



# MARCAR

BIOMMARKERS & MOLECULAR TUMOUR CLASSIFICATION  
FOR NON-GENOTOXIC CARCINOGENESIS



efpia

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# What patients need What Europe needs

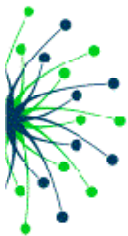
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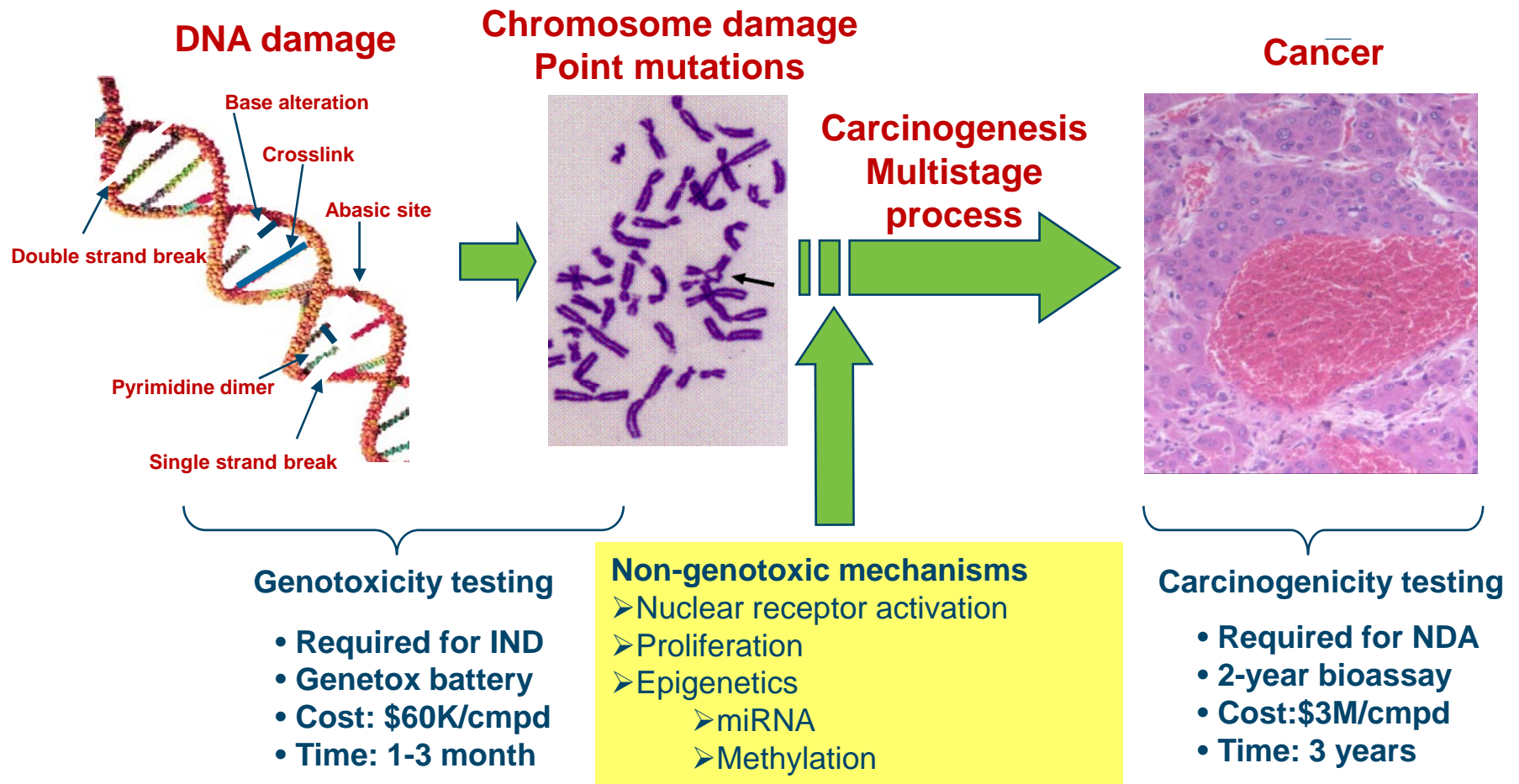
- Improved drug safety
- Increased efficiency of drug development
- Earlier detection of undesired effects of candidate drugs during development

## MARCAR focus on **NON-GENOTOXIC CARCINOGENESIS**

- Identify BIOMARKERS:
  - help predict cancer risk more accurately at a very early stage
  - speed up development & increase drug safety for patients



# Why focus on Non-Genotoxic Carcinogenesis (NGC)?



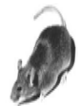
Schematic Figure adapted from Ellinger-Ziegelbauer et al., (2009) *Toxicol Lett.* (Elsevier) 186:36-44 with authors permission

- **No sufficiently accurate or well-validated short-term assays to identify NGC**
- **Need early mechanism-based biomarkers for the design of more predictive tests & improved cancer risk assessment**

# What we will do



**MARCAR will search for biological clues (BIOMARKERS) to enable EARLY preclinical DETECTION of drug-induced tumor formation**



*2 year rodent cancer bioassay supporting clinical development and drug registration*

## **IDENTIFY BIOMARKERS USING INNOVATIVE TECHNOLOGIES AND MECHANISTIC MODELS:**

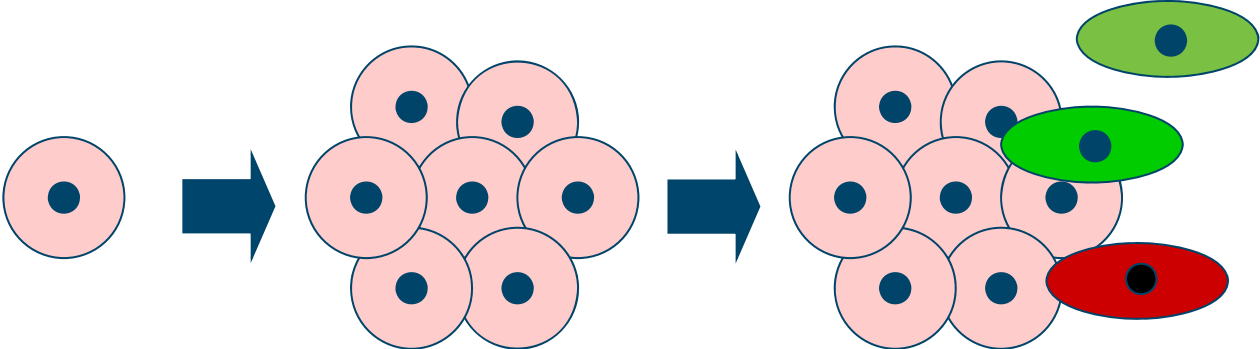
- **EPIGENETICS**; microRNA; BIOINFORMATICS
- MOUSE MODELS CONTAINING HUMAN GENES IMPORTANT FOR NGC
  - TRANSLATIONAL *IN VITRO* CELL-BASED MODELS
  - NON-INVASIVE IMAGING & REPORTER MODELS

**ASSESS PREDICTIVE VALUE OF NOVEL BIOMARKERS & RELEVANCE FOR HUMANS**

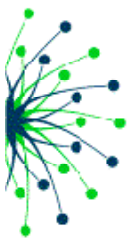
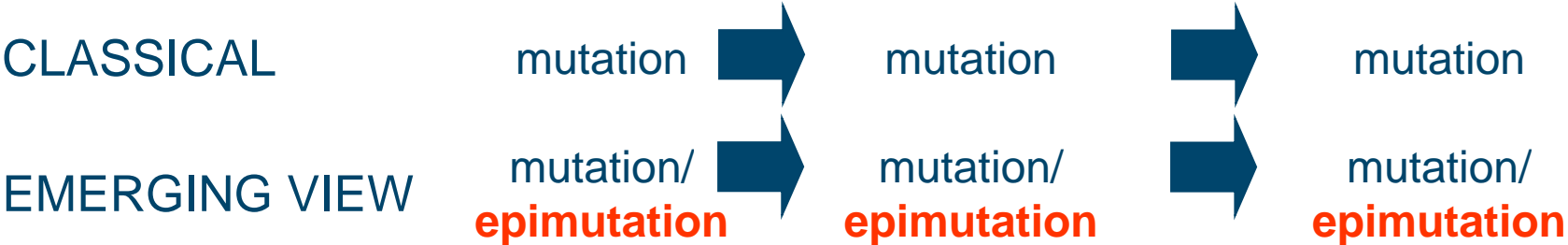
# Emerging importance of epigenetics for cancer biology



INITIATION                      PROMOTION                      PROGRESSION

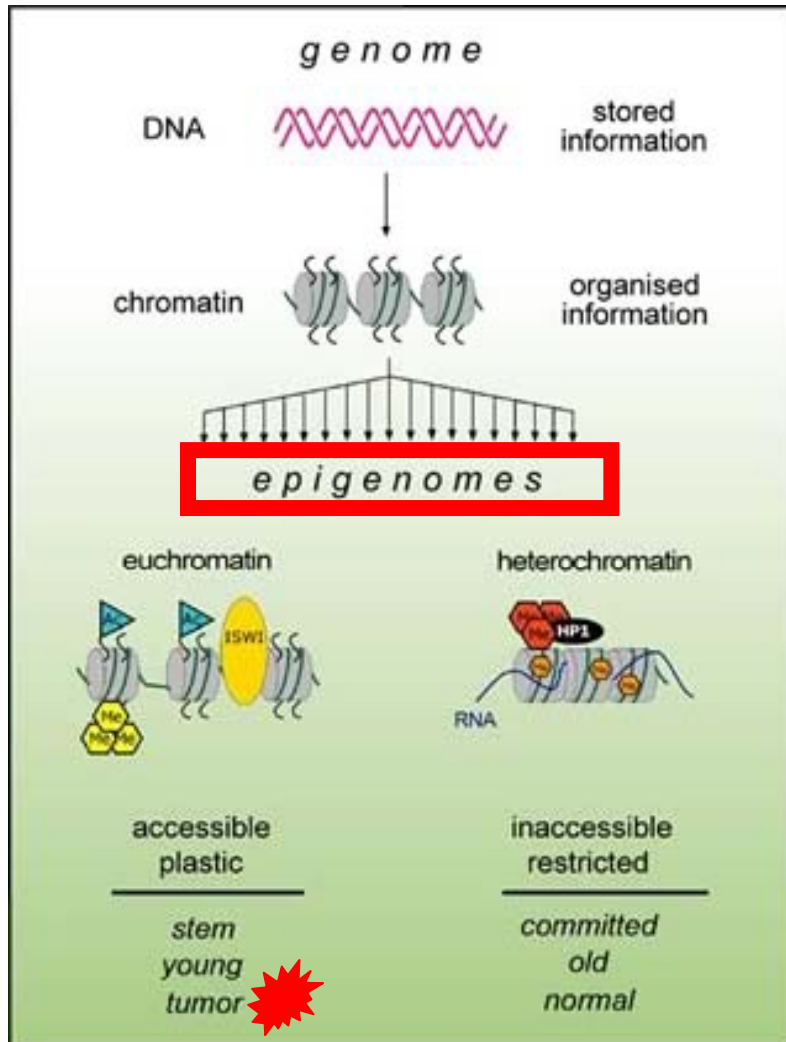


Clonal expansion                      Acquired characteristics  
 •metastasis, angiogenesis

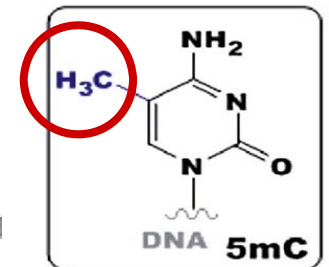
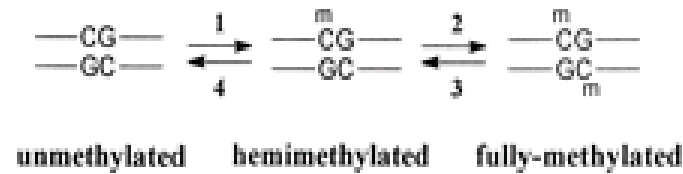


**EPIMUTATIONS:  
 = Paradigm shift in established models for tumour progression**

# EPIGENETIC CODES – opportunity to identify novel biomarkers for early cancer detection



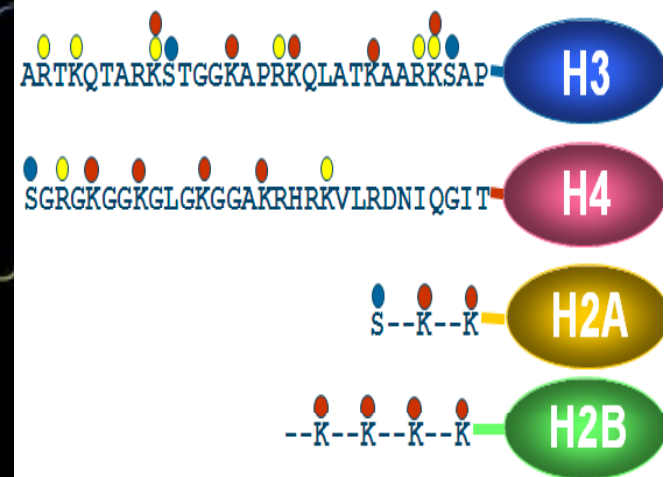
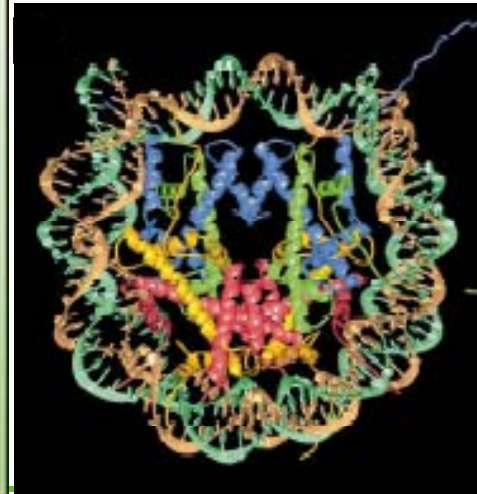
GENETIC CODE (A,C,T,G)



5-methylcytosine



EPIGENETIC CODES



[www.epigenome.eu](http://www.epigenome.eu)

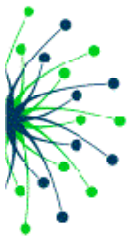
Epigenome Network of Excellence (NoE)  
EU Framework 6 Programme (FP6)

Luger et al., (1997)  
Nature 389:251-260

# Expected outcomes & benefits to patients



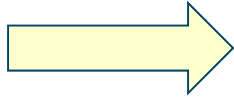
- Improved scientific basis for risk assessment of carcinogenic potential of novel medicines
- Novel early biomarkers for more reliably predicting later cancer development
- Increased efficiency of drug development - fewer delays/attrition during late-phase
- Improved preclinical safety assessment prior to clinical trials & potential translation of early cancer biomarkers to humans



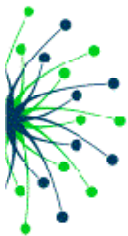


# Added value of consortium



- New knowledge + biomarkers can be more rapidly derived by sharing consortium expertise, technologies, models, and archives
  - Extensive collection of archived samples and data from preclinical studies (3Rs) 
  - Opportunity for continuous regulatory input (via advisory board) on cancer risk assessment & biomarker qualification
  - Access to unique molecular profiling technologies and mechanistic models

Genotoxic carcinogens	2-Nitrofluorene Dimethylnitrosamine Aflatoxin B1 N-Nitrosomorpholine C.I Direct Black 4-(methylnitrosamino)-1-(3-pyridyl)-1-butanone 2-Acetylaminofluorene N-Nitrosopiperidine Methylenedianiline
Non-genotoxic carcinogens	Methapyriene HCl Thioacetamid Diethyl-stilbestrol Wy-14643 Piperonyl-butoxide Methylcarbamate Acetamide Dehydroepiandrosterone Ethionine Acetaminophen Cyproterone acetate Phenobarbital
Non-hepatocarcinogens	Cefuroxime Nifedipine Propranolol Clonidine Prazosin Ibuprofen Allyl alcohol 1,4-Dichlorobenzene

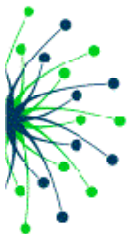




# Results/achievements



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- New technologies established for monitoring drug-induced perturbation of epigenome and microRNAs in rodent tissues (e.g. INTEGRATED GENOME-WIDE MOLECULAR PROFILING)
  - Identification of novel early candidate biomarkers for non-genotoxic carcinogenesis (e.g. NON-CODING RNAs & CANCER SIGNALLING PATHWAYS)
  - Novel insight into human relevance of candidate biomarkers using mouse models (e.g. CAR & PXR NUCLEAR RECEPTORS)



# Time and money



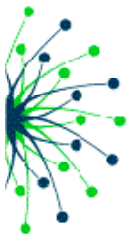
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## Financing

- IMI funding: € 6.049.578
- EFPIA in kind contribution: € 5.155.604
- Other contributions: € 2.114.051
- Total project cost: € 13.319.233

## Timing:

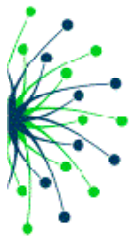
- Starting date: 01.01.2010
- Duration: 60 months



# Participants



- EFPIA
  - Novartis (*Coordinator*)
  - UCB Pharma SA
  - Bayer Schering Pharma AG
  - Lundbeck
  - Boehringer Ingelheim
- SME
  - CXR Biosciences Ltd., UK
- EMEA & FDA scientists
  - via ad-hoc advisory board
- Academic
  - University of Dundee, UK (*Managing Entity of IMI beneficiaries*)
  - Medizinische Universität, Vienna, Austria
  - Medical Research Council, Edinburgh, UK
  - Eberhard Karls Universität, Tübingen, Germany
  - Natural & Medical Sciences Institute, Reutlingen, Germany
  - Institut National de la Santé et de la Recherche, Montpellier, France



Further information

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[www.IMI-MARCAR.EU](http://www.IMI-MARCAR.EU)

[www.imi.europa.eu](http://www.imi.europa.eu)

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