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# **Realidades y Esperanzas**

una reflexión muy personal sobre el cáncer...

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### Disclosures

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### **Conflict of Interest Statement**

This presentation reflects my personal opinion, and not that of the sponsor of this activity. Its main objective is to stimulate independent scientific discussion and does not intend to promote a specific product or indication.



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The hominin wall at the Natural History Museum of Utah in Salt Lake City. (Image credit: Copyright NHMU/Credit: Mark Johnston)

#### "Life is a self-sustaining chemical system capable of Darwinian evolution"

astrobiology.nasa.gov

The hominin wall at the Natural History Museum of Utah in Salt Lake City. (Image credit: Copyright NHMU/Credit: Mark Johnston)









- Cancers arise from single cells.
- Loss of key cellular attributes, (tumor suppressor genes), are a key driver event in the development of cancer
- Inheritance could play a role in cancer susceptibility.
- *Chromosomal (genomic) instability* as key hallmark of cancer

(1862 - 1915)

 Genetic information could be contained in *distinct packages* (genes) that are linearly arranged along chromosomes

### The Clonal Evolution of Tumor Cell Populations

Acquired **genetic lability** permits stepwise selection of variant sublines and underlies tumor progression. (*P Nowell*)



# From a biologic and evolutionary point of view, cancer is an almost unavoidable or inevitable outcome (event)

Cancer as a reactive process, is nothing more than an attempt to survive and therefore reach immortality...

#### Endogenous and Exogenous Mutational Processes are key in the Evolutionary Trajectory of a Tumor



- (A) An age-related mutational process operates throughout the evolution of a lung tumor. A smoking-induced C > A mutation in TP53 (p.R158L) leads to the outgrowth of a major tumor clone. Later in tumor evolution, APOBECmediated mutagenesis results in a mutation to PIK3CA (p.E545K), which leads to a subclonal expansion.
- (B) The evolution a glioblastoma tumor that has undergone treatment with Temozolomide (TMZ). Notably, TMZ leads to mutations in CDKN2A and RB1 in separate subclones, both of which lead to subclonal expansions.



Cancer: Speeding up Darwin's evolution

> Vendramin R, et al. The EMBO Journal, Vol: 40, Issue: 18. 30 August 2021, DOI: (10.15252/embj.2021108389)



Article

# Spatial genomics maps the structure, nature and evolution of cancer clones





Lomakin A, et al. Nature November 2022. https://doi.org/10.1038/s41586-022-05425-2



#### Impact of Personalized Approaches in Clinical Practice











#### Impact of Personalized Approaches in Clinical Practice





#### **MOLECULAR PROFILING / DIAGNOSIS**

#### Impact of Personalized Approaches in Clinical Practice











#### **MOLECULAR PROFILING / DIAGNOSIS**

\* This algorithm is responsibility of the author, is presented for didactic purposes only and may not necessarily be in accordance to approved therapies in certain countries. Please refer to your applicable approved indications.

#### Proposed Standard of Care Treatment Algorithm for Patients with Advanced NSCLC – 2022\*



Algorithm by CH Barrios updated and modified from Karen Kelly

### **Genomics and Cancer Therapy: Hope or Hype!**



Haslam et al, Ann Oncol, 2021 Diestmann R, et al, Ann Oncol, 2021

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#### Further increase in sample size does not increase gene identification



- Positive correlation between sample size and identified driver gene mutations across 33 cancers
- Sequencing as many specimens as possible for all cancer types is becoming inefficient.

#### Near saturation may achieved with 600-5000 samples/tumor



Lawrence SL, et al. Nature, 2014.

# Genes (sequencing) do not hold all the answers...

There is (much) more to cancer than gene abnormalities

Presenter's slide. CH Barrios, 2022.







### **Genomic Signatures**

- Exogenous or endogenous exposures and defective DNA repair/maintenance generate *mutational signatures* including base substitutions, small insertions and deletions, rearrangements and chromosome copynumber changes. (i.e., multiple mutational processes, each generating a characteristic mutational signature)
- Identified: 49 single-base-substitution, 11 doublet-base-substitution, 4 clusteredbase-substitution and 17 small insertion-anddeletion signatures.
- Some signatures are associated to unknown causes.

#### **Mutational Signatures in HR+ and HR- Breast Cancers**



# Response to neoadjuvant chemotherapy in HR+ tumors: multivariable analysis and differences in pCR rate



Denkert C, et al. Ann Oncol, Feb 2021.

#### **Mutations, Co-Mutations and Modifiers**

Computational Analysis of RWD (N=40.903)

**a** ALK - aNSCLC (*N* = 1,053) **b** BRAF - aNSCLC (*N* = 885) **c** EGFR - aNSCLC (*N* = 2,459) **d** KRAS - aNSCLC (*N* = 3,827)



Anchor genes are selected to be genes with available targeted therapies and are shown in the middle of each panel together with their prognostic HRs for OS. Modifiers are genes with significant anchor-modifier interactions (two-sided Wald test P value <0.05 and FDR <0.05). The size of a modifier gene's circle indicates what fraction of patients with the anchor gene mutated also have mutation in the modifier gene. The modifier's color indicates its positive (blue for HR <1) or negative (red for HR >1) impact on the survival of patients who have the anchor gene mutation. Liu R, et al. Nature Medicine, June 30th, 2022



Tumor-resident intracellular microbiota promotes metastatic colonization in breast cancer



Fu A, et al. Cell April 2022.



Figure 1. The vision of personalised healthcare: data, technology and analytics provide deeper insights into disease to accelerate research and development and improve patient outcomes.

#### • Information Technology Revolution

• Big Data, Artificial Intelligence, EMR, Digital follow-up, etc.

#### Personalized Care based on deeper genomic analysis/technology

• CRISPR - gene-editing, mutation-independent cellular programs, etc.

#### • Improvements in Therapy

- Immunotherapy will continue to evolve
  - Cell Therapies (CAR-T)
  - Combinations, patient selection, predictive biomarkers
  - Microbiome
- Therapy based on Monoclonal Antibodies Derivatives
  - Bi-specific antibodies
  - New Antibody-Drug conjugates
  - Combinations

#### Improvements in diagnostics and staging

- Liquid Biopsies and ctDNA diagnostics revolution
  - Early diagnosis of cancer (may be even Prevention...)
  - Better monitoring of disease evolution

## **Take Home Messages**

- **Beware of complexity**! Substantial heterogeneity is the rule for most prevalent tumors.
- Note that "genomics and genes do not hold all the answers" and that cancer biology and evolution allows for a number of yet to be characterized biological intervening and enabling dimensions very likely to be of clinical significance.
- We are just scratching the surface in our understanding of these characteristics.

## **Take Home Messages**

- Personalized Medicine based on mutated drivers is important and rational, but is only one of the possible strategies.
- It is naïve to think that we will be able to face the totality of the complexity of cancer with a single strategy.
- Combinations are required and will be the rule...





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