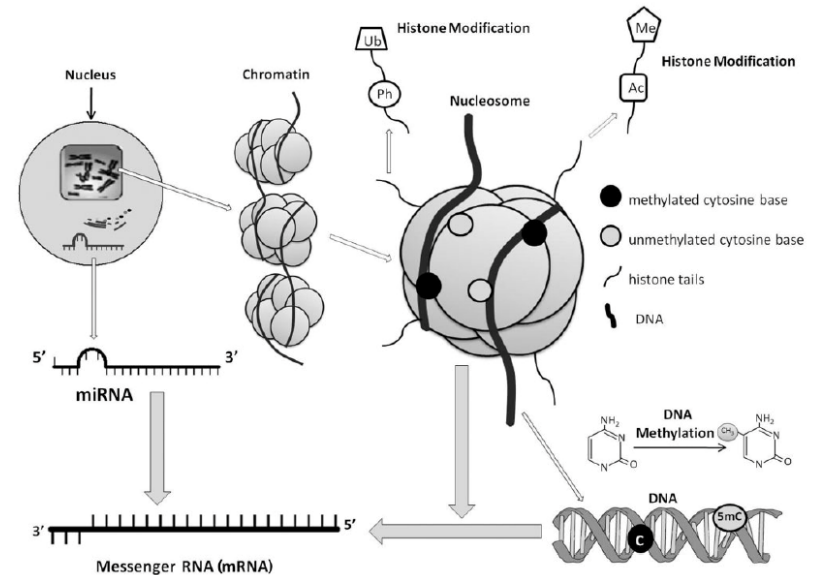
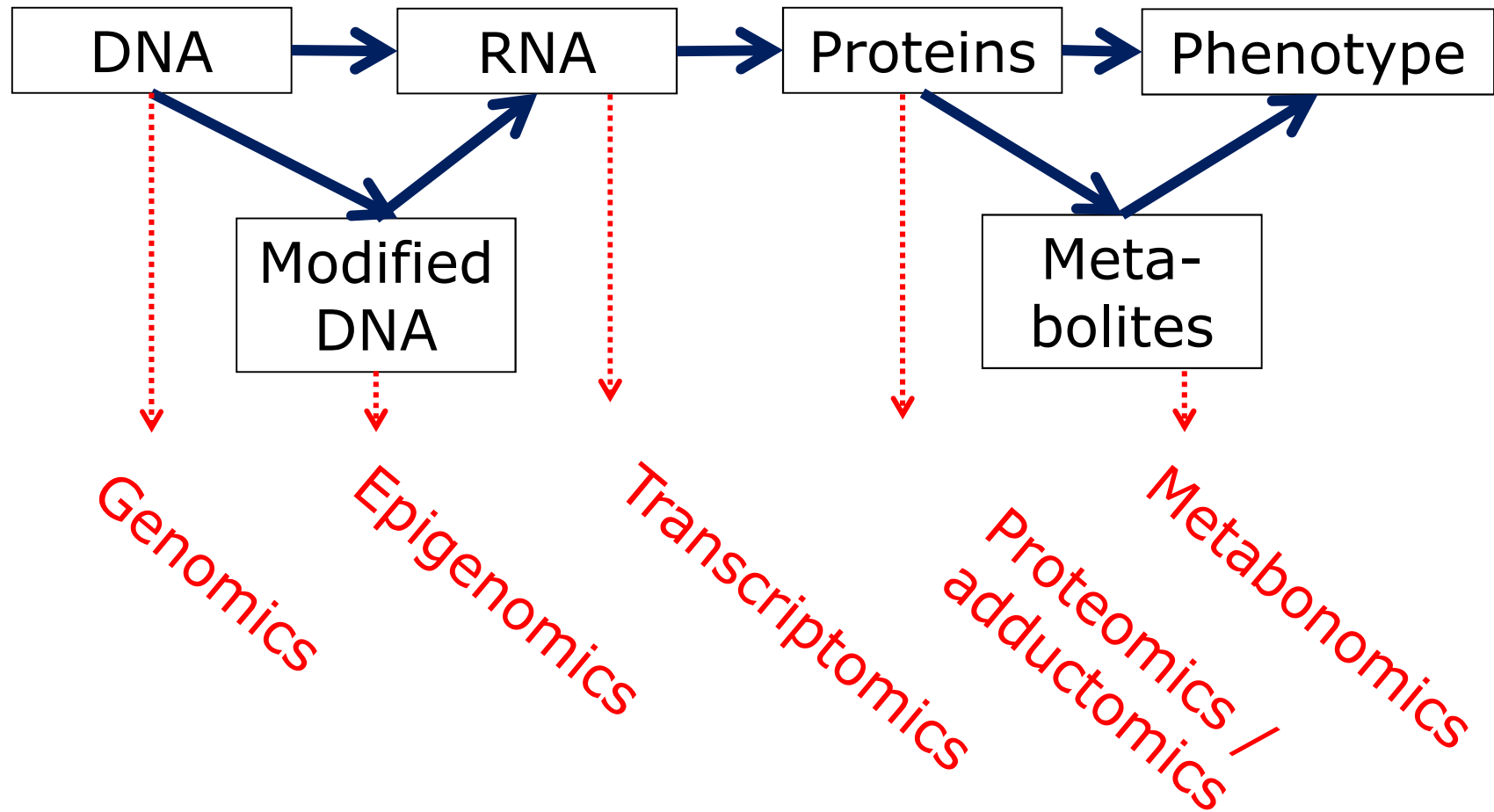


# Epigenetics

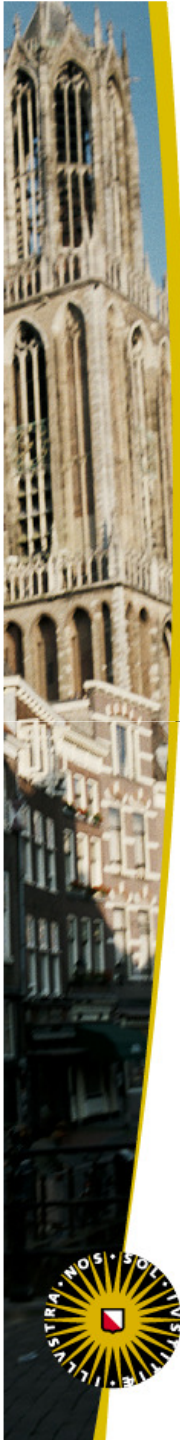
- Stable changes in gene expression and chromatin organization that are independent of the DNA sequence itself and can mitotically inherited over cell divisions
- Epigenetic modifications include:
  - DNA methylation,
  - histone modifications (e.g. acetylation, methylation, ubiquitylation or sumoylation,,),
  - non-coding RNAs (e.g. miRNA),
  - .....
- Methylation
  - Transcriptional repression (CpG)
  - Genomic stability (telomeres)
  - Repress latent viral components
  - Transcriptional activation (gene body)
- *More stable than RNA, proteins*



# Molecular Analyses



Increased relevance (closer to phenotype) but increased epidemiological and analytical challenges

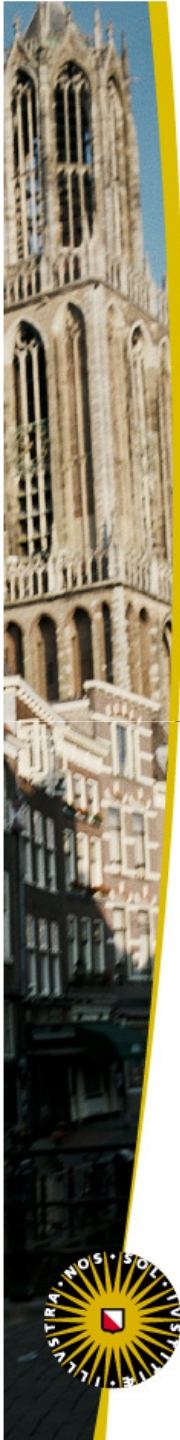




## Epigenetics - Methylation

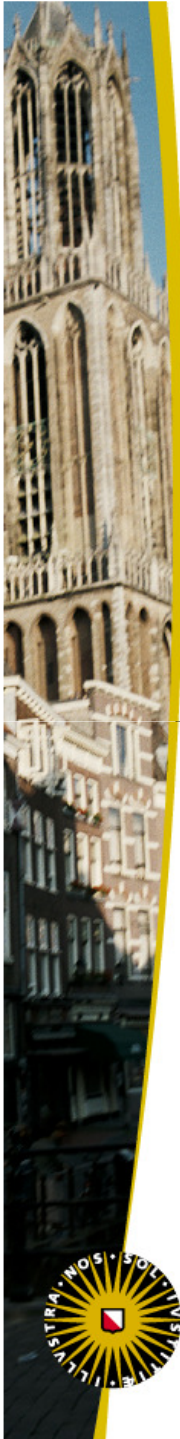
- In recent years, momentum has also been building around the study of epigenetic events in the context of aging and other diseases
  - Autoimmune disorders
  - Cancer
  - Cardiovascular disease
  - Diabetes / metabolic syndrome
  - Neurodegenerative disorders,





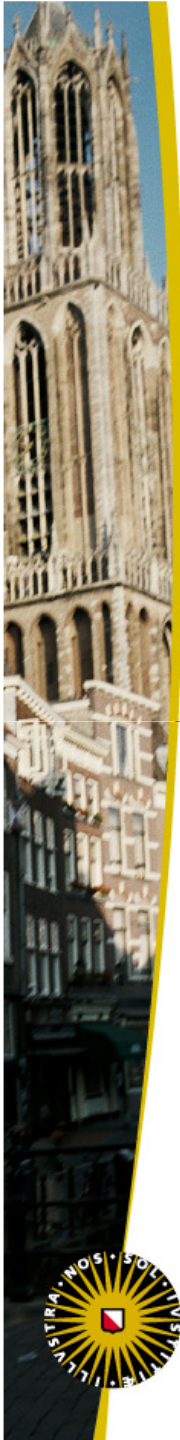
## Plasticity of the epigenome

- Twin studies have demonstrated that epigenetic discordance between identical siblings increases over time; particularly as more time is spent apart [Fraga et al.,
  - Stochastic events that occur over time,
  - preprogrammed,
  - adaptive response to various exposures encountered,
  - during life (or a combination of these events).
- There is additional mounting evidence that environmental or dietary exposures or maternal signals such as stress that are sustained during the developmental stages, in particular during prenatal development, can result in epigenetic errors or modifications that propagate throughout somatic tissues



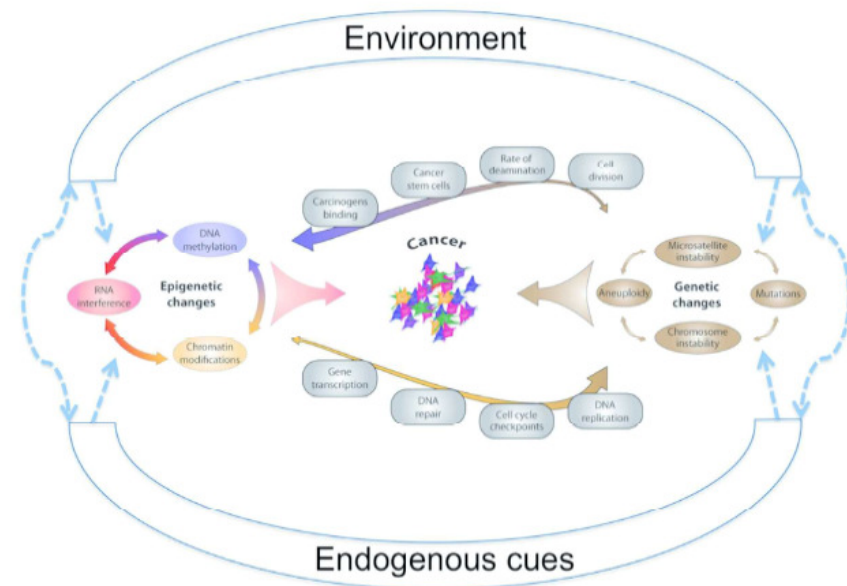
## Environmental factors related to epigenetic deregulation

- Tobacco smoke
- Arsenic
- Cadmium
- Nickel
- Ionizing
- UV radiation
- Air pollution
  
- Human hepatitis B virus
- Hepatitis C virus
- Human papillomavirus
- Helicobacter pylori

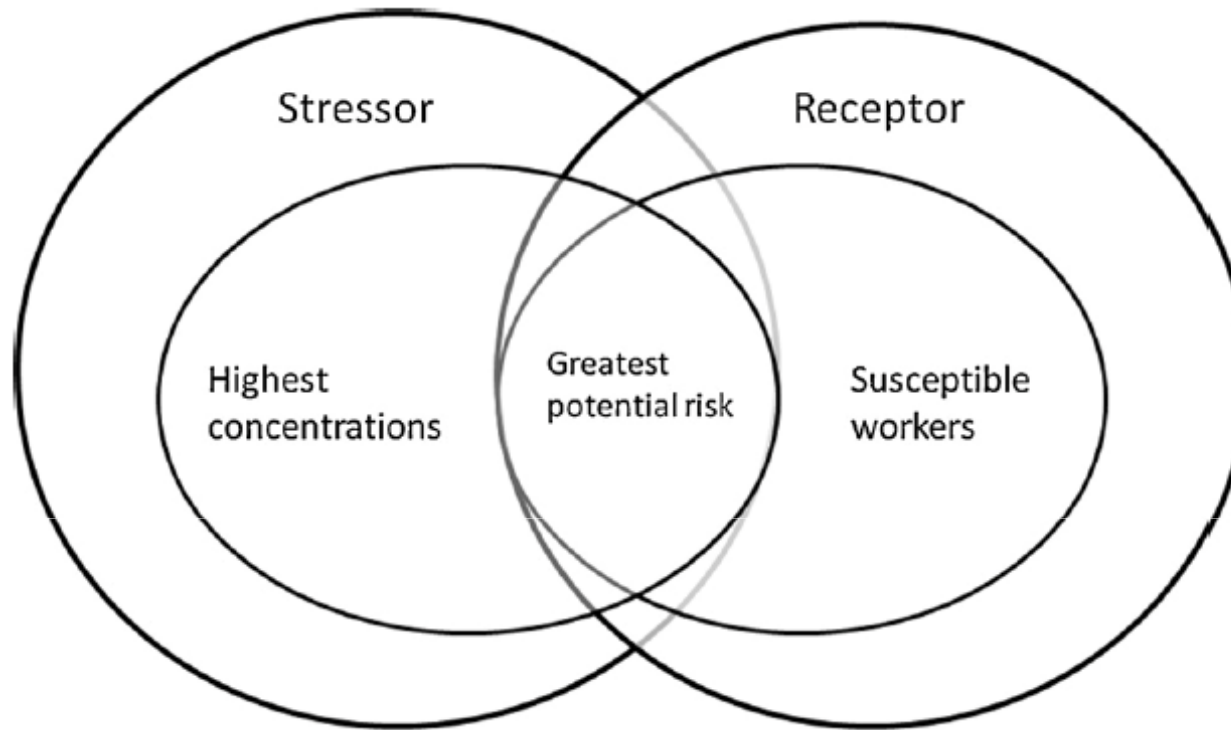


## Epigenetics - Methylation

- Tissue heterogeneity
  - epigenetic signature varies by cell and tissue
- Reversed causality
- Platforms
  - Next generation sequencing
  - Array- based



## Use of biomarkers in occupational health



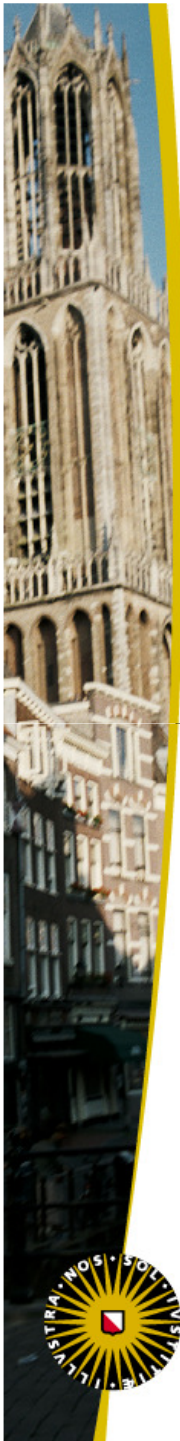
Source: Adapted from Reiter [2009]

- Medical screening
- Diagnostic monitoring



# Use of Biomarkers in Occupational Health; Research

	Research	Practice
Exposure	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as independent and dependent variables</li> <li>• Assess exposure</li> </ul>	<ul style="list-style-type: none"> <li>• Biological monitoring</li> <li>• Genetic monitoring</li> <li>• Assess control effectiveness</li> <li>• Health surveillance</li> </ul>
Effect	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as dependent variables</li> <li>• Useful in developing systems biology approaches</li> </ul>	<ul style="list-style-type: none"> <li>• Medical screening</li> <li>• Health surveillance</li> <li>• Early warning</li> <li>• Genetic monitoring</li> <li>• Fitness for duty</li> </ul>
Susceptibility <b>epimutations</b>	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Investigate new effect modifiers</li> <li>• Develop system biology approach</li> </ul>	<ul style="list-style-type: none"> <li>• Target high risk groups</li> <li>• Develop risk communication</li> <li>• Genetic screening</li> </ul>







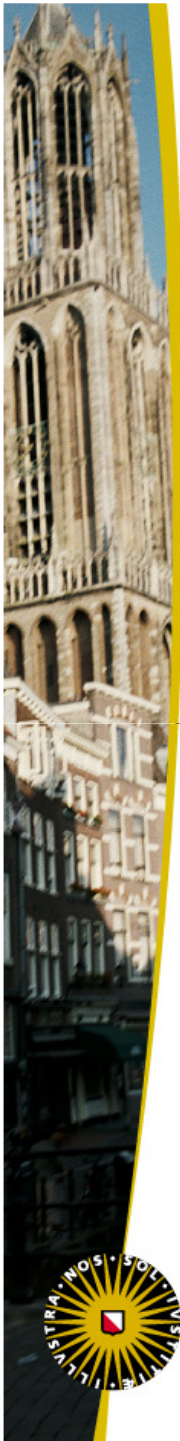
## Use of Biomarkers in Occupational Health; In practice

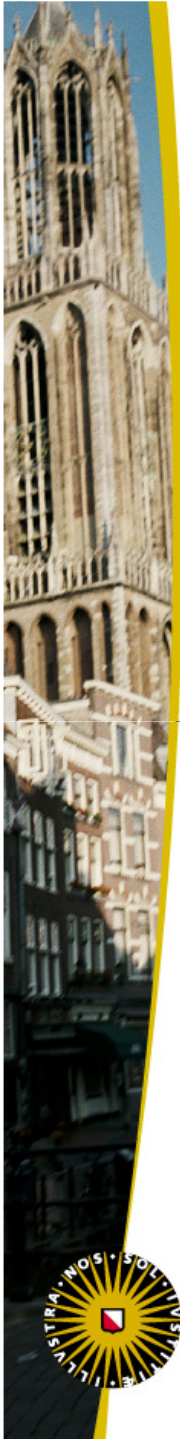
- Exposure assessment
  - Means to assure the effectiveness of risk management and prevention
- Health surveillance
  - Respiratory , cardiovascular conditions
- Identify high risk groups
  - Develop risk communication



# Use of Biomarkers in Occupational Health; Research

	Research	Practice
Exposure	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as independent and dependent variables</li> <li>• Assess exposure</li> </ul>	<ul style="list-style-type: none"> <li>• Biological monitoring</li> <li>• Genetic monitoring</li> <li>• Assess control effectiveness</li> <li>• Health surveillance</li> </ul>
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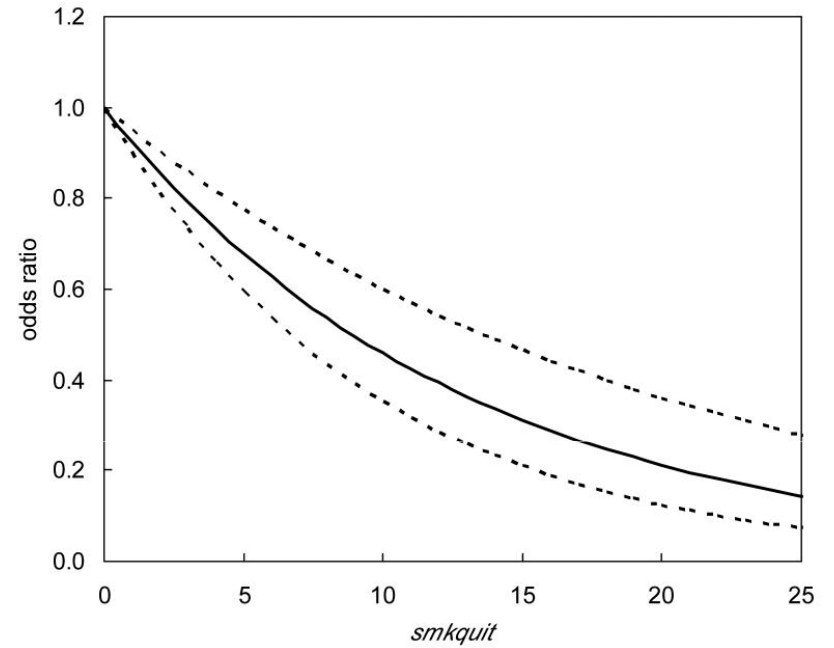
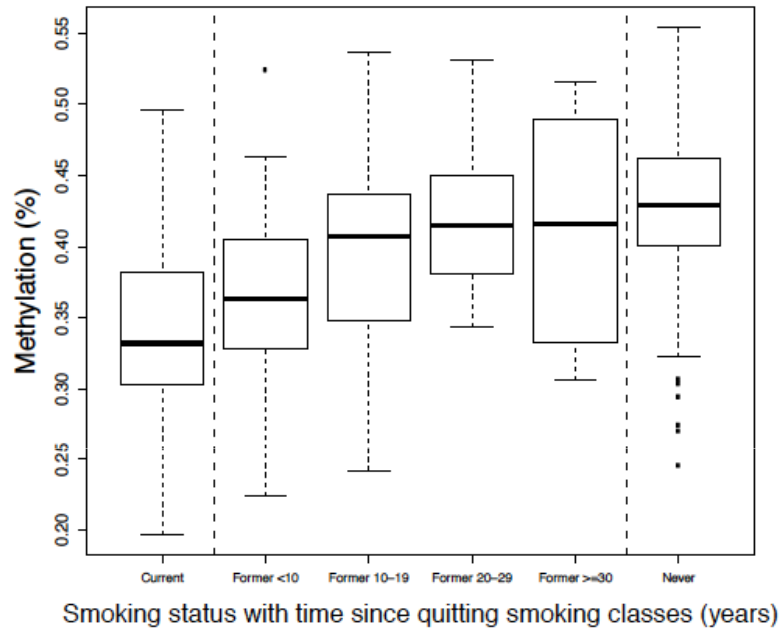


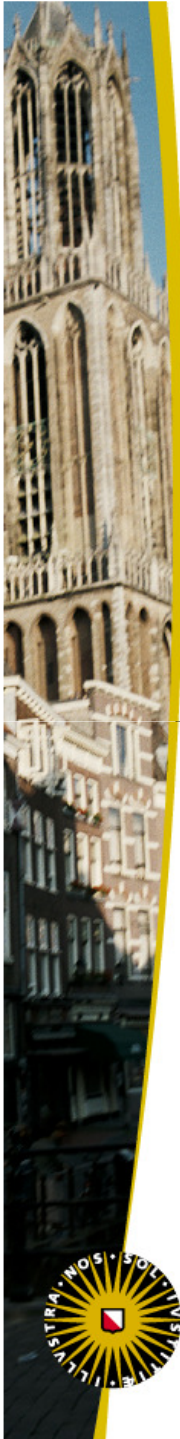
## Epigenetics and exposure; Epigenetic changes and smoking

- Population
  - EPIC-Italy n=620
  - NOWAC n=343
- Detailed information on smoking
- PBMCs
- Methylation
  - Illumina Humanmethylation 450K chip


# Biological effective dose?

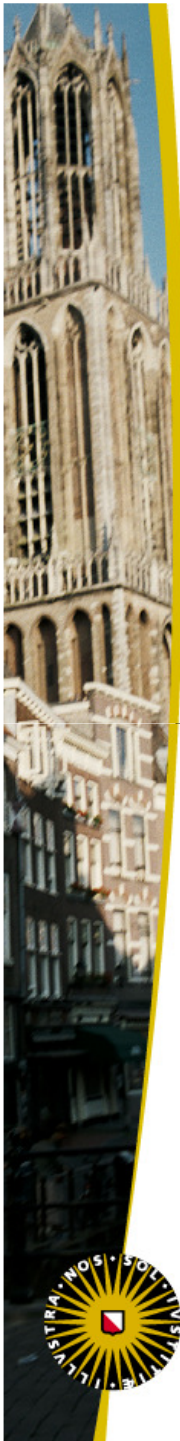
Neuberger et al., 2007





# Use of Biomarkers in Occupational Health; Research

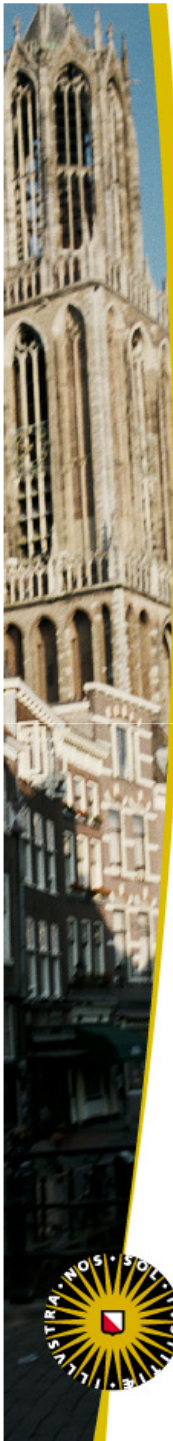
	Research	Practice
Exposure  Sensitivity? Specificity?	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as independent and dependent variables</li> <li>• Assess exposure</li> </ul>	<ul style="list-style-type: none"> <li>• Biological monitoring</li> <li>• Genetic monitoring</li> <li>• Assess control effectiveness</li> <li>• Health surveillance</li> </ul>
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## Effect markers – Epigenetic mechanisms of known carcinogens

Human carcinogen	Epigenetic mechanism	Model		
		Human samples	Cell lines	Animal model
<b>Part C : Arsenic, Metals, Fibres, and Dusts and Part D: Radiation</b>				
<b>Arsenic</b>				
	DNA methylation	*		
			*	
				*
	Histone marks		*	
	miRNA		*	
<b>Cadmium</b>				
	DNA methylation		*	
				*
	miRNA	*		
<b>Nickel</b>				
	DNA methylation		*	
				*
	Histone marks		*	
				*
<b>Beryllium</b>				
	DNA methylation			*

Un



# Effect markers – Epigenetic mechanisms of known carcinogens

Human carcinogen	Epigenetic mechanism	Model		
		Human samples	Cell lines	Animal model
Asbestos	DNA methylation			*
	miRNA			*
X-radiation	DNA methylation	*		
				*
	miRNA	*		
			*	
Gamma radiation				*
	DNA methylation		*	
	miRNA			*
Smoky coal emissions (for cooking and heating)				
	DNA methylation	*		
Chromium VI compounds				
	Histone marks		*	



Air pollution  
DEE

DNA methylation \*  
miRNA \*

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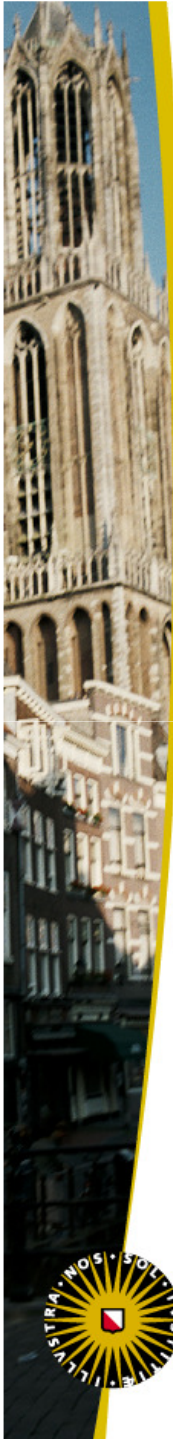
Zdenko et al. unpublished

# Use of Biomarkers in Occupational Health; Research

	Research	Practice
Exposure  Sensitivity? Specificity?	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as independent and dependent variables</li> <li>• Assess exposure</li> </ul>	<ul style="list-style-type: none"> <li>• Biological monitoring</li> <li>• Genetic monitoring</li> <li>• Assess control effectiveness</li> <li>• Health surveillance</li> </ul>
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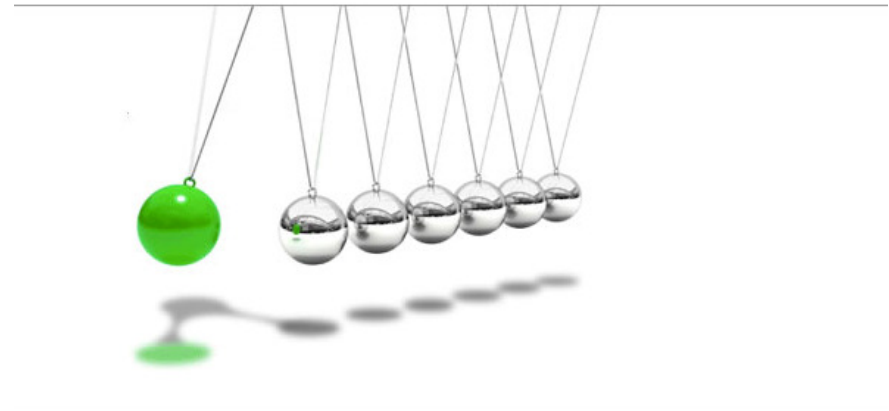






## Cross Omics analyses

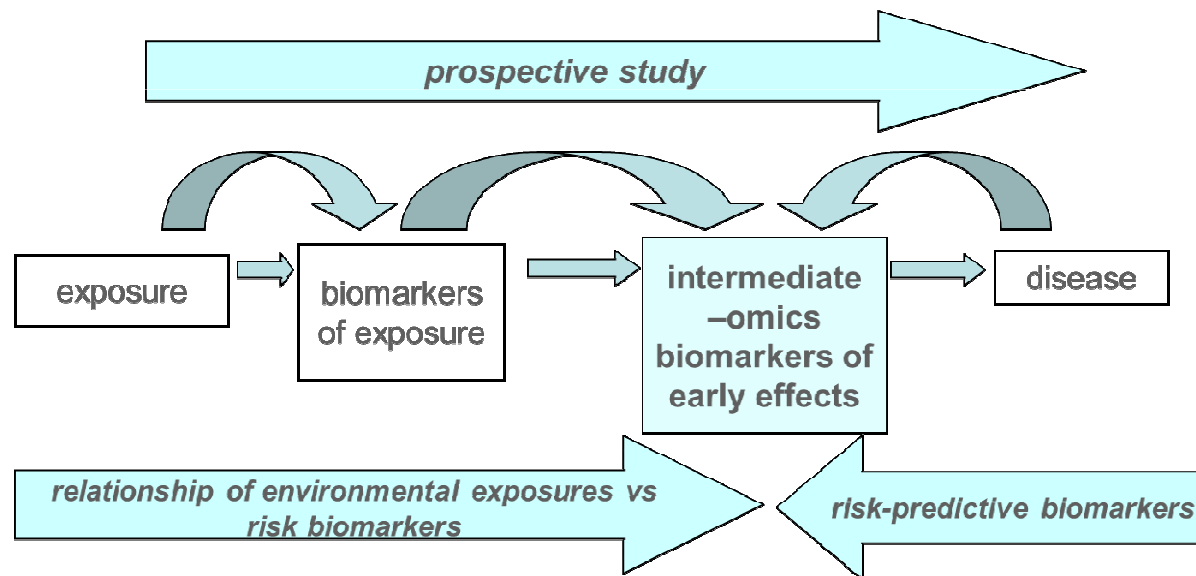
- Concatenating the methylation dataset on gene expression profiles
  - 558 gene expression probes significantly associated with 79 out of the 123 common CpG sites, using a Bonferroni 5% threshold.
  - Differentially significant expression probes were mapped to genes present in DAVID.





# OMICS based biomarkers – Epiphenomena?

- Objectives

- To identify exposure markers
- To find disease classifiers (risk-predictive biomarkers)
- To elucidate the etiology of the exposure – disease association (MITM biomarkers)



# Use of Biomarkers in Occupational Health; Research

	Research	Practice
Exposure  Sensitivity? Specificity?	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Serve as independent and dependent variables</li> <li>• Assess exposure</li> </ul>	<ul style="list-style-type: none"> <li>• Biological monitoring</li> <li>• Genetic monitoring</li> <li>• Assess control effectiveness</li> <li>• Health surveillance</li> </ul>
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epimutations



Universiteit Utrecht



## Susceptibility; Epigenetics adds another layer of complexity to individual variability

- Epigenome may be most vulnerable to environmental insults during periods of extensive epigenetic reprogramming
  - Early stages of embryonic development
  - Childhood (oocyte, hyploid demethylated state)

-> *Employment of fertile women*

-> *Intergenerational equity*
- Pre-exposure testing for individual susceptibility
  - Similar ethical issues as with genetic testing
    - *Privacy & confidentiality*
    - *Undermine individual autonomy*
    - *Create stigma*
    - *Psychological harm*

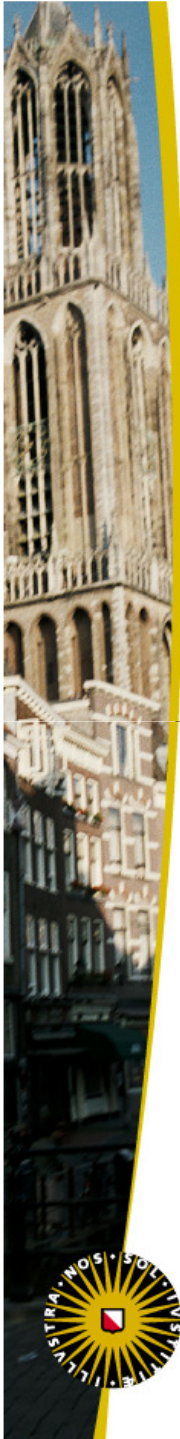


# Use of Biomarkers in Occupational Health; Research

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Susceptibility ✓ Ethics	<ul style="list-style-type: none"> <li>• Identify mechanisms</li> <li>• Investigate new effect modifiers</li> <li>• Develop system biology approach</li> </ul>	<ul style="list-style-type: none"> <li>• Target high risk groups</li> <li>• Develop risk communication</li> <li>• Genetic screening</li> </ul>

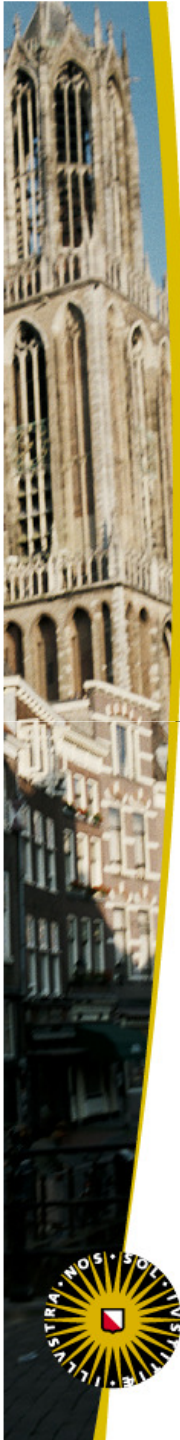
epimutations





## Epigenetica en blootstelling aan chemicaliën; Betekenis voor de bedrijfsgezondheidszorg

- *Epigenetics represents a new class of biological effects of harmful exposures and adds a multi-generational dimension to environmentally-caused adverse effects*
- Epigenetics holds promise for elucidating mechanisms of disease
  - Integral part of the systems biology approach
- May potentially lead to environmental fingerprints
  - Specificity, dynamics
- May potentially provide stable markers of early effect
  - Epiphenomena ~ validation
- May provide markers of susceptibility
  - Ethics



**Biomarkers are not a panacea for preventing all worker illnesses and injuries, but focused on particular issues, their use can contribute to improved protection**

Schulte, 2012



U