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247. Heme-targeting

agents HSP2 and CycT

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Lijie Cao, Hao Wen,
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2020; 11(24):
7368-7377.
doi:10.7150/jca.50752
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In this research paper we discuss the properties of cancer mortality, the primary data sources, and the availability and quality of data between countries and over time, and the ensuing difficulties with interpretation. We also provide some selected descriptions of cancer mortality patterns by sex

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research over the period 2011-2015. In 2014, cancer research represents over 4 % of all re-search performed globally, reflecting the huge amount of

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09:42. Cancer is among
the leading causes of

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Of Cancer death worldwide, accounting for 8.2 million deaths in 2012 according to the World Health Organization. According to the same source, it is expected that annual cancer cases will rise from 14 million in 2012 to 22 million by 2030. Research on cancer development, treatment and prevention, as well as

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Of Cancer in this area, is essential to reduce the burden of this disease.

Cancer research

In summary, I strongly believe that this paper illustrates extremely well what is wrong with biomedical research. It chooses an isolated association of protein

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and cancer, selects supporting evidence — all reductionist — and ignores both the contrary specific evidence and the extensive evidence concerning other disease outcomes.

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in Cancer Research -
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Diet and cancer | Cancer
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Cancer is defined as
“when the body’s cells
begin to divide without
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Of Cancer
 (“What is cancer?”,
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mutations that lead to
the cell cycle to
proceed, regardless if
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The mutations block the
use of the G1, G2, and
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Cancer is a very complex disease and research geared towards learning more about its origin, progression, and treatments. Medical researches use animals to study cancer for a variety of reasons. Animals have shorter lifespans and more rapid generation times

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compared to humans,
and disease progression
moves forward at a
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results of studies using
animals as models are
obtained quicker³.

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Abstract. Skin cancers
are cancers that affect

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of skin. These include melanoma, basal and squamous cell cancers. The symptoms include skin color change and appearance of skin ulcers, as well as, alterations in the moles that are on the skin. Smoking and ultraviolet radiations are some of the leading causes of skin cancer.

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overview of the
molecular mechanisms
underpinning the
development of cancer
and its treatment.

Written by an
international panel of
researchers, specialists
and practitioners in the
field, the text discusses
all aspects of cancer
biology from the causes,
development and
diagnosis through to the

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level courses in the biology of cancer, for medical students and qualified practitioners preparing for higher exams, and for researchers and teachers in the field

Genomic sequencing technologies have augmented the classification of cancer beyond tissue of origin

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and towards a molecular taxonomy of cancer.

This has created opportunities to guide treatment decisions for individual patients with cancer based on their cancer's unique molecular characteristics, also known as precision cancer medicine. The purpose of this text will be to describe the

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Contribution and need for multiple disciplines working together to deliver precision cancer medicine. This entails a multi-disciplinary approach across fields including molecular pathology, computational biology, clinical oncology, cancer biology, drug development, genetics, immunology, and

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bioethics. Thus, we have outlined a current text on each of these fields as they work together to overcome various challenges and create opportunities to deliver precision cancer medicine. As trainees and junior faculty enter their respective fields, this text will provide a framework for understanding the role

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and responsibility for each specialist to contribute to this team science approach.

It has been recognized for almost 200 years that certain families seem to inherit cancer. It is only in the past decade, however, that molecular genetics and epidemiology have combined to define the

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role of inheritance in cancer more clearly, and to identify some of the genes involved. The causative genes can be tracked through cancer-prone families via genetic linkage and positional cloning. Several of the genes discovered have subsequently been proved to play critical roles in normal growth

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and development. There are also implications for the families themselves in terms of genetic testing with its attendant dilemmas, if it is not clear that useful action will result. The chapters in *The Genetics of Cancer* illustrate what has already been achieved and take a critical look at the future directions of this

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research and its
potential clinical
applications.

“Social Inequities in Cancer” is a compendium of articles that identify barriers and metrics on the topic of modifiable and unnecessary cancer inequalities. Social inequities have long been recognised as a

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strong contributing factor in health and cancer inequalities for the past several decades. Despite progress in cancer treatment, cancer incidence, mortality and survival vary markedly between and within countries. Globalisation, greater life expectancy, emerging analytical technologies, and the scalability of big data

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have revolutionized the vantage point from which social inequities can be studied. The focus of these articles is inequalities as they relate to cancer, with the inequalities ranging from the community to the global scale.

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Cancer cell biology research in general, and anti-cancer drug development specifically, still relies on standard cell culture techniques that place the cells in an unnatural environment. As a consequence, growing tumor cells in plastic dishes places a selective pressure that

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of Cancer substantially alters their original molecular and phenotypic properties. The emerging field of regenerative medicine has developed bioengineered tissue platforms that can better mimic the structure and cellular heterogeneity of in vivo tissue, and are suitable for tumor bioengineering research.

Microengineering

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technologies have resulted in advanced methods for creating and culturing 3-D human tissue. By encapsulating the respective cell type or combining several cell types to form tissues, these model organs can be viable for longer periods of time and are cultured to develop functional properties

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This approach recapitulates the dynamic role of cell–cell, cell–ECM, and mechanical interactions inside the tumor. Further incorporation of cells representative of the tumor stroma, such as endothelial cells (EC) and tumor fibroblasts, can mimic the in vivo

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tumor microenvironment.

Collectively, bioengineered tumors create an important resource for the in vitro study of tumor growth in 3D including tumor biomechanics and the effects of anti-cancer drugs on 3D tumor tissue. These technologies have the potential to overcome

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current limitations to
genetic and histological
tumor classification and
development of
personalized therapies.

This book explores in
depth the relation
between physical
activity and cancer

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control, including primary prevention, coping with treatments, recovery after treatments, long-term survivorship, secondary prevention, and survival. The first part of the book presents the most recent research on the impact of physical activity in preventing a range of cancers. In the second part, the

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association between physical activity and cancer survivorship is addressed. The effects of physical activity on supportive care endpoints (e.g., quality of life, fatigue, physical functioning) and disease endpoints (e.g., biomarkers, recurrence, survival) are carefully analyzed. In addition, the determinants of

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physical activity in cancer survivors are discussed, and behavior change strategies for increasing physical activity in cancer survivors are appraised. The final part of the book is devoted to special topics, including the relation of physical activity to pediatric cancer survivorship and to palliative cancer care.

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"Cancer research is at a crossroads.

Traditionally, cancer has been thought of as a disease of gene mutation, where the stepwise accumulation of cancer gene mutations is the key, and the identification of common gene mutations has been considered to be essential for

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diagnosis and treatment.

Despite extensive research efforts and accumulated knowledge on cancer genes and pathways, the clinical benefits of this traditional approach have been limited.

Recently, cancer genome sequencing has revealed an extensive amount of genetic heterogeneity where the

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long-expected common mutation drivers have been difficult, if not impossible, to identify. These realities ultimately challenge the conceptual framework of current cancer biology. This book introduces a new concept of genome theory of cancer evolution, in an attempt to unify the field. Many

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important and representative, but often confusing, questions and paradoxes are critically analyzed. By comparing gene- and genome-based theories, the hidden flaws of many popular viewpoints are addressed. This discussion is intended to initiate a much-needed critical re-evaluation of current cancer

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Since the invention of nanomedicine decades ago, considerable progresses have been made, especially with cancer as a target.

Nanoparticles have been proven to be powerful imaging tools or potent agents for cancer diagnosis, treatment and prevention. Active

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research spread from fundamental research to clinical investigations. This topic intends to cover several important aspects in this field including nanocarrier development, gene delivery, intrinsically active nanoparticles, tumor microenvironment, immunology, and toxicity.

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