



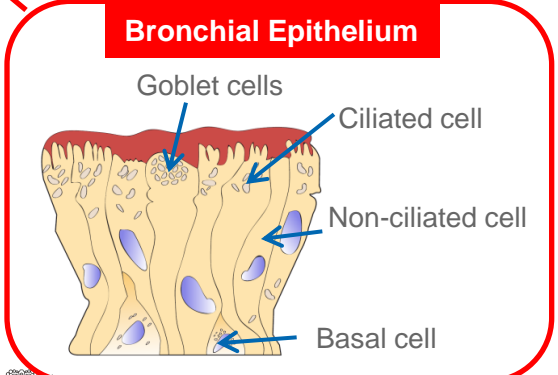
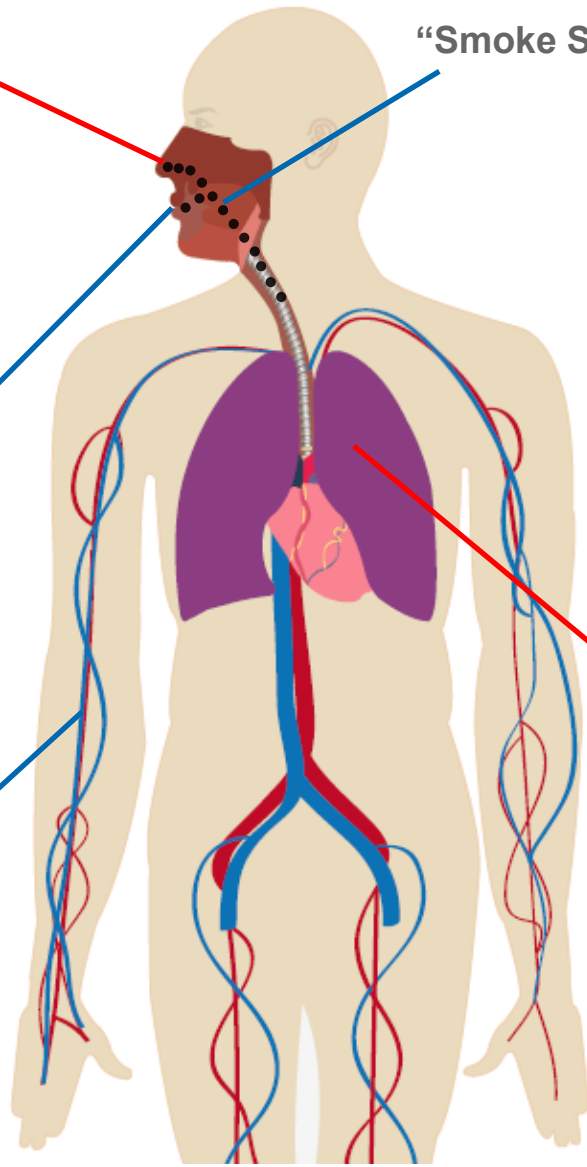
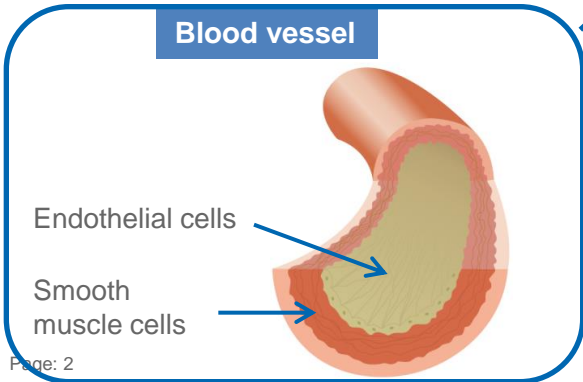
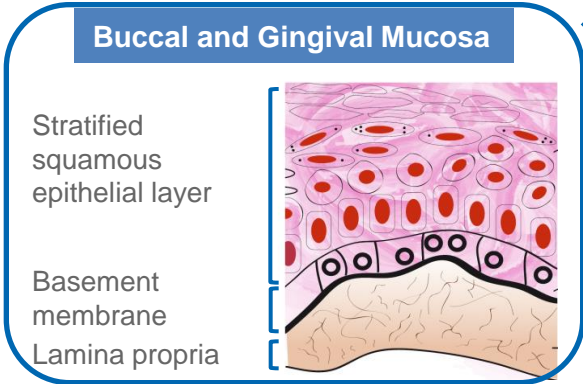
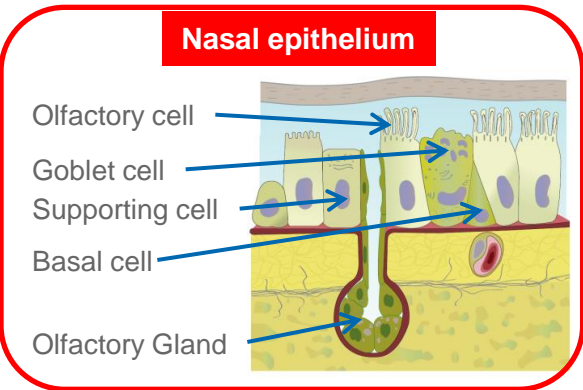
PMI RESEARCH & DEVELOPMENT

**ASSESSING THE EFFECTS OF REPEATED  
CIGARETTE SMOKE-EXPOSURE  
USING HUMAN ORGANOTYPIC SYSTEMS  
REPRODUCING THE RESPIRATORY TRACT IN VITRO**

Carole Mathis

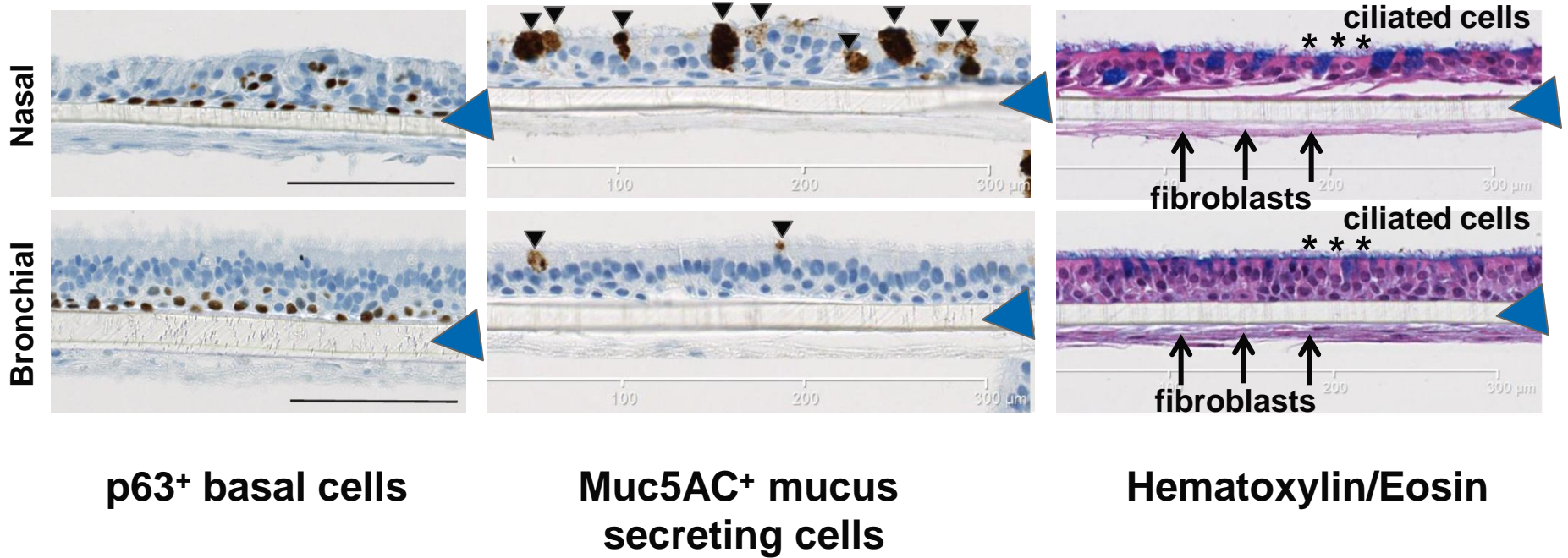
**Philip Morris International R&D, Philip Morris Products S.A.,  
Neuchâtel, Switzerland**

# Recapitulation of In Vivo Biology by Organotypic Systems



# Human Organotypic Cultures of Primary Bronchial and Nasal Epithelial Cells

## COMPARISON AT THE MORPHOLOGICAL LEVEL (ANALYSIS DONE ON UNTREATED TISSUES)



(POROUS MEMBRANE ◀ )



PMI RESEARCH & DEVELOPMENT

# Whole cigarette smoke/aerosol exposure system (Vitrocell®)

Smoking Machine



Humidification System

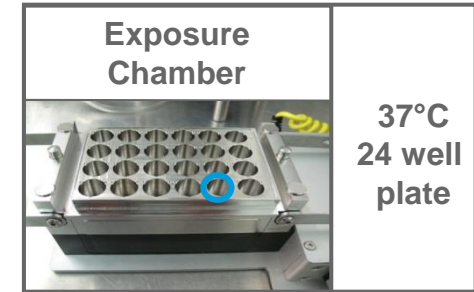
Vitrocell 24/48



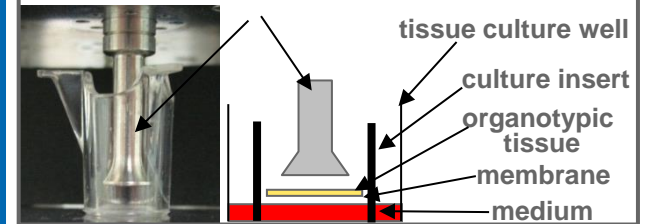
Climatic chamber  
Dilution system  
(8 dilutions @ 6 replicates)



Microbalance

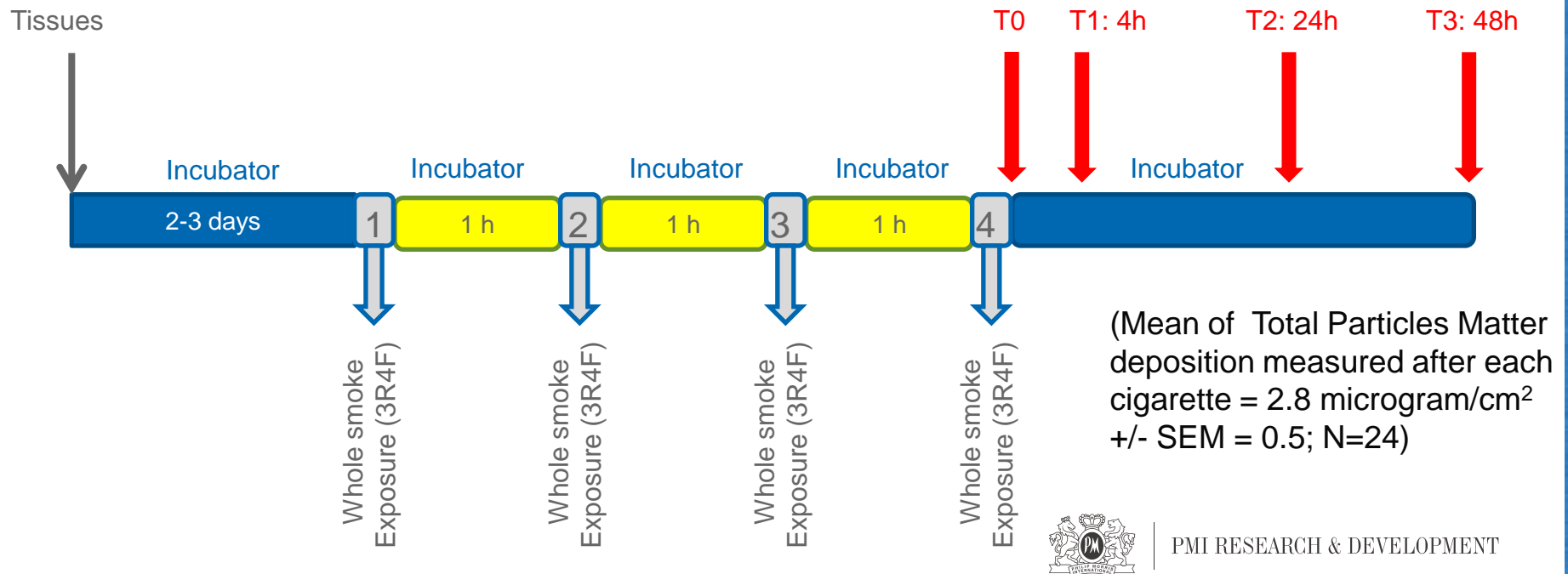


Position of the aerosol inlet above the AIR-100 tissue insert



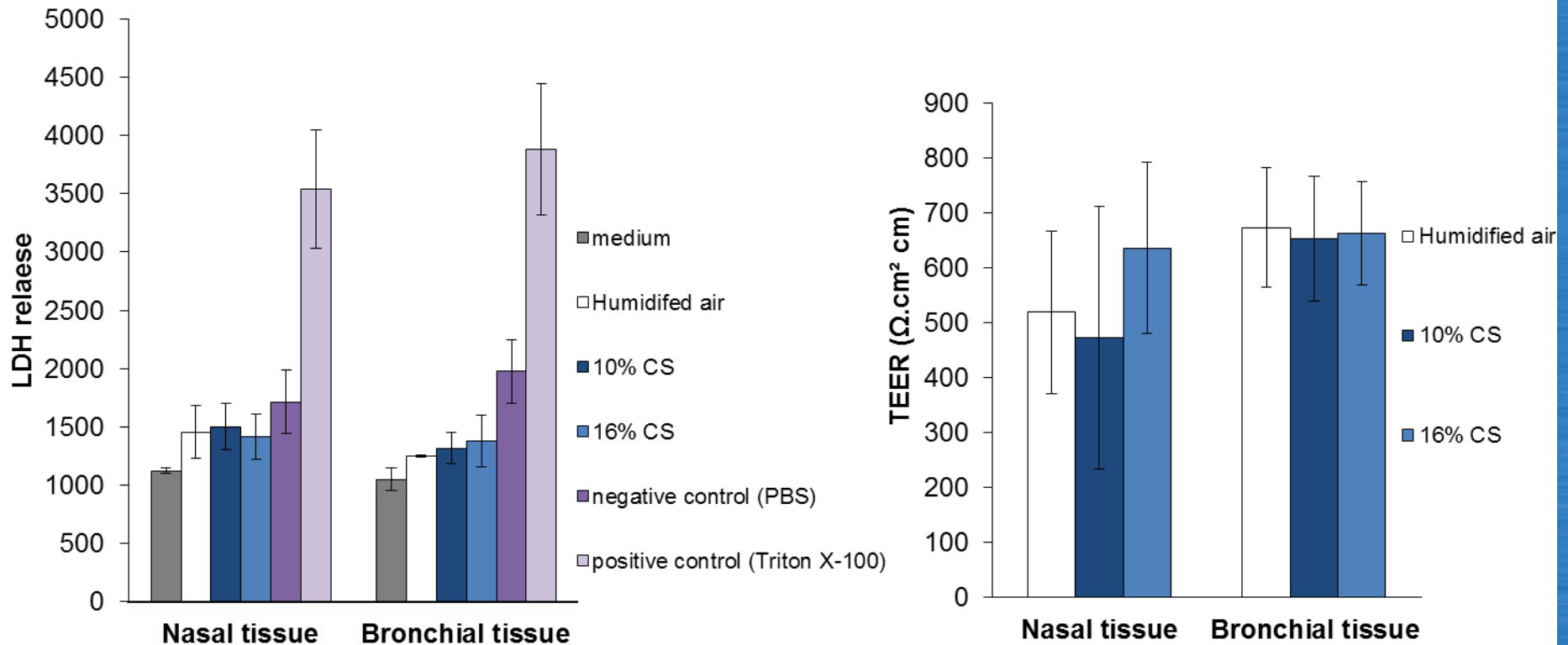
# Whole Smoke Repeated Exposure of Organotypic Cultures of Human Primary Bronchial and Nasal Epithelial Cells

Experimental Design			POST-EXPOSURE TIME			
HUMAN TISSUES	CONDITIONS	ENDPOINTS	0h	4h	24h	48h
BRONCHIAL MUCILAIR™	SHAM	Cytotoxicity (LDH assay)			X	X
		Membrane Integrity (TEER)			X	X
NASAL MUCILAIR™	CS 10%	Gene Expression Profiles (Affymetrix)	X	X	X	X
	CS 16%	MicroRNA Expression Profiles (GeneChip)	X	X	X	X
		Histology (Alcian blue, H&E)			X	
		Immunohistology (p63, Ki67, B-Tubulin, Muc5AC)			X	



# Whole Smoke Repeated Exposure of Organotypic Cultures of Human Primary Bronchial and Nasal Epithelial Cells

The two doses of CS applied on both nasal and bronchial tissue cultures do not induce high toxicity or impair tissue integrity allowing the capture of systems biology endpoints.



24h after exposure



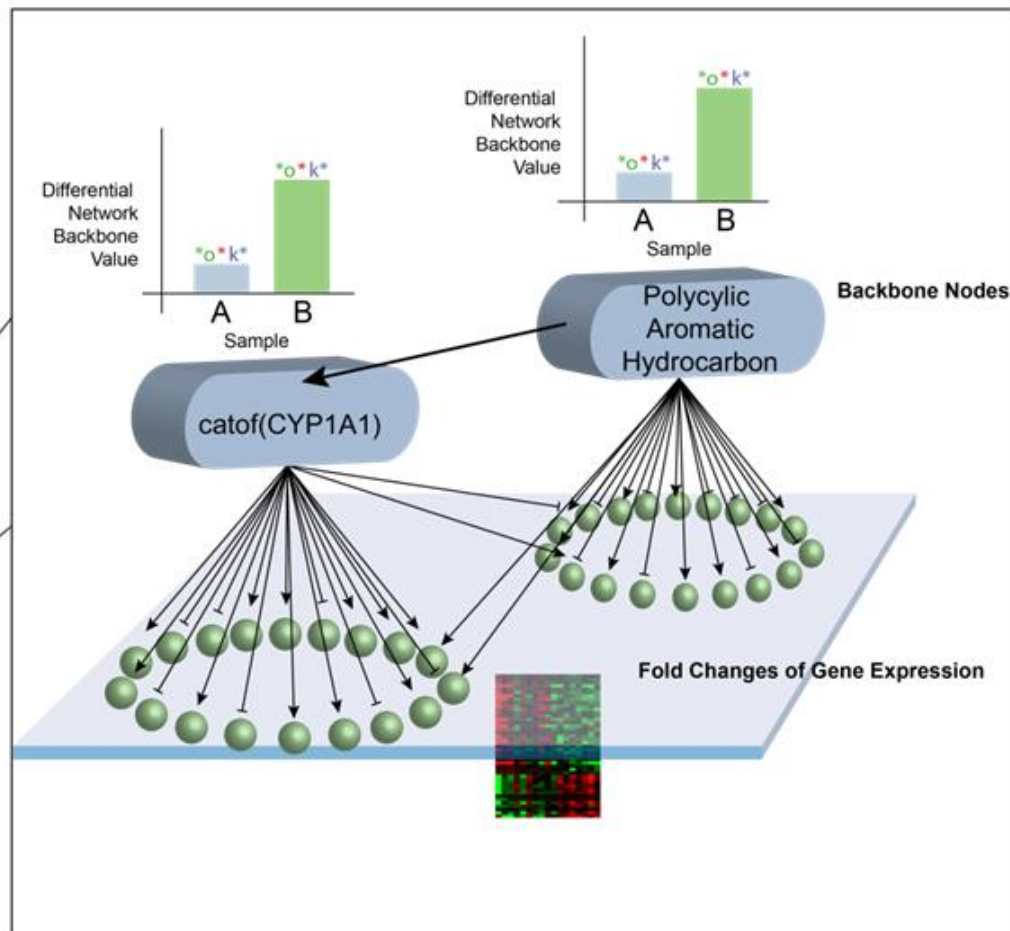
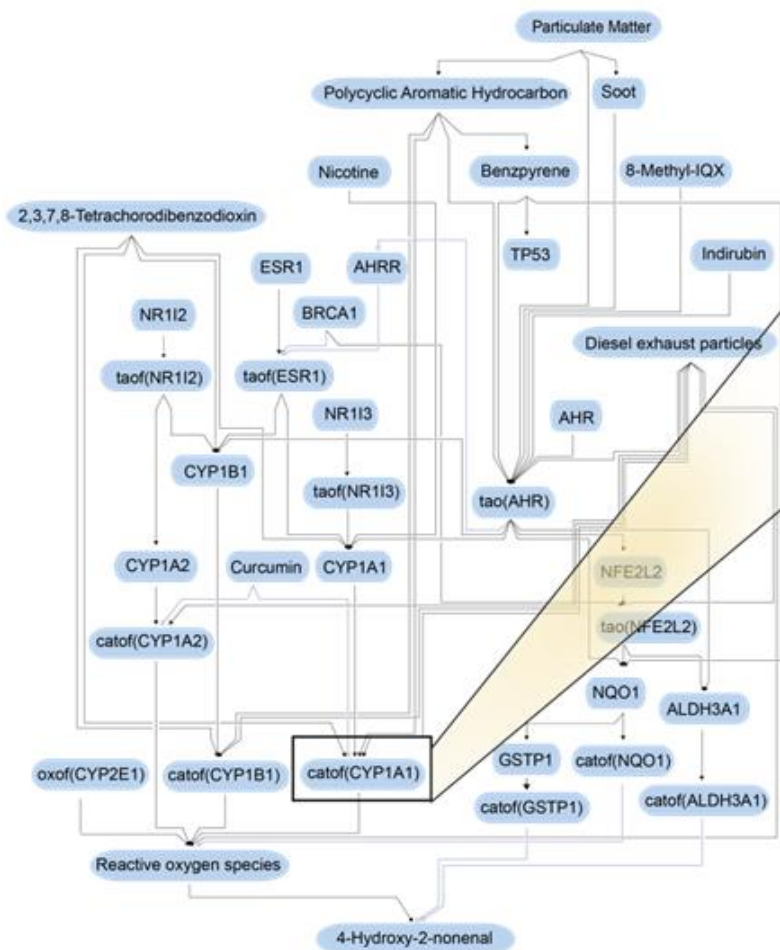
PMI RESEARCH & DEVELOPMENT



# Xenobiotic Metabolism Network Model and Biological Perturbation Assessment



**Xenobiotic Metabolism Network**

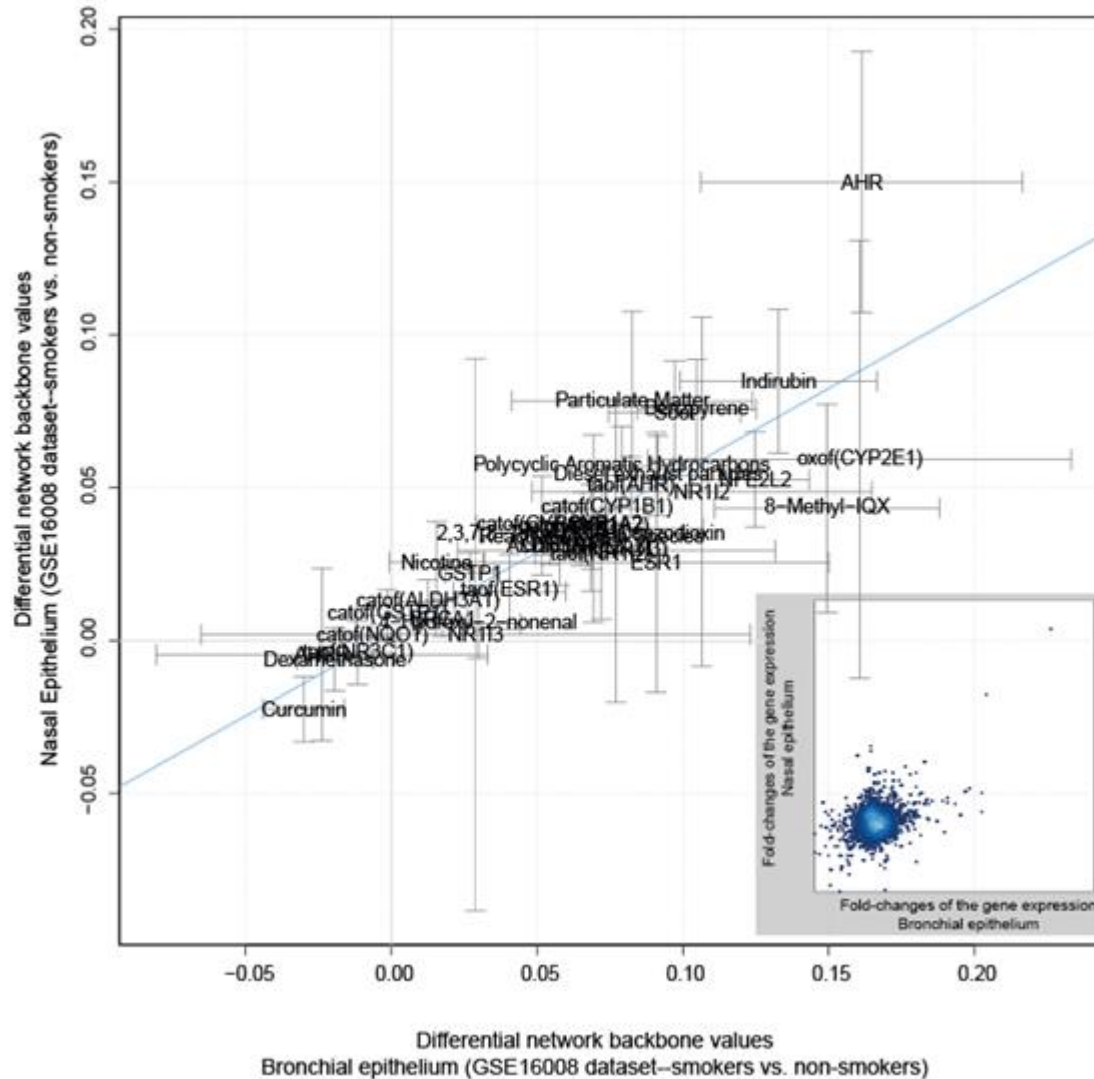


(Ref. Iskandar A. R. et al. BioMed Research International - In press)



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# Nasal as a Surrogate for Bronchus – IN VIVO

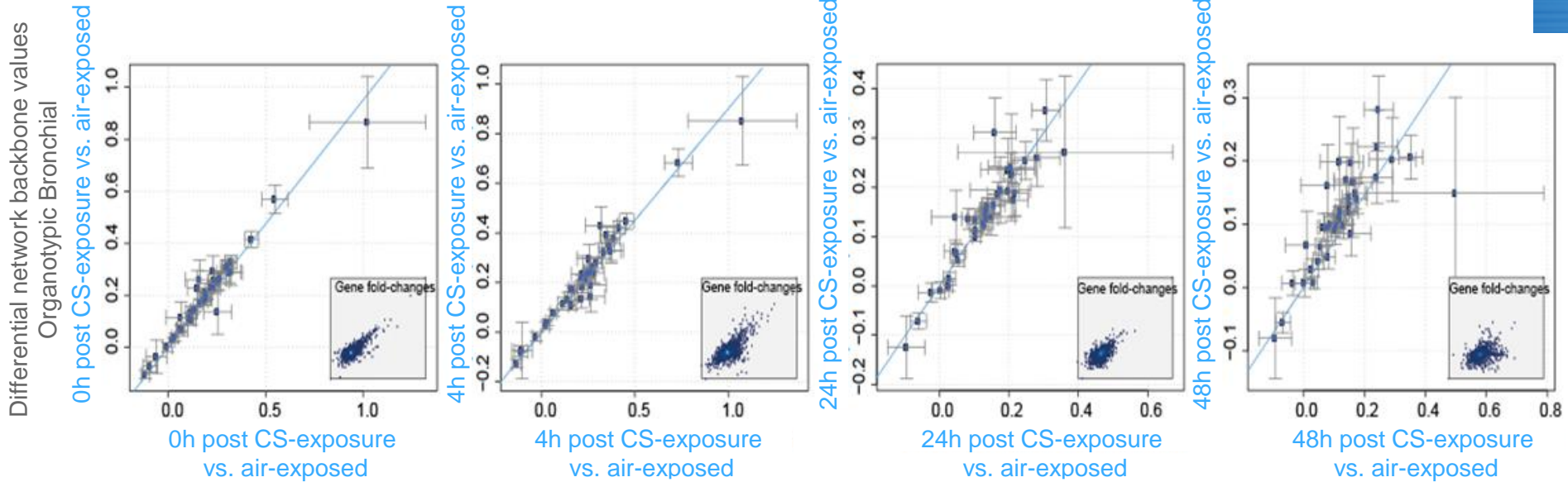


(Ref. Iskandar A. R. et al. BioMed Research International - In press)





# Nasal as a Surrogate for Bronchus – IN VITRO



Differential network backbone values – Organotypic Nasal

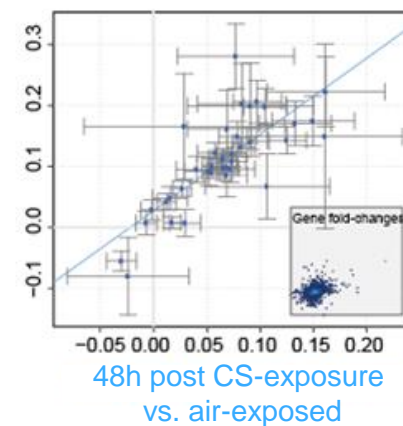
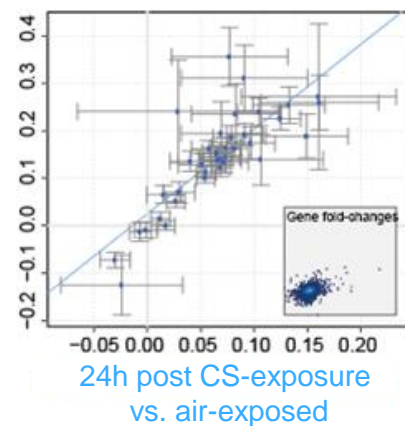
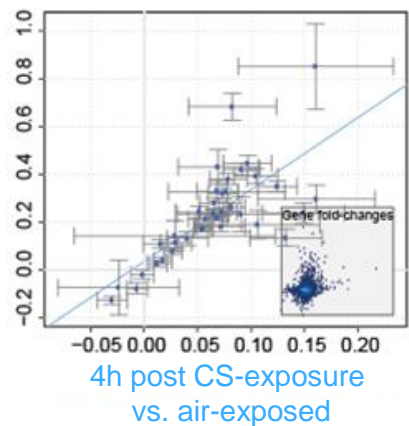
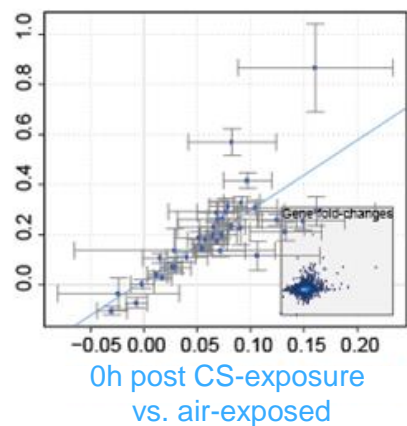
(Ref. Iskandar A. R. et al. *BioMed Research International* - In press)



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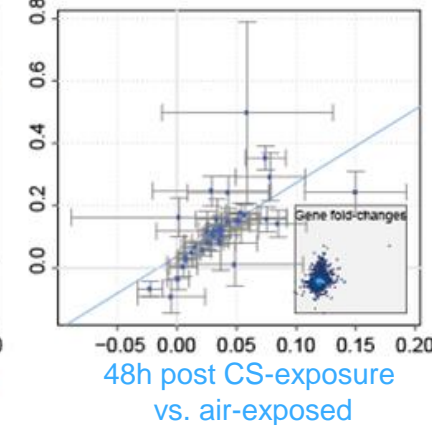
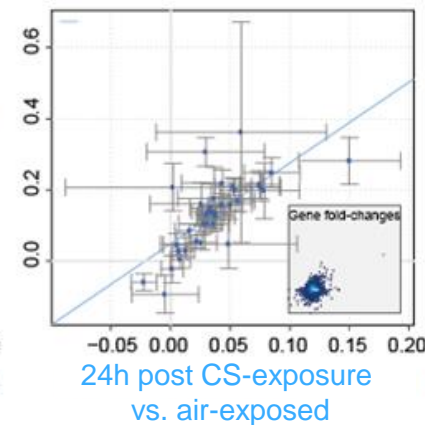
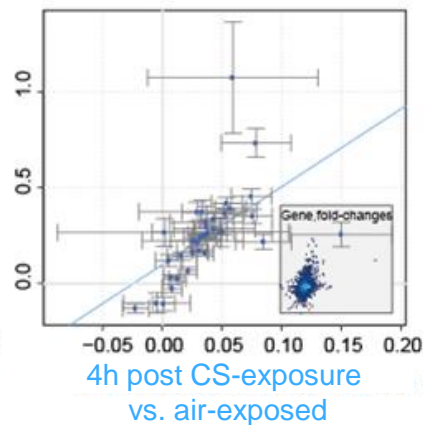
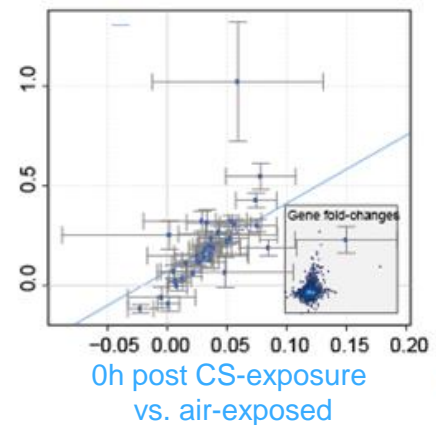
# In Vivo/In Vitro Comparison in the Xenobiotic Metabolism Network Model

Differential network backbone values  
Bronchial Epithelium (Dataset GSE16008)  
Smokers vs. non smokers



Differential network backbone values – Organotypic bronchial

Differential network backbone values  
Nasal Epithelium (Dataset GSE16008)  
Smokers vs. non smokers



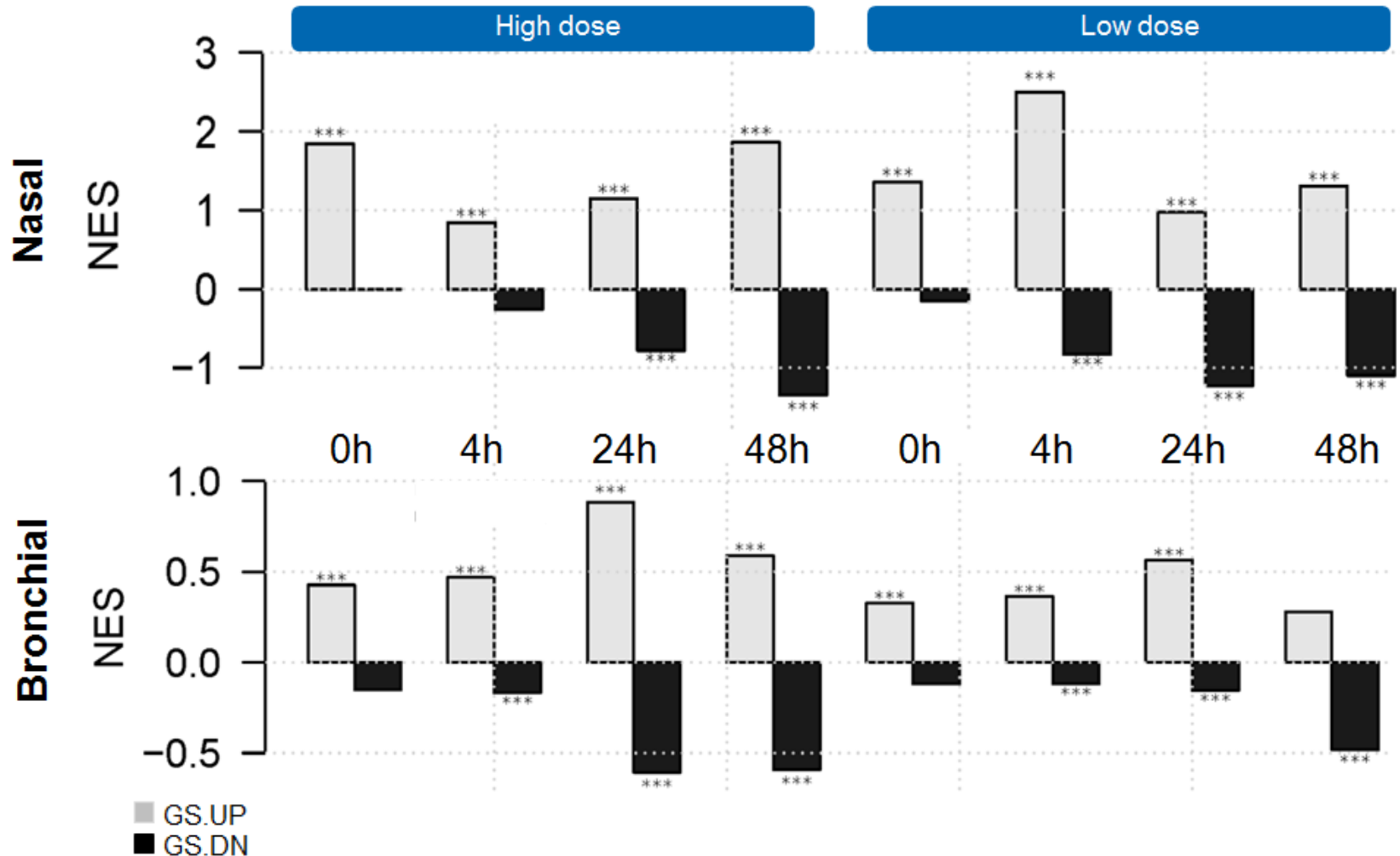
(Ref. Iskandar A. R. et al. BioMed Research International - In press)



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# In Vivo/In Vitro Comparison

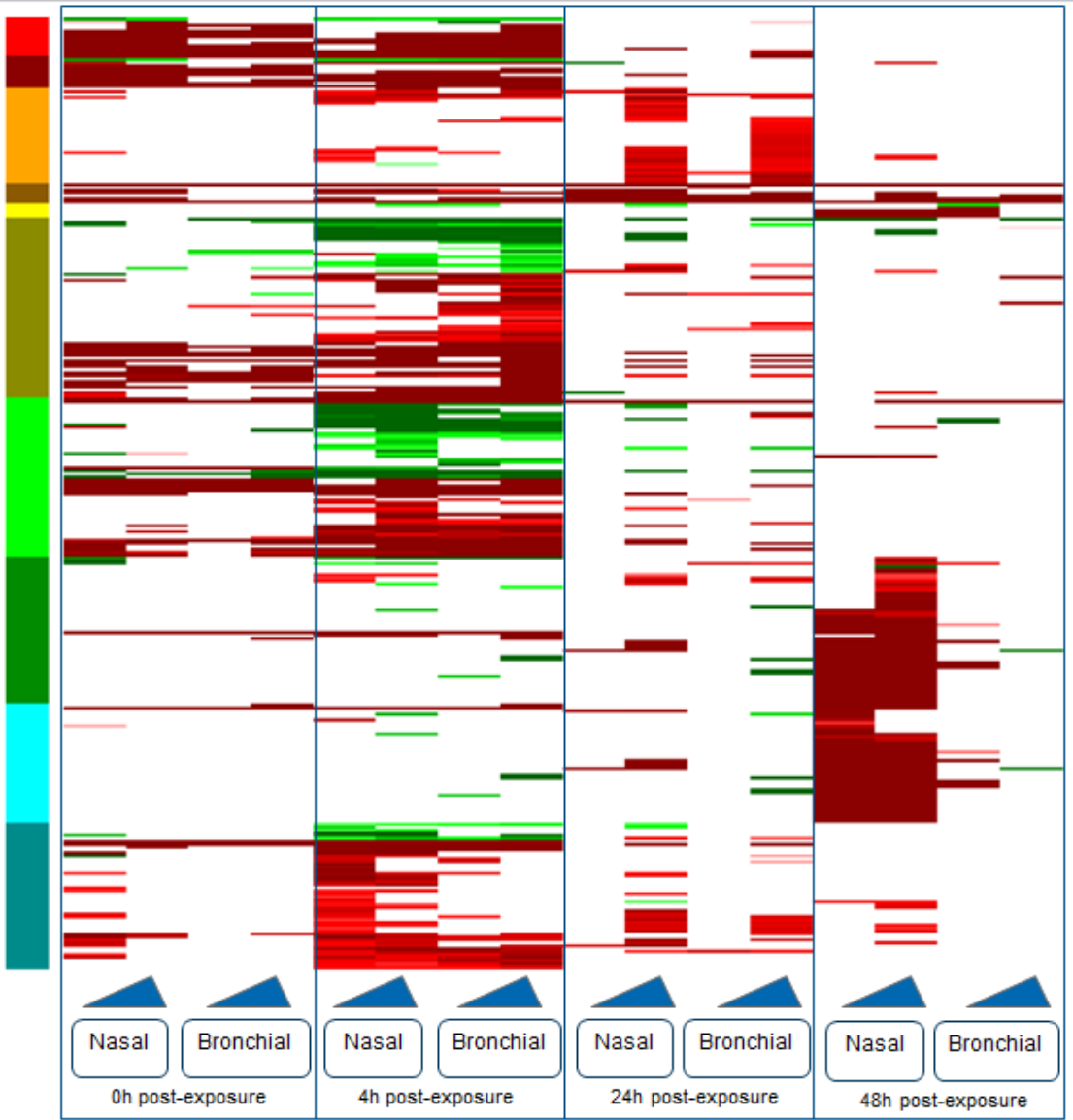
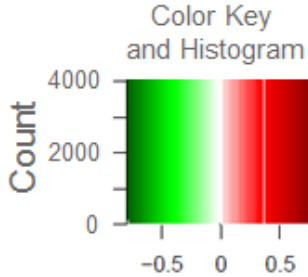
## GENE SET ANALYSIS



# Top 10 Canonical Pathways Perturbed Over Post-Exposure Time and Over Dose in both Nasal and Bronchial Exposed Tissues

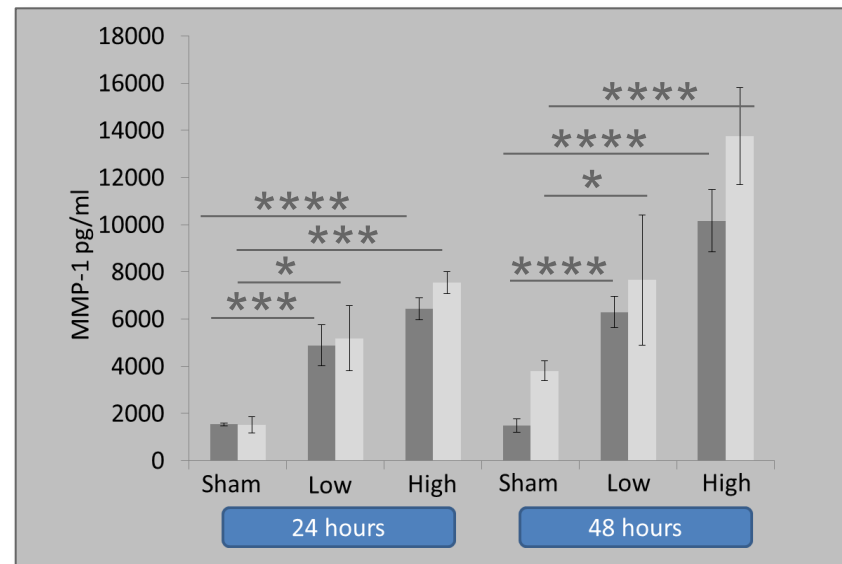
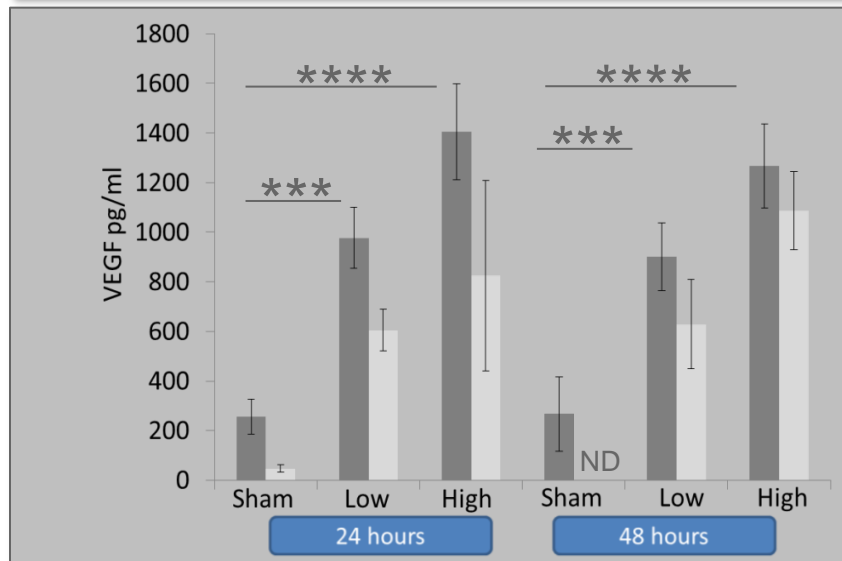
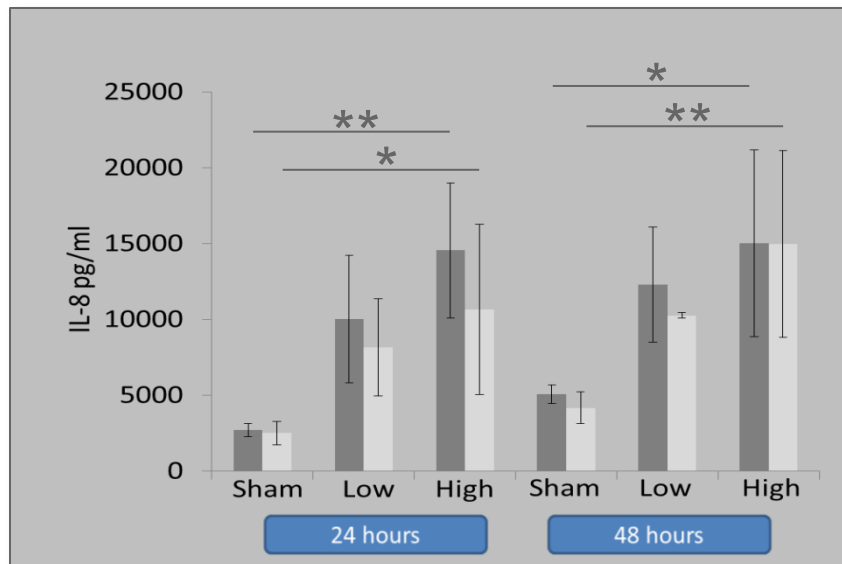
## GENE SET ENRICHMENT ANALYSIS

- AP-1 transcription factor network
- Validated transcriptional targets of AP1 family members Fra1 and Fra2
- Proteasome
- Metabolism of xenobiotics by cytochrome P450
- Aurora B signaling
- MAPK signaling pathway
- Direct p53 effectors
- Genes involved in Cell Cycle, Mitotic
- Genes involved in Cell Cycle
- Genes involved in Metabolism of RNA



The color key and histogram show the distribution of gene counts across the pathways. The color key shows a gradient from green (negative enrichment) to red (positive enrichment), with a peak at 0. The histogram shows the count of genes for each pathway, with a peak around 2000 genes.

# Dose- and Time-dependent Release of Pro-inflammatory Markers in both CS-exposed Bronchial and Nasal Tissues



BRONCHIAL  
NASAL





# CONCLUSIONS

## CS DOSE- AND TIME-DEPENDENCY EFFECTS



- Activation of AP-1 pathway or of the xenobiotic metabolism at early post-exposure time points (0h and 4h).
- Similar dose- and time-dependent regulation of the release of pro-inflammatory markers (VEGF, IL-8 and MMP-1).

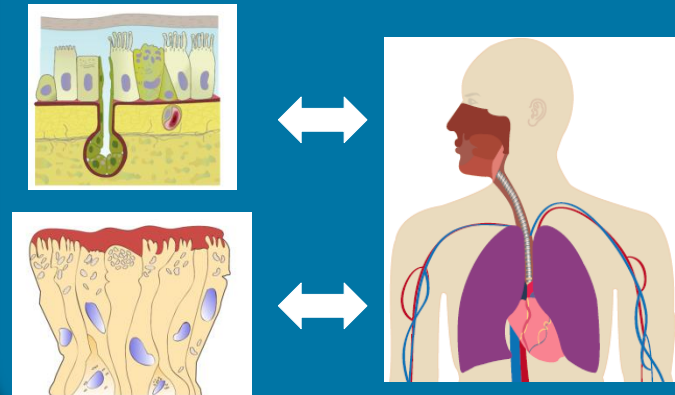


- Up-regulation of genes involved in cell cycle only in nasal tissue culture 48h after exposure  
➔ recovery process specific to the nasal epithelium?

## NASAL AS A SURROGATE FOR BRONCHUS



## COMPARISON IN VIVO / IN VITRO



# Acknowledgment

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