

SmartNanoTox

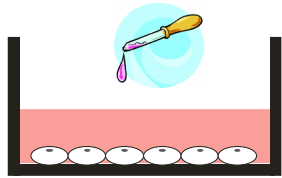
SmartNanoTox Project Online Conference, 24 June 2020

Mechanisms of toxicity related to surface activity, and shape-induced cell-particle interaction

Luc Ferrari

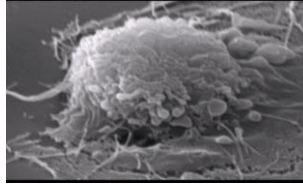
Institut Jean Lamour, UMR CNRS 7198, Université de Lorraine, Nancy, France

Cells, cytotoxicity

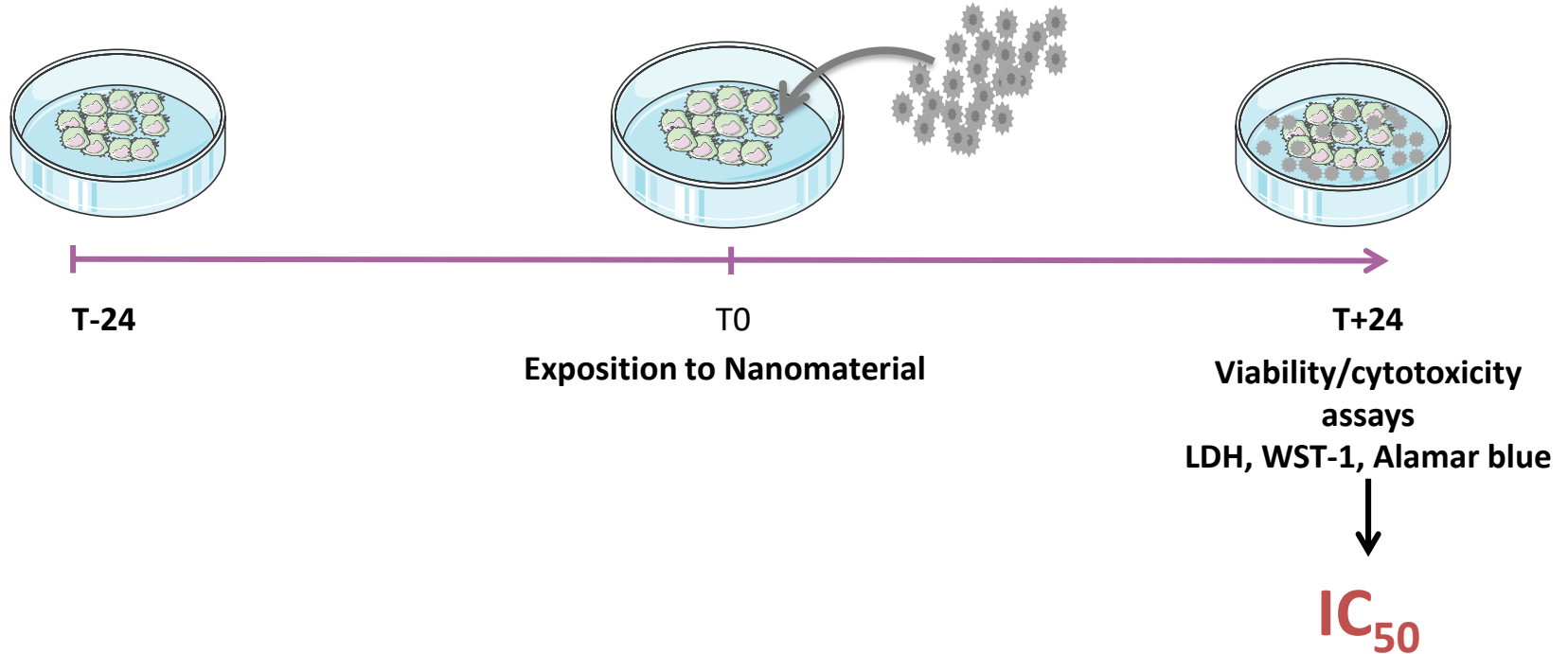
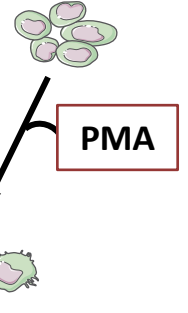


In vitro, submerged

NR8383, rats

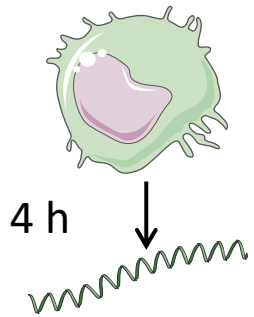


THP-1, human

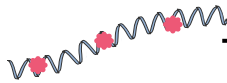


Transcriptomic protocol, simplified

$\frac{1}{4}$ IC₅₀ Nanomaterial

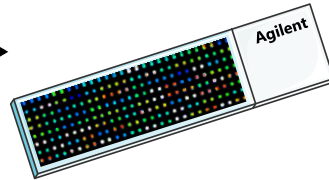


mRNA



cRNA

DNA chip



Data normalisation



Data analysis



Binding to
cyanin-3

Hybridization

Benjamini Hochberg test
 p -value < 0,001
FC > 1,51

Regulated
pathways
 p -value < 0,05
or
 P -value < 0,001

Proteomic protocol, simplified

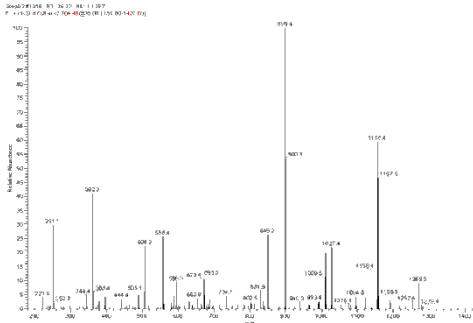
$\frac{1}{4}$ IC₅₀ Nanomaterial



24 h



Proteins



Mass Spectrometry

Data analysis

Ingenuity Pathway Analysis

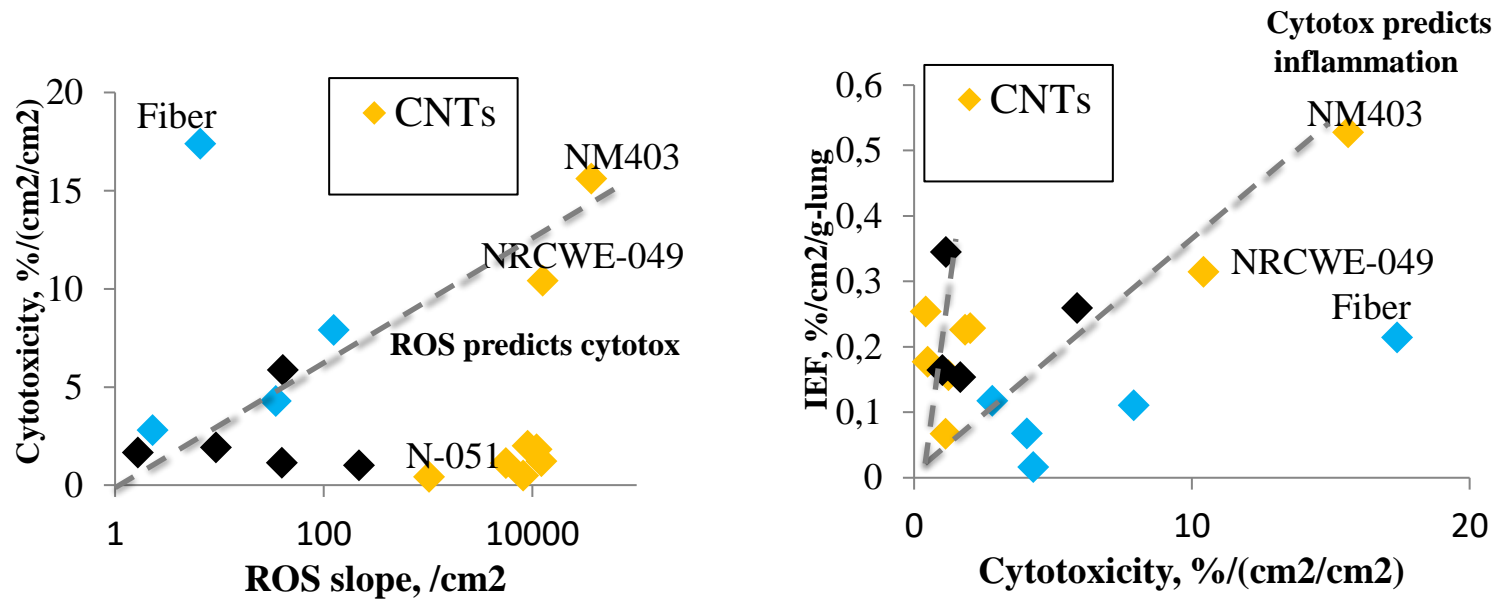


Functional protein association networks



Biomolecular and biomedical Research (Ireland)
H. Cassidy et D. Gomez

ROS–Cytotoxicity/Cytotoxicity-Inflammatory Correlations



Not all of the observed cytotoxicity (reduction in viability) is due to nanomaterial-induced ROS



Cytotoxicity (viability) predicts acute lung inflammation (mouse), but only for some of the nanomaterials

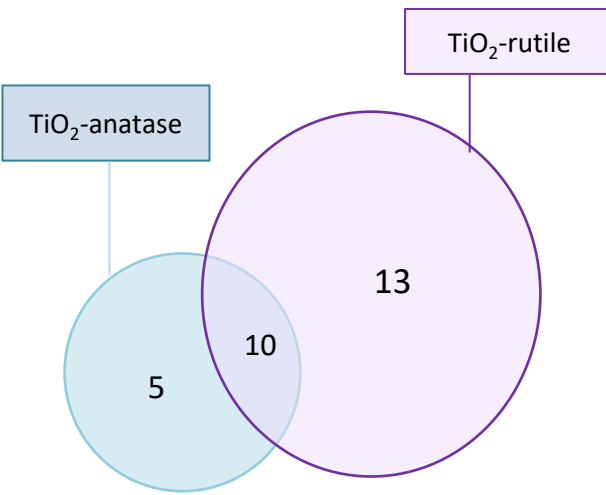


No activity pattern available

Normal regulation

Response		Enrichment score	
Functional categories	Canonical pathways	ZnO NM110	ZnFe ₂ O ₄ NRCWE-021
Stress response	• Mitochondrial dysfunction		
	• Oxidative phosphorylation		
	• mTOR signaling		
	• NRF2-mediated oxidative stress response		
	• Sirtuin signaling		
Cell cycle/proliferation	• PI3K/AKT signalling		
	• VEGF signaling		
Protein synthesis/modification	• EIF2 signaling		
	• Regulation of eIF4 and P70S6K signaling		
	• Protein ubiquitination pathway		
	• Unfolded protein response		
Cell mobility	• Paxillin signaling		
	• Integrin signaling		
Lipid homeostasis	• Superpathway of cholesterol biosynthesis		
Cell modelling	• Actin cytoskeleton signaling		
Cancer	• Cancer drug resistance by drug efflux		
Metal exposure response	• Iron homeostasis signaling pathway		

TiO₂, anatase vs rutile, dysregulated genes in NR8383

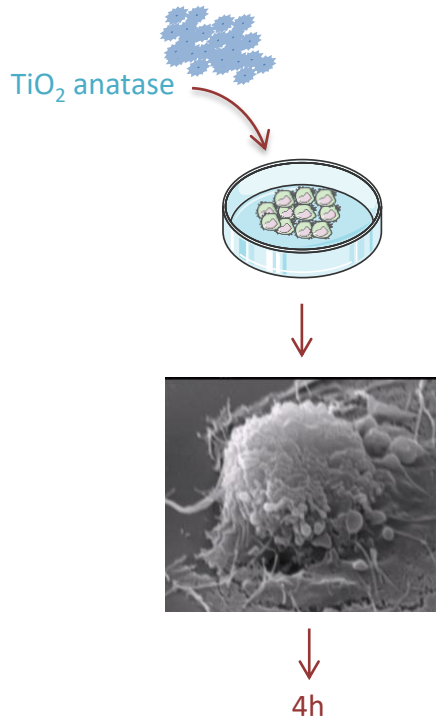


p-valeur < 0,001, FC > 20

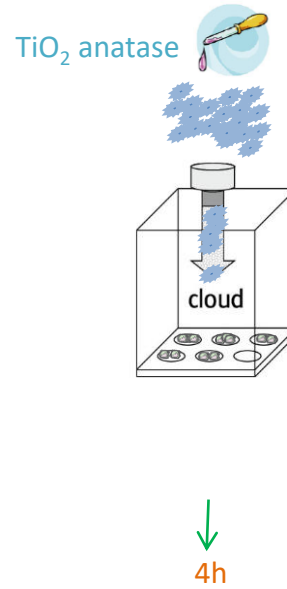
Genes	Genes name	FC (exposure to anatase)	FC (exposure to rutile)
<i>Cxcl2</i>	Chemokine (C-X-C motif) ligand 2	63.36	71.32
<i>Slpi</i>	Secretory Leukocyte Peptidase Inhibitor	51.27	31.19
<i>Slpil3</i>	Secretory Leukocyte Peptidase Inhibitor 3	49.57	28.27
<i>Tac4</i>	Tachykinin 4	42.53	56.20
<i>Tbkbp1</i>	TBK1 Binding Protein 1	23.24	24.43
<i>Phf19</i>	PHD Finger Protein 19	20.03	20.33
<i>Pla2g2d</i>	Phospholipase A2 Group IID	-31.56	-56.16
<i>Dynlt3</i>	Dynein Light Chain Tctex-Type 3	-31.57	-83.03
<i>Cybb</i>	Cytochrome B-245 Beta Chain	-36.61	-85.31
<i>Calcr</i>	Calcitonin Receptor	-36.77	-100.62

TiO₂ : 3 cm²/cm²

Submerged

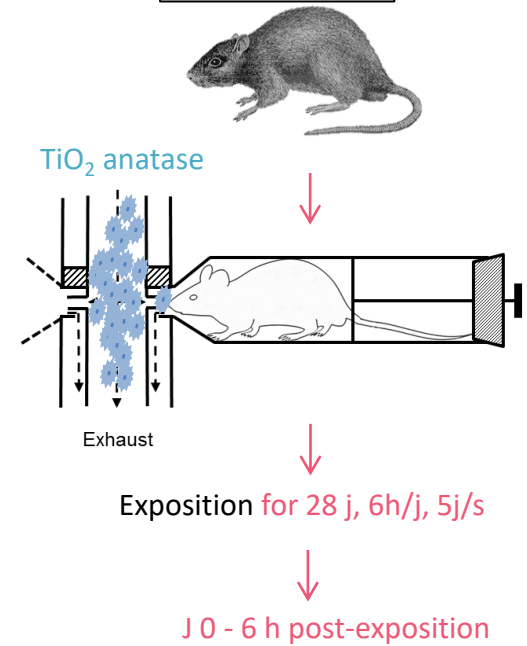


Air-Liquid Interface

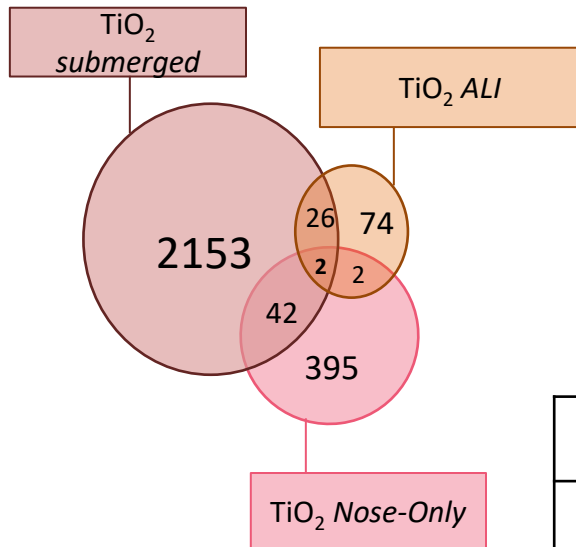


Transcriptomic

Nose-only



L Gaté
C Seidel
L Chezeau

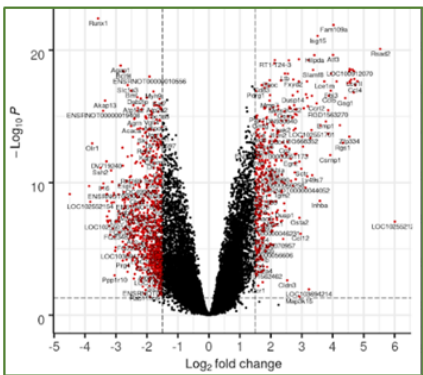


p-value < 0,05, FC > 1,5

Gene ID	FC <i>sub</i>	FC <i>ALI</i>	FC <i>nose</i>	Protein	Biological fonction
<i>Ccl4</i>	3,92	2,12	1,76	C-C Motif Chemokine Ligand, 4	inflammatory response
<i>Cxcl2</i>	63.36	3,16	3.72	C-X-C Motif Chemokine Ligand 2	chimiotoxicity, inflammatory response
<i>Ccl3</i>	3.56	—	1.94	C-C Motif Chemokine Ligand, 3	inflammation
<i>Mmp7</i>	2.72	—	2.66	Metalloproteinase 7	cell division, inflammation
<i>Ccl7</i>	—	1,53	4,8	C-C Motif Chemokine Ligand, 7	inflammatory response

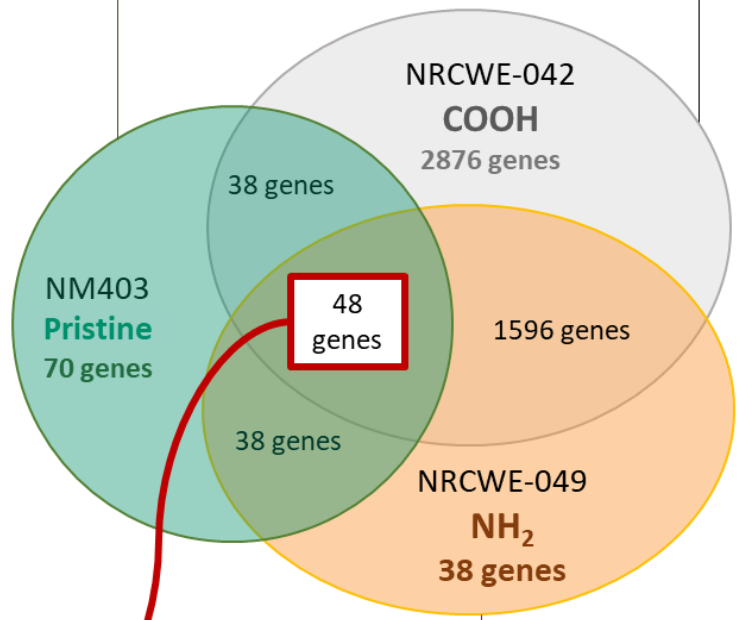
Functionalisation effects on the regulated answer to CNTs, in NR8383

NR8383
Time: 4 h
Doses: 1/4 IC₅₀



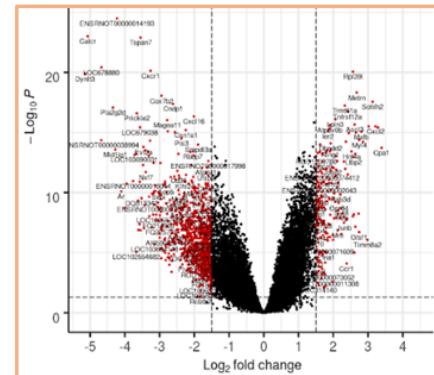
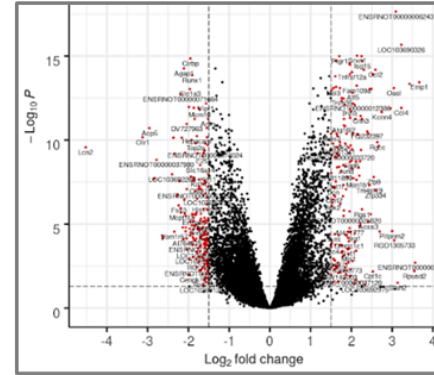
Oxidative stress

Traduction



Commons
Cytoskeleton; Ribosomes
Inflammation

Cytokines production



■ Passed both cut-offs and up-regulated ■ Failed to pass both cut-offs

Functionalisation effects on the regulated response to NMs, in NR8383

Dysregulated pathways after NMs exposure of NR8383		
NM403	NRCWE-042	NRCWE-049
<ul style="list-style-type: none"> • Role of Braca1 in DNA damage <ul style="list-style-type: none"> • Atm signaling • Vitamin-C Transport • Cell cycle: G2/M DNA Damage Checkpoint Regulation • Nrf2 mediated oxidative stress response 	Common	
	<ul style="list-style-type: none"> • Mitochondrial dysfunction <ul style="list-style-type: none"> • EIF2 signaling • Sirtuin signaling • Oxidative phosphorylation 	
	Different	
	Regulation of eIF4 and p70S6K signaling	mTOR signaling

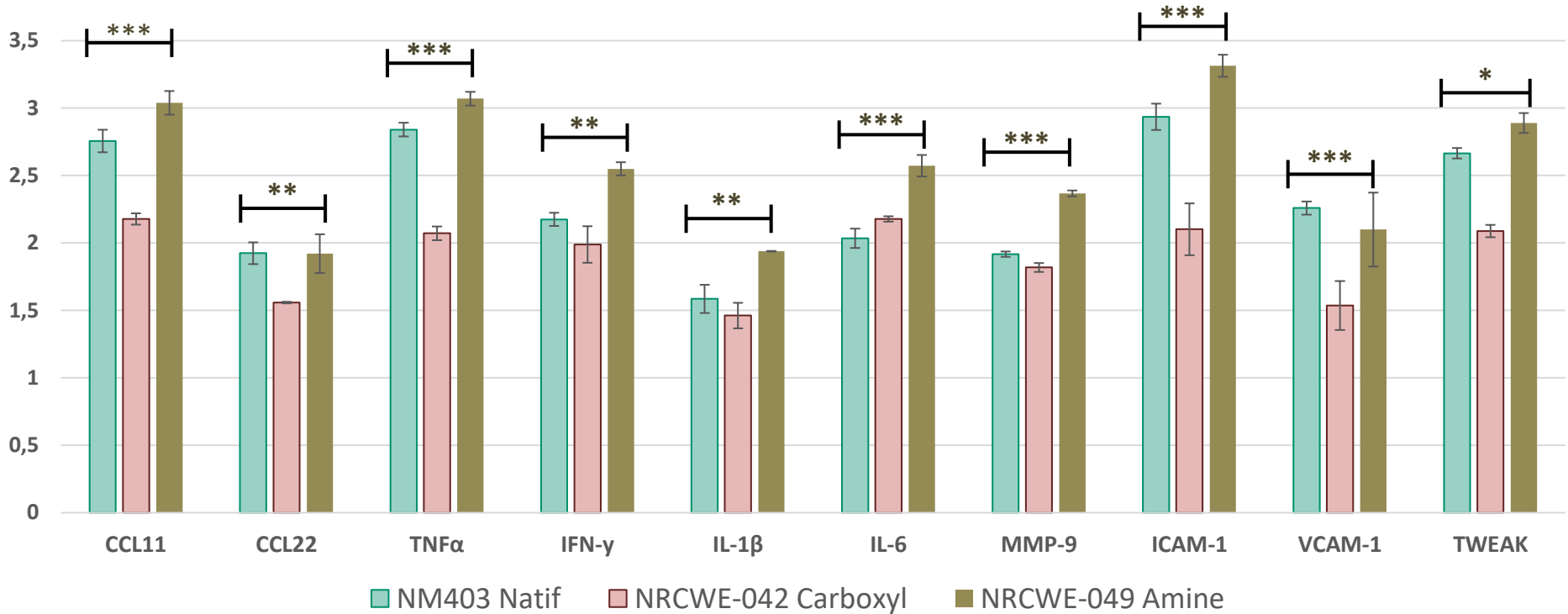
NM403
Pristine
DNA damage
Oxidative stress

NRCWE-042
Fonction : COOH
ER Stress
Protein synthesis

NRCWE-049
Fonction : NH₂
ER Stress
Protein synthesis

Cytokines expression in NR8383 after exposure to NM

(Relative Protein amount)



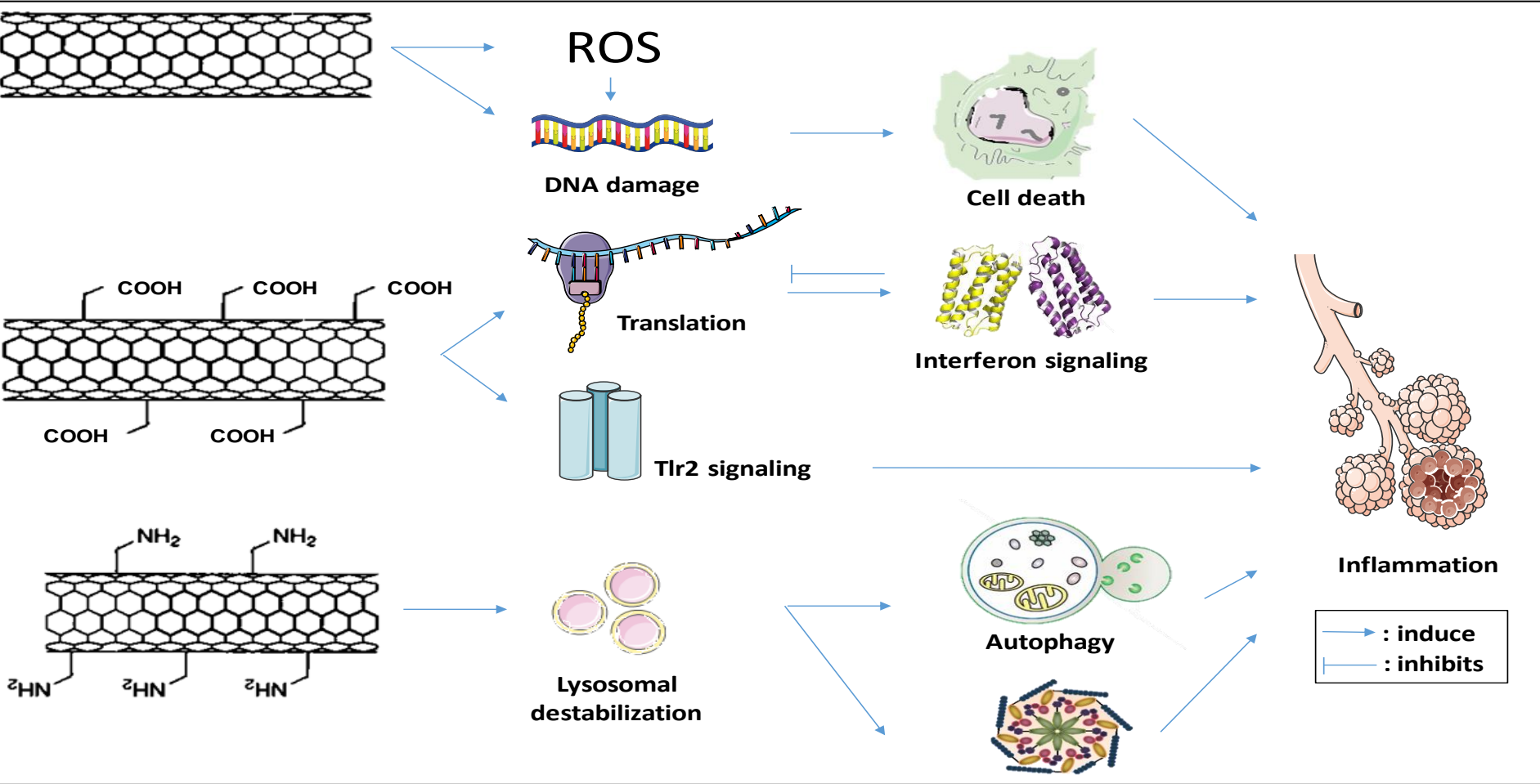
NR8383

Time: 24 h

Doses: $\frac{1}{4}$ IC₅₀

* p<0,10 ** p<0,05 *** p<0,01

Functionalisation effects on the regulated response to NMs, in NR8383



SW vs MW effects on the regulated answer to NMs, in NR8383

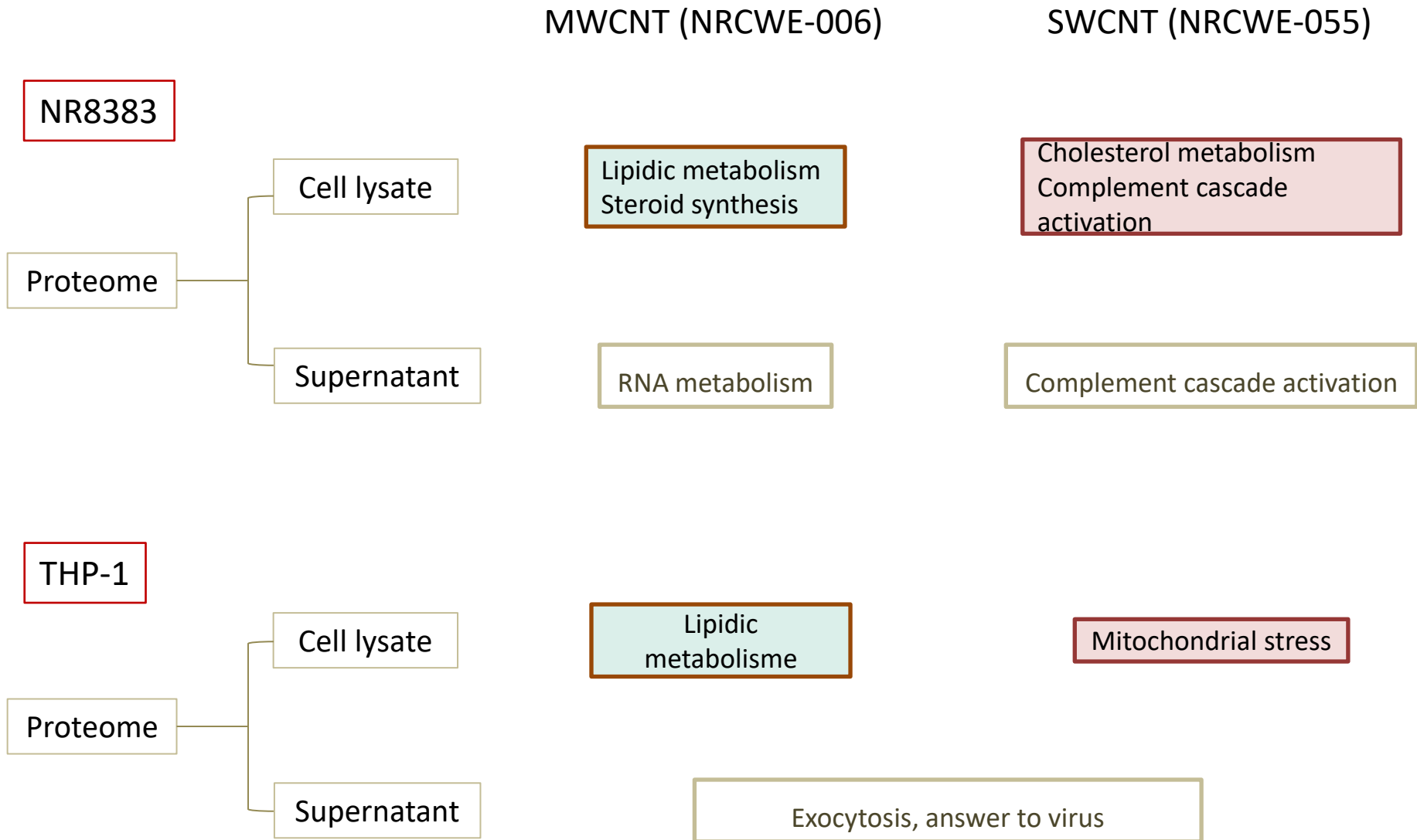
Time = 4h
 Doses: SWCNT = 11 cm²/cm²
 MWCNT = 1 cm²/cm²

- 3 Enrichment score 6

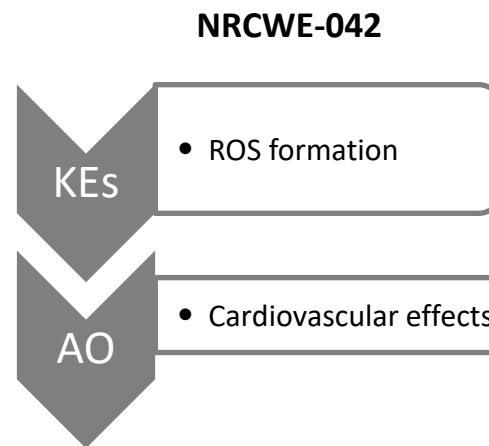
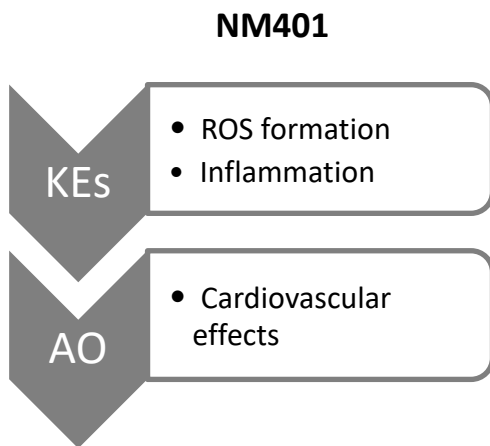
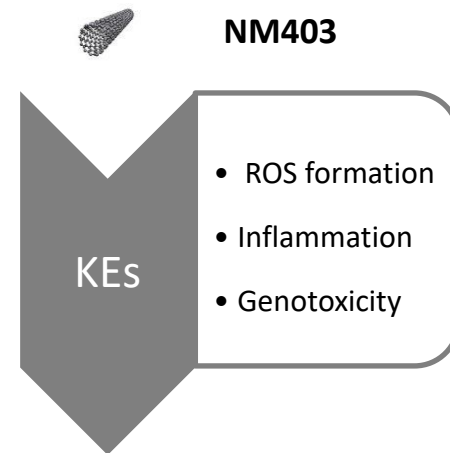
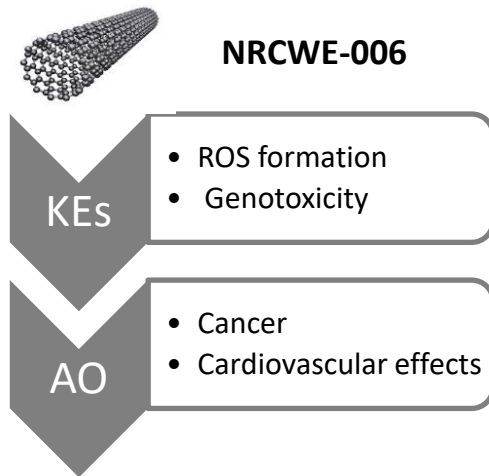
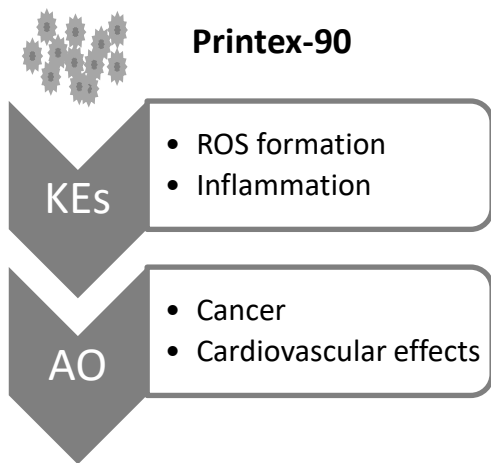


Cells	Canonical pathways			
	SWCNT (NRCWE-055)		MWCNT (NRCWE-006)	
THP-1 Human	• Mitochondrial Dysfunction		• Sirtuin Signaling Pathway	
	• Oxidative Phosphorylation		• mTOR Signaling	
	• Sirtuin Signaling Pathway		• Regulation of eIF4 and p70S6K Signaling	
	• EIF2 Signaling		• EIF2 Signaling	
	• mTOR Signaling		• Sumoylation Pathway	
NR8383 Rat	• EIF2 Signaling		• EIF2 Signaling	
	• Mitochondrial Dysfunction		• Protein Ubiquitination Pathway	
	• Oxidative Phosphorylation		• Sirtuin Signaling Pathway	
	• Sirtuin Signaling Pathway		• Oxidative Phosphorylation	
	• mTOR Signaling		• Mitochondrial Dysfunction	

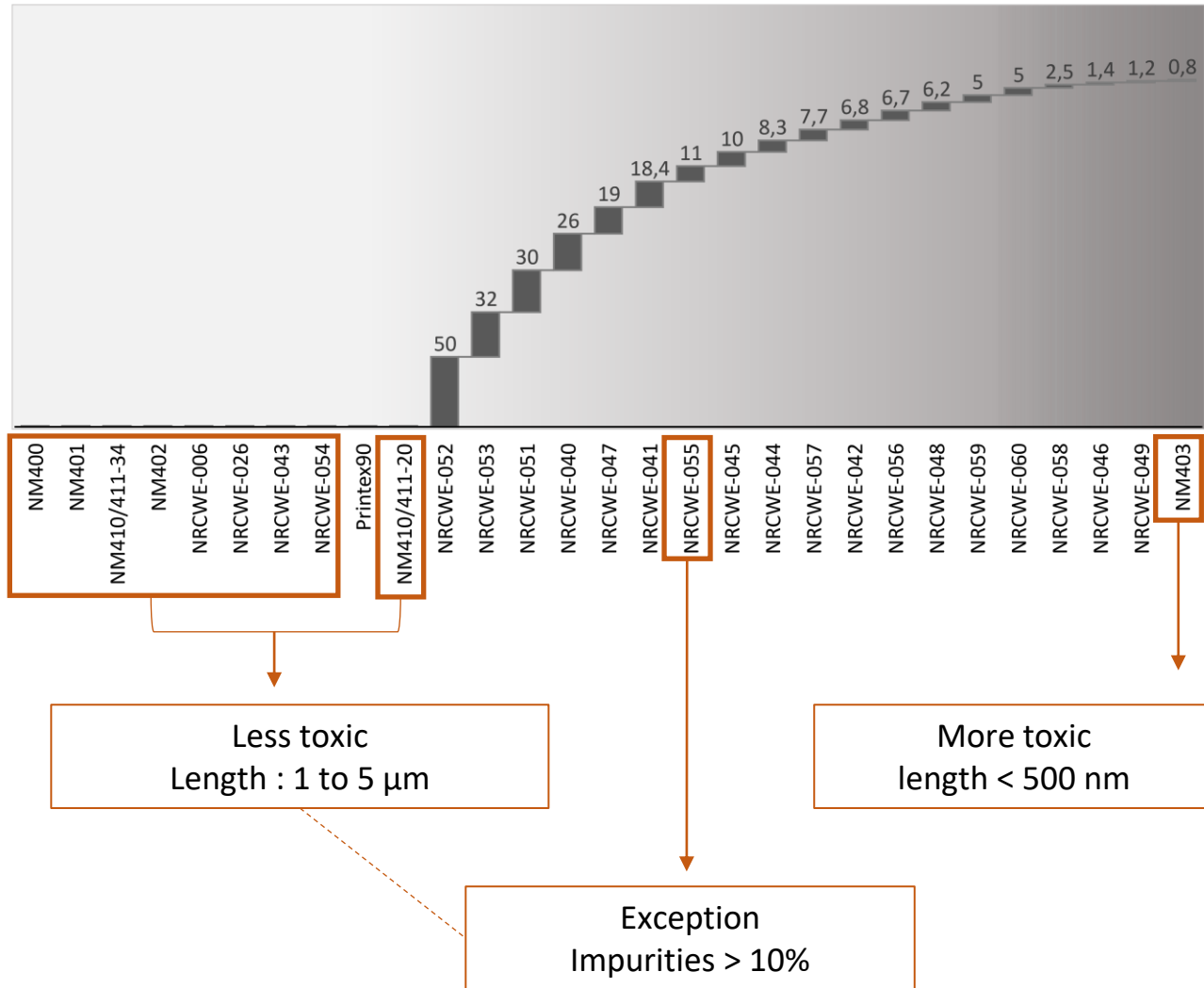
Proteomic primary analysis of SW vs MW effects in NR8383



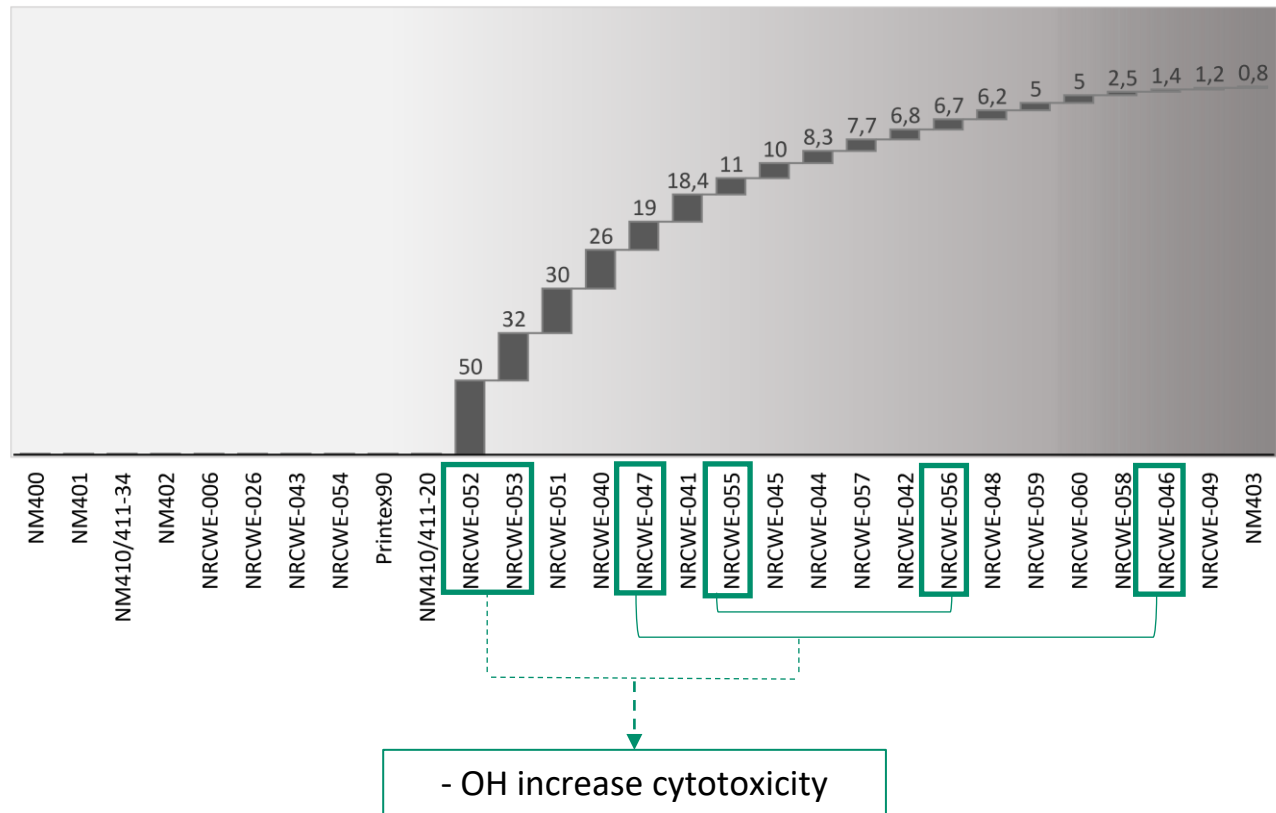
KEs and AOs for some MWCNT



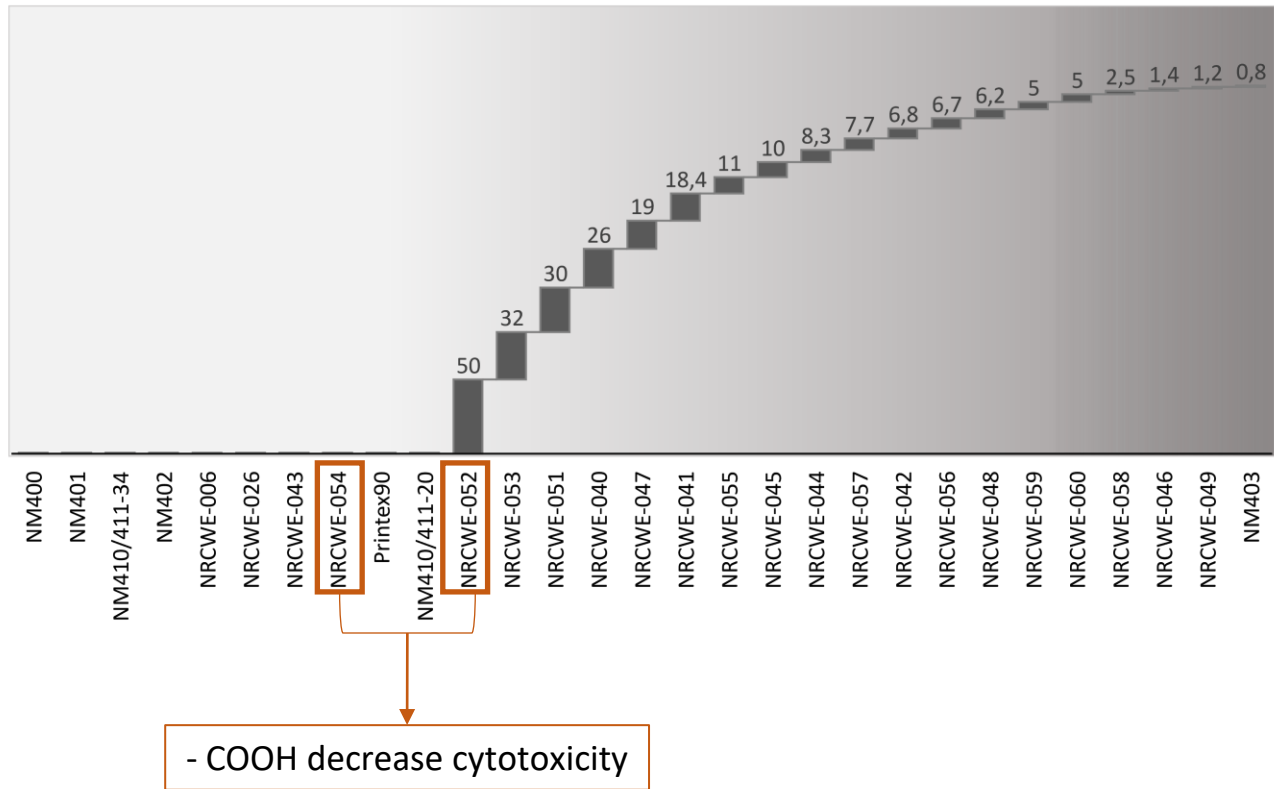
CNT length effect on IC₅₀, NR8383, WST-1



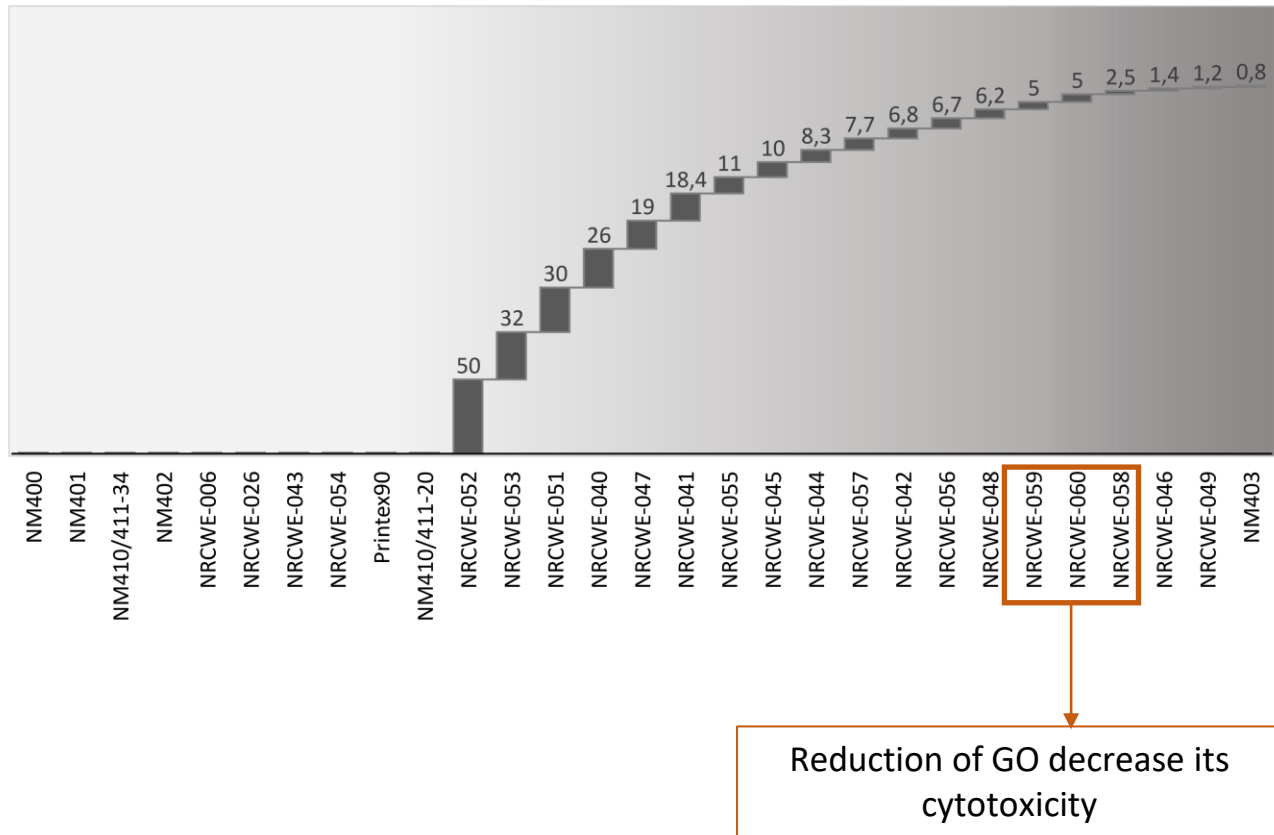
CNT hydroxylation effect on IC₅₀, NR8383, WST-1



CNT carboxylation effect on IC₅₀, NR8383, WST-1



Reduction of graphene oxide effect on IC₅₀, NR8383, WST-1



Reduction status of GO:

NRCWE-058 < NRCWE-060 < NRCWE-059

Thank you for your attention!