XXIII Encuentro de Cooperación Farma-Biotech

28 de noviembre de 2023

#### Thyromethyl, a method for the diagnosis of thyroid cancer based on epigenetic biomarkers



#### Mario Fernández Fraga







# Content

- 1. The Institution
- 2. The Product
  - a) Target Indications
  - b) Innovative mechanisms of action
  - c) Differential features facing the market
  - d) Current status of development
  - e) IPR protection
  - f) Pitfalls & Risks to be considered
- 3. Partnering Opportunities

#### The Institution

CSIC

cipado de Asturias

# Cancer Epigenetics and Nanomedicine

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# Endocrinología, nutrición, diabetes y obesidad

Elías Delgado Edelmiro Menéndez Torre Carmen Lambert Goitia Elsa Vera



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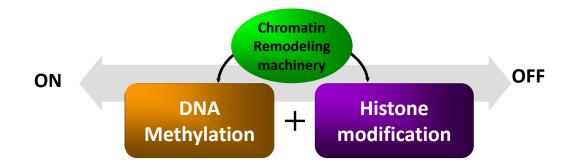
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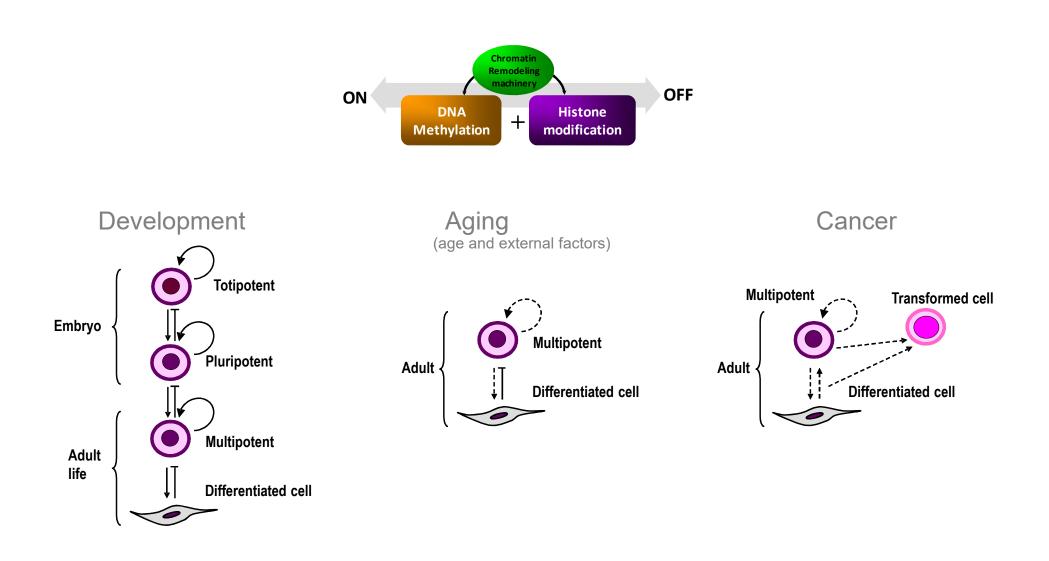
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### **Epigenetics**

"The study of stable genetic modifications that result in changes in gene expression and function without a corresponding alteration in DNA sequence"

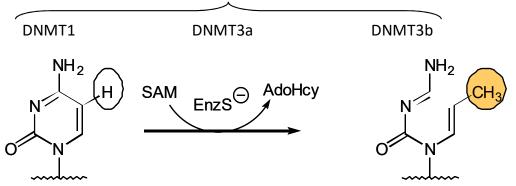


## **Epigenetics**



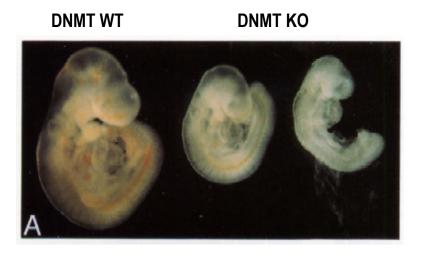
- DNA methylation is a process by which methyl groups are added to the DNA molecule.
- The reaction is catalyzed by a family of proteins called DNA methyltransferases.

DNA methyltransferase familly of proteins



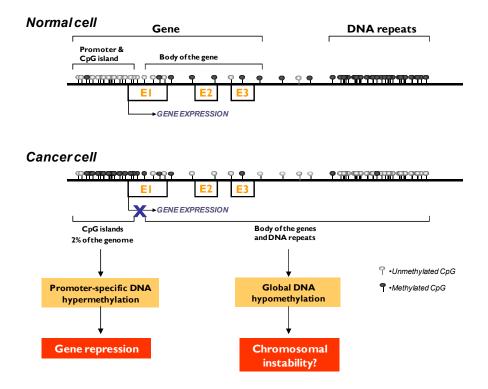
Fraga et al., 2009. Curr Opin Immunol.

- Genomic DNA methylation is essential for life in mammals
- Mice deficient in DNA methyltransferases die during the first stages of embryonic development.



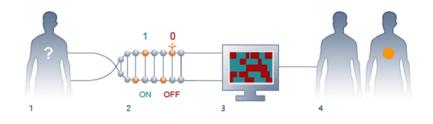
Li et al., 1992, Cell

- Global DNA hypomethylation
- CpG island promoter hypermethylation
- Aberrant patterns of DNA methylation are specific to tumor type and subtype

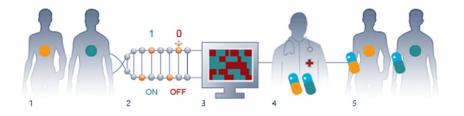


 Aberrant DNA methylation in cancer has clinical applications in diagnosis and pharmacodiagnosis

#### • **Diagnostic tests** (screening, monitoring, classification)



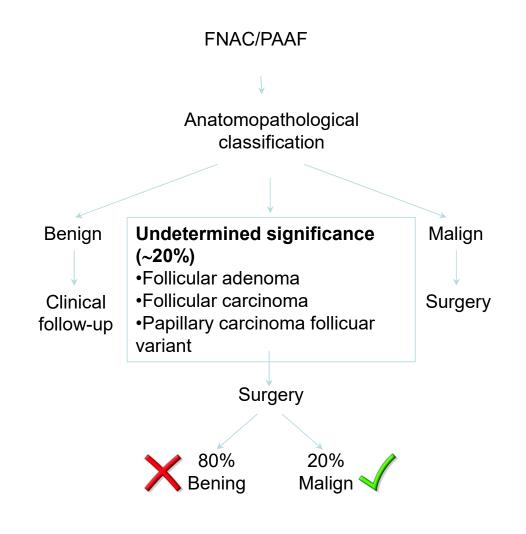
•Pharmacodiagnostic tests (responder/non-responder)



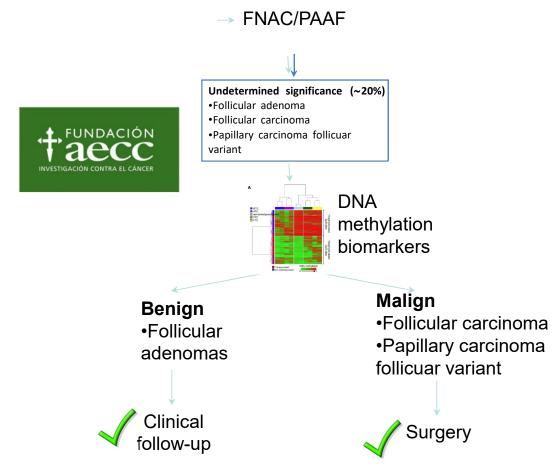
Adapted from www.epigenomics.com/

- Surgical resection of 80% of the thyroid nodules with undetermined anatomopathological significance is <u>unnecessary</u>.
- DNA methylation biomarkers to help anatomopathological classification

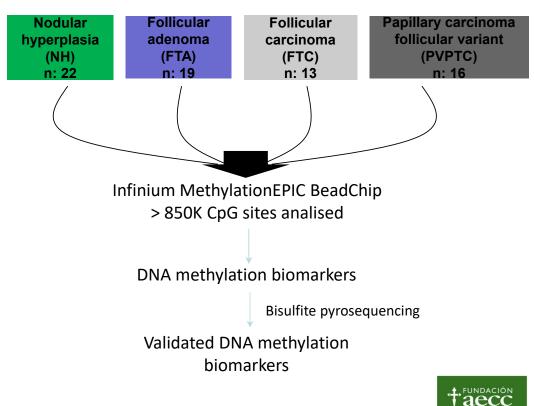
- Fine needle aspiration cytology (FNAC/PAAF) is commonly used in the diagnosis thyroid cancer.
- Lesions with undetermined cytology are surgically removed. However, surgery biopsies reveal that 80% of them are benign and, consequently, that surgery was unnecessary.



- Improvement of the diagnostic yield of fine needle aspiration cytology (FNAC) samples by the detection of DNA methylation markers, and its application for the early diagnosis of Follicular Thyroid Carcinoma
- AECC grant.

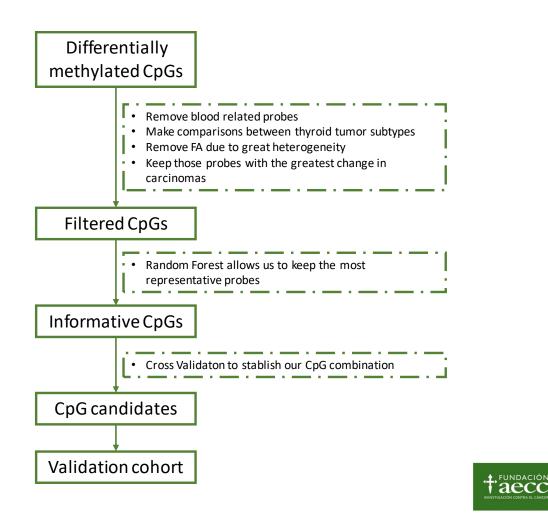


- Discovery cohort: Nodular hyperplasia (n=22), Follicular adenoma (n=19), Follicular carcinoma (n=13), Papillary carcinoma follicular variant (n=16).
- Validation cohort: Nodular hyperplasia (n=84), Follicular adenoma (n=39), Follicular carcinoma (n=26), Papillary carcinoma follicular variant (n=36).

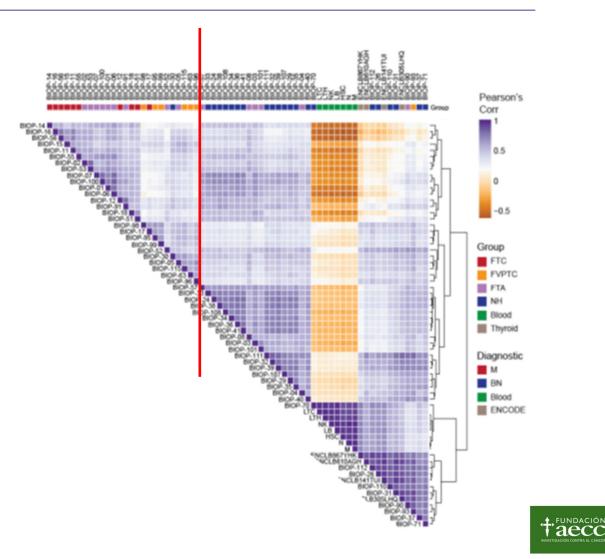


#### Experimental design

- A two-step data analysis strategy
- Classical approach to identify differentially methylated CpG sites and random forest approach to identify a minimal set of informative CpG sites

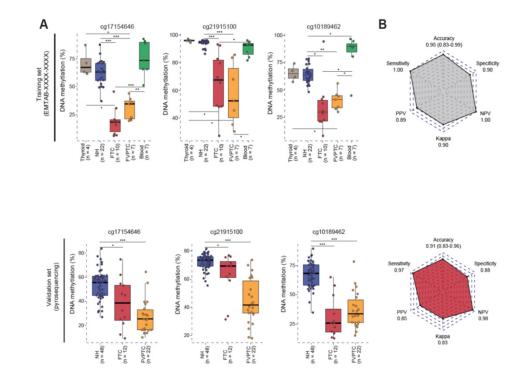


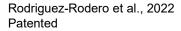
### DNA methylation biomarkers in thyroid cancer



- DNA methylation patterns distinguish malignant and benign samples
- Difficult to implement in the routine clinical practice

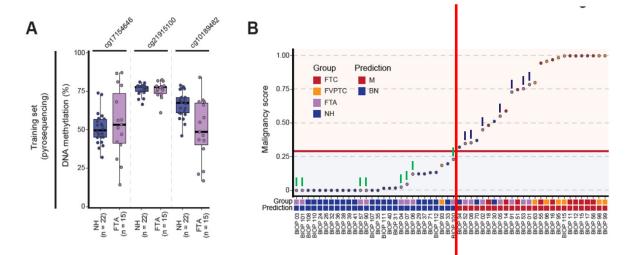
- Identification of the most informative CpG sites using random forest classification.
- Validation of these CpG sites in an independent cohort (Methylation arrays)
- Validation of these CpG sites in an independent cohort (Pyrosequencing)







- Follicular adenomas present a heterogenous behavior.
- A malignant score can stratify thyroid lesions accordingly to risk of malignancy.





Validation of the epigenetic classification system of thyroid lesion malignancy in Thyroid nodules (FNACs)

FNAC-21 FNAC-06 FNAC-08 FNAC-08 FNAC-04 FNAC-05

FNAC-1

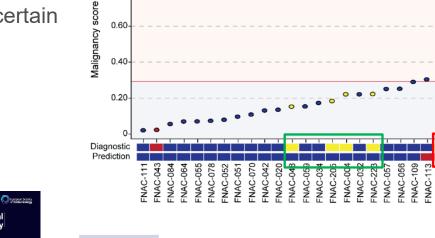
NAC-222-NAC-222-

FNAC-07

FNAC-1



• 9 Uncertain



Diagnostic

Uncertair

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Prediction

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BN



JOURNAL ARTICLE

1.00

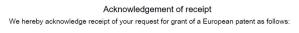
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# Classification of follicular-patterned thyroid lesions using a minimal set of epigenetic biomarkers Get access >

Sandra Rodríguez-Rodero, Paula Morales-Sánchez, Juan Ramón Tejedor, Andrés Coca-Pelaz, Cristina Mangas, Alfonso Peñarroya, Iván Fernández-Vega, Luís Fernández-Fernández, Carmen M Álvarez-López, Agustín F Fernández ... Show more

*European Journal of Endocrinology*, Volume 187, Issue 3, Sep 2022, Pages 335–347, https://doi.org/10.1530/EJE-22-0012 **Published:** 19 July 2022 Article history ▼





Submission number	300441836
Application number	EP22382165.3
File No. to be used for priority declarations	EP22382165
Date of receipt	25 February 2022
Your reference	EP1641.1719
Applicant	CONSEJO SUPERIOR DE INVESTIGACIONES CIENTIFICAS
Country	ES
<b>T</b> 241 -	

Title EPIGENETIC BIOMARKERS FOR THE DIAGNOSIS OF THYROID CANCER



- Public and private hospitals who treat patients with thyroid tumors
- In Spain about 50,000 PAAFs are performed every year. Therefore, about 10,000 patients per year could benefit from the use of this product.

• In our hospital, about 450-500 PAAFs per year.

*Epigenetic vs Genetic biomarkers* 

- Plasticity and Environmental Response
- Dynamic and Temporal Changes
- Better Reflection of Current Cellular Function
- Potential to Identify Modifiable Risk Factors

## **ThyroMethyl**

- Simplicity: 60-70 targets vs 3
- Sensitivity, specificity, etc.
- Availability of Pyrosequencers in public hospitals. E.g. Tryroidprint requires the purchase of costly equipment
- Valorization of national product. None of the similar products in the market have been developed in Spain
- Connections with the Spanish Public Health System, ISCIII, Public Hospitals, CIBER, etc. The product will be used by clinicians throughout Spain who have participated in its development.

TLR5: Validated with clinical samples

Biomarker identification (machine learning).

Versatility in multiple technologic environments (arrays, pyroseq)

Validation of these results with additional biospies with external cohorts

Proof of concept in a relevant clinical setting (PAAFs)

Standardizing protocols for extracting DNA from PAAF biopsies

Standardizing protocols for subsequent DNA methylation analyses, implementation in the clinical practice

Expand the metodology in novel clinical trials focused in the implementation of diagnostic biomarkers of tyroid cancer cohorts

Develop molecular extraction kits for the implementation of a standarized conditions across laboratories.

Implementation of an easy-to-use web interface to analyze data from the prospective cases.