form as:
Mol Pharm. 2012 April 2; 9(4): 914-921. doi:10.1021/mp200458e.

Exploiting evolution to treat drug resistance: Combination therapy and the double bind

David Basanta¹, Robert A. Gatenby¹, and Alexander R. A. Anderson¹

¹Integrated Mathematical Oncology, H. Lee Moffitt Cancer Center and Research Institute, Tampa, FL 33612, USA

Abstract

Although many anticancer therapies are successful when initially administered, the evolutionary dynamics that often, resistance is an inevitable outcome. Research evolutionary double bind could be an effective way to bind two therapies are used in combination such that more susceptible to the other. In this paper we present framework of a double bind to study the effect that suggest a synergistic effect between a p53 cancer vac recapitulates the latest experimental data and provide on the commensalistic relationship between the tumor

Keywords

Evolutionary Game Theory; Evolutionary Double Bind Immunotherapy; Chemotherapy; Combination therapy

1.2 Introduction

Cancer is an evolutionary disease [1]. One

Perspectives In Cancer Research

Lessons from Applied Ecology: Cancer Control Using an Evolutionary Double Bind

Robert A. Gatenby,¹ Joel Brown,² and Thomas Vincent¹

¹Departments of Radiology and Integrative Mathematical Oncology, Moffitt Cancer Center, Tampa, Florida; ²Department of Ecology and Evolutionary Biology, University of Illinois, Chicago, Illinois and ³Department of Aerospace Engineering, University of Illinois, Urbana, Illinois

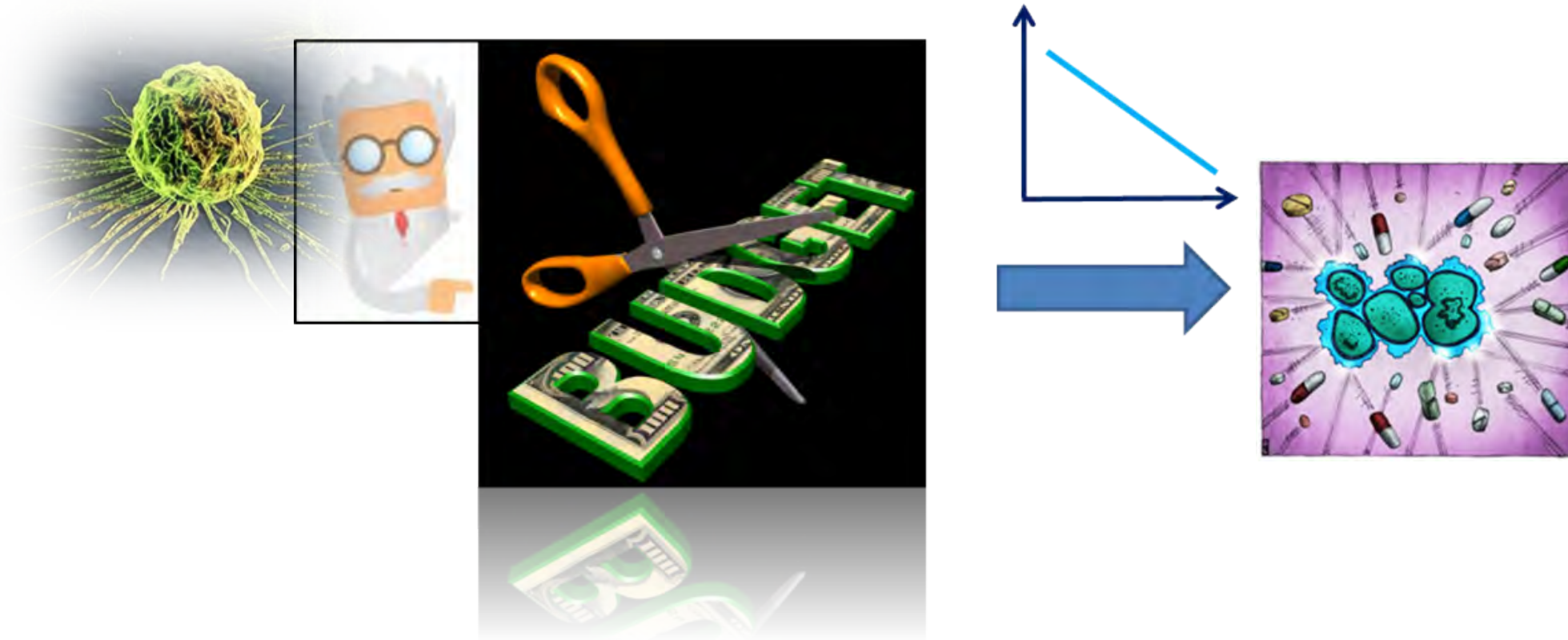
Abstract

Because the metastatic cascade is largely governed by the ability of malignant cells to adapt and proliferate at the distant tissue site, we propose that disseminated cancers are analogous in many important ways to the evolutionary and ecological dynamics of exotic species. Although goals can be decimated through the application of chemical toxins, this strategy virtually never achieves robust control as evolution of resistant phenotypes typically permits population recovery to pretreatment levels. In general, biological strategies that introduce predators, parasitoids, or pathogens have achieved more durable control of pest populations even after emergence of resistant phenotypes. From this we propose that long term outcomes from any treatment strategy for invasive pests, including cancers, is not limited by evolution of resistance, but rather by the phenotypic cost of that resistance. If a cancerous cell's adaptation to therapy is achieved by upregulating oncogenic metabolism or a redundant signaling pathway, the required investment in resources is small, and the original malignant phenotype remains essentially intact. As a result, the cancer cell's initial high level of fitness is little changed and unconstrained proliferation will resume once resistance evolves. Robust population control is possible if resistance to therapy requires a substantial and costly phenotypic adaptation that also significantly reduces the organism's fitness in its original niche: an evolutionary double bind. [Cancer Res 2009;69(12):3999-3002]

failures to proliferate in a distant organ. For example, in one study 87% (14) of injected cells survived to invade into the extravascular tissue space of a distant organ where they remained viable for prolonged periods of time, often several years. Only 10/25 (4) of these surviving cells grew into clinically evident metastases. Although some of the impacted cells formed clinically insignificant microtumors, the vast majority did not survive.

These results indicate that development of metastases is largely dependent on the complex, dynamic interactions between the phenotypic properties of the circulating tumor cell and micro-environmental conditions in the tissue at the metastatic site. Only tumor cells that are pre-adapted to or able to adapt to the local "ecological" conditions of the "foreign" landscape within the distant organ can form a metastasis. This finding is consistent with the long held "seed and soil" concept [7, 8] and indicates that Darwinian dynamics at the distant site play a fundamentally important role in the metastatic cascade. A feedback loop exists between the tumor cells' new environment and the ecological and evolutionary responses of the tumor cell's phenotype to its novel circumstances.

In the past 200 years, foreign species have been introduced into a wide range of habitats by human activities as well as random natural processes [9]. Clearly, metastatic cancers and invasive pests are both complex and highly diverse processes, and there are many obvious and subtle differences. However, we propose that there are sufficient similarities that general principles from the evolutionary ecology of invasive pest species may provide insights into treatment strategies for metastatic cancers. [10]. For example,



Evolutionary double bind

In an evolutionary double bind two therapies are used in combination such that evolving resistance to one leaves individuals more susceptible to the other.

Controversy and Consensus

First Strike–Second Strike Strategies in Metastatic Cancer: Lessons from the Evolutionary Dynamics of Extinction

Robert A. Gatenby, Jingsong Zhang, and Joel S. Brown

DOI: 10.1158/0008-5472.CAN-19-0807 [Check for updates](#)

[Article](#)

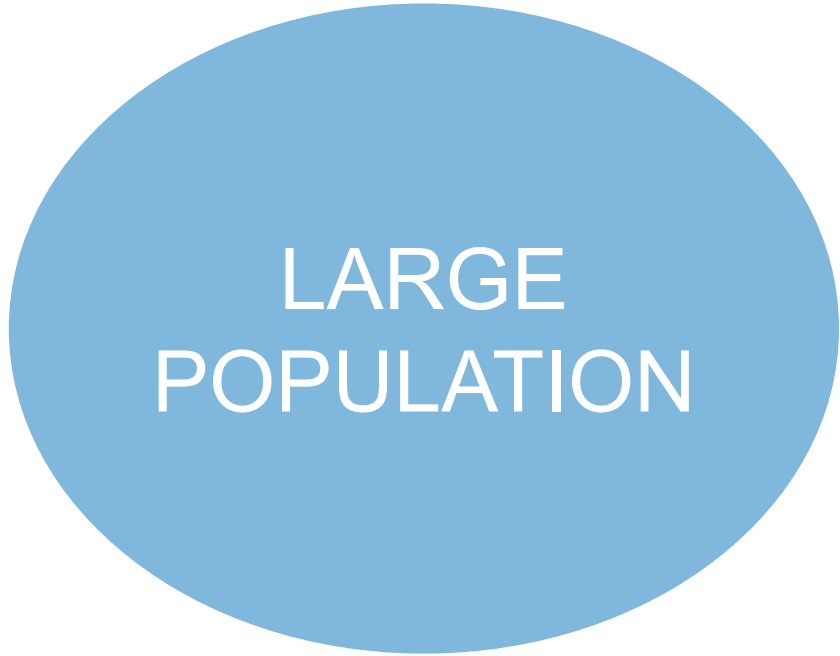
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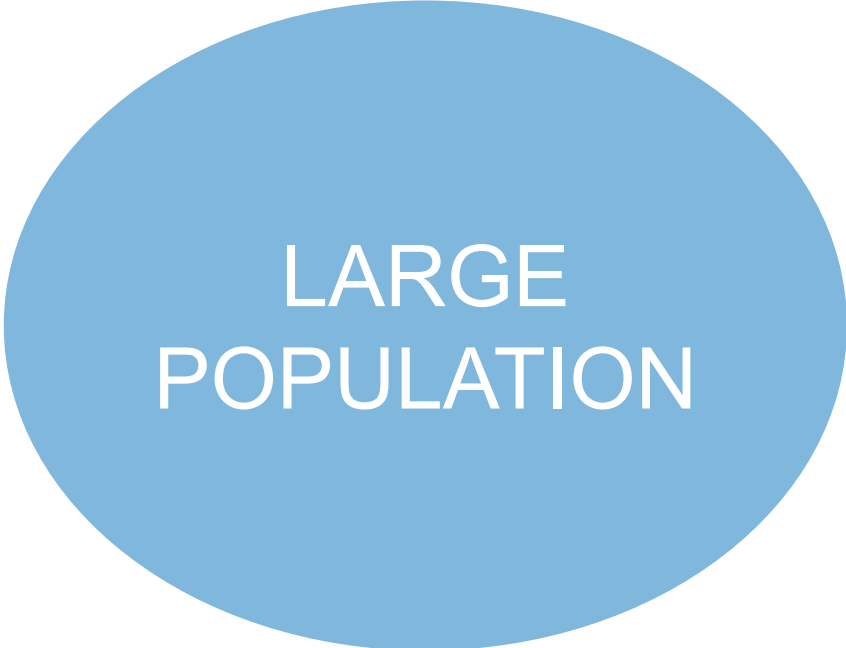


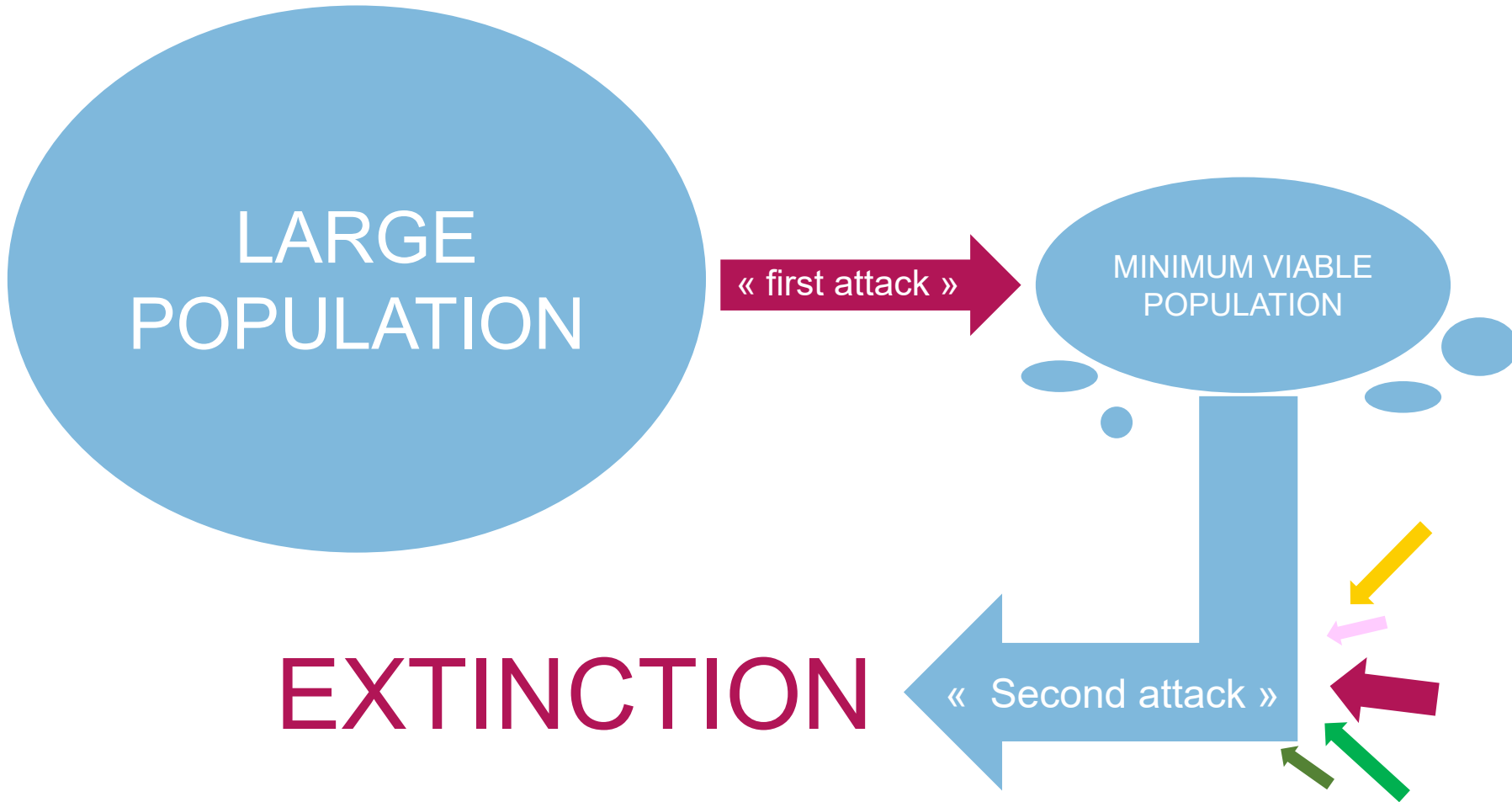
Evolutionary dynamics of Extinction



EXTINCTION







LARGE
POPULATION

« first attack »

MINIMUM VIABLE
POPULATION

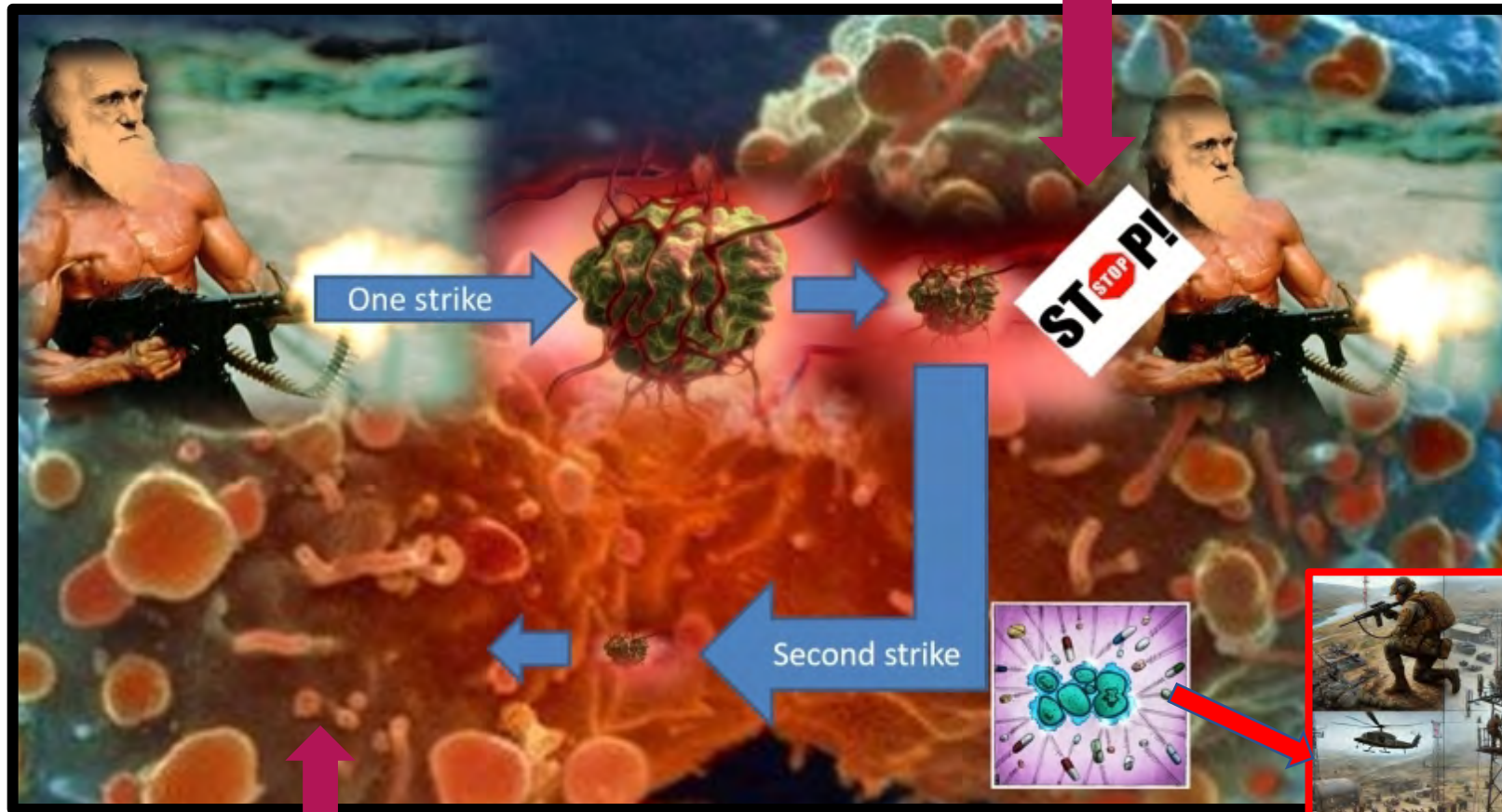
EXTINCTION

« Second attack »

First strike-second strike



Change the therapy even if it works



Continue the therapy



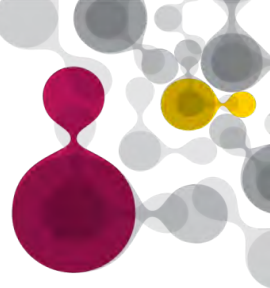
Cancer is problematic for human health mainly because it often evolves so fast that it is able to outrun our defence systems.



By reducing cancer evolution rate by 2 or 3 fold, most of our tumours would become harmless due to the cancer would appear beyond our present life expectancy.



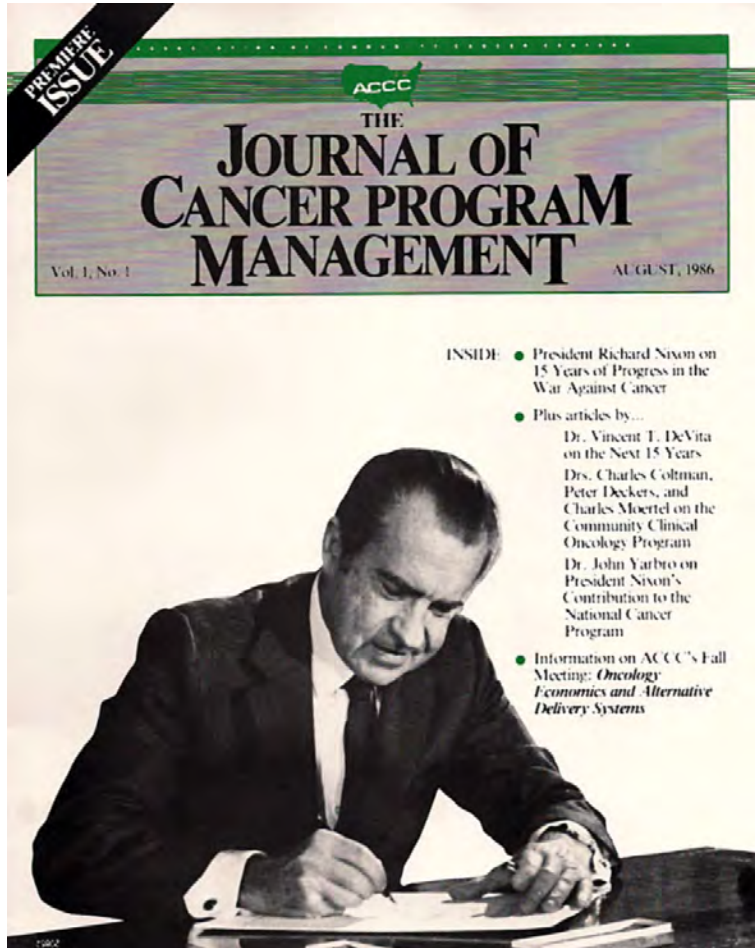
How Do We Slow Down Evolution?



- Reducing the mutation rate since it reduces the diversity in the population of cells
- Slow down the division rate
- Slow evolution by reducing the population size of the tumor and/or by reducing the fitness differences among cells (since differences in survival and reproduction are what drive evolution)
- Manipulating the environment to encourage cancer cells to evolve in the direction of a slower life history rendering them for instance dormant



More than 50 years after President Richard Nixon declared a war on cancer, victory remains elusive...



President Richard Nixon signs the National Cancer Act, Dec. 23, 1971, launching a \$1.6 billion federal crusade to conquer cancer. (AP)



COMMENTARY

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EVOLUTION, MEDICINE, & PUBLIC HEALTH

A new perspective on tumor progression

Evolution via selection for function

Frédéric Thomas^{1,2}, James DeGregori³, Andriy Maruyki⁴, Antoine M. Dujon^{1,4}, Beata Ujvari⁵, Jean-Pascal Cappi¹, Robert Gatenby⁶ and Aurora M. Nedelcu¹

¹CRIS/CANCOPI MRESEC (CRIS) Department, University of Montpellier, CNRS, IRD, Montpellier, France; ²Department of Biochemistry and Molecular Genetics, University of Colorado Anschutz Medical Campus, Aurora, CO, USA; ³Department of Cancer Physiology, Hillier Medical Cancer Center and Research Institute, Tampa, FL, USA; ⁴School of Life and Environmental Sciences, Deakin University Geelong, Centre for Invasive Ecology, Warrnambool, VIC 3276, Australia; ⁵Toulouse Bioinformatics Institute, University of Toulouse, INSA, CNRS, INRAE, Toulouse, France; and ⁶Department of Biology, University of New Brunswick, Fredericton, New Brunswick, Canada

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ABSTRACT
Tumorigenesis is commonly attributed to Darwinian processes involving natural selection among cells and groups of cells. However, progressing tumors are those that also achieve an appropriate group phenotypic composition (GPC). Yet, the selective processes acting on tumor GPCs are distinct from that associated with classical Darwinian evolution (i.e. natural selection based on differential reproductive success) as tumors are not genuine evolutionary individuals and do not exhibit heritable variation in fitness. This complex evolutionary scenario is analogous to the recently proposed concept of 'selection for function' involved for the evolution of both living and non-living systems. Therefore, we argue that it is inaccurate to assert that Darwinian processes alone account for all the aspects characterizing tumorigenesis and cancer progression; rather, by producing the genetic and phenotypic diversity required for creating novel GPCs, these processes fuel the evolutionary success of tumors that is dependent on selection for function at the tumor level.

KEYWORDS: cancer; evolution; natural selection; function; group; phenotype; composition

THE PREMISE
Following the pioneering work of Cairns [1] and Nowell [2], tumorigenesis has been generally viewed as underpinned by a classical Darwinian evolutionary process (i.e. somatic evolution) primarily governed by natural selection among mutant clones differing in fitness (i.e. survival and reproduction), starting from the emergence

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Thérapies fondées sur le GPC : pour quels types de tumeurs ?

- Tumeurs dont la complexité structurelle et fonctionnelle limite l'efficacité des approches traditionnelles ciblant les cellules individuelles, e.g. cas de cancers localement avancés mais non métastatiques, où les interactions complexes entre la tumeur et son microenvironnement rendent une résection chirurgicale complète difficile
- Pour les cancers ayant développé une résistance aux traitements cytotoxiques, la thérapie basée sur le GPC peut cibler les réseaux fonctionnels qui soutiennent la résilience de la tumeur.
- Les tumeurs présentant une forte hétérogénéité intratumorale, contenant de nombreuses sous-populations cellulaires coopératives, pourraient également bénéficier de cette approche, car elle perturbe les interactions qui soutiennent leur adaptabilité.
- cas où les amas de cellules tumorales circulantes (CTC) contribuent à la dissémination métastatique, cibler le GPC de ces amas pourrait limiter leur capacité à échapper au système immunitaire et à coloniser d'autres sites. En revanche, pour les tumeurs plus petites et localisées, les approches traditionnelles (par exemple, la chirurgie suivie de radiothérapie ou de chimiothérapie) peuvent être plus appropriées.

La stratégie thérapeutique que nous proposons cible le GPC de la tumeur, plutôt que de se concentrer uniquement sur les cellules cancéreuses individuelles. Cette approche viserait à déstabiliser les réseaux fonctionnels de la tumeur qui soutiennent sa survie et sa progression.

Voici une stratégie proposée en plusieurs volets :

1. Déstabilisation des réseaux fonctionnels : En ciblant des nœuds clés dans les réseaux fonctionnels de la tumeur (comme ceux impliqués dans l'angiogenèse, l'évasion immunitaire ou le remodelage de la matrice extracellulaire), on pourrait démanteler le système de soutien de la tumeur. Par exemple :

1. Des **agents anti-angiogéniques** (comme le bevacizumab) pourraient priver la tumeur de nutriments essentiels.
2. Les **inhibiteurs des points de contrôle immunitaires** (comme les inhibiteurs de PD-1/PD-L1, tel que le pembrolizumab) pourraient restaurer la surveillance immunitaire, renforçant ainsi la capacité du système immunitaire à attaquer la tumeur.
3. Les **inhibiteurs du remodelage de la matrice** empêcheraient la propagation de la tumeur en maintenant la structure extracellulaire.

2. Thérapie séquentielle et adaptative : La mise en place d'un plan de traitement séquentiel pourrait empêcher la tumeur de stabiliser une GPC résistante. En alternant des thérapies ciblant différentes fonctions (par exemple, le métabolisme, l'évasion immunitaire), cette approche pourrait maintenir la tumeur en perpétuelle adaptation, jusqu'à la piéger dans un état non oncogénique.

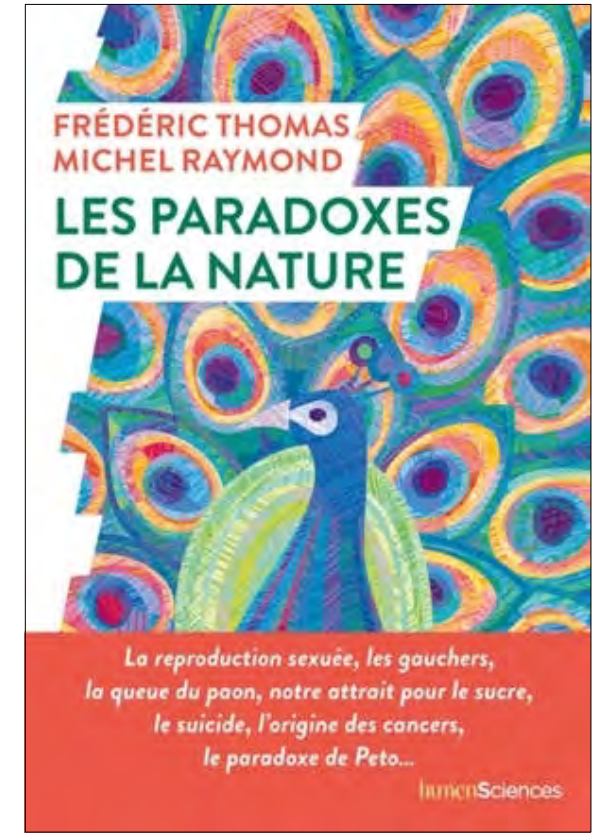
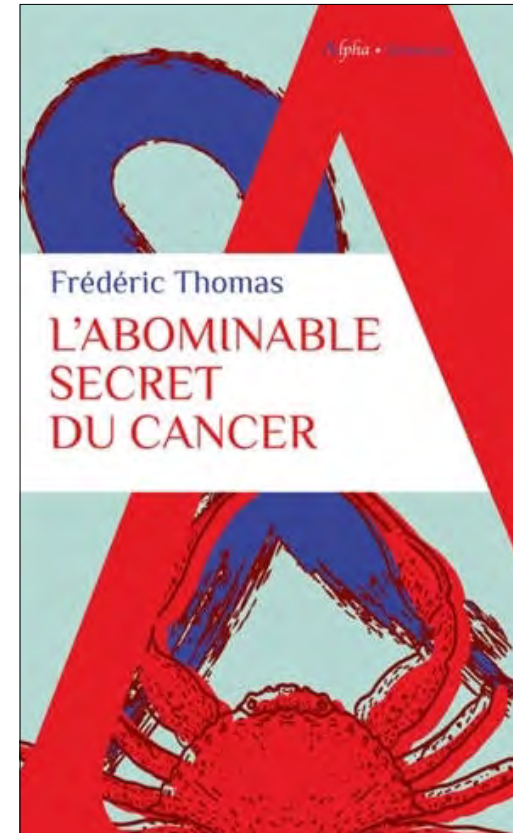
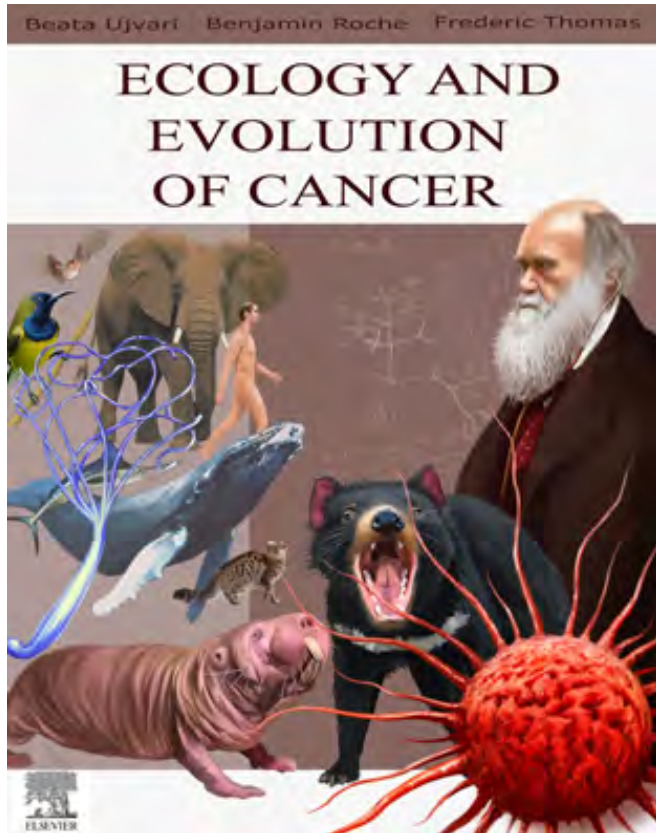
3. Modulation du microenvironnement : Modifier le microenvironnement de la tumeur pour créer des discordances avec son GPC pourrait encore déstabiliser la tumeur. Par exemple, des thérapies qui stabilisent le pH extracellulaire pourraient perturber l'environnement acide souvent exploité par les cellules cancéreuses pour survivre, et le Losartan pourrait être utilisé pour réduire la densité des fibroblastes associés au cancer et améliorer la délivrance des médicaments.

4. Favoriser les clones non-oncogéniques : Introduire des conditions qui privilégient les phénotypes non oncogéniques pourrait orienter l'évolution de la tumeur vers un état moins agressif. Par exemple, altérer les voies métaboliques pour favoriser la phosphorylation oxydative par rapport à la glycolyse pourrait diminuer la capacité d'adaptation et l'agressivité de la tumeur.

5. Surveillance en temps réel et ajustements : En utilisant des biopsies liquides et la transcriptomique spatiale, il serait possible de suivre l'évolution de la GPC et d'ajuster les traitements en temps réel. Cette approche adaptative permettrait de choisir des combinaisons de traitements et des moments précis pour prendre de vitesse l'évolution de la tumeur.

Cette stratégie s'appuie sur la dynamique évolutive de la tumeur, en visant à la fois à déstabiliser les GPC oncogéniques et à promouvoir l'émergence de phénotypes moins agressifs, avec pour ambition de transformer le cancer en une condition chronique et gérable.

Thank you for your attention...



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