



GASTRO
JournalClub

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UNIMORE
UNIVERSITÀ DEGLI STUDI DI
MODENA E REGGIO EMILIA

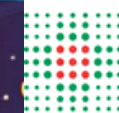


L'importanza della ricerca in Oncologia

10-11 OTTOBRE 2019 - ROMA

VOI Donna Camilla Savelli Hotel - Via Garibaldi, 27

**Dual targeting of
histone methyltransferase G9a
and DNMT1 in Gastric Cancer**



SERVIZIO SANITARIO REGIONALE
EMILIA-ROMAGNA
Azienda Ospedaliero - Universitaria di Modena

OUTLINE

BACKGROUND

- **Epigenetics**
- **Epigenetics in cancer**
- **Focus on G9a and DNMT1**
- **Dual targeting of G9a and DNMT1 in HCC e CCA**
- **Epigenetics and Gastric Cancer**

RESEARCH PROJECT

Epigenetics

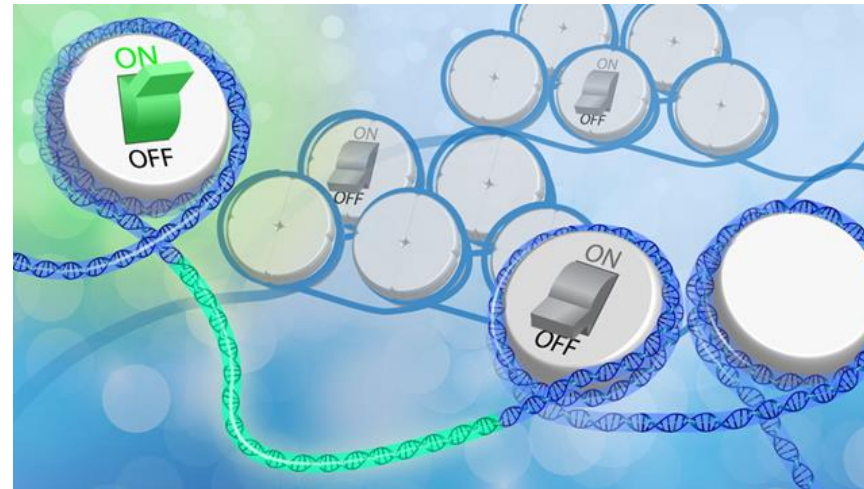
from transgenerational inheritance to disease



Initial definition: “The casual interactions between genes and their products, which bring the phenotype into being” (C.H. Waddington, 1939)

Current definition: “Heritable changes in gene expression that occur independent of changes in the primary DNA sequence”

Epigenetics in cancer. Estellar NEJM 2008.



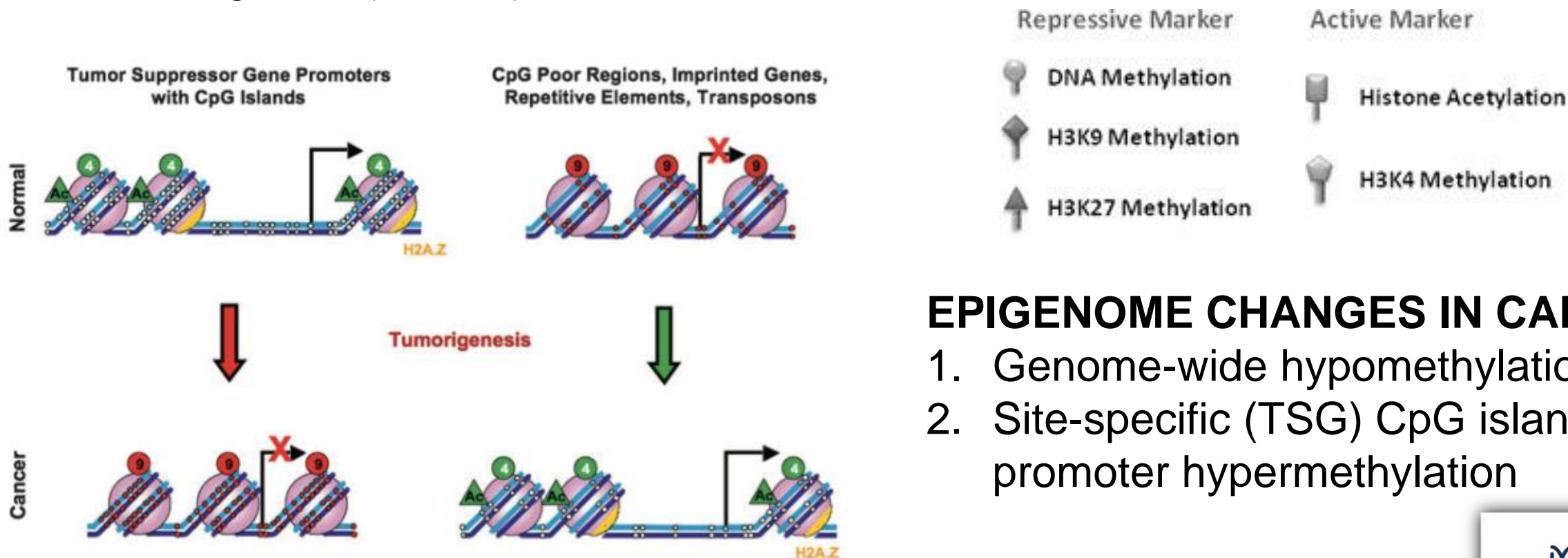
Choices you make can change your genes, and those of your kids

Time cover, 18th January 2010

Epigenetics and cancer – 1/3

EPIGENETIC MECHANISMS:

- DNA methylation (DNMT1, DNMT3A, DNMT3B)
- Histone (H2A, H2B, H3, H4) modifications: methylation (HMT) and acetylation
- Non coding RNAs (miRNAs)



EPIGENOME CHANGES IN CANCER:

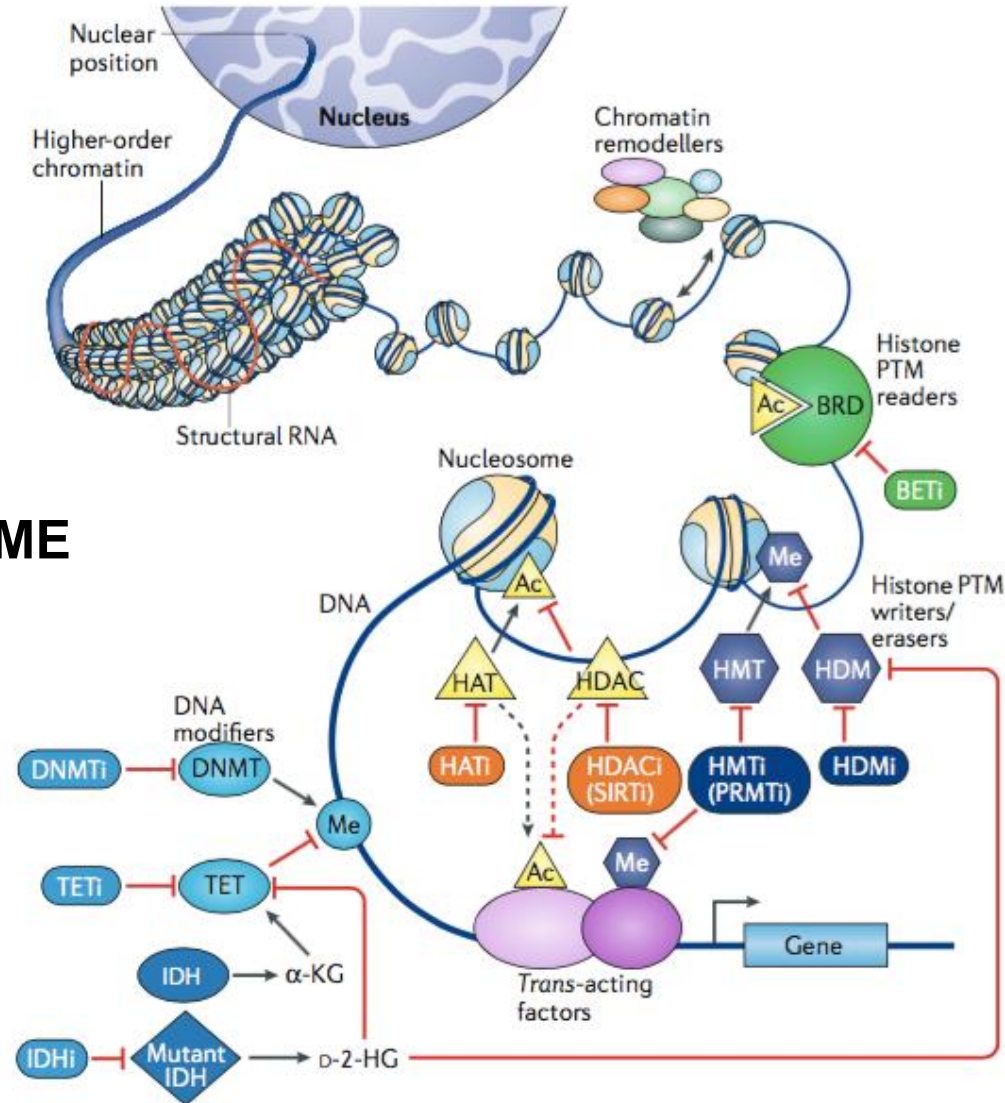
1. Genome-wide hypomethylation
2. Site-specific (TSG) CpG island promoter hypermethylation

Epigenetics in cancer. Sharma S et al. Carcinogenesis 2010.



Epigenetics and cancer – 2/3

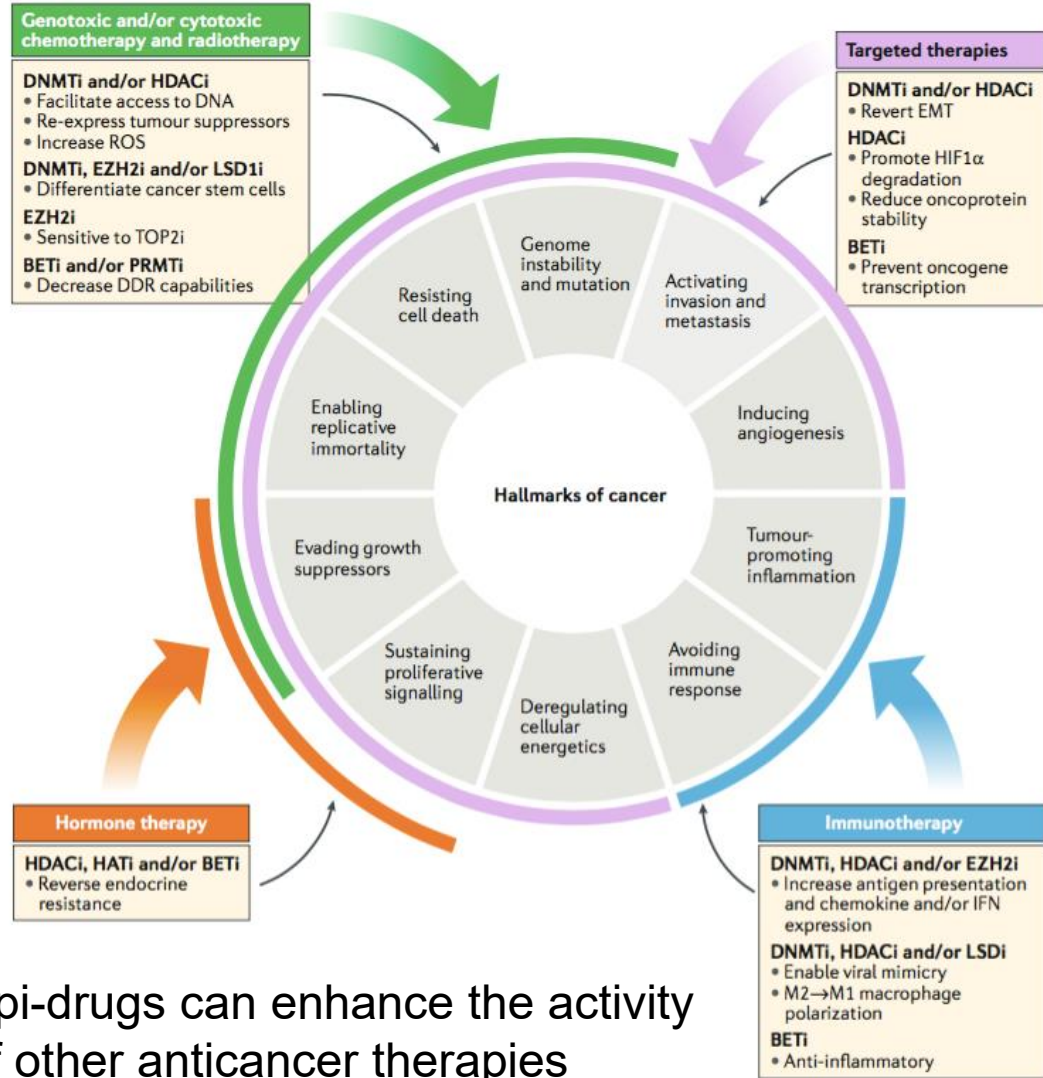
TARGETING THE EPIGENOME



Combining epigenetic drugs with other therapies for solid tumours – past lessons and future promise. Morel D et al. Nature Reviews Sept 2019.

Epigenetics and cancer – 3/3

Morel D et al. Nature Reviews Sept 2019.



EPI-DRUGS

Drugs	classification	Approved Year	Indicated disease	ORR
Azacitidine ²⁰	DNMT inhibitor	2004	MDS	17.9%
Vorinostat ³²	HDAC inhibitor	2006	CTCL	30%
Decitabine ³⁵	DNMT inhibitor	2006	MDS	42%~54%
Romidepesin ³⁴	HDAC inhibitor	2009	TCL	34%
Ruxolitinib ³⁶	JAK1/2 inhibitor	2011	Myelofibrosis	30%
Belinostat ³³	HDAC inhibitor	2015	PTCL	25.8%
Panobinostat ²⁵	HDAC inhibitor	2015	MM	NA

Epigenetic targeting drugs potentiate chemotherapeutic effects in solid tumor therapy. Li J et al. Sci Rep 2017

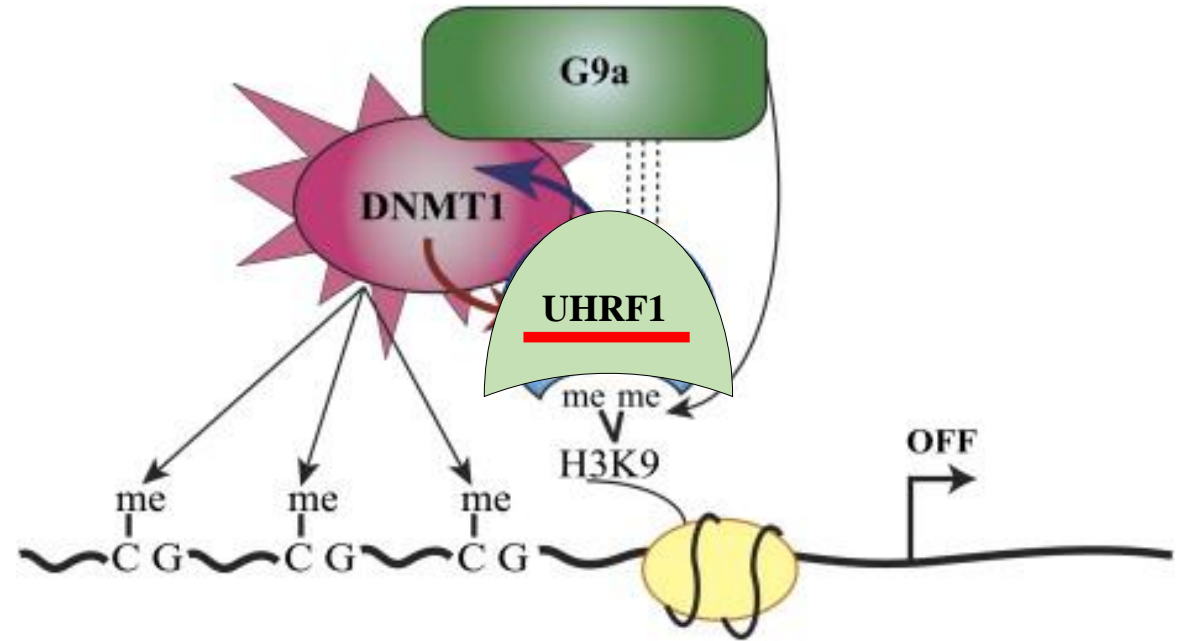
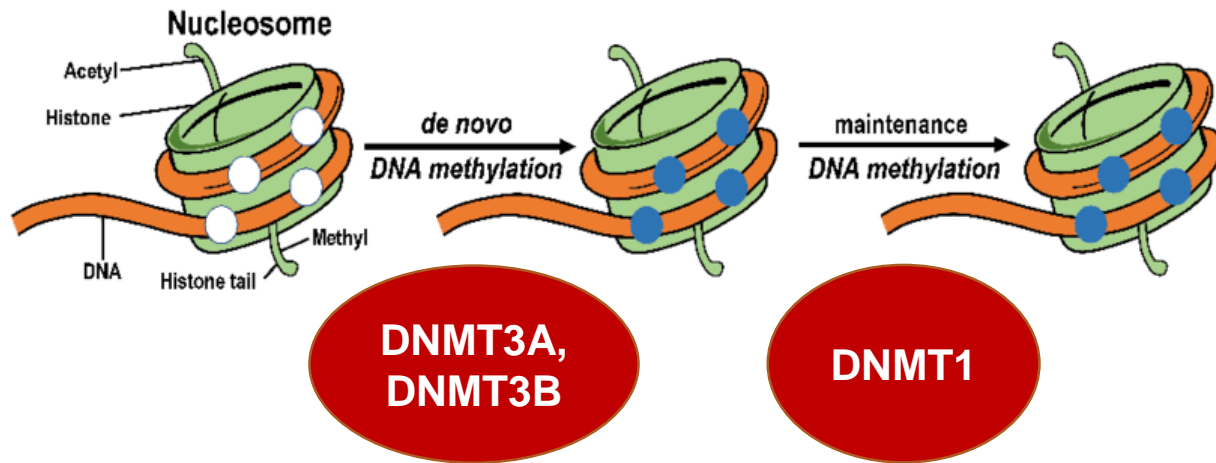
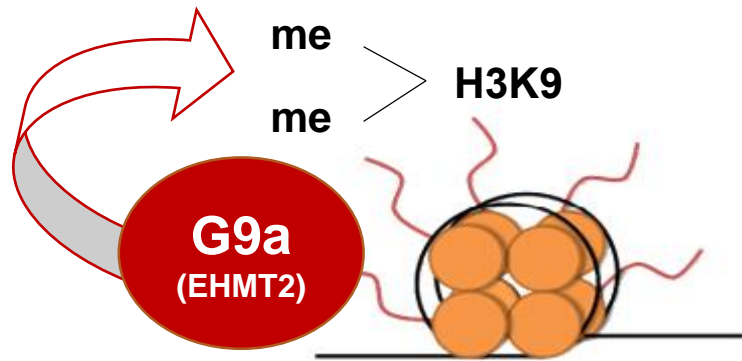


Azacitidine

Decitabine

Epi-drugs can enhance the activity of other anticancer therapies to counteract the hallmarks of cancer

Focus on G9a and DNMT1



Direct interaction between DNMT1 and G9a coordinates DNA and histone methylation during replication

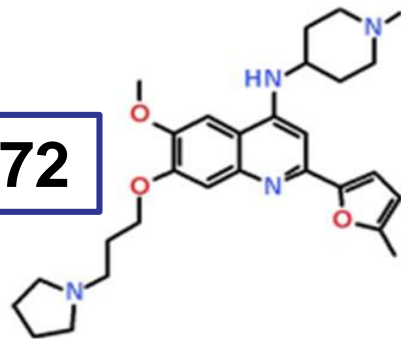
Estève *et al.* Genes Dev. 2006.

Dual Targeting of Histone Methyltransferase G9a and DNA-Methyltransferase 1 for the Treatment of Experimental Hepatocellular Carcinoma

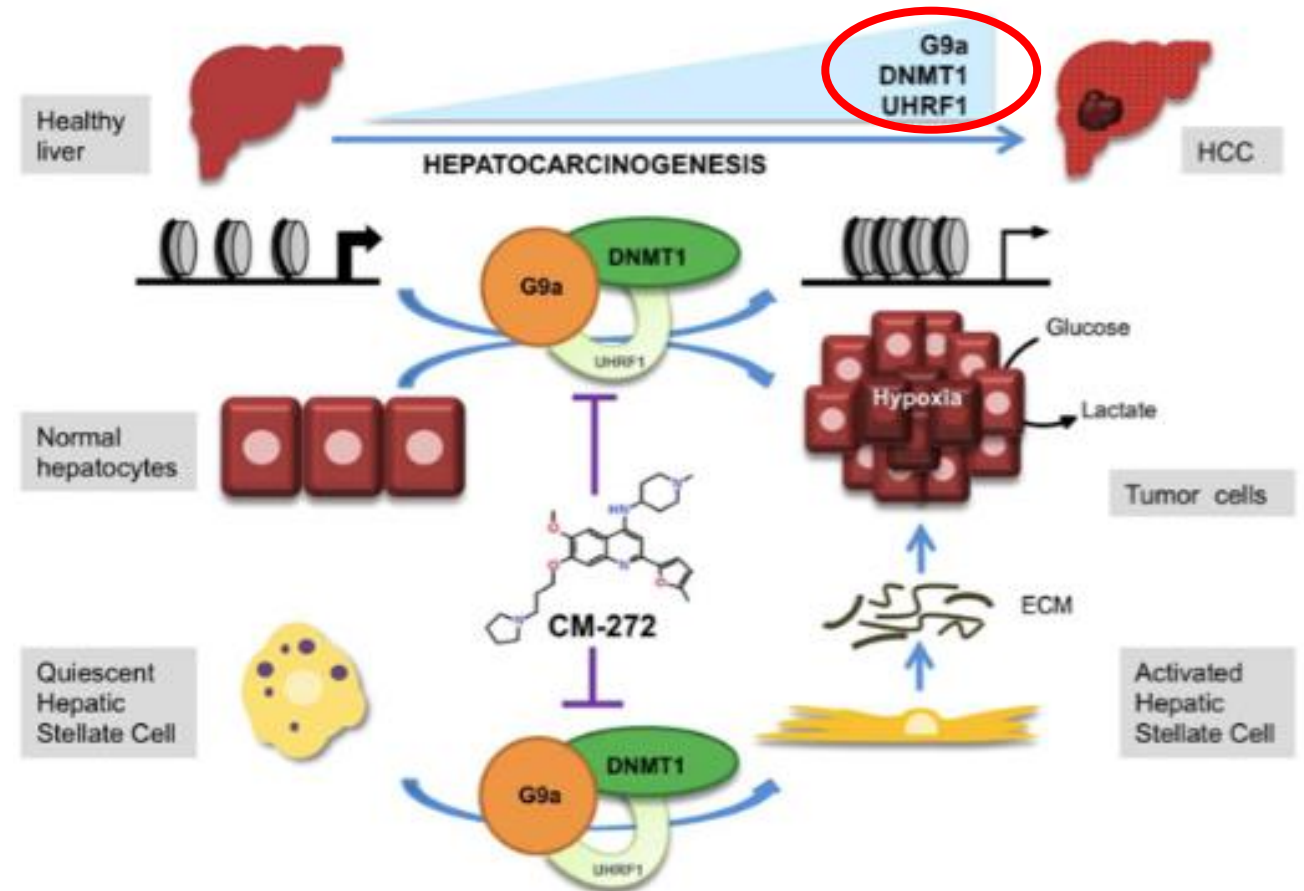
Marina Bárcena-Varela,¹ Stefano Caruso,² Susana Llerena,^{3,4} Gloria Álvarez-Sola,^{1,4} Iker Uriarte,^{1,4} M. Ujue Latasa,¹ Raquel Urtasun,¹ Sandra Rebouissou,² Laura Alvarez,¹ Maddalen Jimenez,¹ Eva Santamaría,^{4,7} Carlos Rodríguez-Ortigosa,^{1,4,7} Giuseppe Mazza,⁵ Krista Rombouts,⁵ Edurne San José-Eneriz,^{6,7} Obdulia Rabal,⁸ Xabier Agirre,^{6,7} Maria Iraburu,⁹ Alvaro Santos-Laso,^{9,10} Jesus M. Banales,^{4,9,10} Jessica Zucman-Rossi,¹⁰ Felipe Prósper,^{6,7} Julen Oyazabal,⁸ Carmen Berasain,^{1,4,7*} Matias A. Ávila,^{1,4,7*} and Maite G. Fernández-Barrena^{1,4,7*}

- First-in-class **dual** and **reversible** inhibitor targeting **G9a** & **DNMT1**.
- Substrate competitive (H3K9 and DNA).

Lead compound **CM-272**



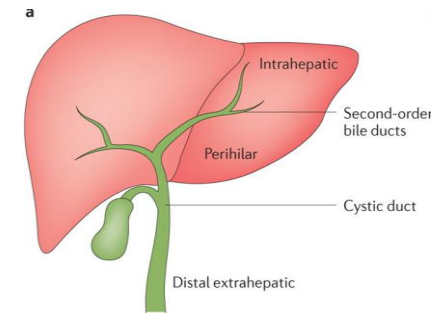
Dual targeting of G9a and DNMT1 in HCC



Dual targeting of G9a and DNMT1 in CCA

PS-043-Dual targeting of G9a and DNMT1 for the treatment of experimental cholangiocarcinoma

[Leticia Colyn](#), [Gloria Álvarez-Sola](#), [Maria U Latasa](#), [Iker Uriarte](#), [Marina Bárcena-Varela](#), [Maria Arechederra](#), [Maddalen Jimenez](#), [Sergio Morini](#), [Simone Carotti](#), [Julen Oyarzabal](#), [Felipe Prosper](#), [Matteo Canale](#), [Andrea Casadei Gardini](#), [Maria Iraburu Elizalde](#), [Jesus Urman](#), [Chaobo Chen](#), [Francisco Javier Cubero](#), [Leonard J Nelson](#), [Bruno Sangro](#), [María Luz Martínez-Chantar](#), [Jesús María Banales](#), [Jose Marin](#), [Carmen Berasain](#), [Maite G Fernandez-Barrena](#), [Matías A Avila](#)

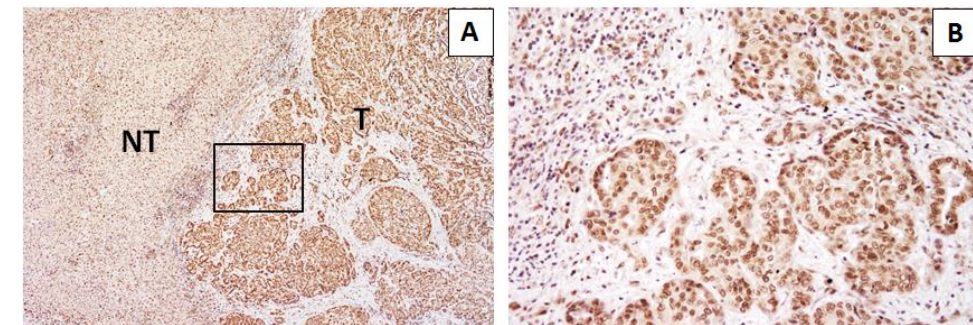


Expression levels of epigenetic modifiers G9a, DNMT1 and UHRF1 significantly increased in tumor tissue

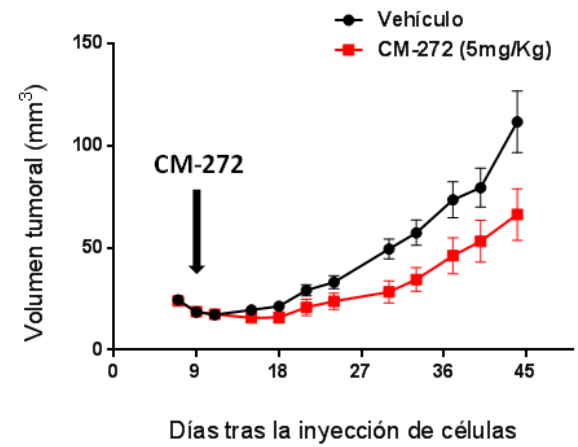
Synergistic antiproliferative effect over CCA cells by simultaneous inhibition of G9a and DNMT1.

CM-272

- Strong antiproliferative activity on different CCA cell lines
- Synergistic effect with different chemotherapeutics in the inhibition of CCA cells proliferation: Cisplatin, Mcl-1 Inhibitor and ErbB Inhibitors.
- In vivo potent inhibition of CCA growth



G9a expression in iCCA (T= tumoral tissue, NT= non tumoral tissue)



Epigenetics and Gastric Cancer 1/5

Oncogene (2009) 28, 184–194

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www.nature.com/onc

ORIGINAL ARTICLE

Hypoxic silencing of tumor suppressor *RUNX3* by histone modification in gastric cancer cells

SH Lee^{1,3}, J Kim^{2,3}, W-H Kim² and YM Lee¹

Hypoxia → ↑ **G9a (HMT)** → ↑ H3K9me → ↓ **RUNX3 (TSG)**

Cancer Res. 2005 Jun 1;65(11):4809-16.

Loss of RUNX3 expression significantly affects the clinical outcome of gastric cancer patients and its restoration causes drastic suppression of tumor growth and metastasis.

Wei D¹, Gong W, Oh SC, Li Q, Kim WD, Wang L, Le X, Yao J, Wu TT, Huang S, Xie K.



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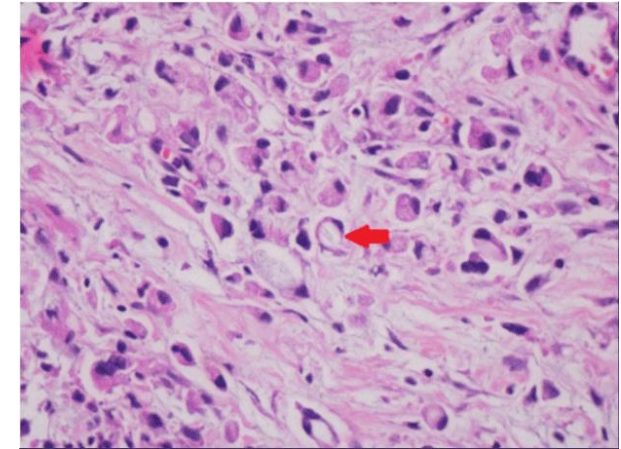
Epigenetics and Gastric Cancer 2/5

Mol Med Rep. 2013 Sep;8(3):942-8. doi: 10.3892/mmr.2013.1566. Epub 2013 Jul 2.

Expression of DNMTs and genomic DNA methylation in gastric signet ring cell carcinoma.

He M¹, Fan J, Jiang R, Tang WX, Wang ZW.

↑ ↑ ↑ **DNMT1** in SRC tissue compared with matched mucosal tissue.



Positive expression of DNMT1 ↔ N+ and late TNM stages of SRC

Epigenetics and Gastric Cancer 3/5

The FASEB Journal • Research Communication

Regulation of UHRF1 by miR-146a/b modulates gastric cancer invasion and metastasis

Lin Zhou,^{1,2} Xiaodi Zhao,¹ Yanan Han,¹ Yuanyuan Lu,¹ Yulong Shang, Changhao Liu, Ting Li, Zhian Jin, Daiming Fan,³ and Kaichun Wu³

UHRF1 expression in GC and adjacent nontumor tissues

Tissue	n	Expression level of UHRF1				P
		-	+	++	+++	
Nontumor	72	37	24	7	4	<0.01
GC	106	17	31	41	17	

The χ^2 test was used to evaluate the significance of differences between the two groups.

. Association of UHRF1 expression in the tumor tissues with demographic and clinicopathologic characteristics in 106 patients with GC

Category	n	Expression of UHRF1 (n)				P	Correlation coefficient
		-	+	++	+++		
Sex					0.427	0.007 ^{ns}	
Male	67	9	23	24	11		
Female	39	8	8	17	6		
Age (yr)					0.351	0.135 ^{ns}	
<60	58	10	19	23	6		
≥60	48	7	12	18	11		
Differentiation					<0.05	0.286 ^{**}	
Well	8	5	1	2	0		
Moderate	46	7	16	17	6		
Poor	52	5	14	22	11		
Stage					<0.01	0.334 ^{***}	
I-II	34	11	12	8	3		
III-IV	72	6	19	33	14		
Lymph node metastases					<0.05	0.322 ^{***}	
0	25	8	10	6	1		
≥1	81	9	21	35	16		
Metastases to other organs					<0.05	0.249 [*]	
Present	7	0	1	2	4		
Absent	99	17	30	39	13		

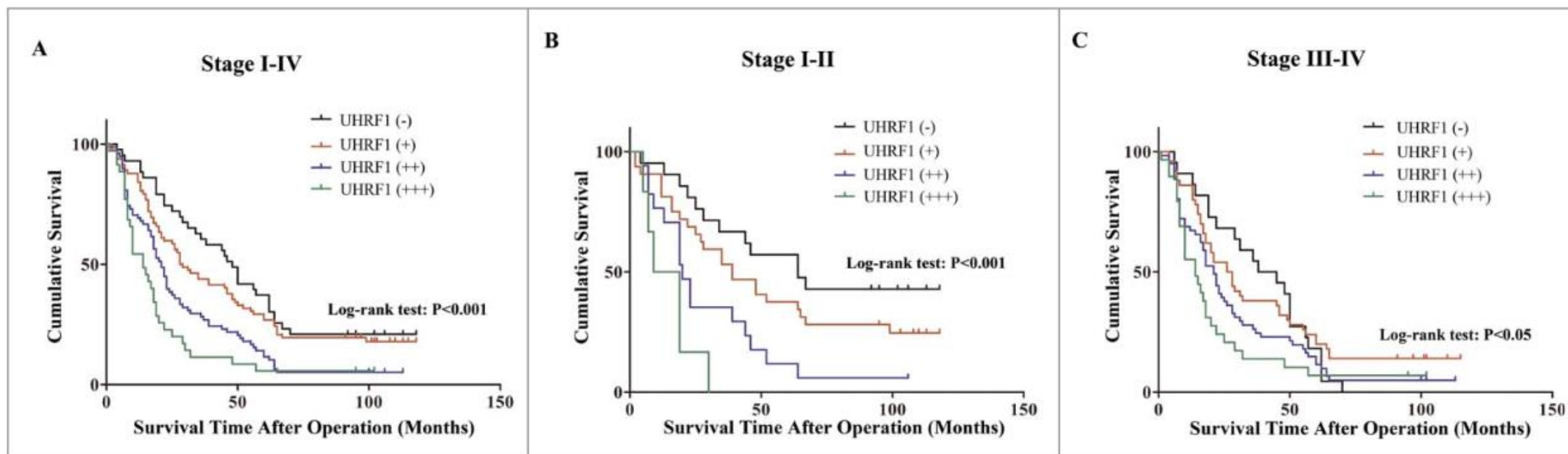


Epigenetics and Gastric Cancer 4/5

UHRF1 promotes proliferation of gastric cancer via mediating tumor suppressor gene hypermethylation

Lin Zhou^{1,4,*}, Yulong Shang^{2,4}, Zhi'an Jin^{3,4}, Wei Zhang^{4,4}, Chunlei Lv¹, Xiaodi Zhao², Yongqiang Liu¹, Naiyi Li¹, and Jie Li^{1,2,4*}

Kaplan–Meier survival curves of GC patients with different level of UHRF1 expression stratified by the TNM stage of the tumor (log-rank test).



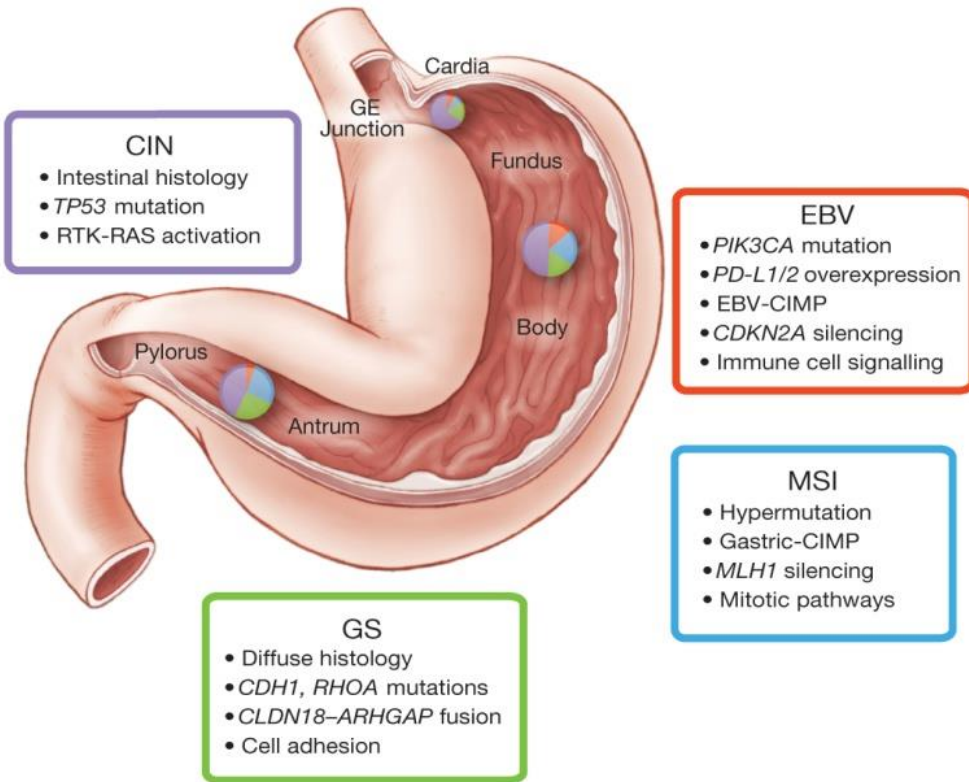
Hypermethylation of 7 TSG (CDKN2A, RUNX3, CDX2, DOXO4, PPARG, BRCA1 e PML)

Epigenetics and Gastric Cancer 5/5

nature

Comprehensive molecular characterization of gastric adenocarcinoma

The Cancer Genome Atlas Research Network*



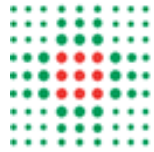
EBV- positive tumours had a **higher prevalence of DNA hypermethylation** than any cancers reported by TCGA

All EBV-positive tumours clustered together and **exhibited extreme CIMP** (CpG island methylator phenotype)

All EBV-positive tumours assayed displayed **CDKN2A (p16INK4A) promoter hypermethylation**, but lacked the **MLH1** hypermethylation

Key features of gastric cancer subtypes

RESEARCH PROJECT



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EMILIA-ROMAGNA
Azienda Ospedaliero - Universitaria di Modena

Collection of human Gastric Cancer (GC) samples and clinico-pathological data (Jan 2000 – Feb 2019) → 100 samples

Preparation of tissue samples for histological examination

Inclusion Criteria

Patients with histological diagnosis of **resectable or advanced gastric cancer**

Availability of **surgical (or bioptic) specimens** for the analysis

No chemo- and/or radiotherapy prior to surgery

Written informed consent

Exclusion Criteria

Patients with uncertain histological diagnosis of gastric cancer

Chemo- and/or radiotherapy prior to surgery

Refusal of consent

INITIAL PHASES



cima

CENTRO DE INVESTIGACIÓN MÉDICA APLICADA
UNIVERSIDAD DE NAVARRA

IHC Analysis
of G9a, DNMT1 and UHRF1



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SUBSEQUENT PHASES



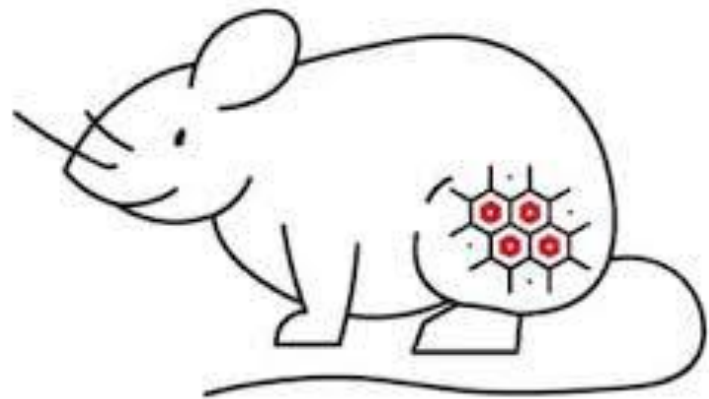
IN VITRO (GC Cell lines)

Evaluation of **G9a, DNMT1 and UHRF1** expression

Evaluation of **CM-272** effect

Evaluation of potential synergistic effect with other anticancer agents

Evaluation of **CM-272** effect



IN VIVO (mouse xenograft models)



Aims

Primary objective:

Expression evaluation of G9a, DNMT1 and UHRF1 in GC patients

- Provide a better characterization of methylation in GC
- Identify potential novel therapeutic targets providing a background for further investigations in order to develop more specific therapeutic strategies (*Epi-drugs*)

Secondary objective:

Correlation of profile expression of G9a, DNMT1 and UHRF1 in GC **with clinico-pathological features** and **survival parameters**

- Identify possible prognostic and predictive factors

Take-home messages

- **Epigenetic mechanisms** are emerging as **attractive therapeutic target** in solid tumors
- **Epi-drugs** are already being tested
- Given the crosstalk between chromatin marks, **simultaneous targeting** of different epigenetic modifiers **may improve therapeutic efficacy**
- **CM-272** as emerged as a **promising** dual (G9a and DNMT1) targeting agent (HCC, CCA)
- **Epigenetic dysregulation** plays a crucial role in **GC development**
- **G9a and DNMT1** seems **potential epi-target** also in the context of **GC**, but few data are available
- Further investigation are needed



Grazie!

Acknowledgments

Prof. Stefano Cascinu
Dr. Andrea Casadei Gardini
Dr.ssa Giulia Orsi
Dr. Andrea Spallanzani
Dr.ssa Kalliopi Andrikou

CIMA, Pamplona:
Prof. Matías Ávila