

Molecular biomarkers of effect in the causality determination process: an important piece of the puzzle?

Valentina Bollati

EPIGET Lab

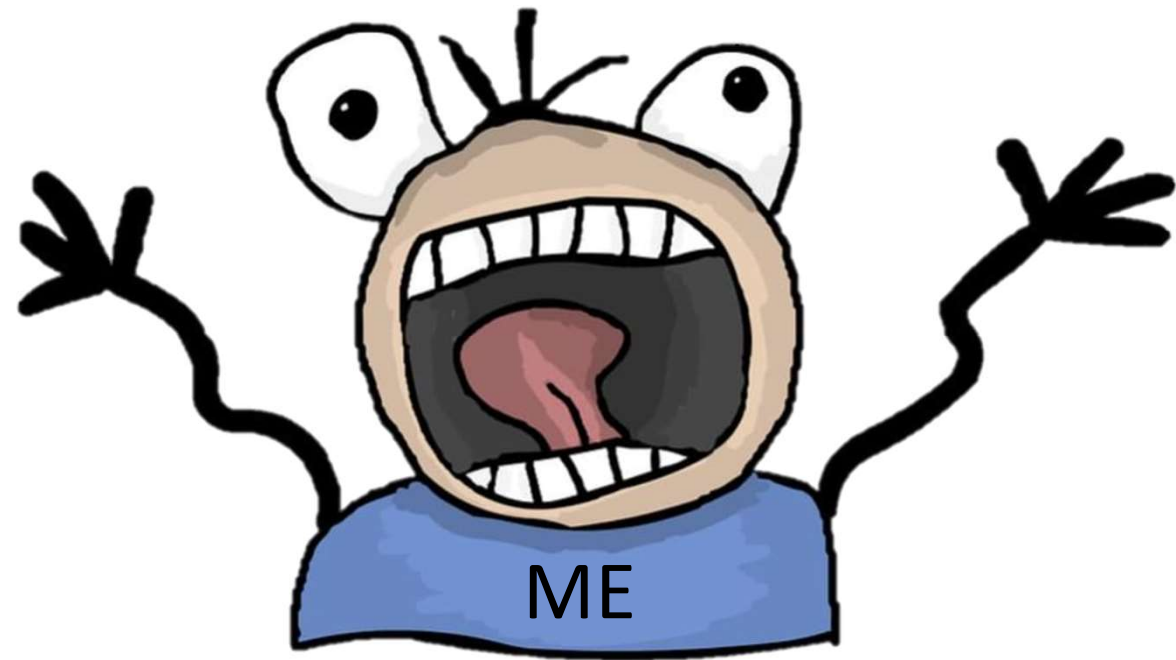
Department of Clinical Sciences and Community Health

University of Milan, Italy



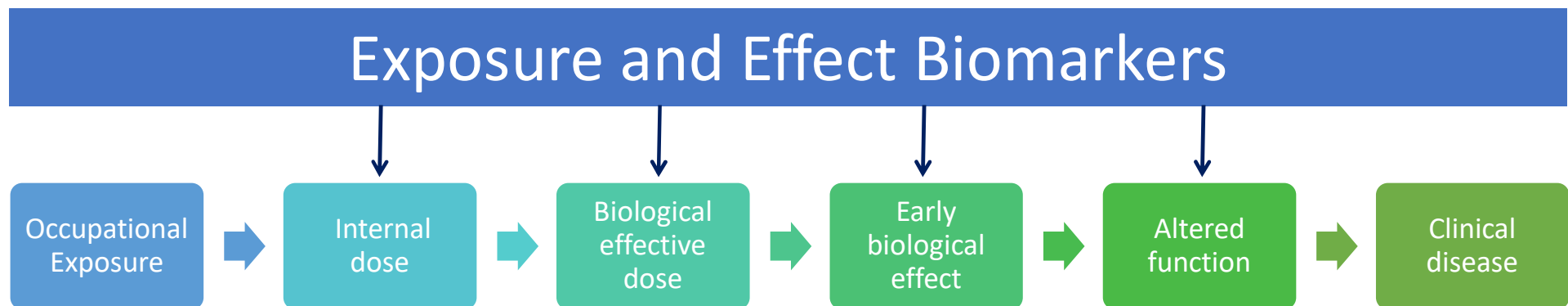
Graphical abstract of my talk

TOPIC OF THIS
PRESENTATION



«Biomarkers of effect»

Biomarkers of effect are the **quantifiable** changes that an individual endures, which **indicates an exposure** to a compound and may indicate a resulting **health effect**.



«Causality determination» in occupational health

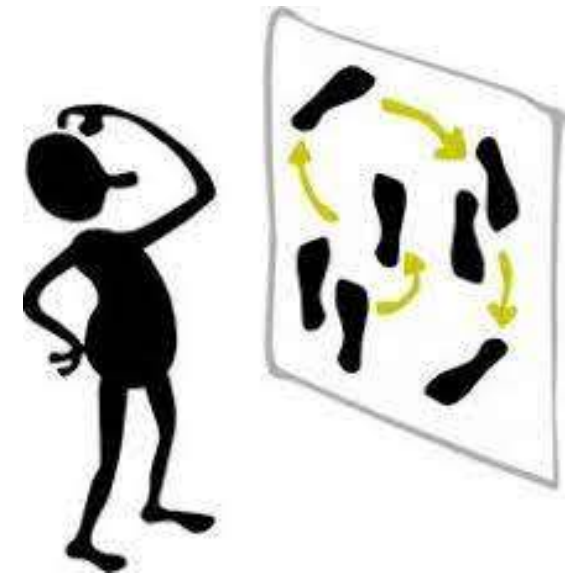
An **injury or illness** is considered by Occupational Safety and Health Administration to be “**work related**” if an event or exposure in the work environment either **caused or contributed** to the resulting condition or significantly aggravated a pre-existing condition.

<https://www.osha.gov>

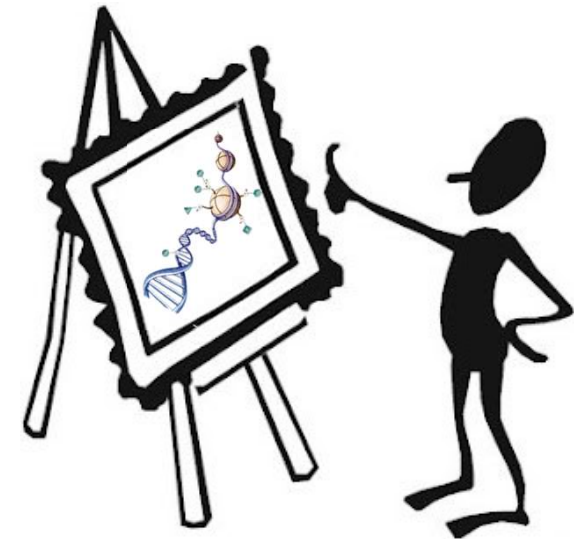


The challenge

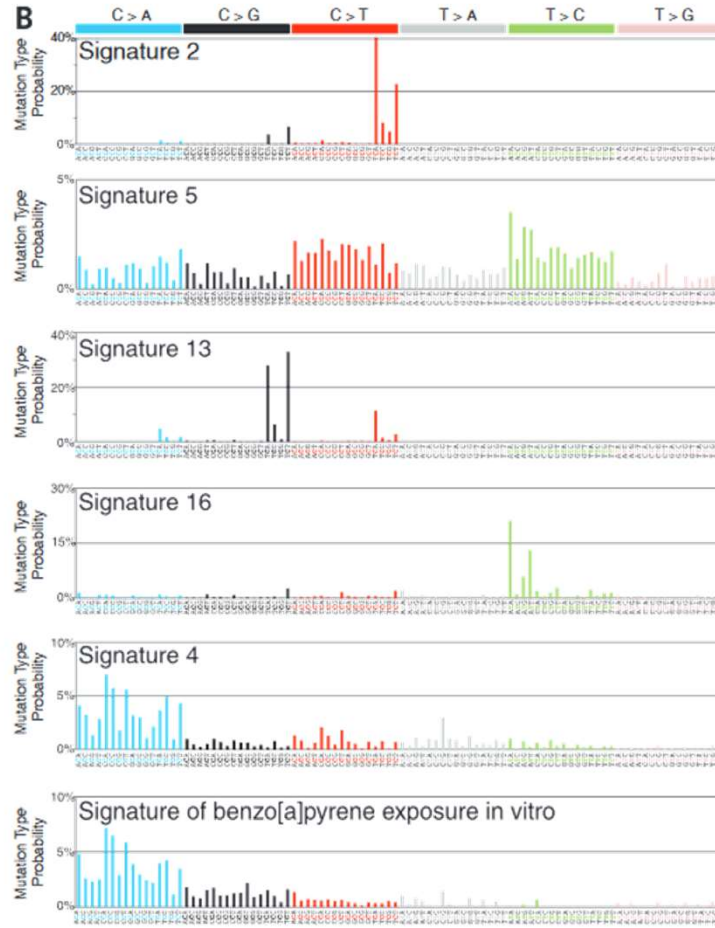
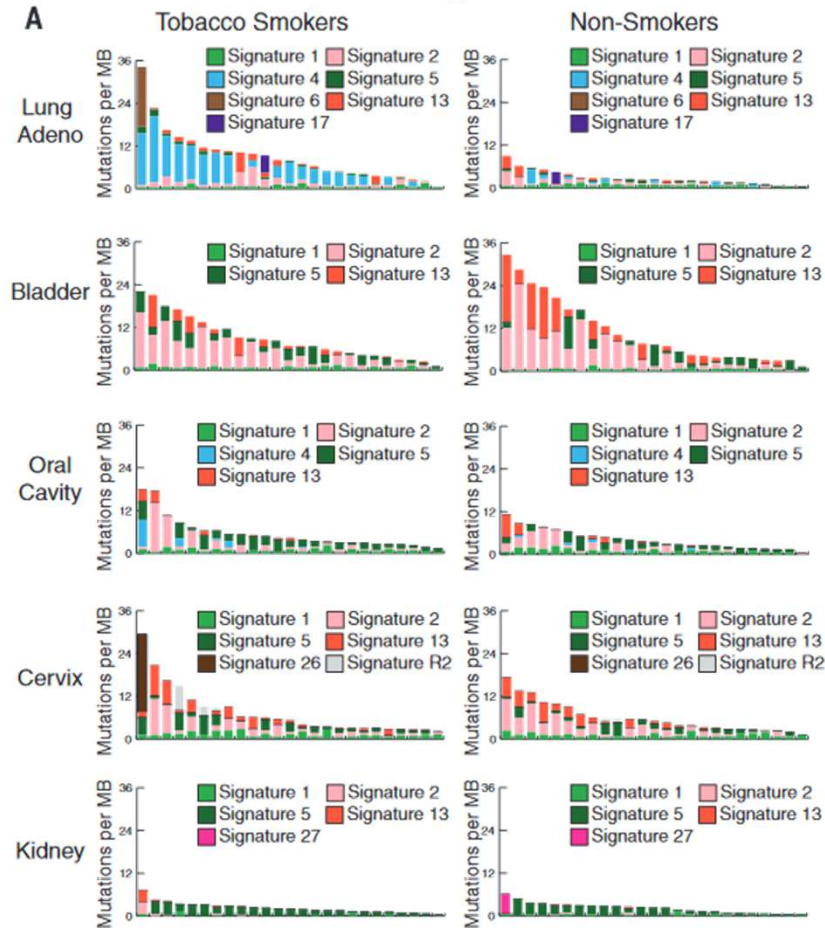
- Establishing a work-related cause of a disease is one of the most challenging aspects of occupational medicine.
- ALL diseases are multicausal
- Occupational diseases can be:
 - mainly caused by factors at work (e.g. mesothelioma)
 - diseases in which work plays a minor role in causation (e.g. back pain).



Environmental events
can be permanently
registered by our cells



Mutational signatures associated with tobacco smoking



Tobacco smoking increases the risk of at least 17 classes of cancer

Genome analyses of cancer tissues in:
 -2,490 tobacco smokers
 -1,063 never smokers

Alexandrov LB, et al. Mutational signatures associated with tobacco smoking in human cancer. *Science*. 2016 Nov;354(6312):618-622.

WEBINAR Settling a debate about theoretical bases of occupational disease



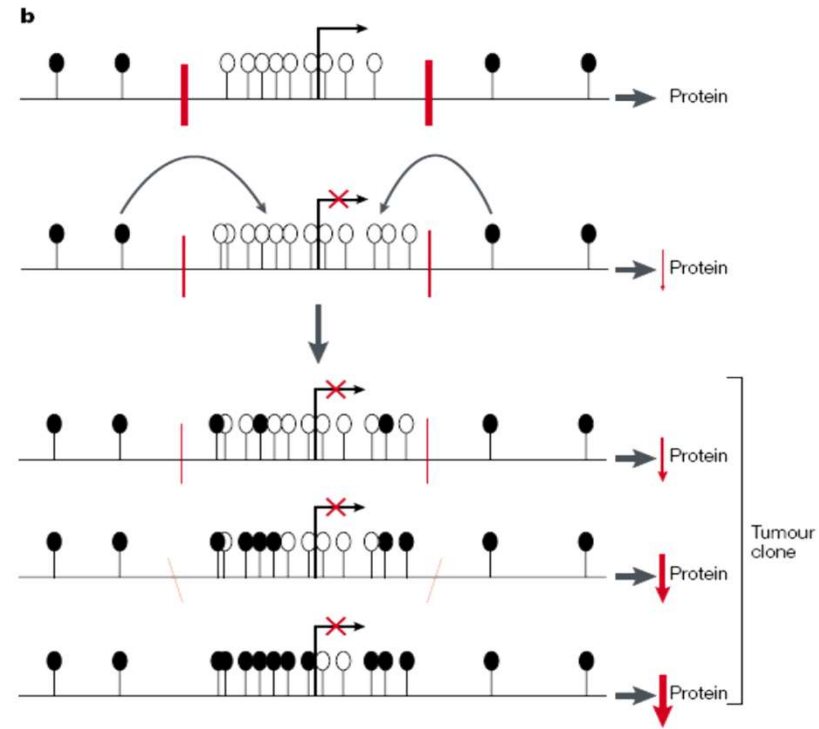
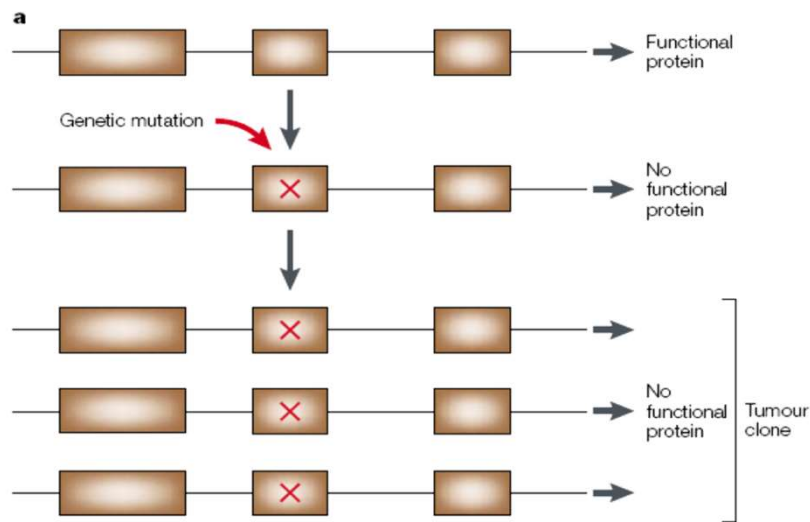
PROs and CONs

- Mutational signature is specific.
- Exposure signature was identified in cancer tissue, in the fully developed disease.
- Mutational signature is not dynamic by definition.

EPIGENETICS



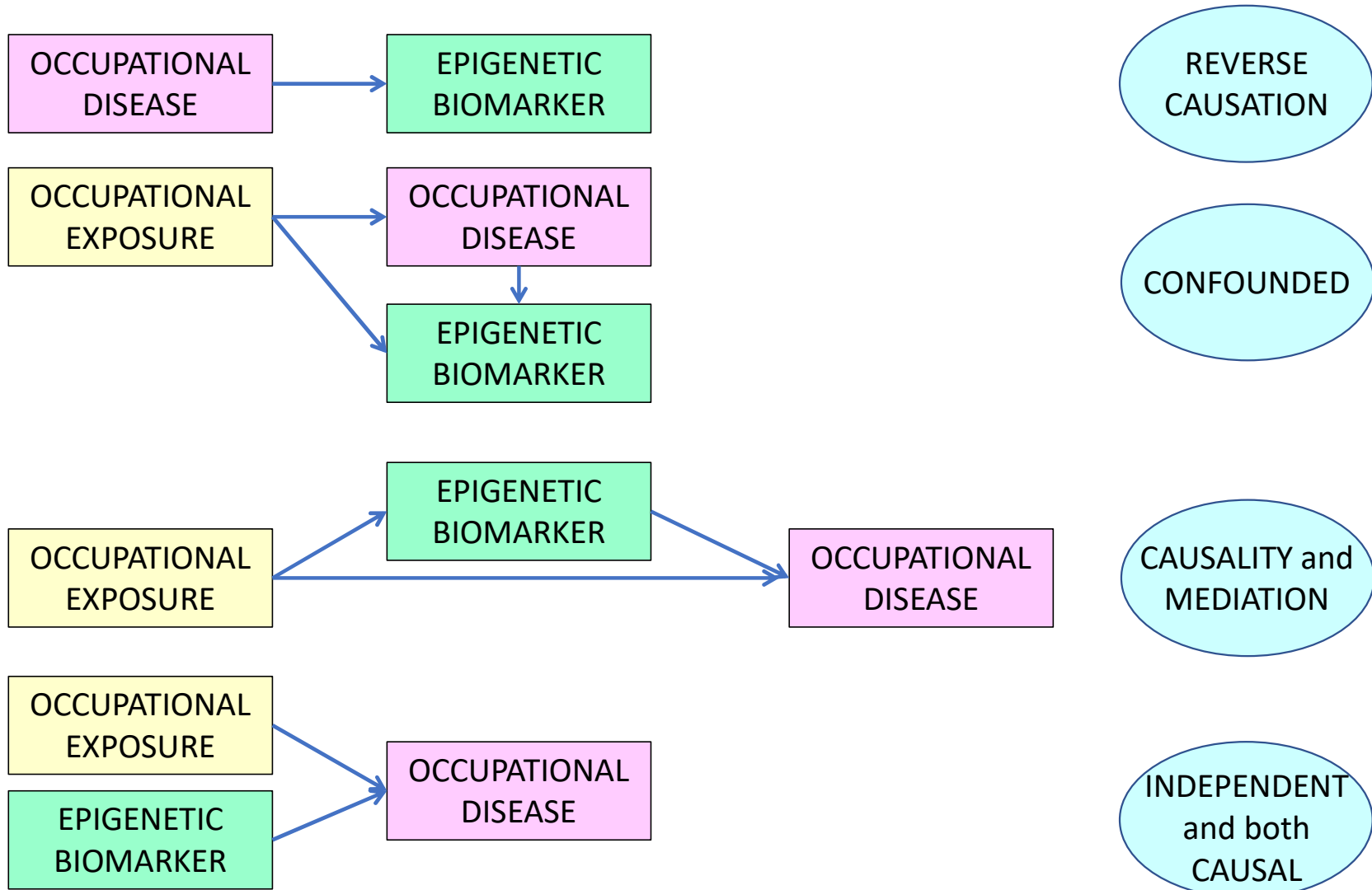
DNA mutations versus “epimutations”



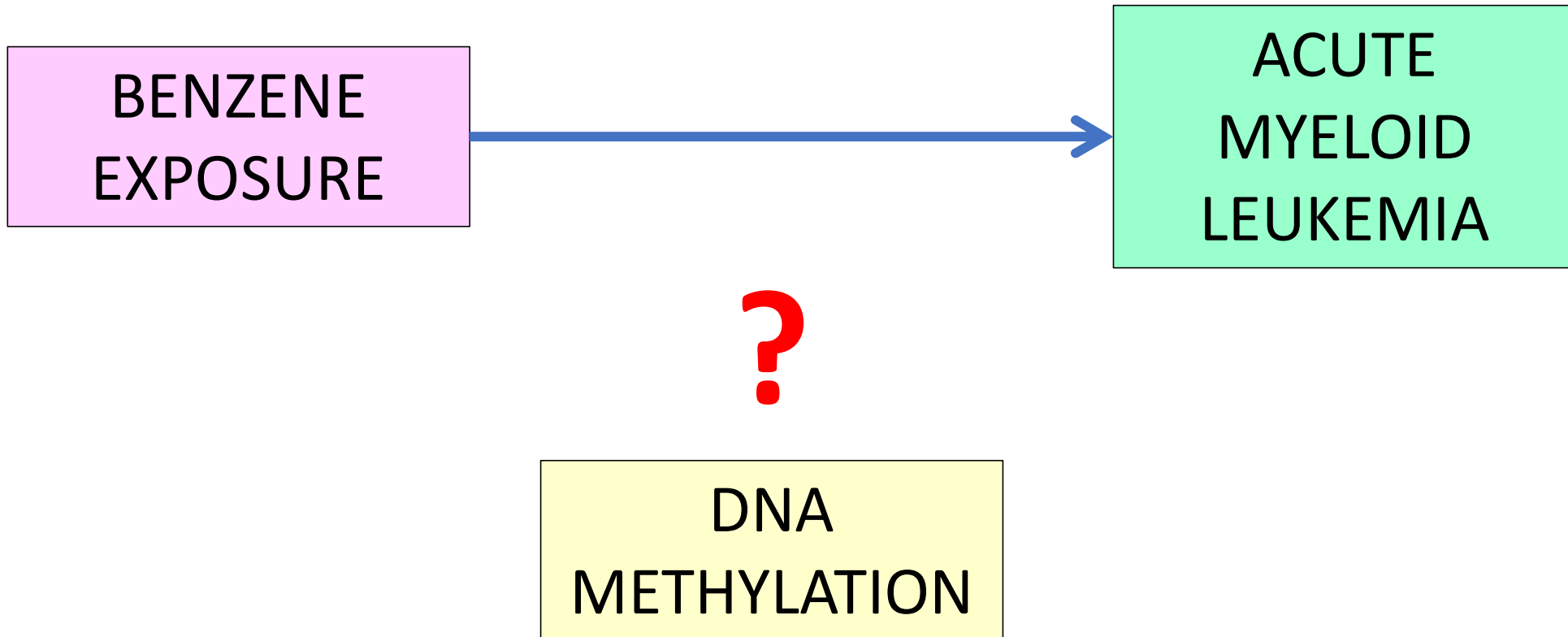
Jones & Baylin Nature Rev 3:415, 2002



The possible scenarios



An example



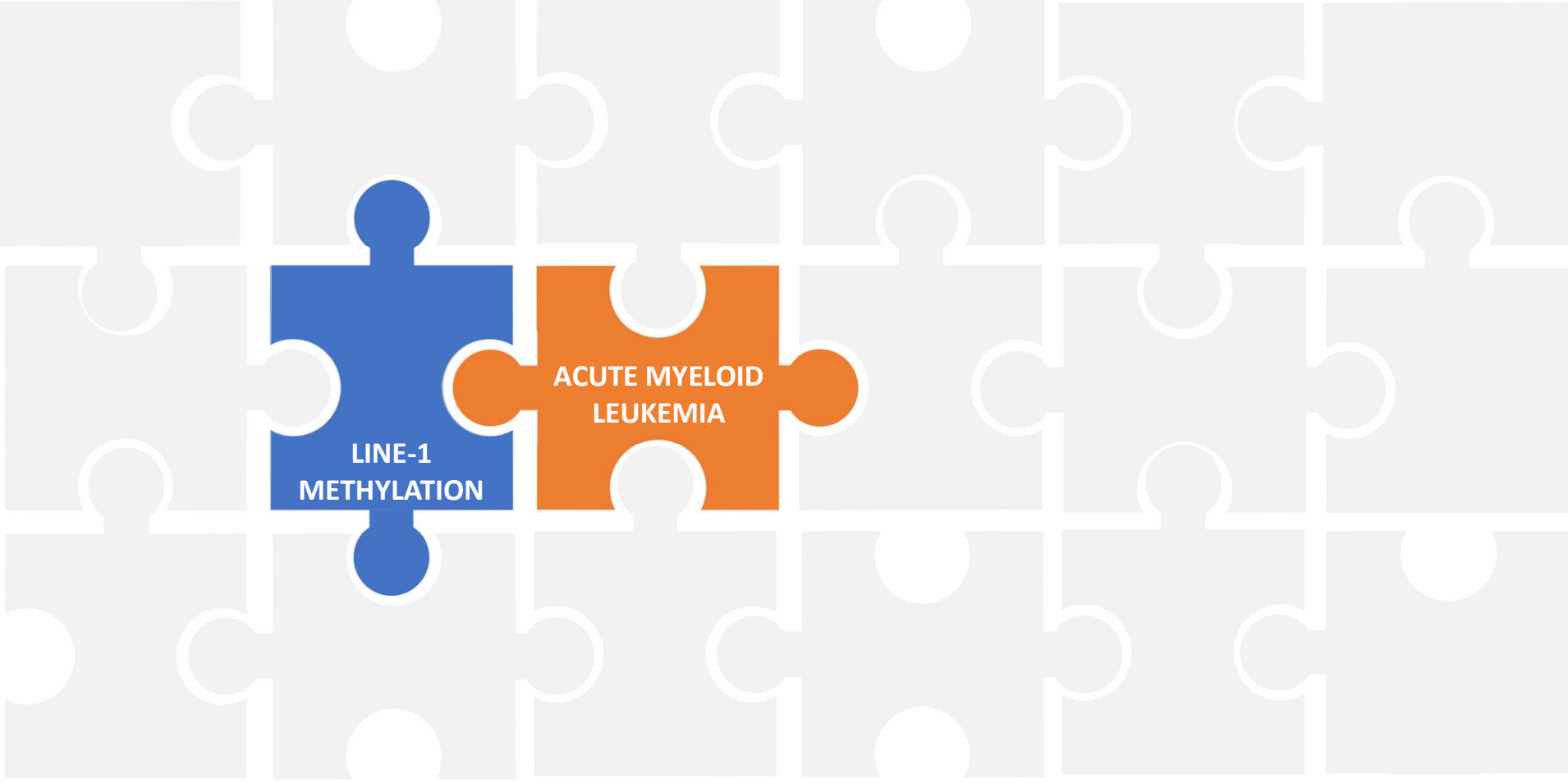
Benzene and Acute Myeloid Leukemia

- Benzene (C_6H_6) is the smallest and most stable aromatic hydrocarbon
- Benzene is an airborne pollutant of industrial and general environment
- Benzene is a well known human leukemogen (group 1 IARC) (IARC, 1987)
- Benzene exposure in human populations is causally related to the development of AML (Savitz and Andrews, 1997).
- An increased risk of leukemia was reported in different occupational settings characterized by high exposure, but conflicting results for low-exposures



DNA methylation and Acute myeloid leukemia

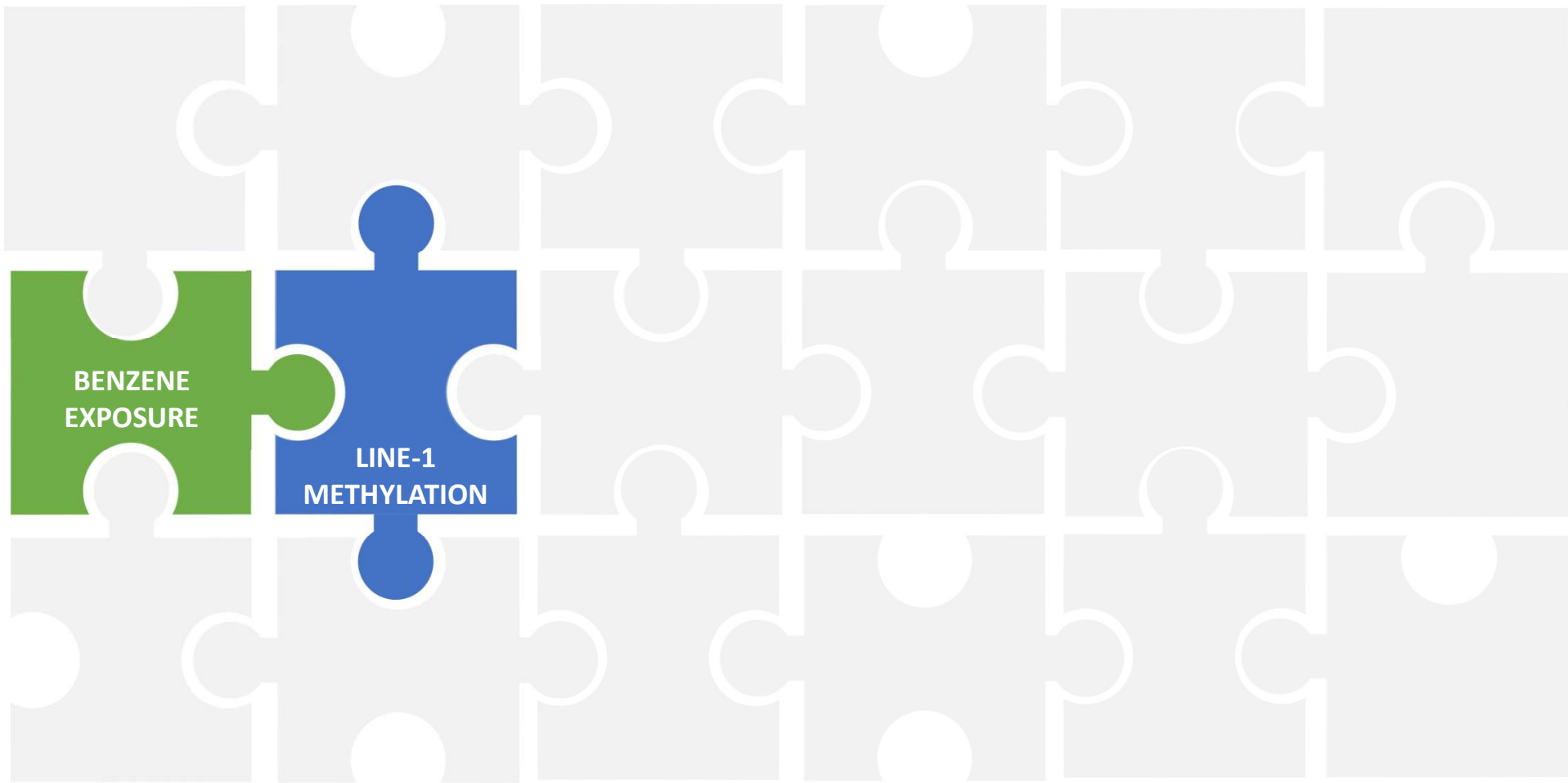
- Leukemia cells exhibit an abnormal methylation pattern.
- The CpG islands in the promoters of some genes become hypermethylated.
- The global genome of cancer cells undergoes hypomethylation, up to 20-60% less than normal cells.
- **LINE-1 methylation** is highly correlated with global methylation, thus undergoes hypomethylation in cancer cells.



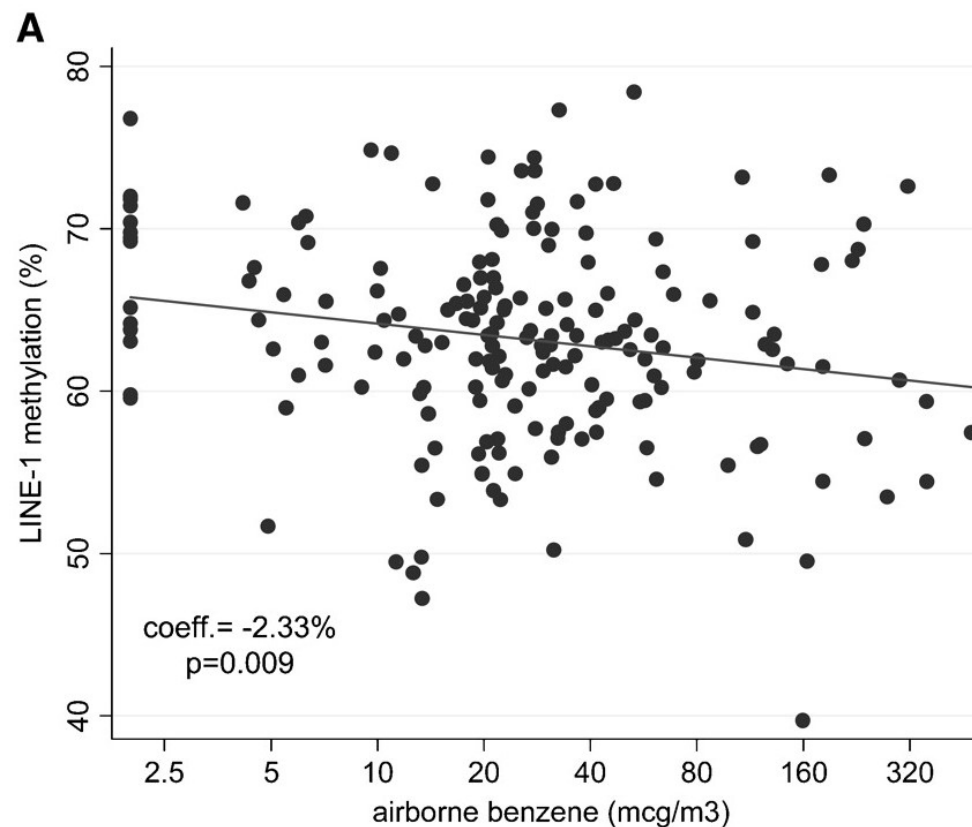
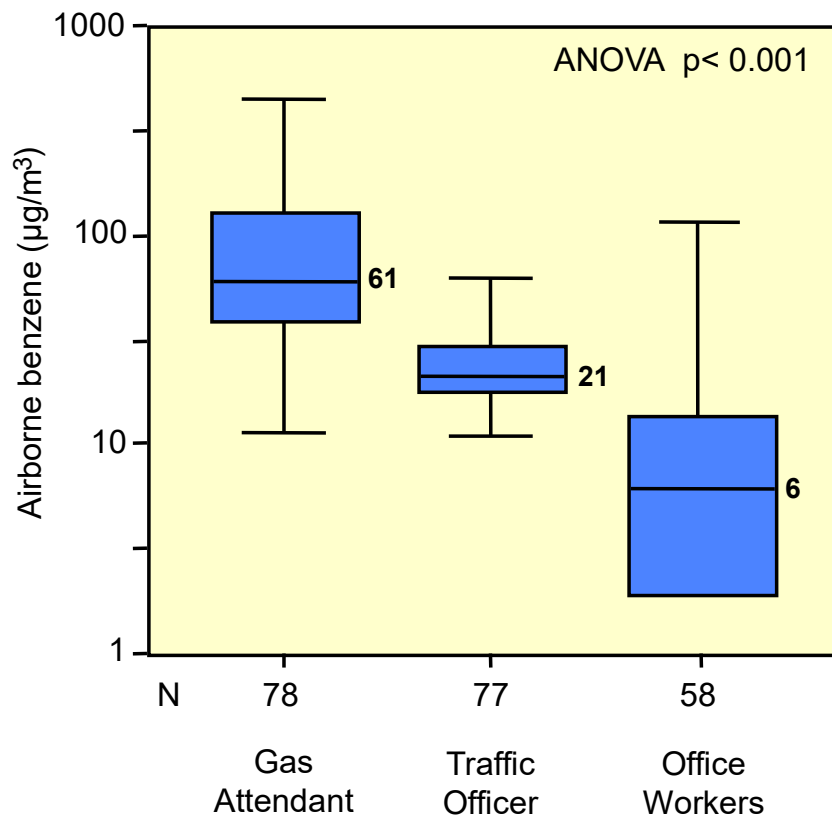
LINE-1
METHYLATION

ACUTE MYELOID
LEUKEMIA



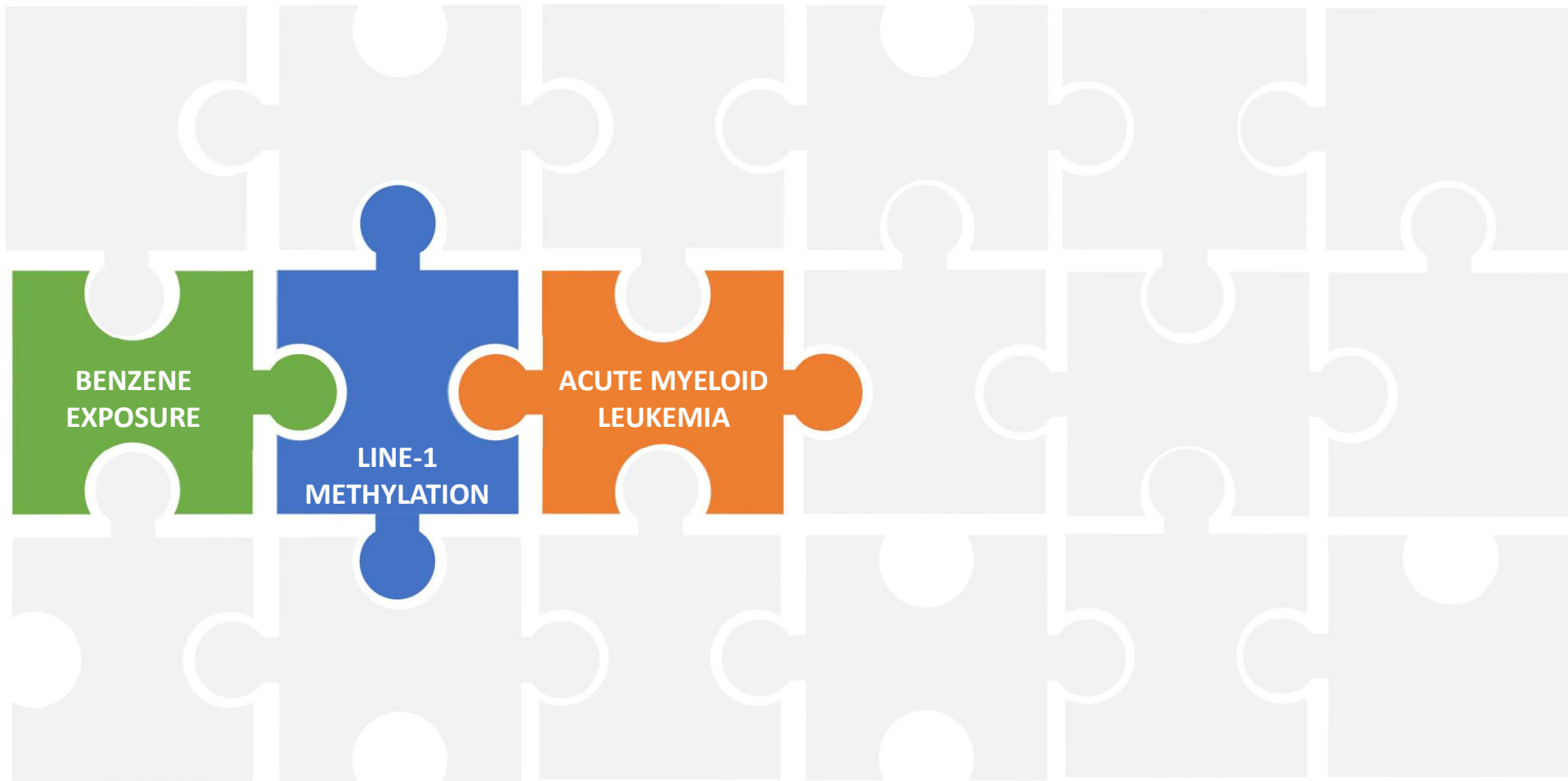


Benzene exposure and LINE-1 methylation



Bollati et al. Cancer Res 2007;67:876-880





HOWEVER...



Not only AML...

Guarera et al. *Clinical Epigenetics* (2015) 7:133
DOI 10.1186/s13148-015-0164-3

Clinical Epigenetics

RESEARCH

Open Access

Gene-specific DNA methylation profiles and LINE-1 hypomethylation are associated with myocardial infarction risk



Simonetta Guarera^{1,2†}, Giovanni Florito^{1,2†}, N. Charlotte Onland-Moret³, Alessia Russo^{1,2}, Claudia Agnoli⁴, Alessandra Allione^{1,2}, Cornelia Di Gaetano^{1,2}, Amalia Mattiello⁵, Fulvio Ricceri⁶, Paolo Chiodini⁷, Silvia Polidoro¹, Graziella Frasca⁸, Monique W. M. Verschuren^{3,9}, Jolanda M. A. Boer⁹, Licia Iacoviello¹⁰, Yvonne T. van der Schouw³, Rosario Tumino⁸, Paolo Vineis^{1,11}, Vittorio Krogh⁴, Salvatore Panico⁵, Carlotta Sacerdote⁶ and Giuseppe Matullo^{1,2*}



ORIGINAL RESEARCH
published: 05 October 2015
doi: 10.3389/fped.2015.01053

Analysis of Global and Local DNA Methylation Patterns in Blood Samples of Patients With Autism Spectrum Disorder

María Victoria García-Ortiz^{1,2,3†}, María José de la Torre-Aguilar^{1,4†}, Teresa Morales-Ruiz^{1,2,3}, Antonio Gómez-Fernández^{1,4}, Katherine Flores-Rojas^{1,4,5}, Mercedes Gil-Campos^{1,4,6*}, Pilar Martín-Borreguero^{1,7}, Rafael R. Ariza^{1,4,3}, Teresa Roldán-Arjona^{1,4,3} and Juan Luis Pérez-Navero^{1,4,3}

WEBINAR Settling a debate about theoretical bases of occupational disease



OPEN ACCESS Freely available online

PLOS ONE

LINE-1 Hypomethylation in Blood and Tissue Samples as an Epigenetic Marker for Cancer Risk: A Systematic Review and Meta-Analysis



Martina Barchitta¹, Annalisa Quattrocchi¹, Andrea Maugeri¹, Manlio Vinciguerra^{2,3*}, Antonella Agodi^{1*}

1 Department GF Ingrassia, University of Catania, Catania, Italy, **2** University College London, Institute for Liver and Digestive Health, Royal Free Campus, London, United Kingdom, **3** Gastroenterology Unit, Department of Medical Sciences, IRCCS Casa Sollievo della Sofferenza, San Giovanni Rotondo, Italy

scientific reports

Check for updates

OPEN Aging-associated distinctive DNA methylation changes of LINE-1 retrotransposons in pure cell-free DNA from human blood

Wardah Mahmood^{1,2}, Lars Erichsen^{1,2}, Pauline Ott¹, Wolfgang A. Schulz², Johannes C. Fischer³, Marcos J. Arauzo-Bravo^{4,5}, Marcelo L. Bendhack⁶, Mohamed Hassan^{7,8} & Simeon Santourlidis^{1,2*}





Not only benzene...

Rapid DNA Methylation Changes after Exposure to Traffic Particles

Andrea Baccarelli^{1,2}, Robert O. Wright^{2,3}, Valentina Bollati¹, Letizia Tarantini¹, Augusto A. Litonjua³, Helen H. Suh², Antonella Zanobetti², David Sparrow⁴, Pantel S. Vokonas⁴, and Joel Schwartz²

¹Laboratory of Environmental Epigenetics, Department of Environmental and Occupational Health, University of Milan & IRCCS Maggiore Hospital, Mangiagalli and Regina Elena Foundation, Milan, Italy; ²Department of Environmental Health, Harvard School of Public Health; ³Channing Laboratory, Department of Medicine, Brigham and Women's Hospital, Harvard Medical School; and ⁴VA Normative Aging Study, Veterans Affairs Boston Healthcare System and the Department of Medicine, Boston University School of Medicine, Boston, Massachusetts

Received: 12 December 2020 | Revised: 11 June 2021 | Accepted: 14 July 2021
DOI: 10.1002/em.22451

RESEARCH ARTICLE



Global and gene-specific promoter methylation, and micronuclei induction in lead-exposed workers: A cross-sectional study

Kan Wang^{1,2} | Yu Meng¹ | Tuanwei Wang¹ | Yuting Tu¹ | Shiyang Gong¹ |
Guanghui Zhang³ | William Au⁴ | Zhaolin Xia¹

Mediterranean Diet and Particulate Matter Exposure Are Associated With LINE-1 Methylation: Results From a Cross-Sectional Study in Women

Martina Barchitta¹, Andrea Maugeri¹, Annalisa Quattrocchi¹, Germana Barone², Paolo Mazzoleni²,
Alfio Catalfo³, Guido De Guidi^{3,4}, Maria Giovanna Iemmolo⁵, Nunzio Crimi⁵ and Antonella Agodi^{1*}

¹Department of Medical and Surgical Sciences and Advanced Technologies "GF Ingrassia", University of Catania, Catania, Italy

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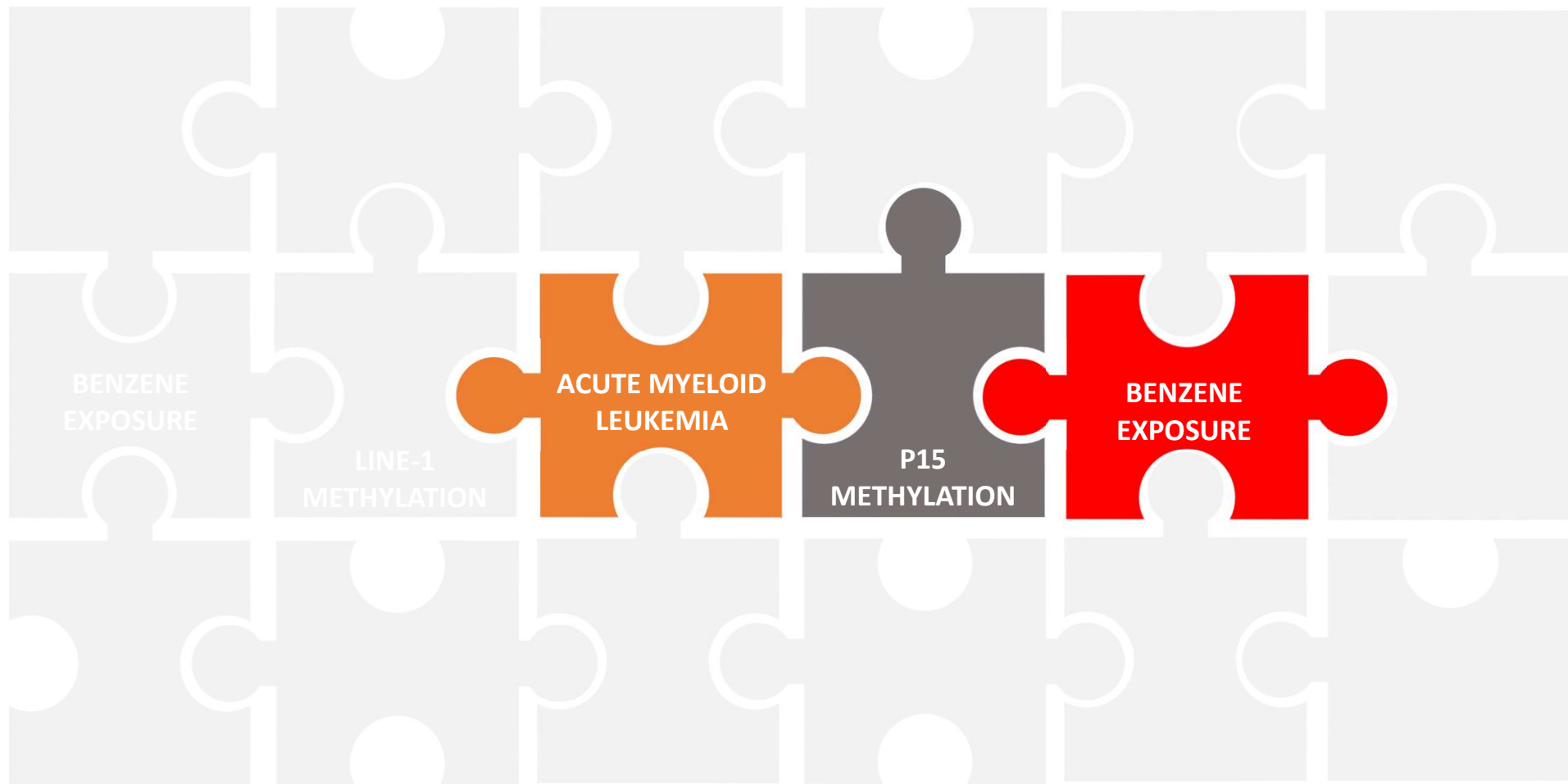
³Department of Chemical Science, Section of Photochemistry and Photobiology, University of Catania, Catania, Italy

⁴Research Centre for the Analysis, the Monitoring and Methodology for Environmental Risk Assessment, University of Catania, Catania, Italy

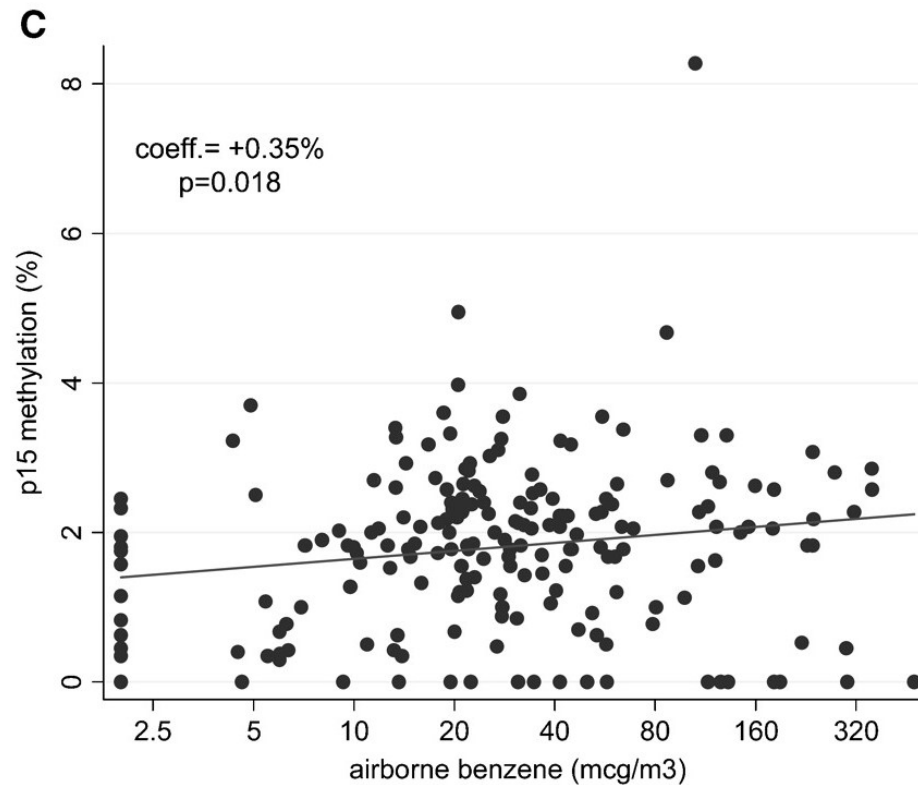
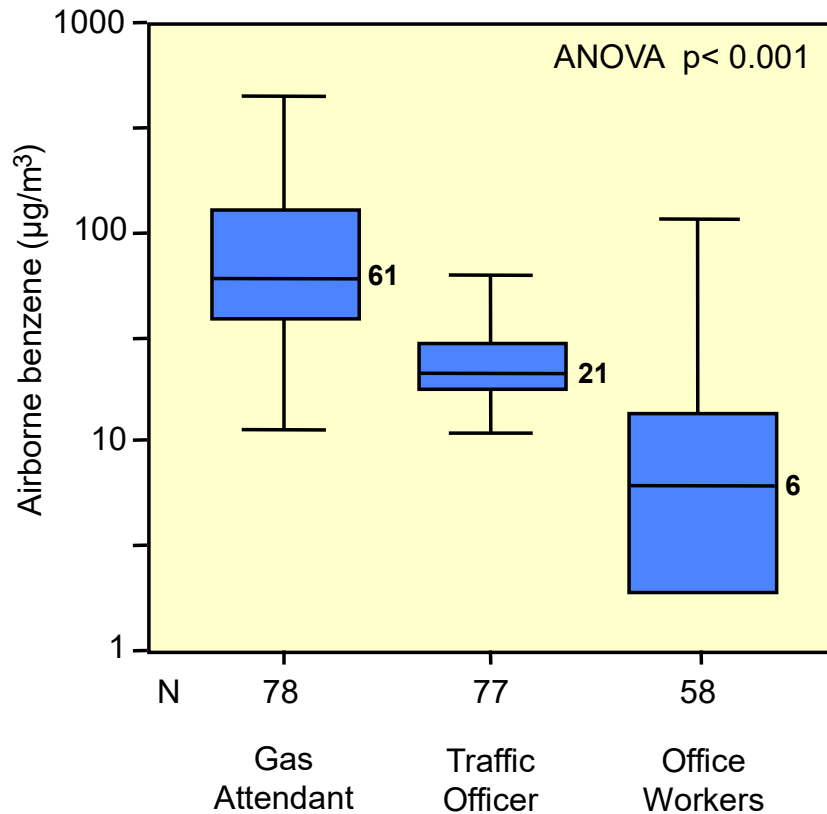
⁵Department of Clinical and Experimental Medicine, University of Catania, Catania, Italy

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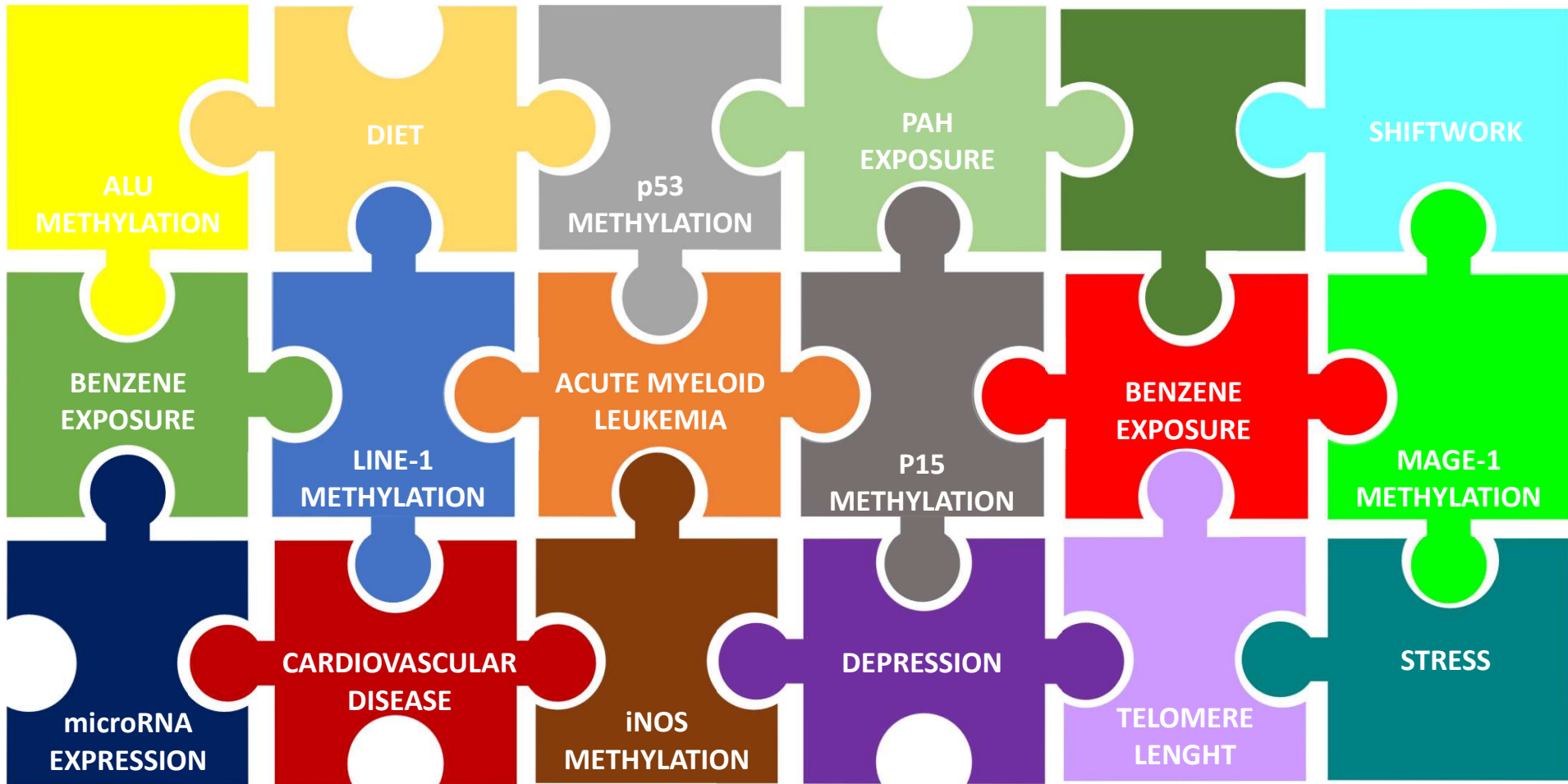


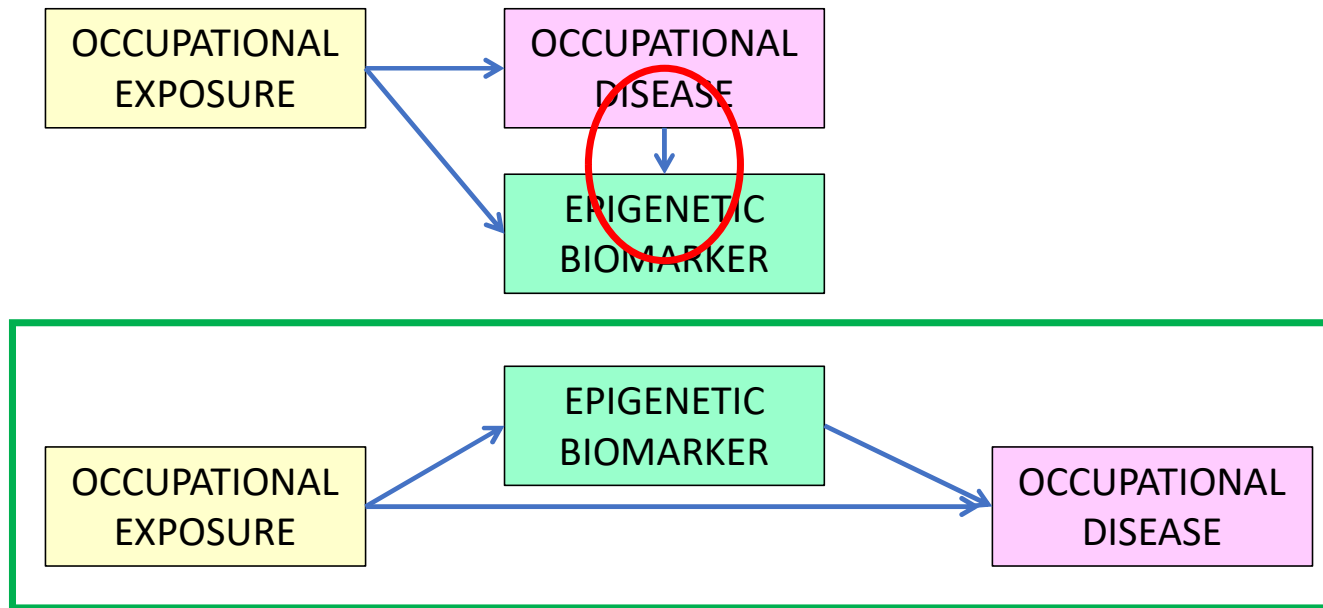
Benzene exposure and p15 methylation



Bollati et al. Cancer Res 2007;67:876-880







NEED of PROSPECTIVE STUDIES

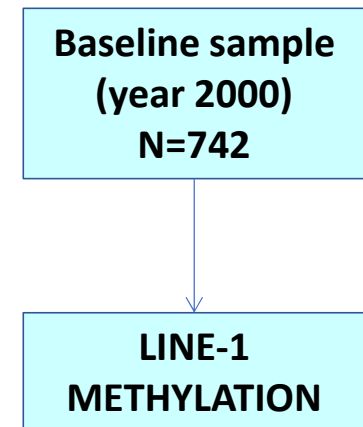
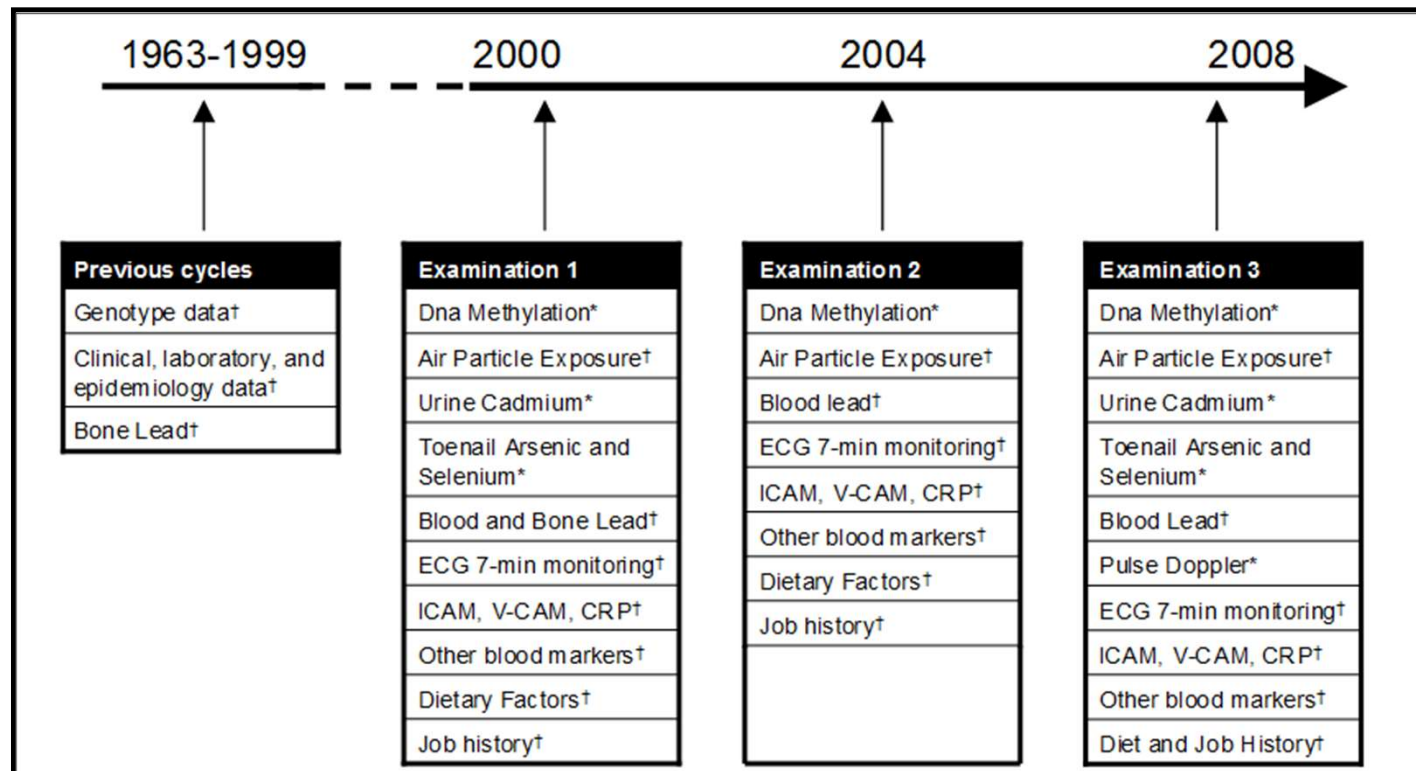


Normative Aging Study (NAS)

- The Normative Aging Study (NAS) is a longitudinal study established by the United States Department of Veterans Affairs in 1963.
- Most participants are veterans from World War II and the Korean War.
- Participants were male residents and free of chronic disease at the time of recruitment.
- Participants were invited to undergo an in-person examination every 3-5 years.



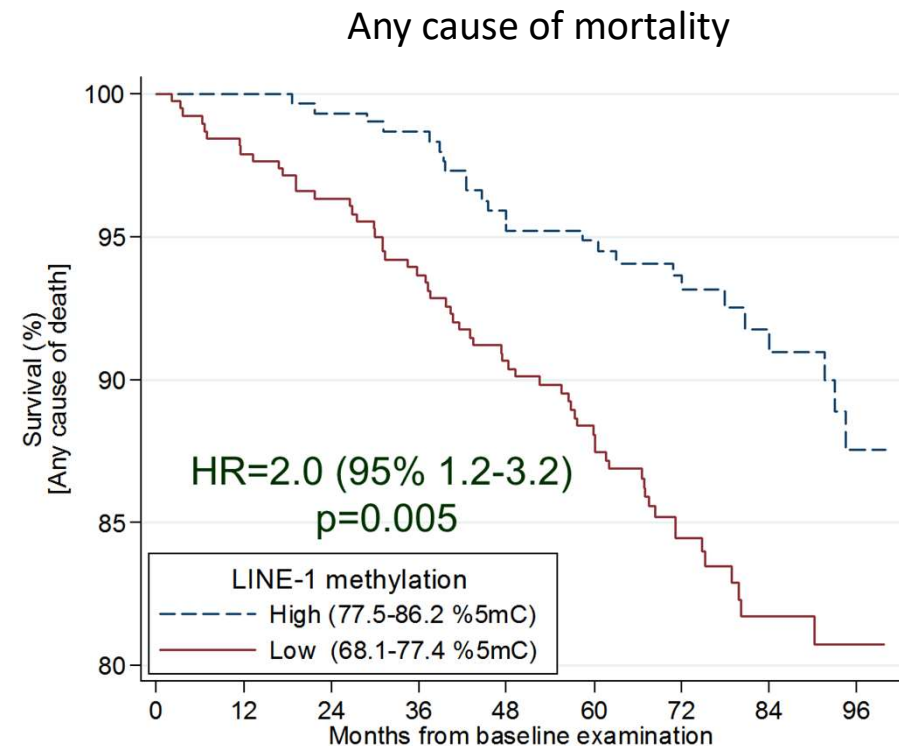
Normative Aging Study (NAS)



Results

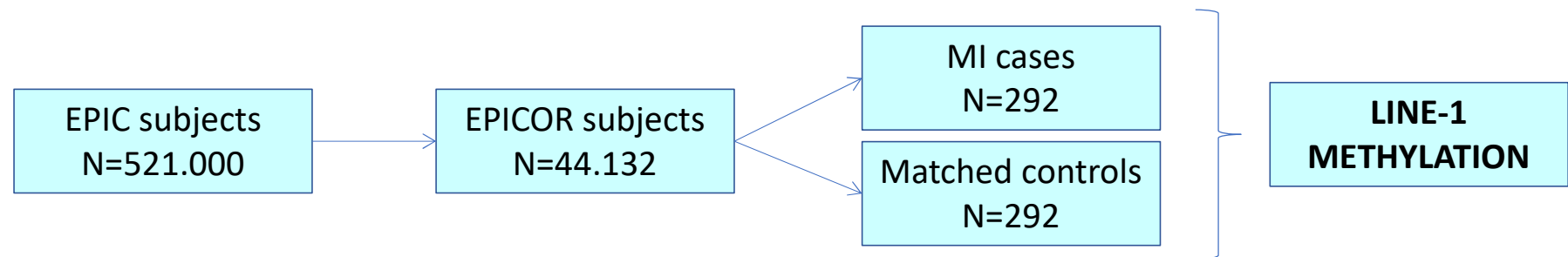
- Individuals with <median LINE-1 methylation show an HR = 3.0 (95% CI 1.3-6.9) of developing cancer (follow-up: mean of 2,894 person-years, 30 cases of cancer).
- Individuals with <median LINE-1 methylation have increased mortality (HR = 3.2, 95% CI 1.3-7.9).

Zhu ZZ, et al. Cancer Causes Control. 2011;22(3):437-47.
Baccarelli A, et al. Epidemiology. 2010 Nov;21(6):819-28.



Italian European Prospective Investigation into Cancer and Nutrition (EPIC) cohort

- The EPIC study is one of the largest cohort studies in the world, with more than half a million (521 000) participants recruited between 1993-1998 across 10 European countries.
- EPIC was designed to investigate the relationships between diet, nutritional status, lifestyle and environmental factors, and the incidence of cancer and other chronic diseases.



Guarrera, S. et al. *Clin Epigenet* 7, 133 (2015)



LINE-1 hypomethylation is associated with myocardial infarction (MI) risk

- MI cases had statistically significant LINE-1 hypomethylation as compared to controls ($\beta = -0.511$, $p\text{-value} = 0.0005$).
- LINE-1 was hypomethylated in cases with shorter time to disease (meta-analysis, men: $P_{\text{TREND}} = 0.0016$, women $P_{\text{TREND}} = 0.026$).
- Findings were replicated in independent samples (EPIC-NL):
 - MI risk associated with LINE-1 lower methylation class in EPIC-NL men (OR = 1.95, $P = 0.043$, ref. group above the median).
 - No difference was found in EPIC-NL women (OR = 1.05, $P = 0.850$).
- **DNA methylation profiles improved case-control discrimination and reclassification when compared with traditional MI risk factors only.**

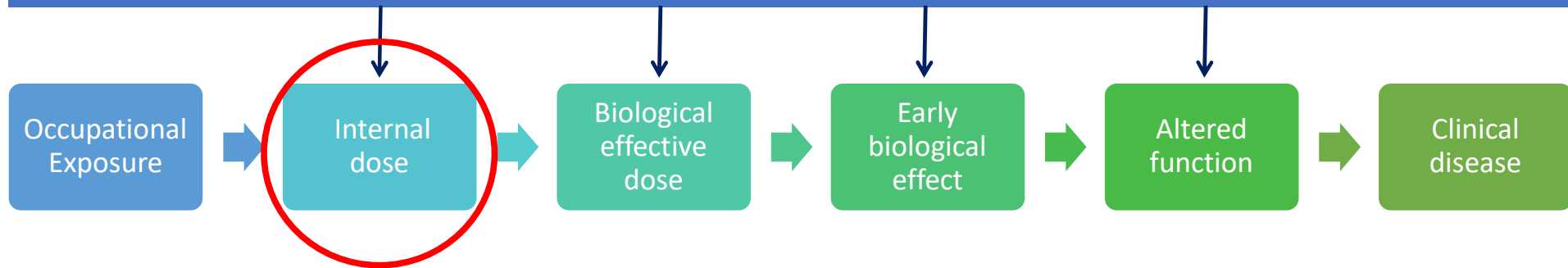
Guarrera, S. et al. *Clin Epigenet* 7, 133 (2015)



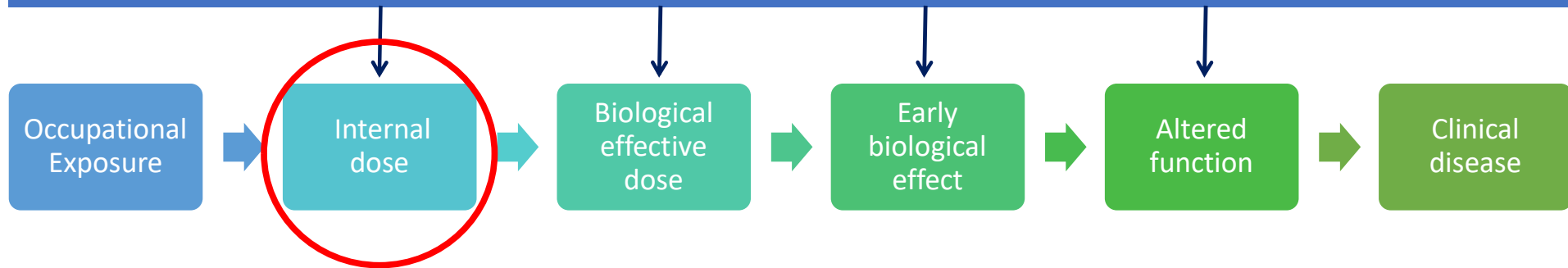
Effect biomarker or NOT?



Exposure and Effect Biomarkers



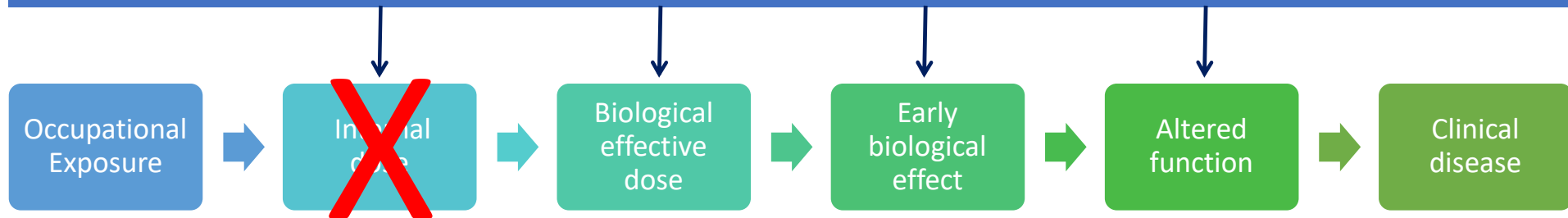
Exposure and Effect Biomarkers



- Often a dose-response is present
- Not specific at all



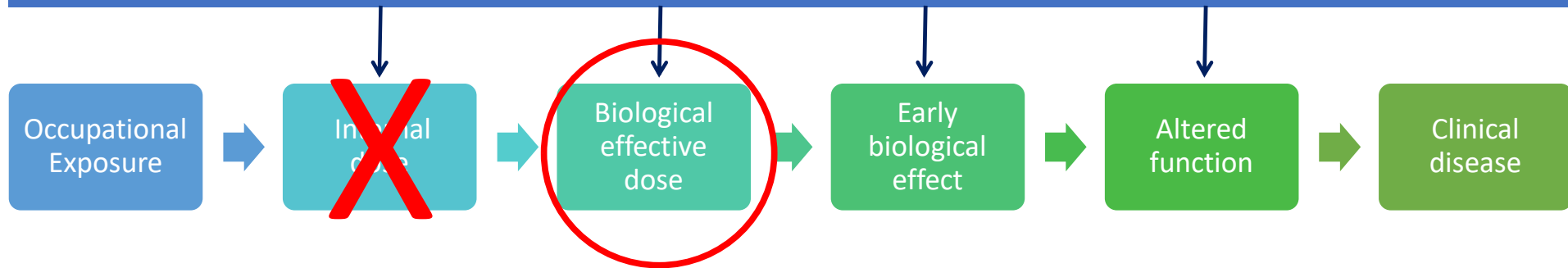
Exposure and Effect Biomarkers



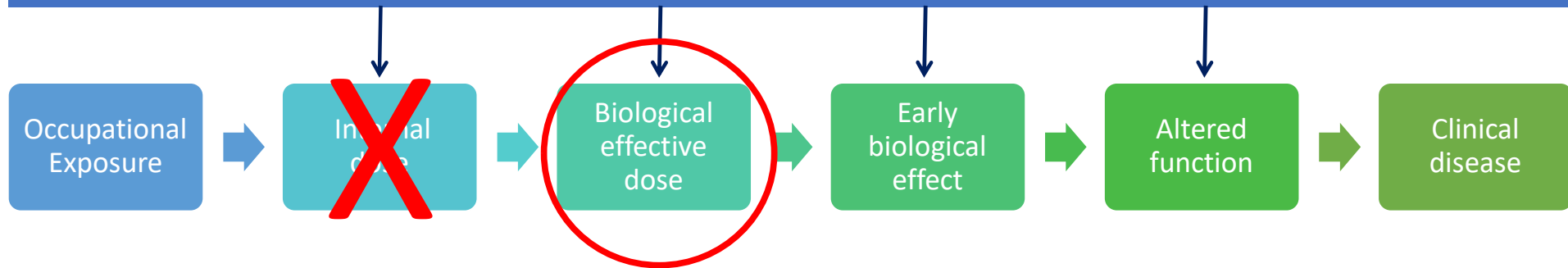
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Exposure and Effect Biomarkers



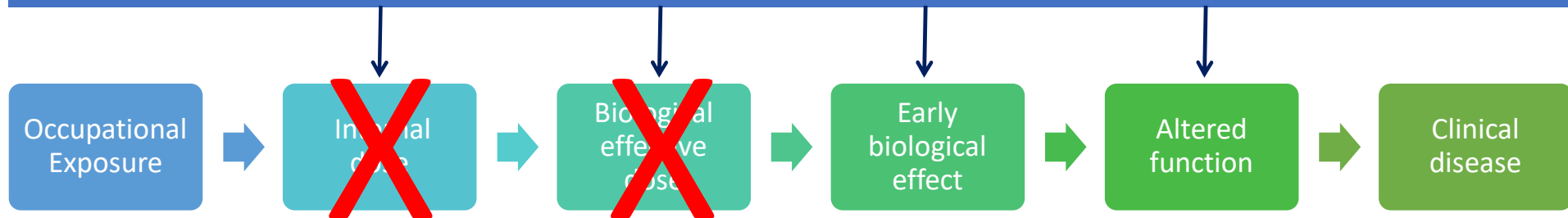
Exposure and Effect Biomarkers



- A threshold of effect is usually not defined



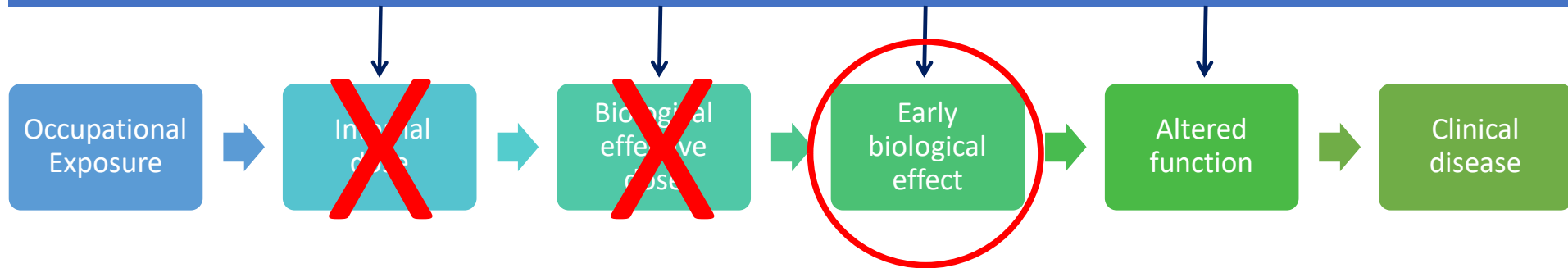
Exposure and Effect Biomarkers



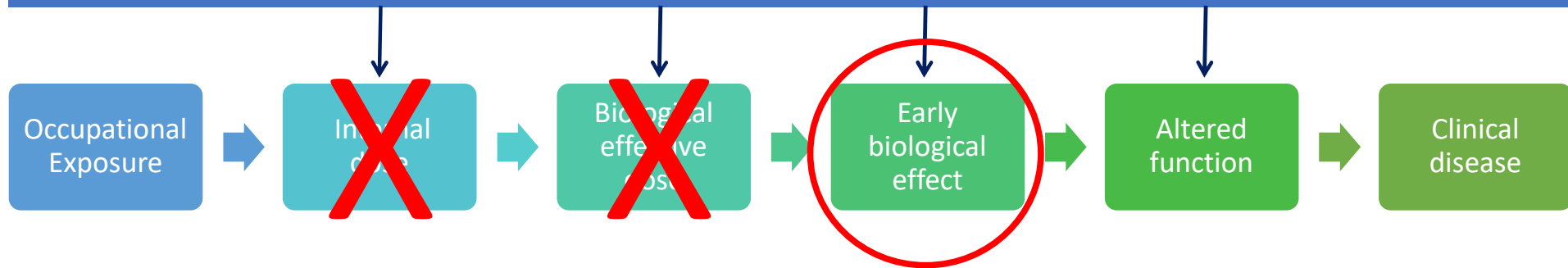
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Exposure and Effect Biomarkers



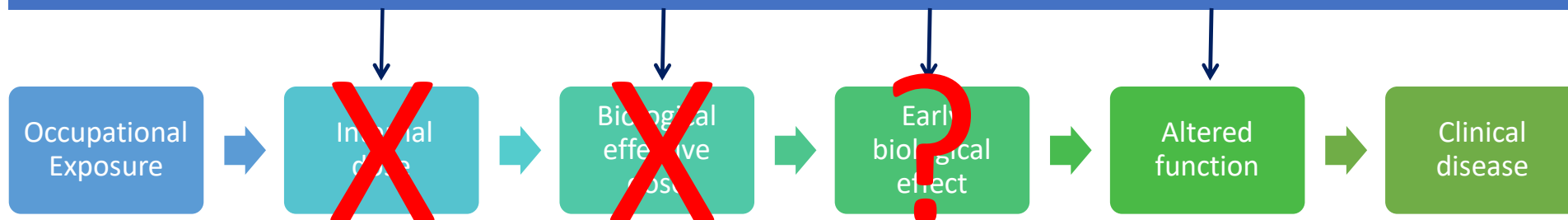
Exposure and Effect Biomarkers



- Biological changes are in common with several exposures
- Not a specific change but rather the response of the organism to changes in its environment, taken as a whole
- It is possible that a single exposure is linked to a PATTERN of modifications



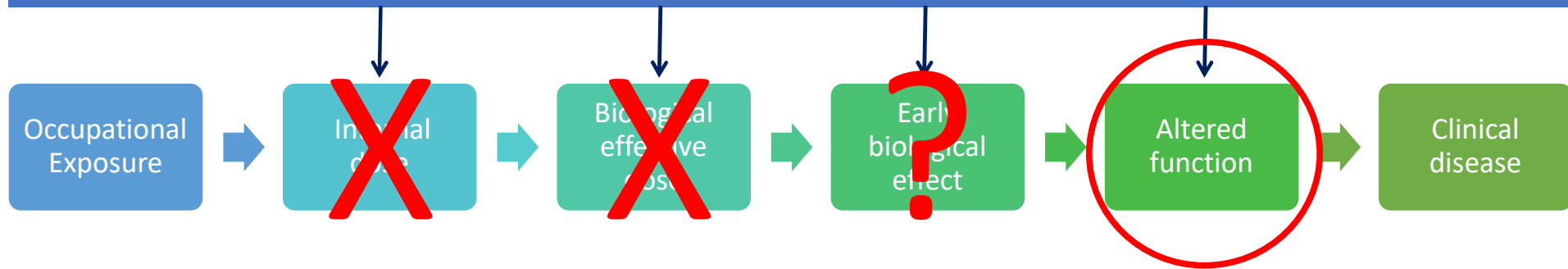
Exposure and Effect Biomarkers



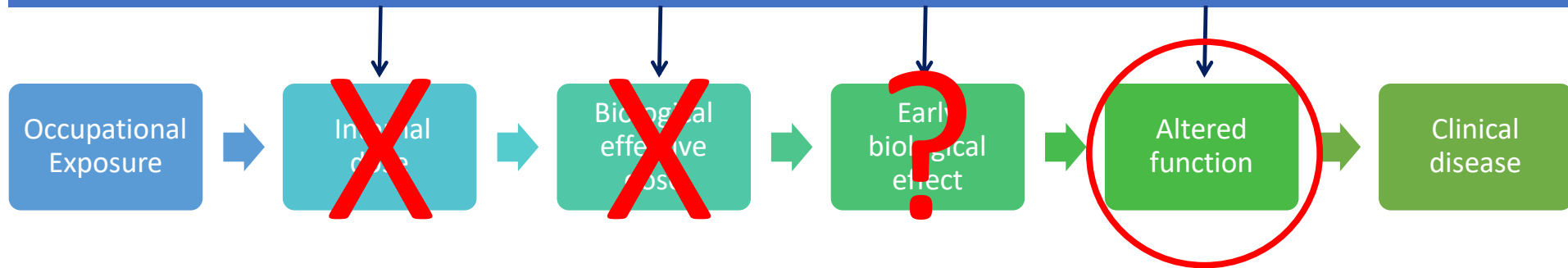
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Exposure and Effect Biomarkers



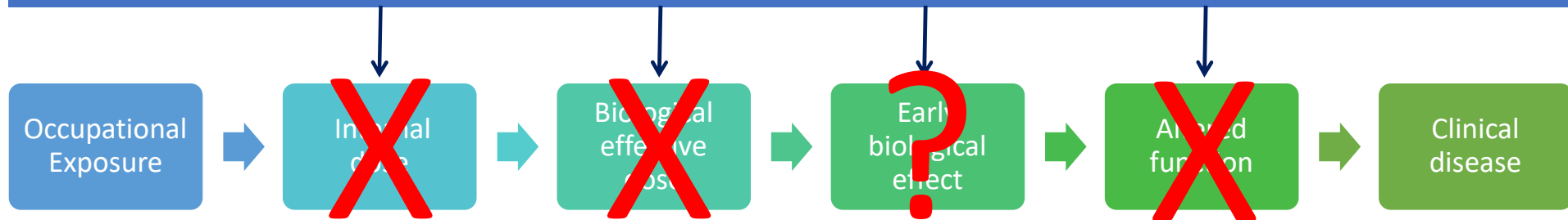
Exposure and Effect Biomarkers



- More than one exposure can converge on pathogenic pathways
- Can be used to estimate a risk of disease, but not to diagnose a disease



Exposure and Effect Biomarkers



Are these markers **USEFUL?**





TARGETED WORKER HEALTH PROMOTION

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Thank
You!

