



PMI RESEARCH & DEVELOPMENT

Integrating multi-omics and high-content screening data to link biological network perturbations with cellular phenotypes to elucidate pathways of toxicity

Ignacio Gonzalez Suarez, Ph.D.

Senior scientist,

Philip Morris International R&D



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Integrating multi-omics and high-content screening data
to link biological network perturbations with cellular
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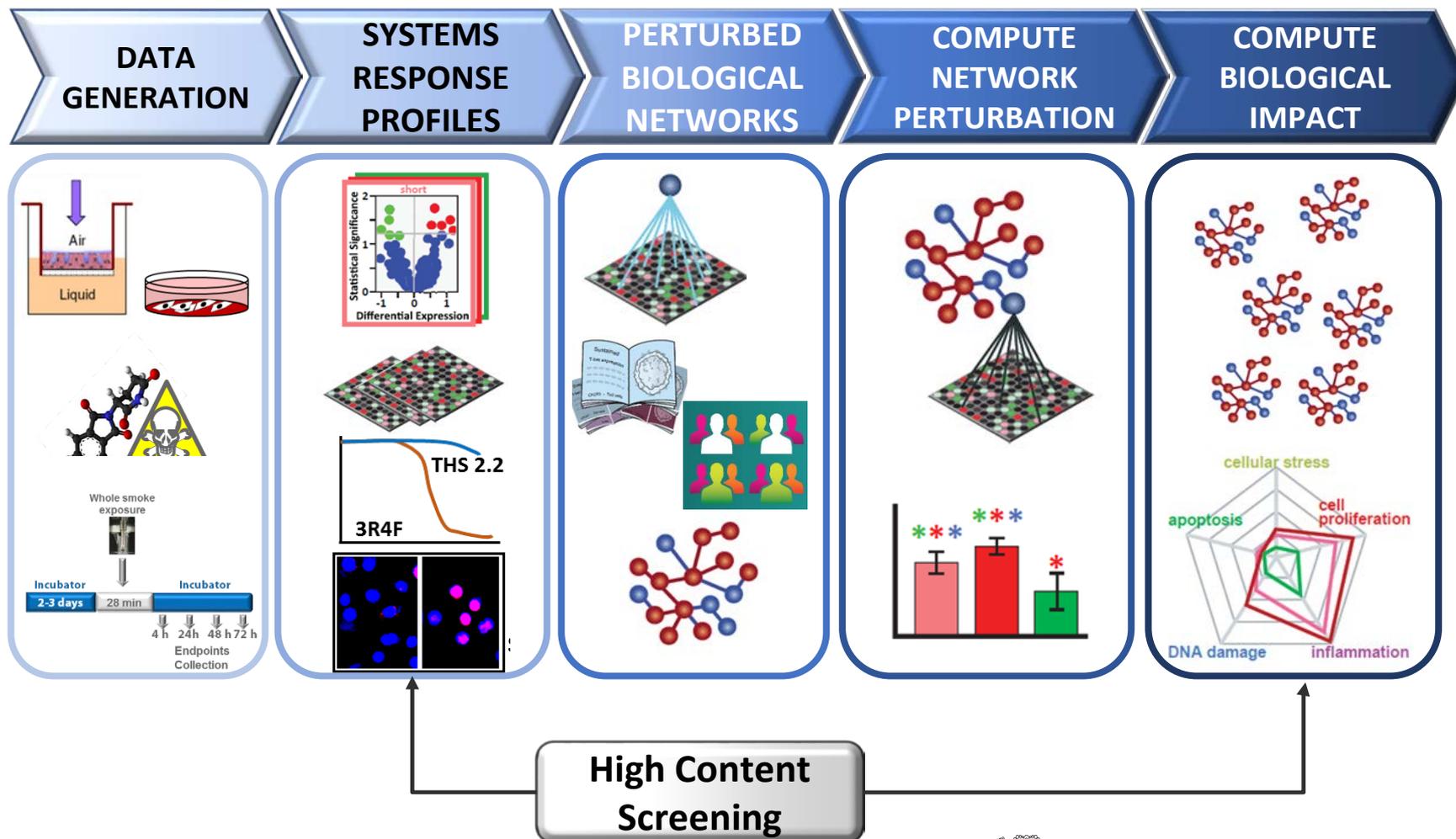
How to apply omics and cellular imaging to toxicological assessment

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Systems Toxicology Assessment workflow at PMI



Toxicological Assessment of Tobacco Products



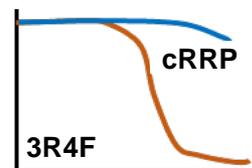
Toxicological Assessment of Tobacco Products

Study Design

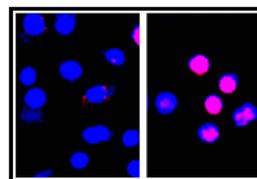
Primary Human
Bronchial epithelial
cells (NHBE)



Exposure

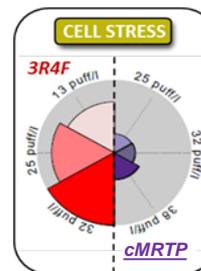


Cell
viability



High
Content
Screening

- Conventional cigarette (3R4F)
- Candidate Reduced Risk Product (cRRP)

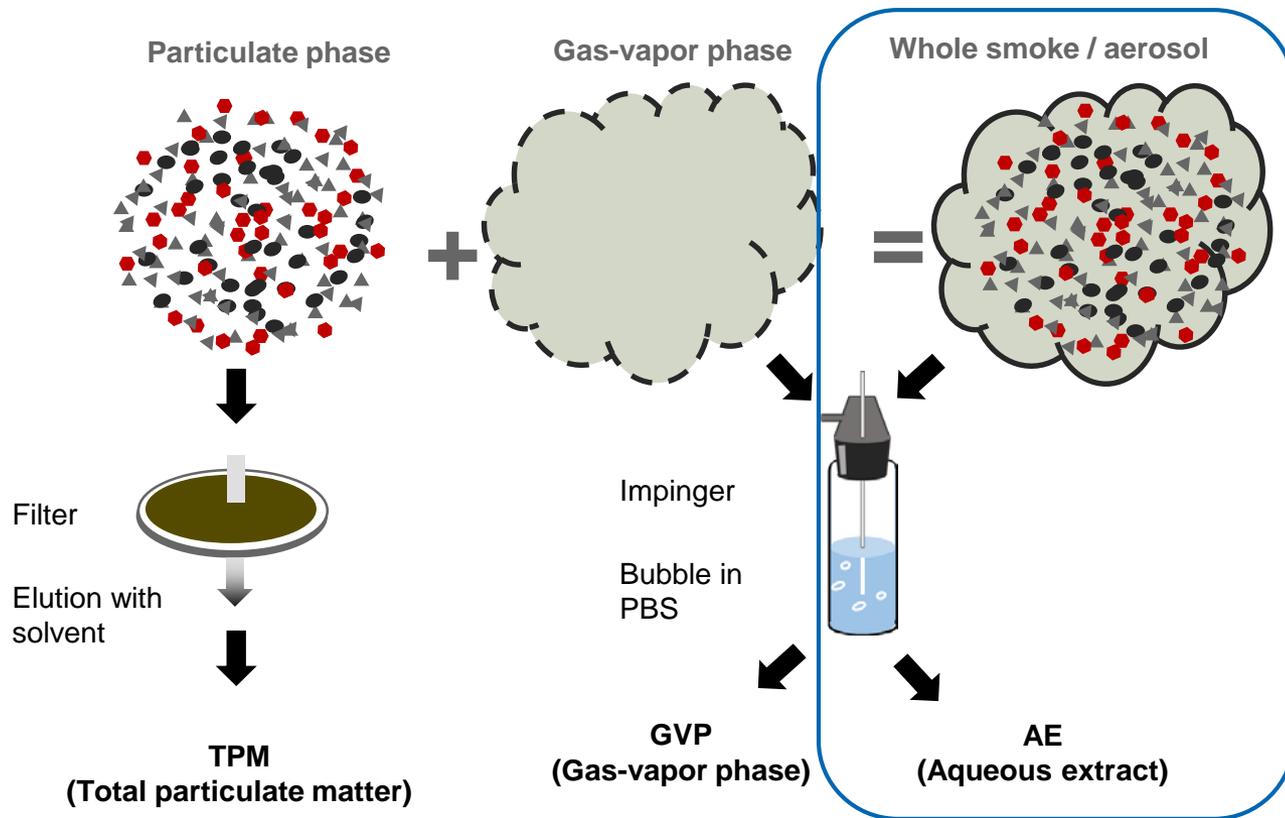


Gene
Expression

What is the Biological impact of a cRRP compared to 3R4F?

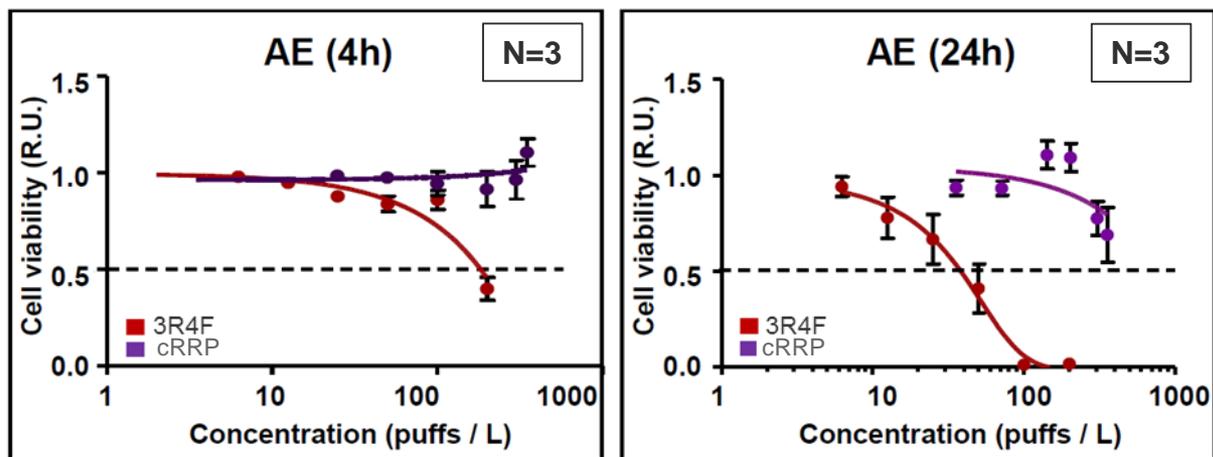
Toxicological Assessment of Tobacco Products

Cigarette smoke / cRRP aerosol



Toxicological Assessment of Tobacco Products

Step I: Cell Viability



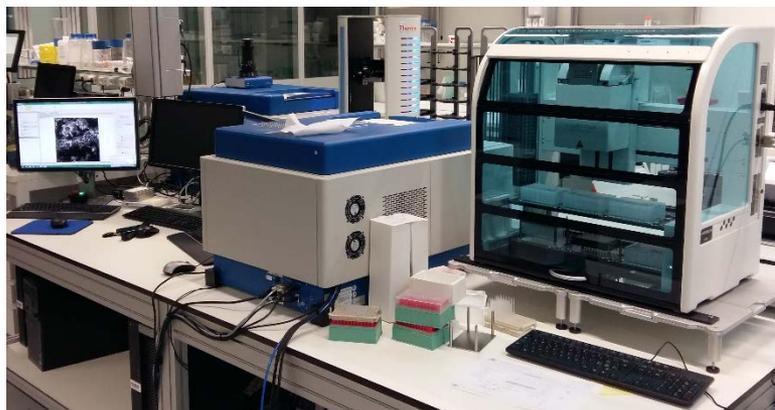
Gonzalez-Suarez et al. Chem. Res. Tox. 2015

- ✓ Dose-dependent decrease in cell viability upon exposure to 3R4F AE.
- ✓ Selected appropriate doses for HCS analysis



Toxicological Assessment of Tobacco Products

Step II: High-Content Screening



- 7 Different Assays
- 14 Toxicological endpoints
- 6 Doses + vehicle
- 3 replicate wells per dose
- 2 time points (4h & 24h)
- ≥ 3 independent experiments

- Cell Count

Proliferation

- Cell cycle

- Phospho-H2AX

DNA Damage

- Phospho cJun

- ROS formation

- GSH content

Cell / Oxidative stress

- Caspase 3/7

- Cytochrome C

- Membrane permeability

- Nuclear size

Apoptosis / necrosis

- Mitochondrial mass

- Mitochondrial potential

Mitochondrial health

- NF- κ B translocation

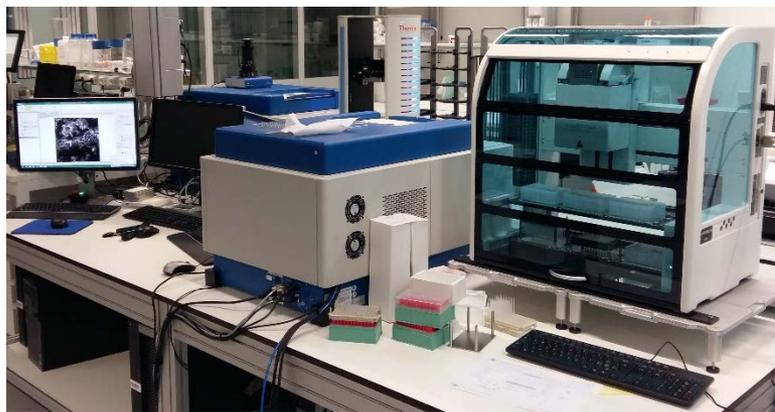
Inflammation

- Gap-Junction

Cell communication

Toxicological Assessment of Tobacco Products

Step II: High-Content Screening



- 7 Different Assays
- 14 Toxicological endpoints
- 6 Doses + vehicle
- 3 replicate wells per dose
- 2 time points (4h & 24h)
- ≥ 3 independent experiments

	3R4F		cRRP	
	4h	24h	4h	24h
Cell Count	-	100	-	200*
DNA Damage (p-H2AX)	200*	200*	-	-
Cell Stress (p-cJun)	-	100	-	200*
ROS Formation	-	100*	-	-
GSH Content	50	100	-	200*
Cell cycle	NA	13	NA	140
Apoptosis (Cytochrome C)	-	100	-	280*
Necrosis (membrane permeability)	100	100	350*	-
Mitochondrial Membrane Potential	-	100	-	280
Mitochondrial Mass	50*	200*	-	-

Gonzalez-Suarez et al. Chem. Res. Tox. 2015

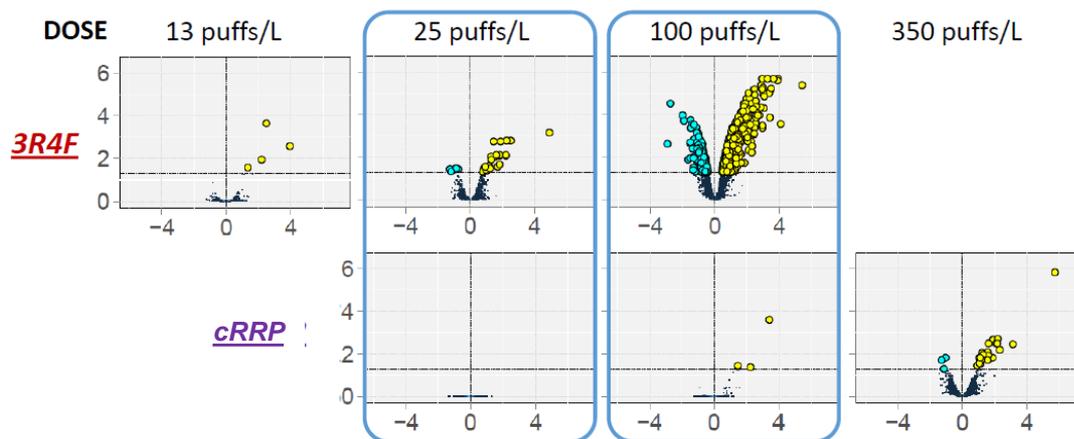
- ✓ Dose-dependent responses in multiple endpoints upon exposure to 3R4F.
- ✓ Selected appropriate doses for Transcriptomics



Toxicological Assessment of Tobacco Products

Gene expression (DEG)

- 2 items: 3R4F & cRRP
- 3 Doses + vehicle
- 1 time points (4h)
- ≥ 3 independent experiments

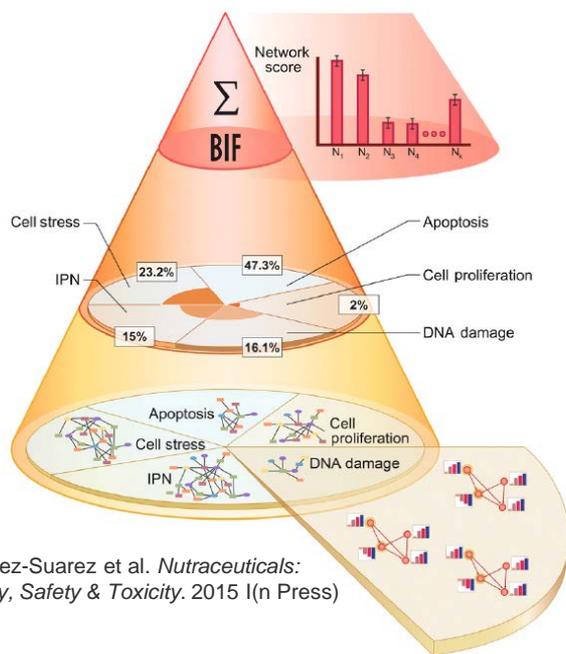


Gonzalez-Suarez et al. *Chem. Res. Tox.* 2015

- ✓ Increased number of DEG in response to 3R4F compared to cRRP.

Toxicological Assessment of Tobacco Products

Gene expression (Network Biology)



Gonzalez-Suarez et al. *Nutraceuticals: Efficacy, Safety & Toxicity*. 2015 I(n Press)

1. Biological Impact



2. Network Level Perturbation



3. Subnetwork Level Perturbation



4. Leading nodes

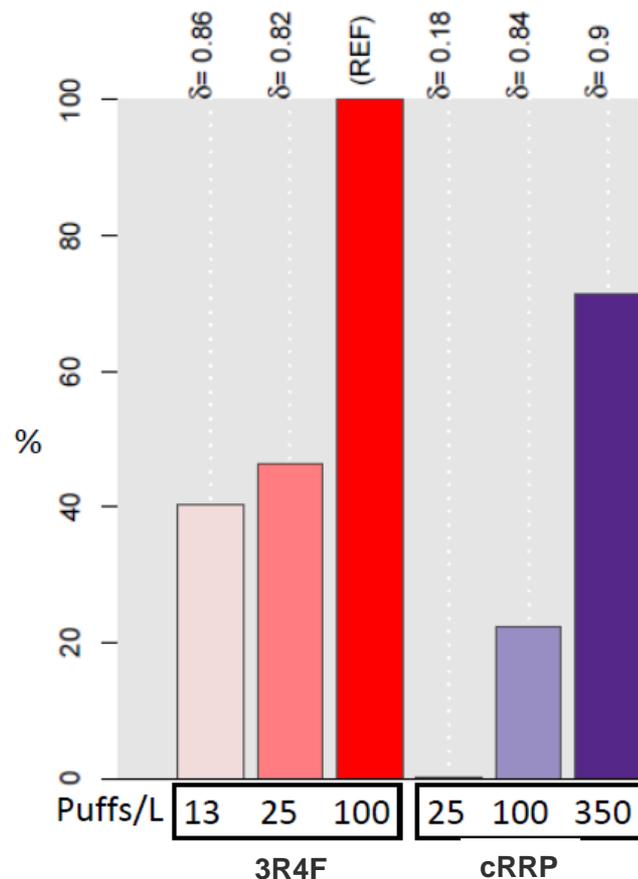
Step-wise approach
Increased Granularity
Increased mechanistic insight

Toxicological Assessment of Tobacco Products

Gene expression: Biological Impact Factor (BIF)

- **Sum of all perturbations** across all biological networks
- **Vehicle control (0%)**
- **Reference value (100%)**
- **Reference:** highest level of overall perturbation
- **δ value (-1 to 1):** compares underlying biology to reference

- ✓ **Dose-dependent responses in 3R4F and cRRP.**
- ✓ **At comparable doses, lower biological impact of cRRP**
- ✓ **δ suggest similar underlying biology between 3R4F and cRRP**

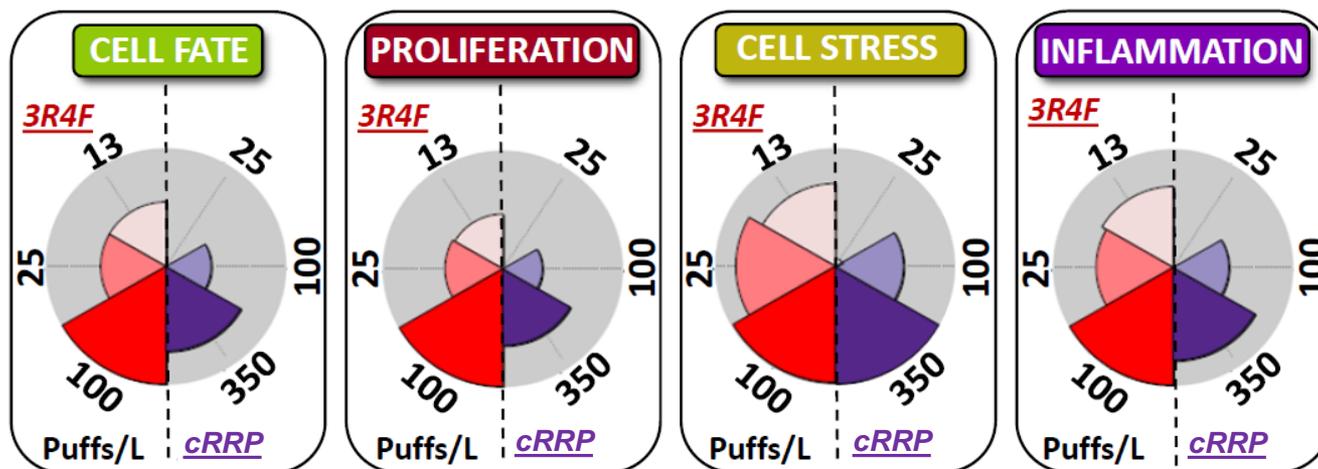


Gonzalez-Suarez et al. *Chem. Res. Tox.* 2015



Toxicological Assessment of Tobacco Products

Gene expression: BIF Mechanistic Components



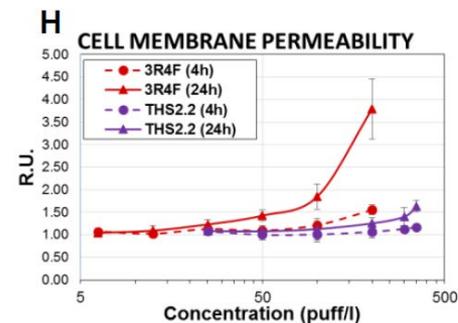
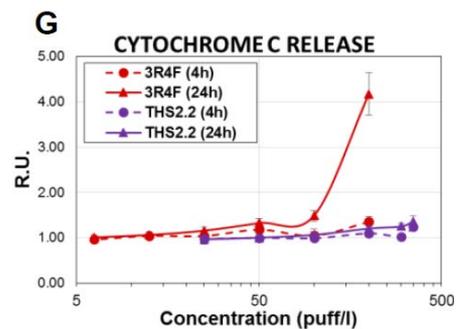
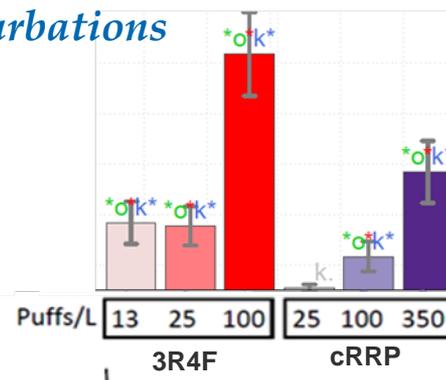
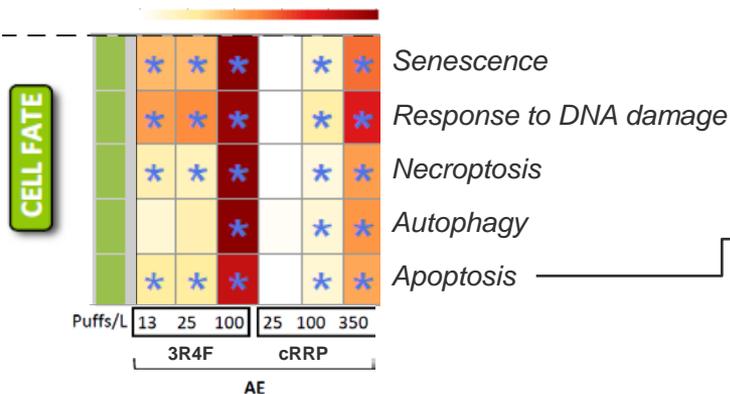
Gonzalez-Suarez et al. *Chem. Res. Tox.* 2015

- BIF values can be separated into its mechanistic components
- Surface area proportional to level of network perturbation
- Vehicle control (0%)
- Values normalized to reference

- ✓ Dose-dependent responses in 3R4F and cRRP.
- ✓ At comparable doses, lower biological impact of cMRTP

Toxicological Assessment of Tobacco Products

Gene expression: Cell Fate Subnetwork Perturbations

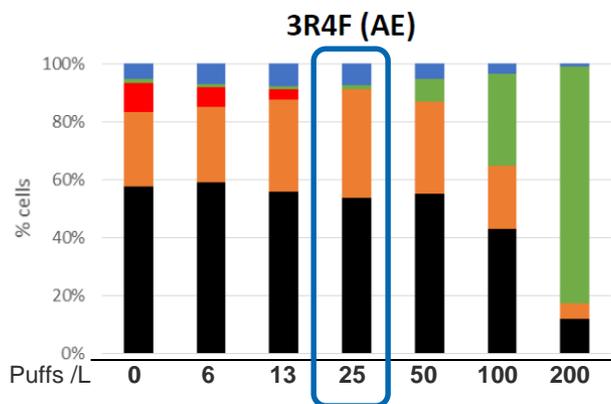
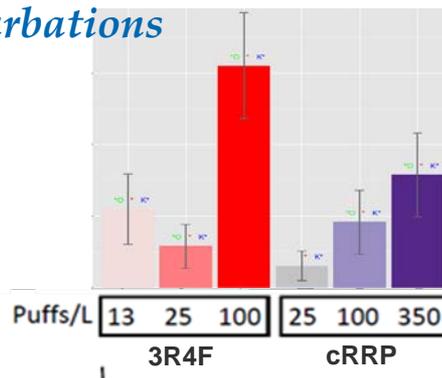
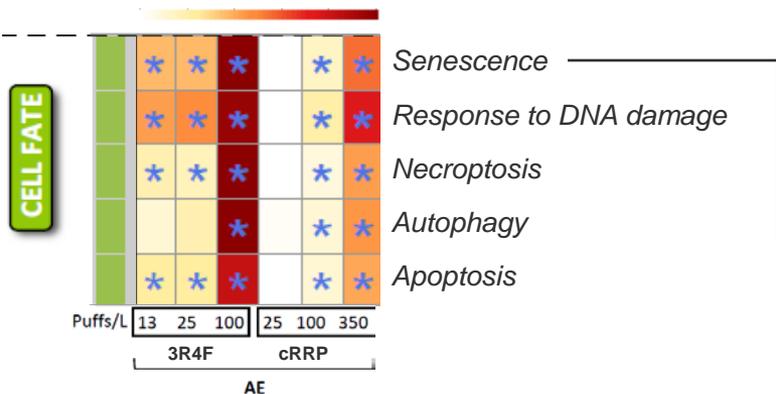


Gonzalez-Suarez et al. *Chem. Res. Tox.* 2015

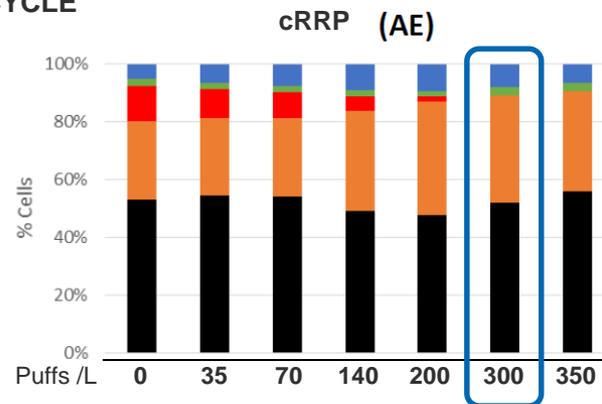


Toxicological Assessment of Tobacco Products

Gene expression: Cell Fate Subnetwork Perturbations



CELL CYCLE



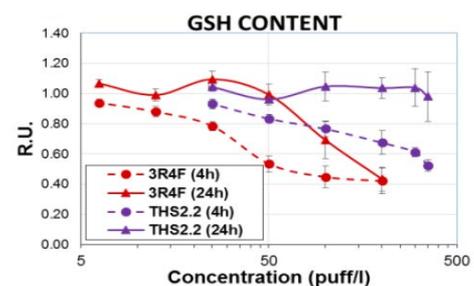
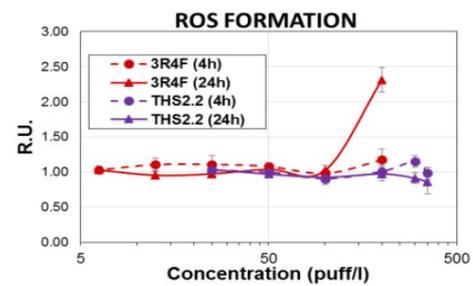
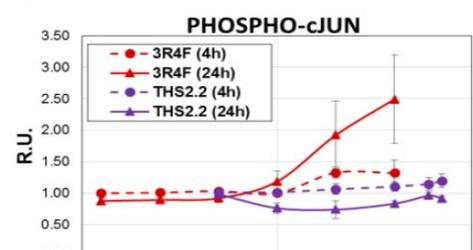
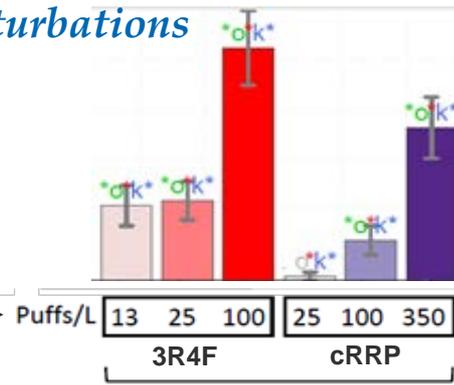
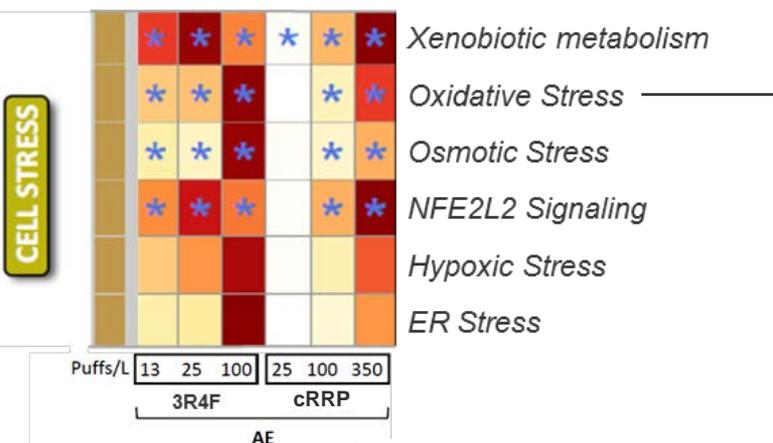
Gonzalez-Suarez et al. *Chem. Res. Tox.* 2015



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Toxicological Assessment of Tobacco Products

Gene expression: Cell stress Subnetwork perturbations



Gonzalez-Suarez et al. Chem. Res. Tox. 2015

Exposure to cRRP has a lower biological impact on NHBE cells compared to 3R4F



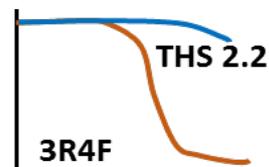
Toxicological Assessment of Environmental toxicants

Study Design

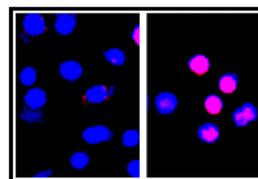
Primary Human
Bronchial epithelial
cells (NHBE)



Exposure



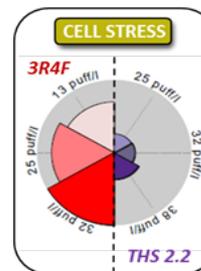
Cell
viability



High
Content
Screening

- Phase I (2012): 15 compounds*
- Phase I (2014): 32 compounds*

*HPHC (harmful & potentially harmful smoke constituents)



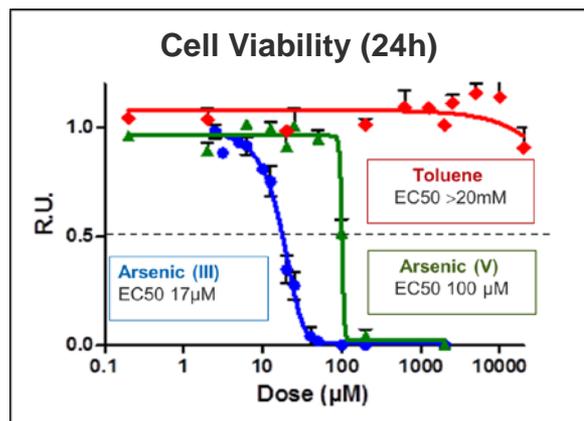
Gene
Expression

What is the Biological impact of these chemicals in NHBE cells?

Toxicological Assessment of Environmental toxicants

Step I: Cell Viability

- 32 toxicants
- 6 Doses + vehicle
- 24h exposure
- ≥ 3 independent experiments



	HPHC	EC50 Value	R ²		HPHC	EC50 Value	R ²
1	Chromium (VI)	4 µM	0.995	17	o-Anisidine	11970 µM	0.968
2	Arsenic (III)	17 µM	0.968	18	2-nitropropane	> 20 mM	-
3	5-Methylchrysene	28 µM	0.961	19	Acetamide	> 20 mM	-
4	Arsenic (V)	100 µM	0.990	20	Acetone	> 20 mM	-
5	Mercury (II)	110 µM	0.999	21	Benzene	> 20 mM	-
6	Selenium (IV)	338 µM	0.982	22	MEK	> 20 mM	-
7	Crotonaldehyde	501 µM	0.994	23	Nitrobenzene	> 20 mM	-
8	Nickel (II)	520 µM	0.999	24	Quinoline	> 20 mM	-
9	Lead (II)	528 µM	0.918	25	Toluene	> 20 mM	-
10	1-Aminonaphthalene	1000 µM	0.964	26	Benz [a] anthracene	> 100 µM	-
11	Naphthalene	1176 µM	0.902	27	Benzo [a] pyrene	> 100 µM	-
12	m-Cresol	2028 µM	0.936	28	Benzo [b] fluoranthene	> 100 µM	-
13	o-Cresol	2170 µM	0.912	29	Benzo [k]fluoranthene	> 100 µM	-
14	p-Cresol	5060 µM	0.900	30	Dibenz [a,h] anthracene	> 100 µM	-
15	Acrilamide	5880 µM	0.981	31	Dibenzo [a,l] pyrene	> 100 µM	-
16	Phenol	6680 µM	0.982	32	Indeno [1,2,3-cd] Pyrene	> 100 µM	-

- ✓ Dose-dependent decrease in cell viability observed in 17 toxicants.
- ✓ Selected appropriate doses for HCS analysis

Toxicological Assessment of Environmental toxicants

High-Content Screening

- 6 Different Assays
- 13 Toxicological endpoints
- 6 Doses + vehicle
- 3 replicate wells per dose
- 2 time points (4h & 24h)
- ≥ 3 independent experiments

✓ 10 toxicants selected for Transcriptomics

✓ 3 Doses selected for Transcriptomics:

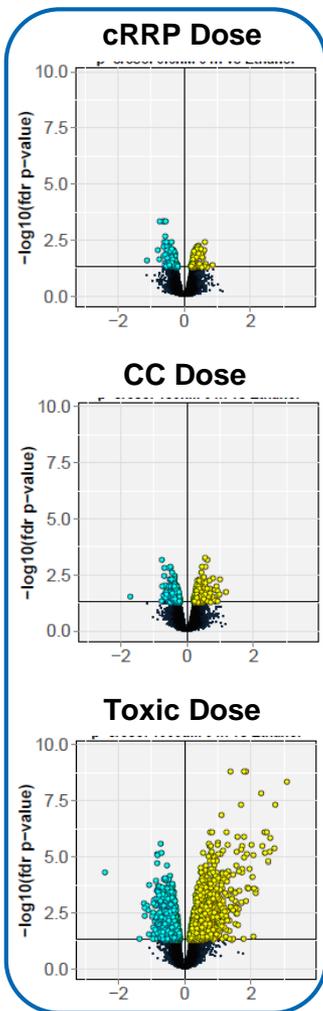
- cRRP dose
- 3R4F (CC) dose
- HCS-toxic dose

HPHC		Cell Loss	DNA Damage	Stress Kinase	GSH Content	Oxidative Stress	Caspase 3/7	Cytochrome C Release	Cell Membrane Permeability	Mitochondrial Membrane Potential	Mitochondrial Mass
5-Methylchrysene	4h	-	-	✓	✓	-	-	-	-	-	-
	24h	✓	✓	✓	✓	-	-	-	-	-	-
Arsenic (III)	4h	-	-	✓✓	✓	-	-	-	-	-	-
	24h	✓	✓	✓✓	✓✓	-	✓	✓✓	✓	✓	-
Lead (II)	4h	-	-	-	✓	-	-	-	-	✓	✓✓
	24h	✓	-	-	✓	-	-	✓	-	✓	✓✓
m-Cresol	4h	-	✓✓	✓	✓	✓	-	-	-	-	-
	24h	✓	✓✓	-	✓✓	-	✓	-	✓	✓	-
Mercury (II)	4h	-	✓✓	-	✓✓	✓	✓	✓✓	✓✓	✓✓	✓✓
	24h	✓	✓✓	-	✓✓	✓	✓✓	✓✓	✓✓	✓✓	✓✓
Naphthalene	4h	-	-	-	✓✓	✓	-	-	-	-	-
	24h	✓	✓	-	✓✓	-	-	-	-	-	-
o-Anisidine	4h	-	✓	-	✓✓	-	-	-	-	-	-
	24h	✓	✓✓	-	✓✓	-	✓✓	-	✓✓	-	✓
o-Cresol	4h	-	✓✓	-	✓✓	✓	-	-	-	-	-
	24h	✓	✓	-	✓✓	-	✓✓	✓	✓✓	-	-
p-Cresol	4h	-	✓✓	✓	✓	-	-	-	-	-	✓
	24h	✓	✓✓	✓	✓✓	-	✓	-	-	-	✓
Selenium (IV)	4h	-	✓✓	-	✓✓	✓	-	✓✓	-	-	✓✓
	24h	✓	✓✓	✓	✓✓	-	-	✓✓	✓✓	-	-
1-aminonaphthalene	4h	-	✓✓	-	✓	✓✓	-	-	✓✓	-	-
	24h	✓	✓✓	✓	✓	✓✓	✓	✓	✓✓	-	-
Chromium (VI)	4h	-	✓	-	✓	-	-	-	-	-	-
	24h	-	✓✓	-	✓	-	✓✓	-	✓✓	-	-
Crotonaldehyde	4h	-	✓✓	-	-	✓✓	✓	-	✓✓	-	-
	24h	✓	✓✓	✓	-	✓	✓	✓✓	✓	✓	-
Acrylamide	4h	-	✓✓	-	✓	-	-	-	-	-	-
	24h	✓	✓✓	✓	✓	-	-	-	✓	-	-
Phenol	4h	-	✓	-	✓	-	-	-	-	-	-
	24h	✓	✓✓	-	✓	-	✓	-	✓	-	✓
Nickel (II)	4h	-	-	-	✓✓	-	-	-	-	-	-
	24h	✓	-	-	✓✓	-	✓	-	-	-	-
Arsenic (V)	4h	-	-	-	✓✓	-	-	-	-	-	-
	24h	✓	-	-	✓	-	-	-	-	-	-

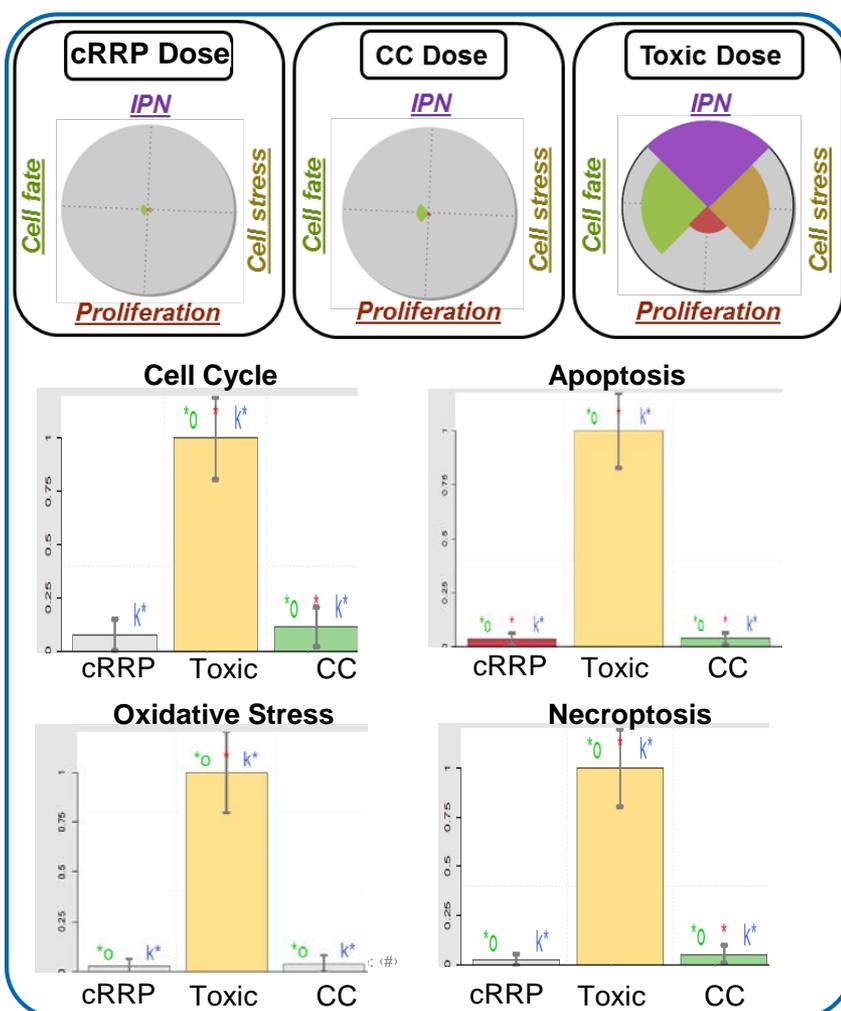


In vitro Toxicological assessment of p-Cresol

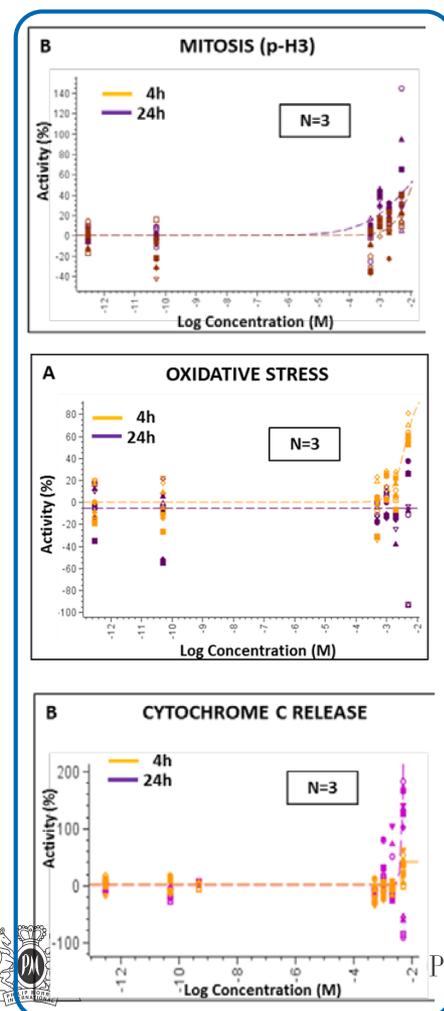
Gene Expression



Network Biology

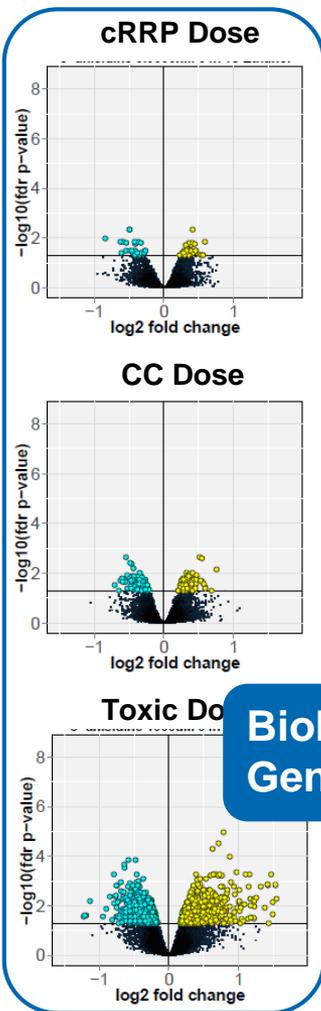


High Content Screening

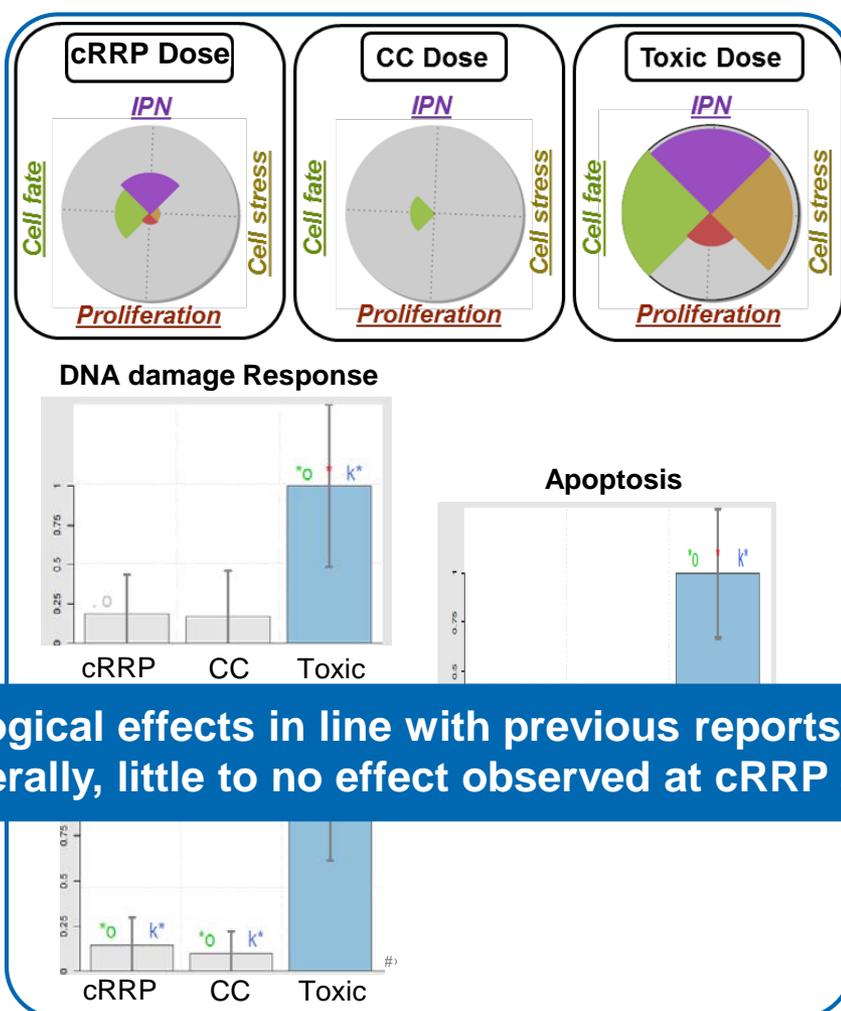


In vitro Toxicological assessment of o-Anisidine

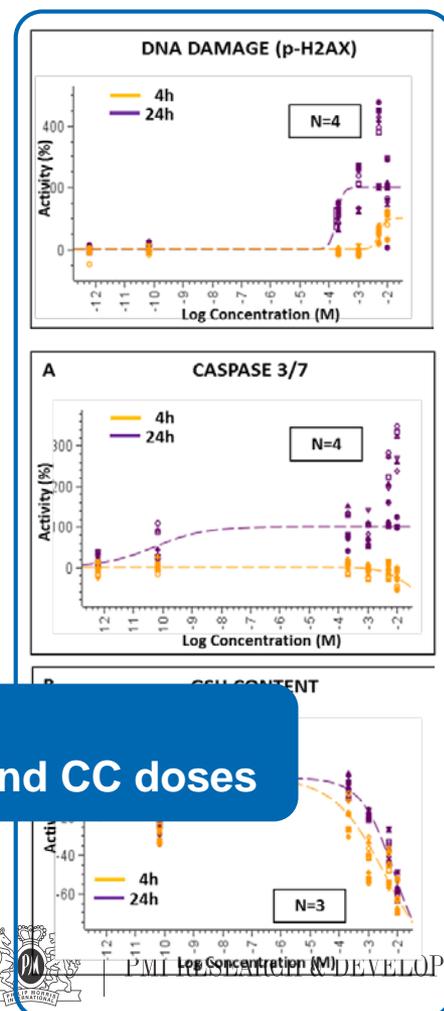
Gene Expression



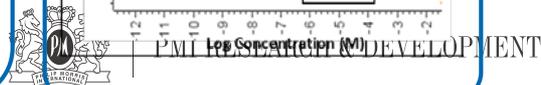
Network Biology



High Content Screening



Biological effects in line with previous reports
Generally, little to no effect observed at cRRP and CC doses



Summary

- The combination of systems biology and high-throughput imaging tools is a valuable approach to investigate molecular mechanisms of toxicity:
 - *Mechanistic insight into toxicity pathways activated upon exposure*
 - *Investigate biological perturbations at sub-cytotoxic exposures*
 - *Systematic and robust assessment*

- Challenges and future directions:
 - *Continuous improvement of Biological Networks*
 - *Incorporation of additional “omics” endpoints*
 - *Increase the number of HCS-based endpoints*
 - *Expand the number of cellular models*

- Ensure transparency and data traceability
- Foster collaboration and data sharing



Acknowledgements

Aerosol Generation Lab:

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- Stephanie Boue

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- Stephan Frentzel

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- Samantha Ellis
- Heather Woodhouse

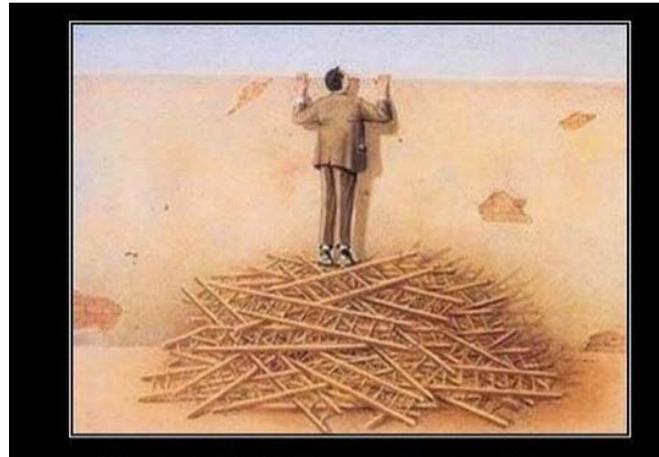
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- Carole Mathis
- Julia Hoeng
- Manuel c. Peitsch

...and many more...



Thank you for your attention and Q&A



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**“It doesn’t matter how many resources you have,
if you don’t know how to use them, they will never be enough”**