

Input of proteomics in nanoparticles toxicology: the example of macrophages responses to mineral nanoparticles

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The challenge of transposing lab. toxicology results into real life

Gold standard: lab animals (healthy life)

Combinatorial explosion of interfering factors (lifestyle, prof. etc...)

Necessity of obtaining molecular mechanisms

Role of high content in vitro approaches

The different toxicities of nanoparticles

(besides the shape/size/agglomeration issues)

Intrinsic toxicity

- examples: asbestos, crystalline silica
 - targeted approaches
 - high throughput screening
-

Helper toxicity

- adsorption As on Fe oxides (Auffan et al. Langmuir 2008)
 - adsorption Cd on amorphous silica (Guo et al. J. Hazard. Mat. 2013)
 - case of diesel exhaust particles
-

Cross-toxicities

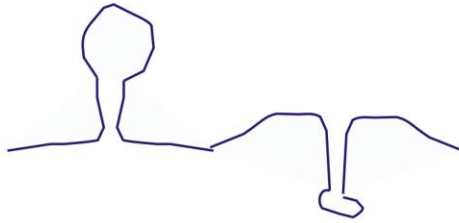
- synergistic toxicities without direct interactions
- **response mechanisms** => vulnerability points => sorting cross toxicities

The price for life: complexity



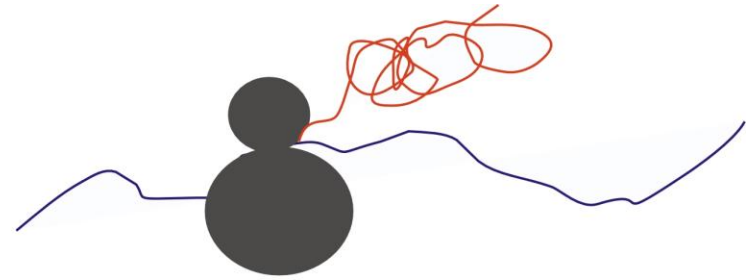
DNA

1500



mRNA

2000

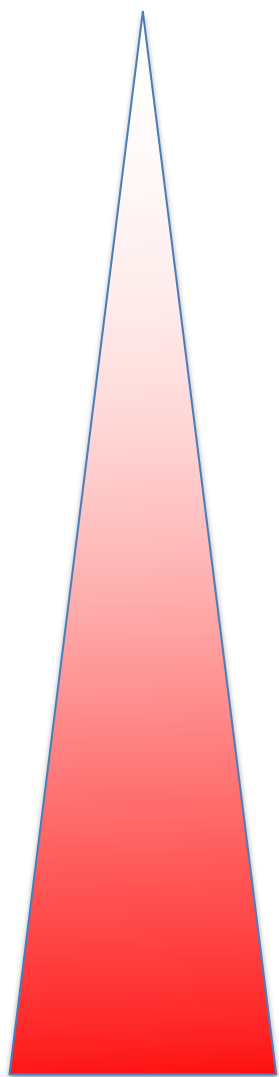


Translation

Protein

2500





Genomics

The world of possibles

Transcriptomics

What is going to happen (maybe)

Proteomics

What is happening now

Metabolomics

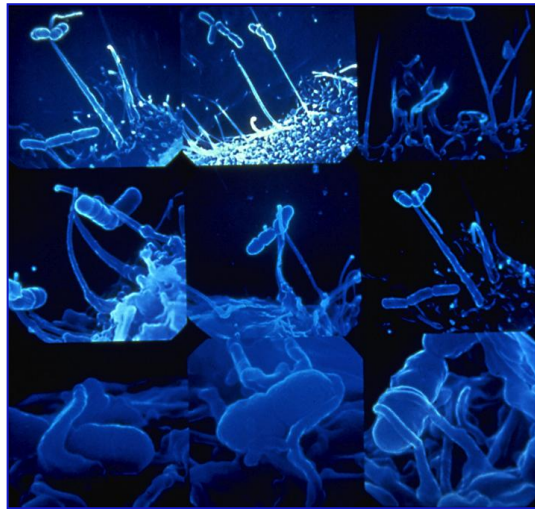
What has happened

Chemical diversity
Dynamic range

Macrophages: first line sentinels, immunity effectors and final scavengers

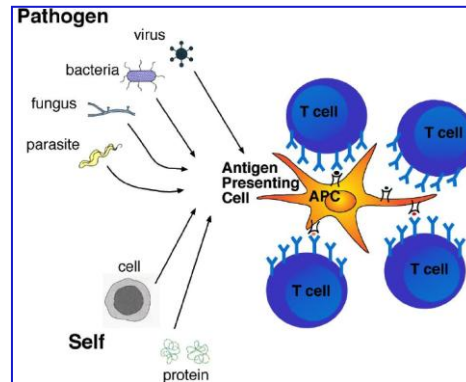
Macrophages

Phagocytosis, destruction of pathogens and abnormal cells

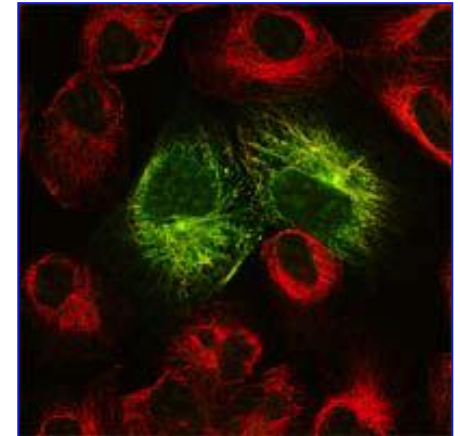


Scavenging of toxic particles
(e.g. altered LDL)

Antigen presentation

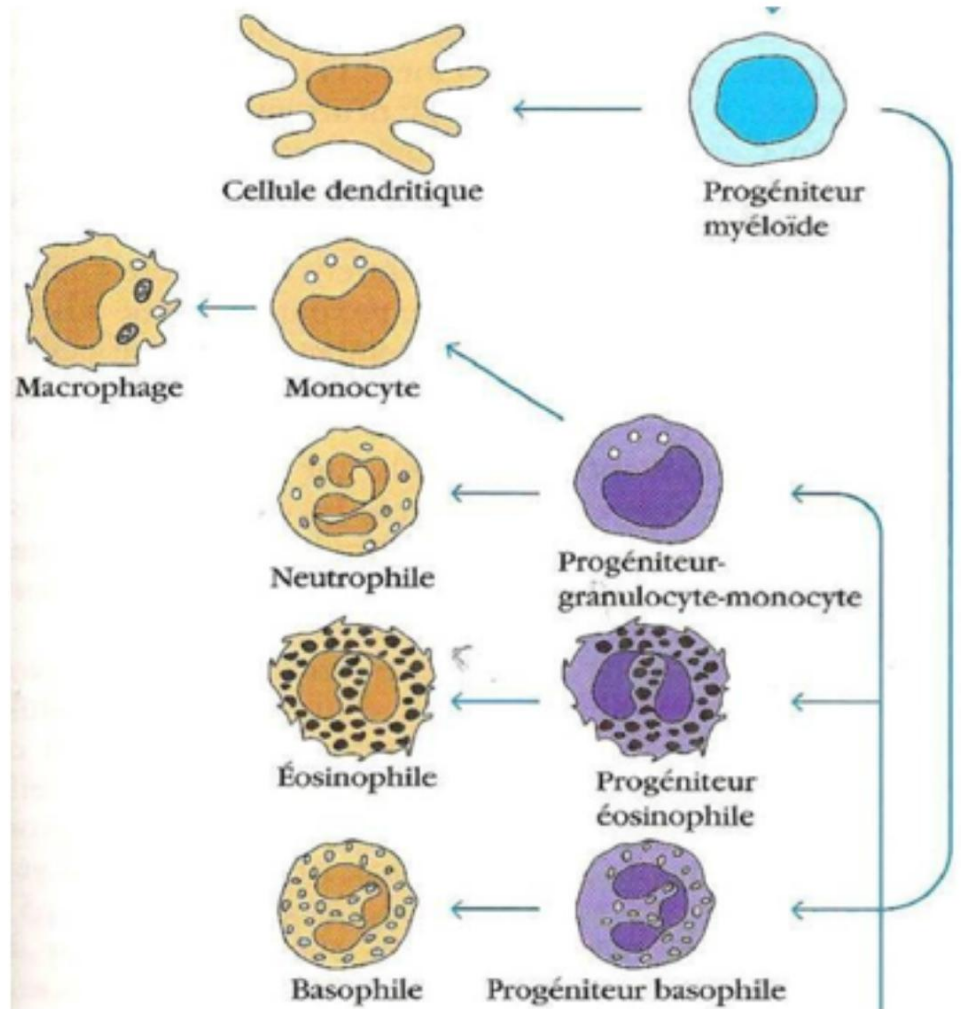
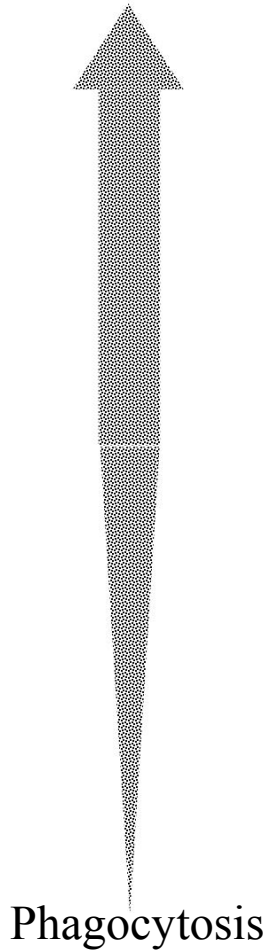
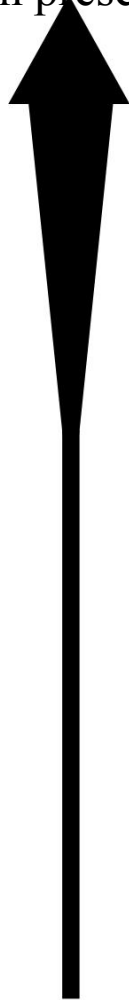
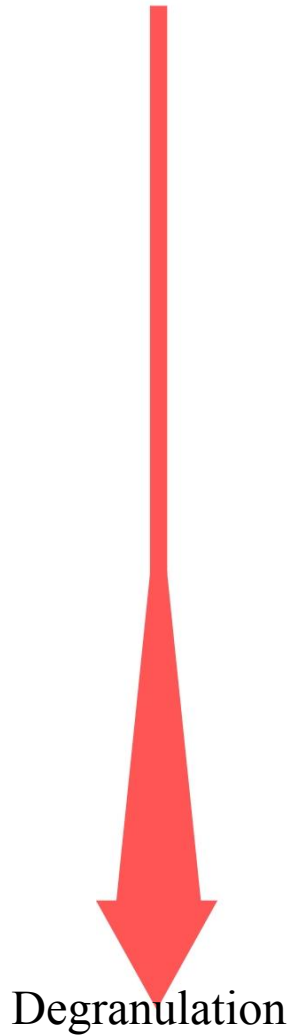


Cytokinic signalling



Inflammation
Tissue healing

Signalisation
Antigen presentation



The nanoparticles investigated: ZnO and CuO

ZnO (30,000 tons/year ww) used in sunscreens, biocidal, UV protection

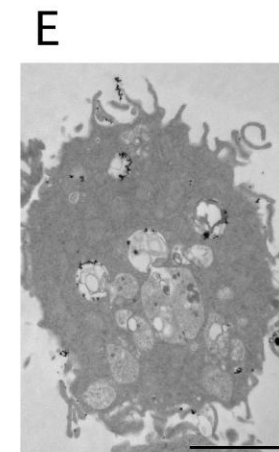
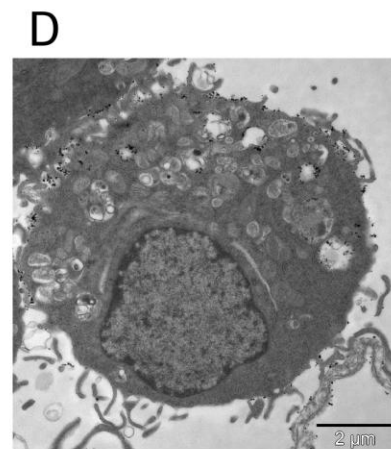
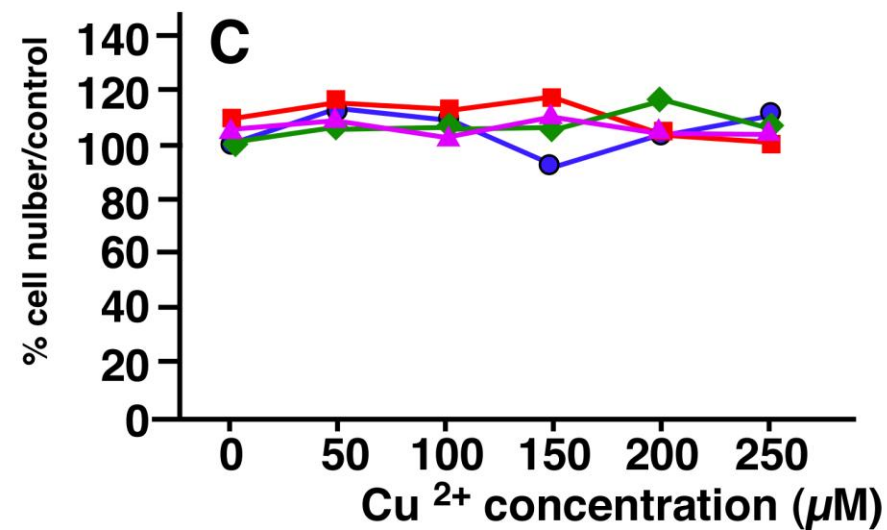
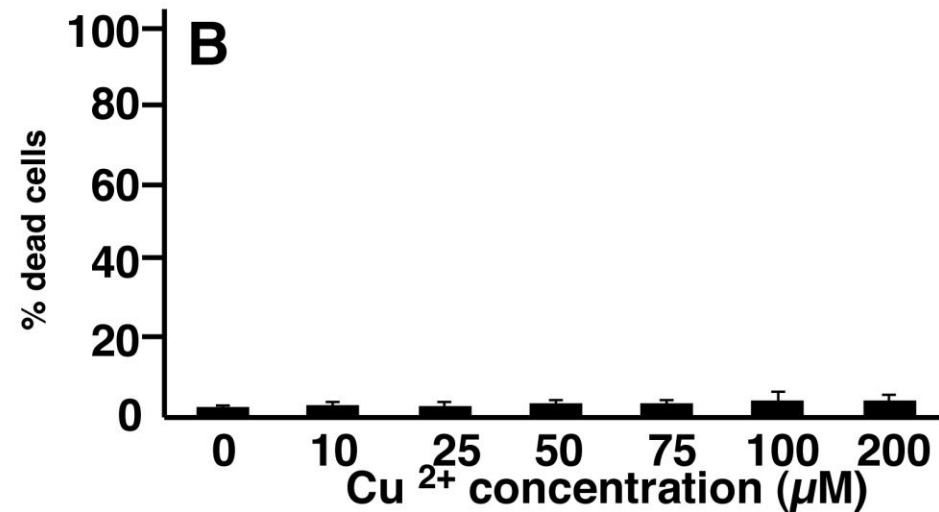
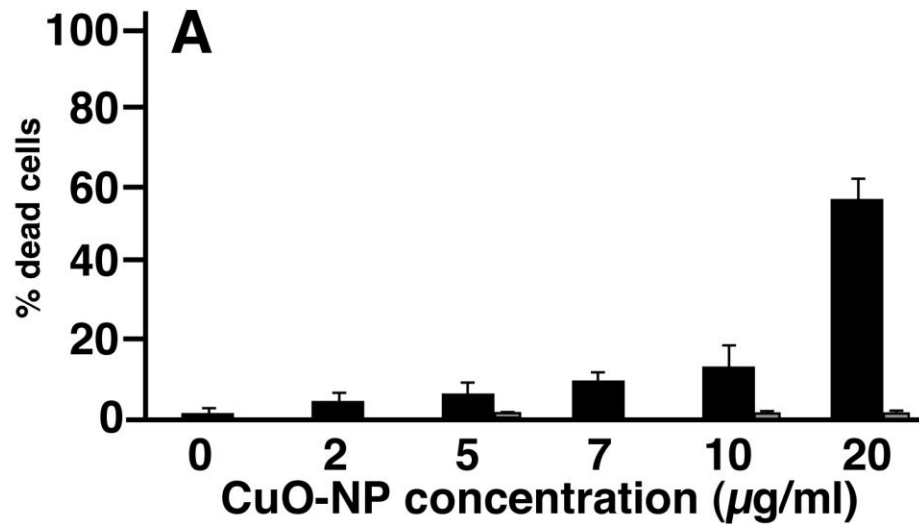
CuO used in water depollution, biocidal, conductive inks, wood treatment

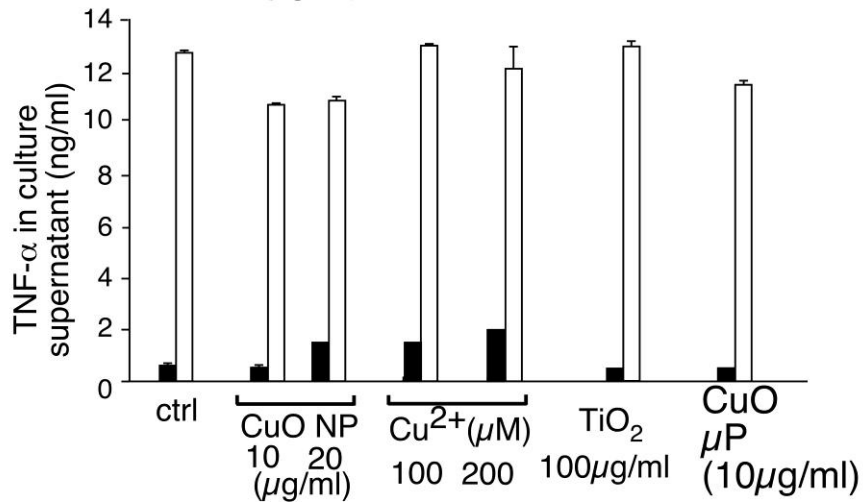
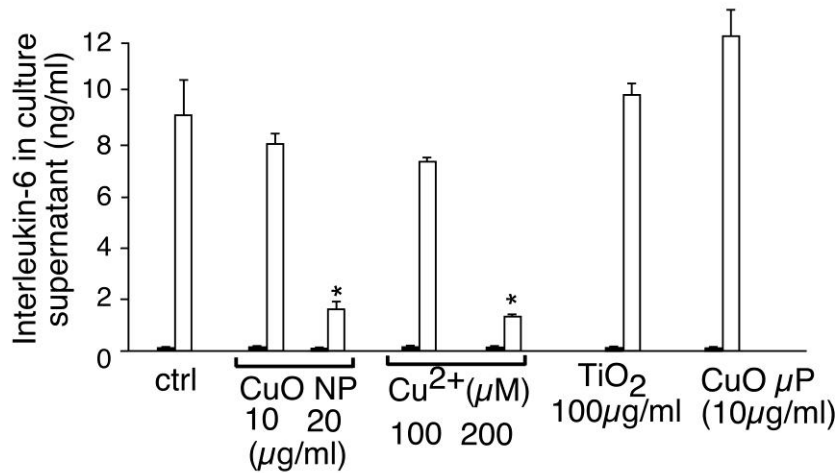
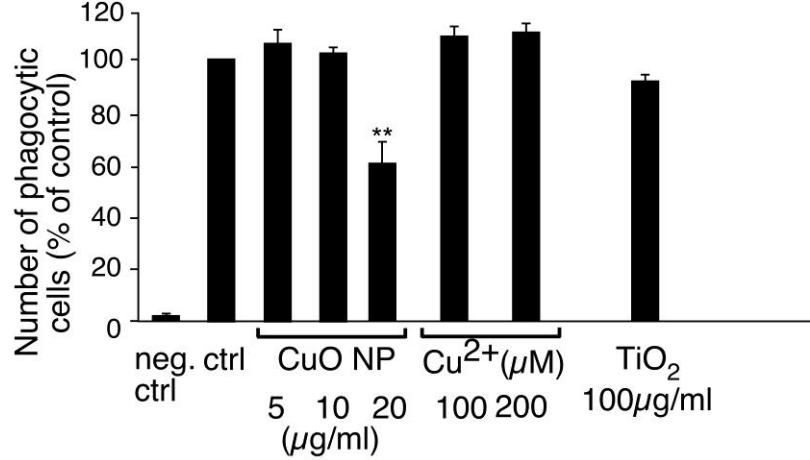
8 (VIII)	9 (VIII)	10 (VIII)	11 (IB)	12 (IIB)
Iron Fe ₂₆ 55.845 8.00284%	Cobalt Co ₂₇ 58.93200 7.3x10 ⁻⁵ %	Nickel Ni ₂₈ 58.6934 0.000161%	Copper Cu ₂₉ 63.546 1.70x10 ⁻⁵ %	Zinc Zn ₃₀ 65.39 4.11x10 ⁻⁵ %
Ruthenium Ru ₄₄ 101.07 6.1x10 ⁻⁸ %	Rhodium Rh ₄₅ 102.90550 1.12x10 ⁻⁶ %	Palladium Pd ₄₆ 106.42 4.5x10 ⁻⁶ %	Silver Ag ₄₇ 107.8682 1.58x10 ⁻⁶ %	Cadmium Cd ₄₈ 112.411 5.3x10 ⁻⁶ %
Osmium Os ₇₆ 198.23 2.26x10 ⁻⁶ %	Iridium Ir ₇₇ 192.217 2.16x10 ⁻⁶ %	Platinum Pt ₇₈ 195.078 4.4x10 ⁻⁶ %	Gold Au ₇₉ 196.96655 6.1x10 ⁻⁶ %	Mercury Hg ₈₀ 200.59 1.11x10 ⁻⁵ %

Very similar parameters: -primary particle size <50nm
-agglomerate size in culture medium ca. 200-250 nm
-similar toxicity (LD20 ca. 10 µg/ml)

Fairly different proteomic responses for ZnO and CuO

Effect of copper on cell survival

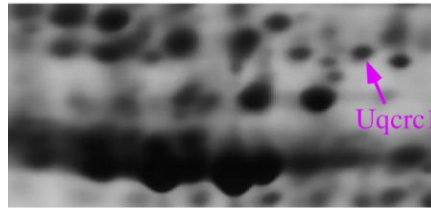




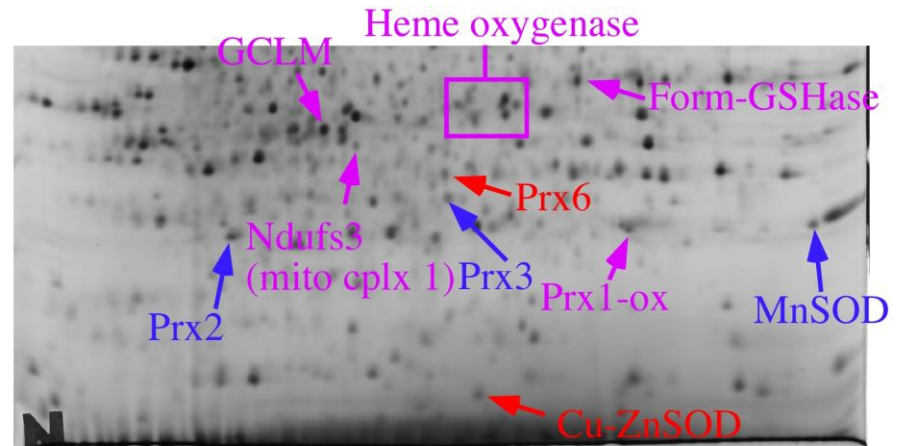
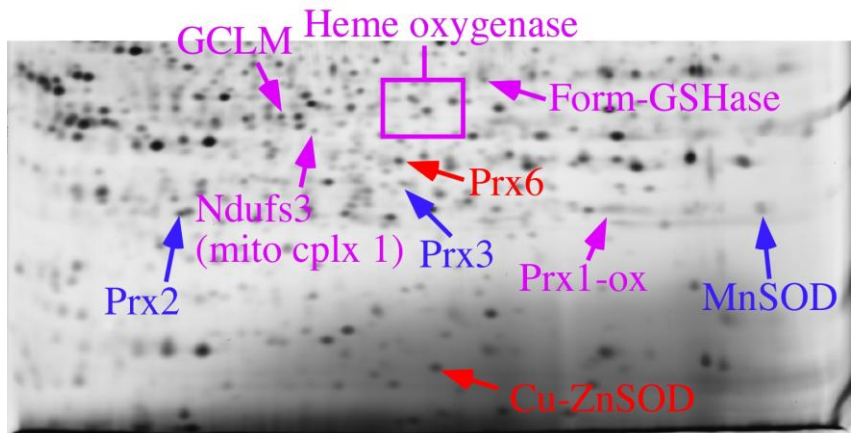
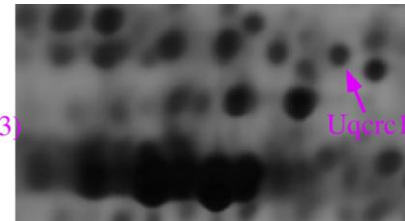
Functional effects of Cu on macrophages

Oxidative and mitochondrial stress response to CuO (1)

Control

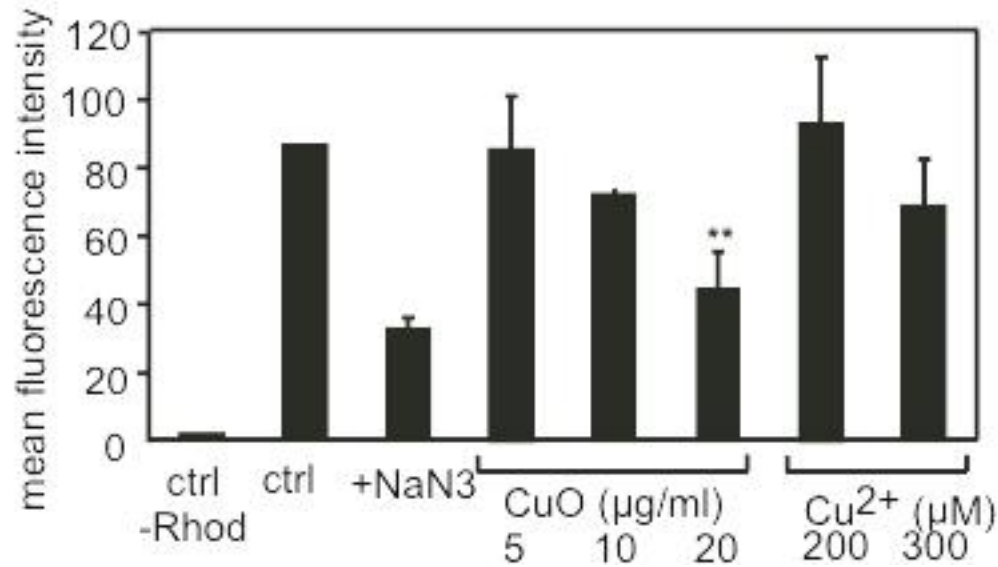


CuO-treated

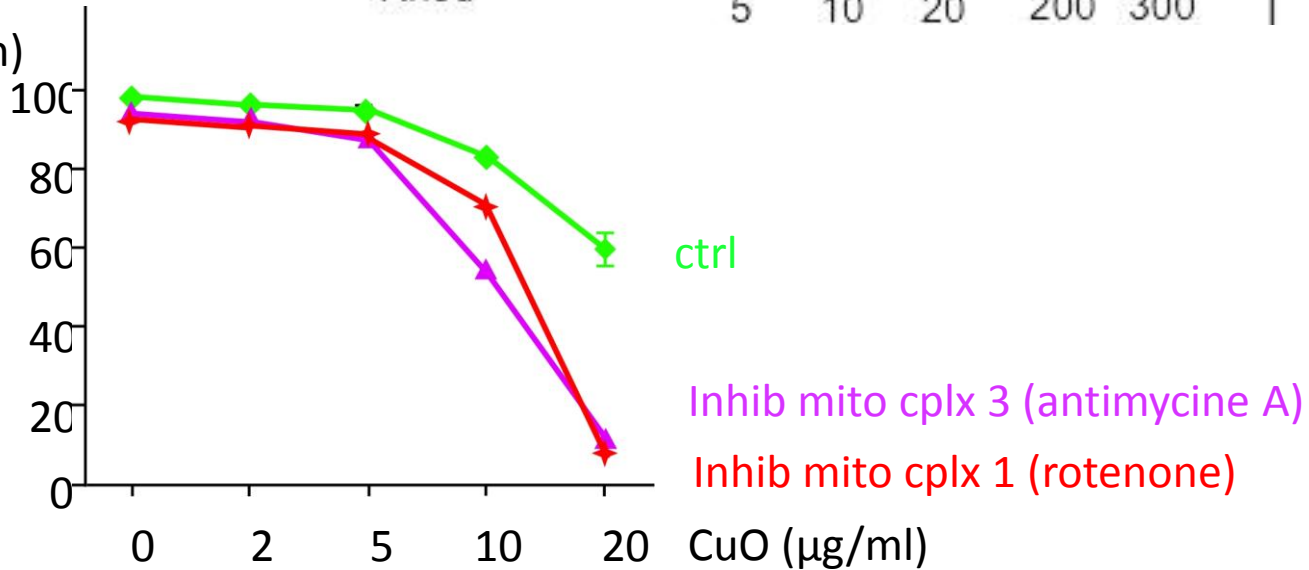


Oxidative and mitochondrial stress response to CuO (2)

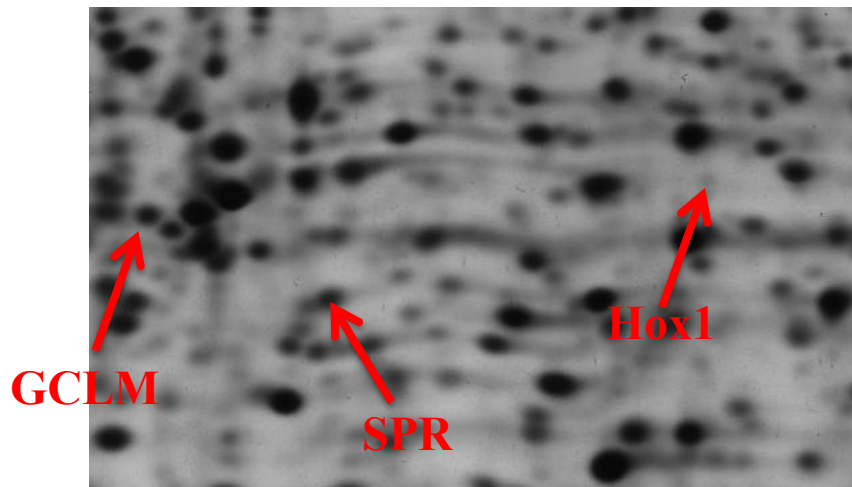
Induction of mitochondrial respiratory complexes



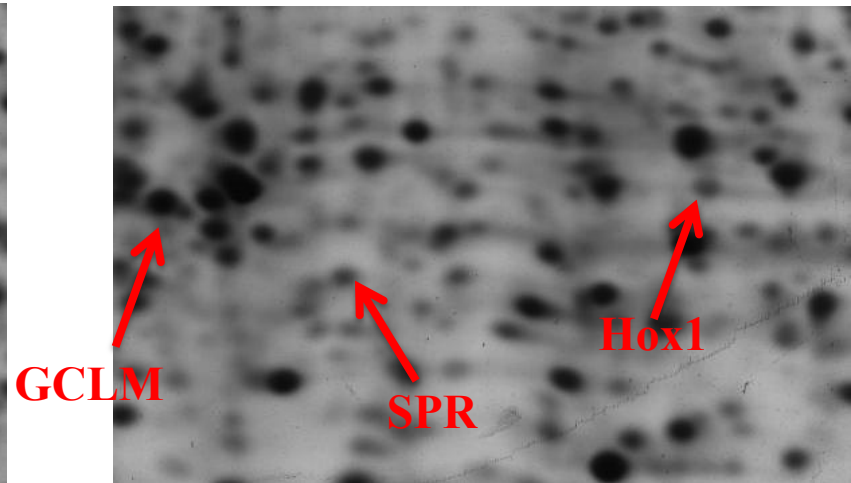
% viable cells
(dye exclusion)



Further validation of proteomic findings

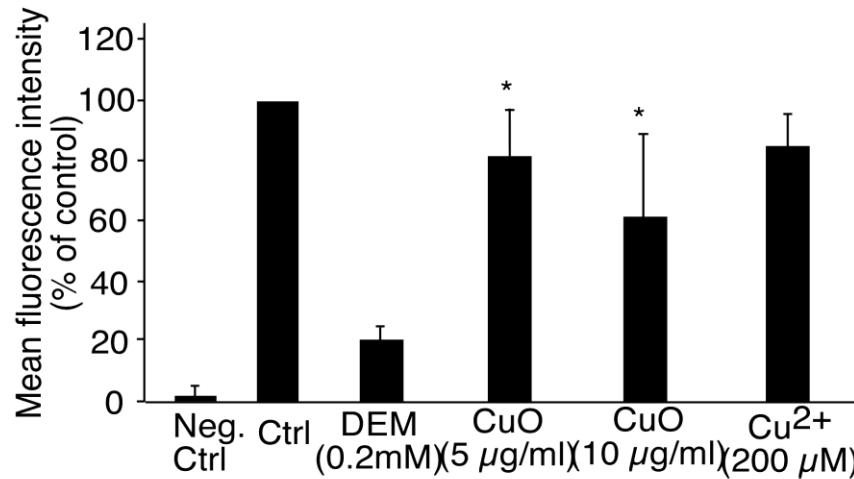


Control



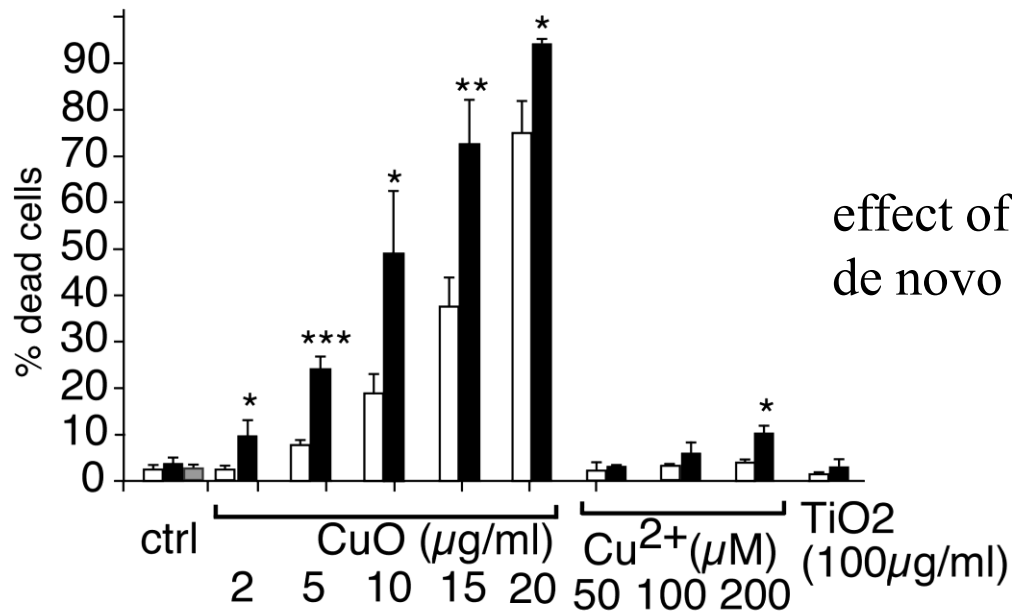
CuO-treated

Role of glutathione in survival to copper oxide nanoparticles (GCLM)



glutathione levels

A

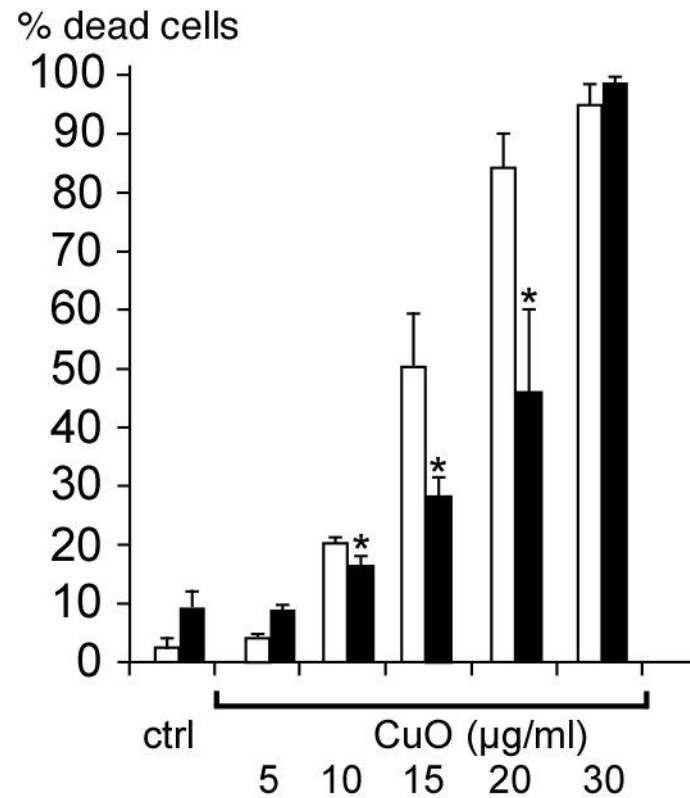


effect of glutathione
de novo synthesis inhibition

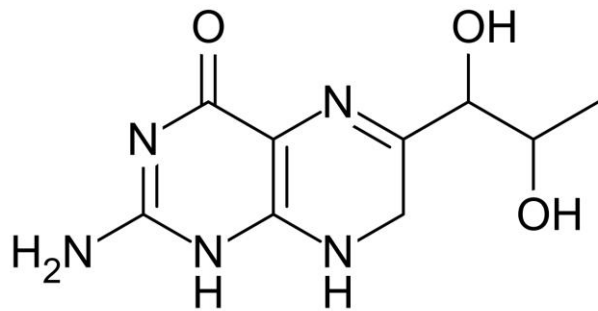
B

Role of heme oxygenase (HOX1) in copper resistance

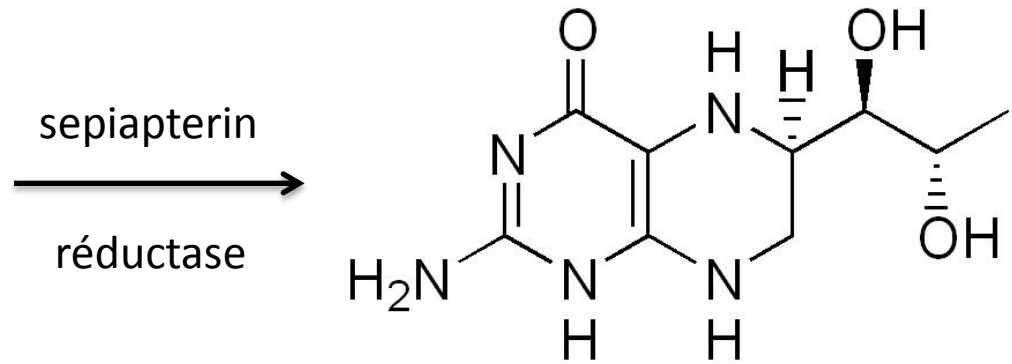
**Pre induction of Hox
with 1 μ M lovastatin
for 6hrs prior to
CuO challenge**



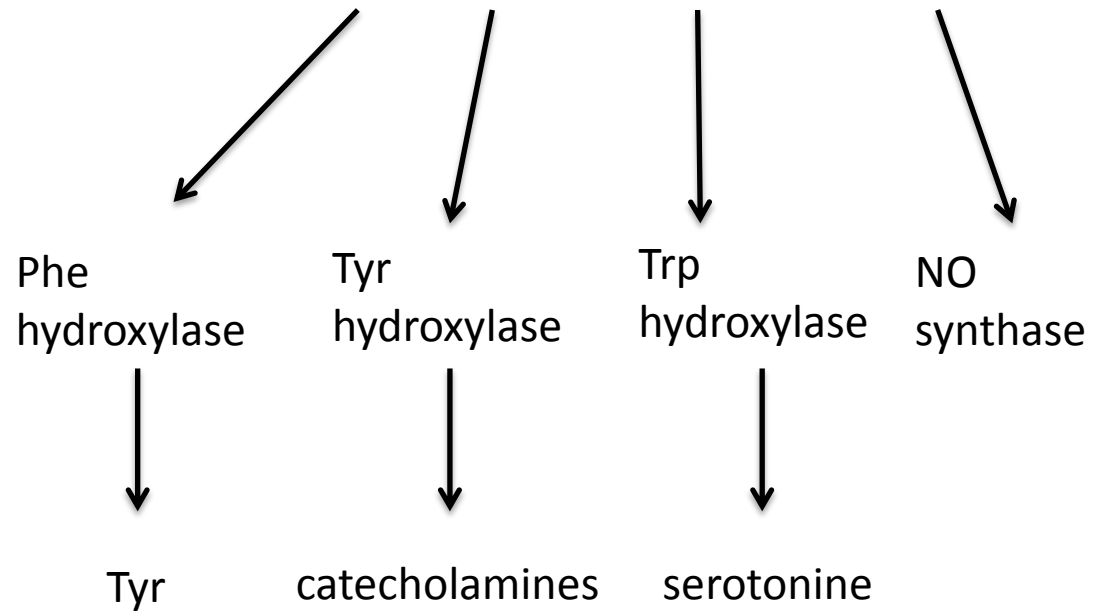
Role of sepiapterin reductase (SPR)



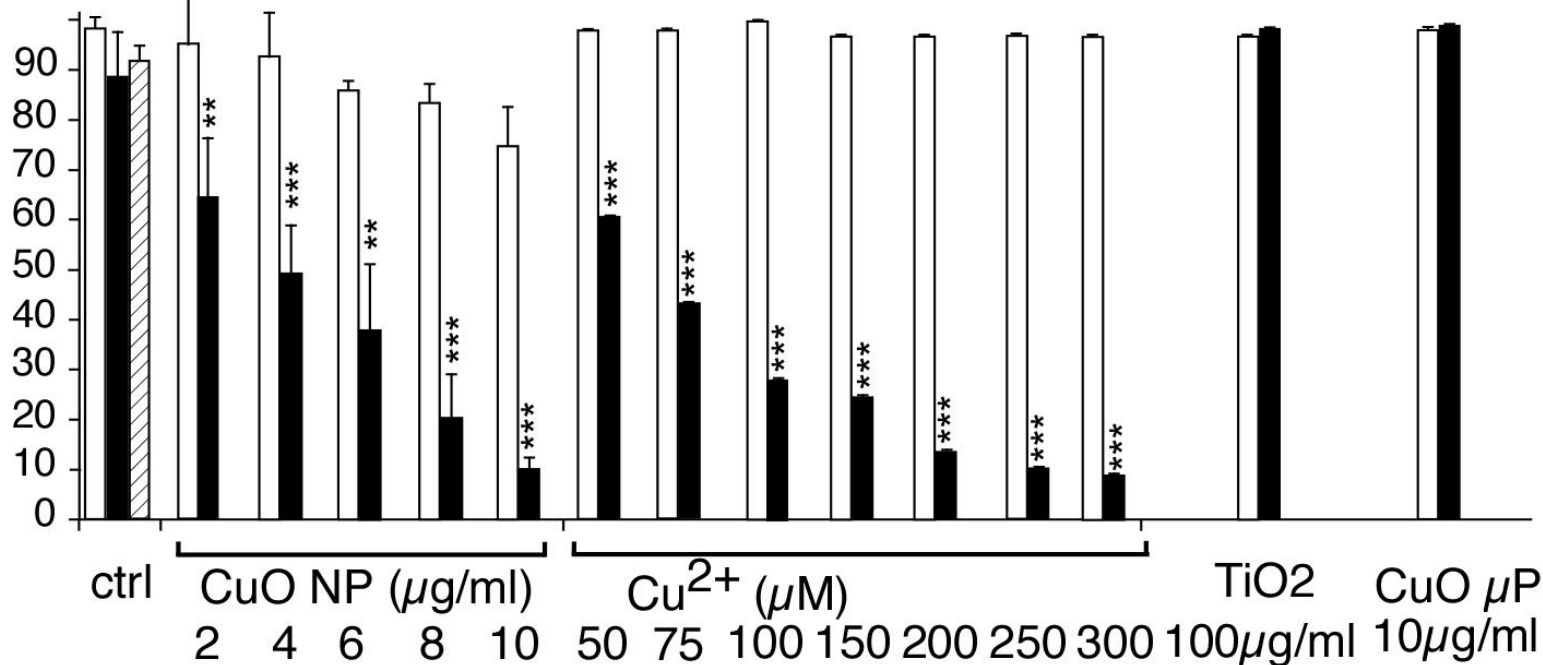
dihydrobiopterin



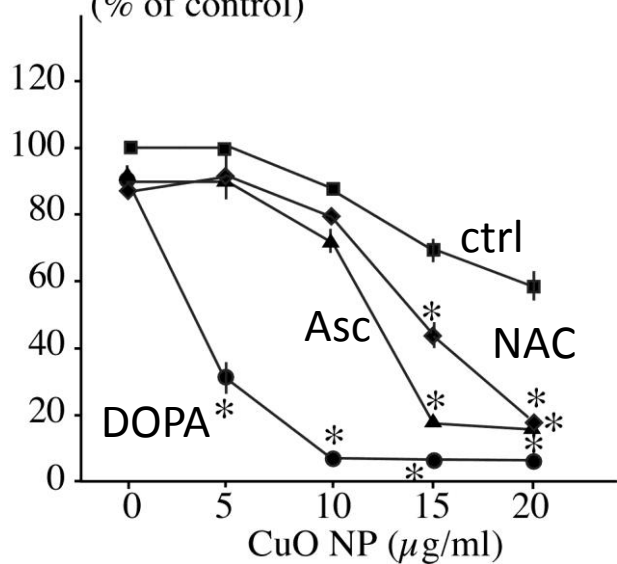
tetrahydrobiopterin



% viable cells



Cell survival
(% of control)



The second nanoparticle investigated: ZnO

ZnO (30,000 tons/year ww) used in sunscreens, biocidal, UV protection

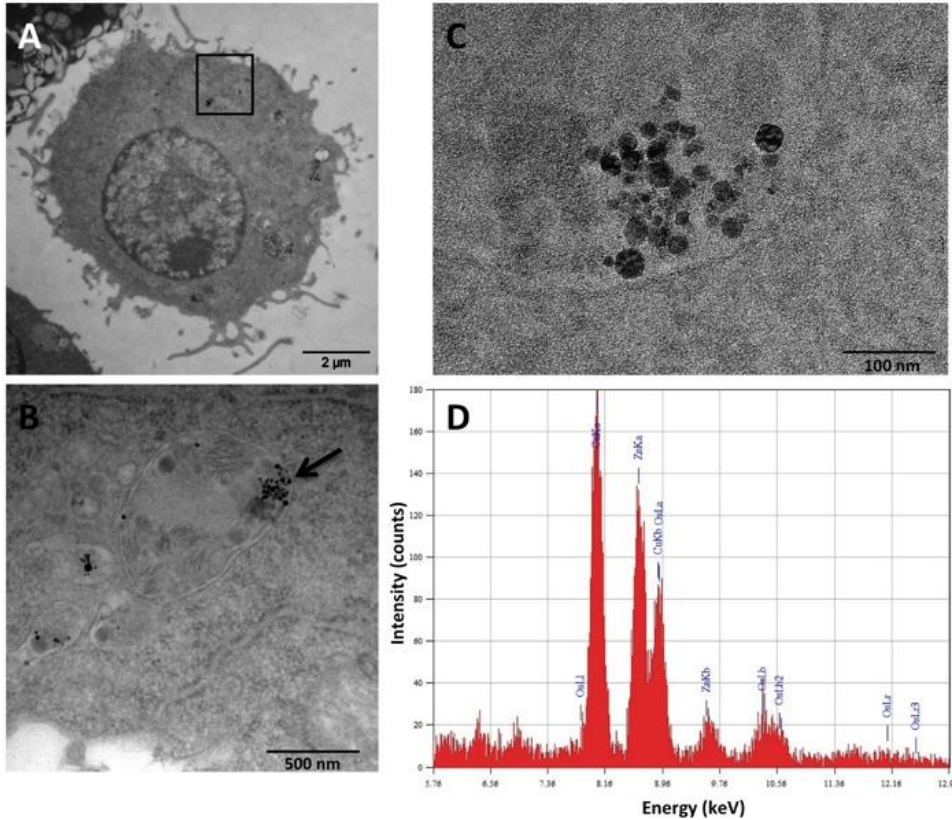
Parameters:

- primary particle size <50nm**
- agglomerate size in culture medium ca. 200-250 nm**
- moderate toxicity (LD20 ca. 10 µg/ml)**

ZnO: causative agent of the metal fume fever (at doses >50mg/m³ air)

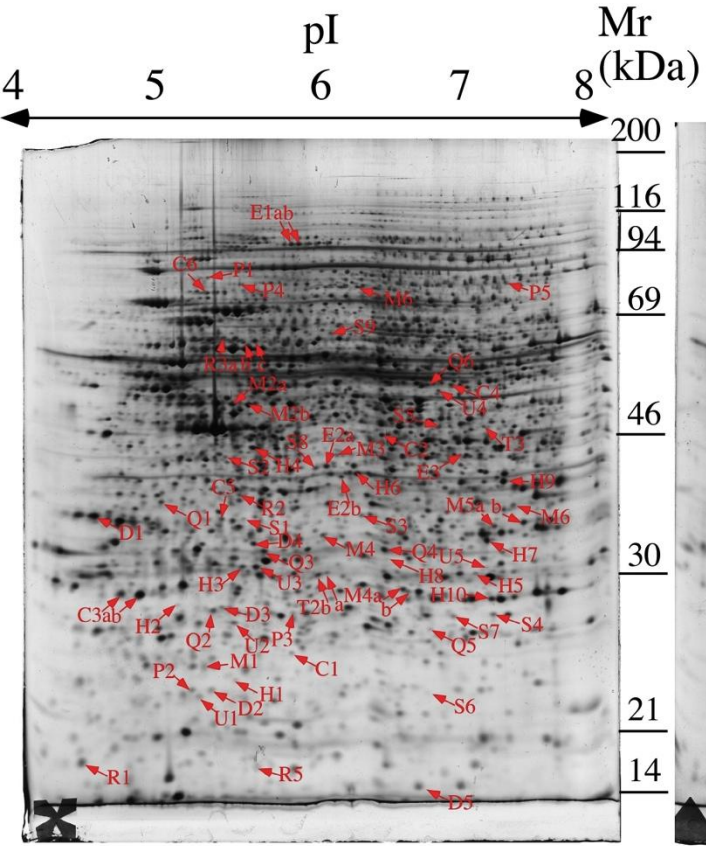
50mg/m³ air => 10 ppm in our culture system

Uptake of zinc oxide by macrophages

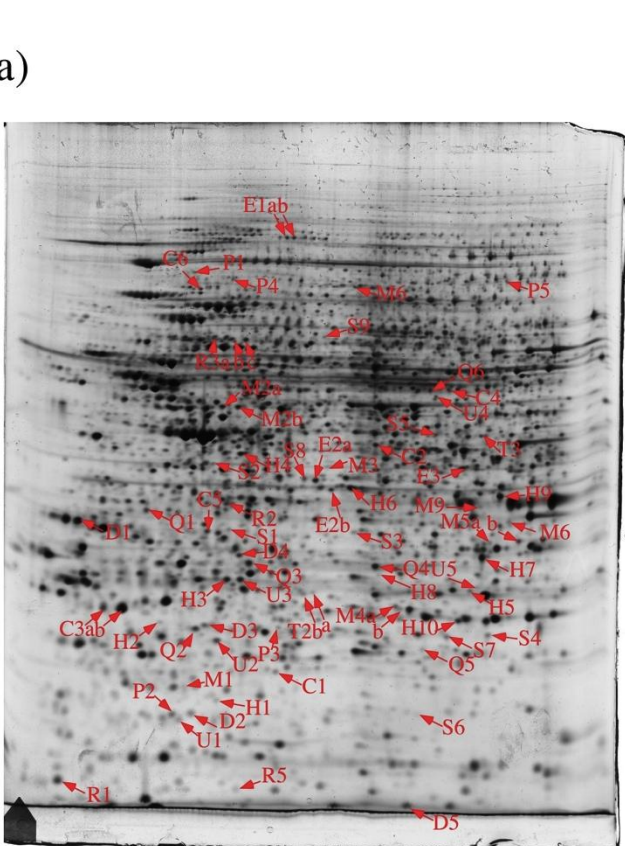


Input in cell culture	Concentration in cell extract	Concentration in cells
Cationic zinc oxide (90 μ M)	27 \pm 7 μ M	135 \pm 35 μ M
Zinc acetate (100 μ M)	37.5 \pm 2.5 μ M	187.5 \pm 12.5 μ M
Zinc oxide (100 μ M)	30 \pm 1 μ M	150 \pm 5 μ M

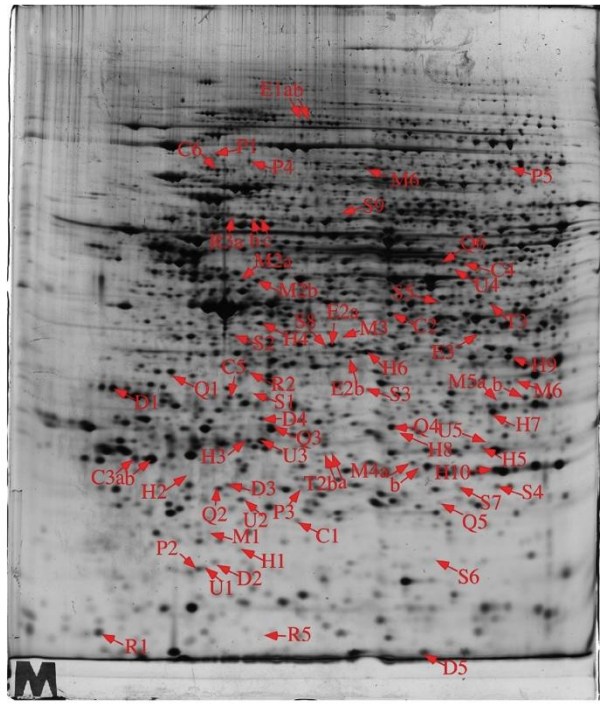
Proteomic analysis of J774 cells in response to ZnO nanoparticles



Ctrl



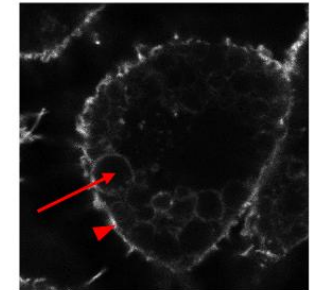
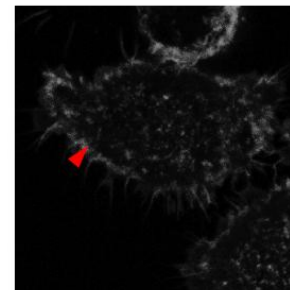
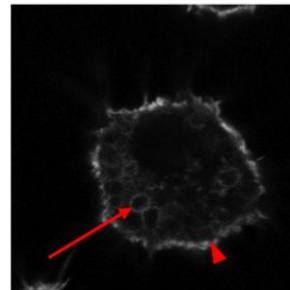
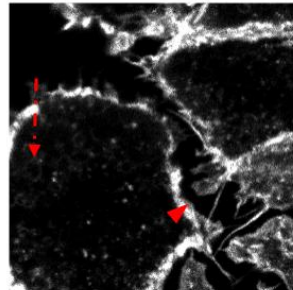
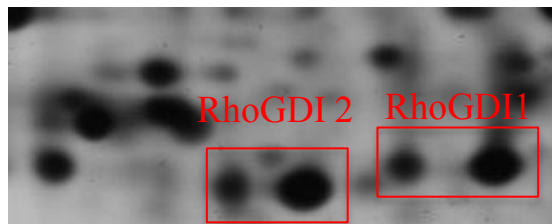
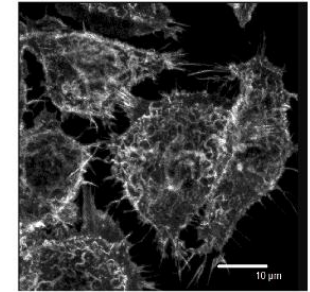
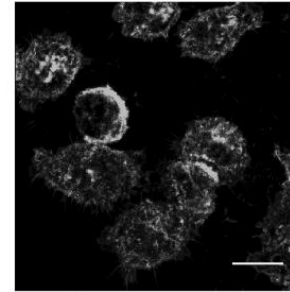
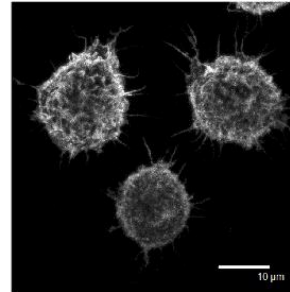
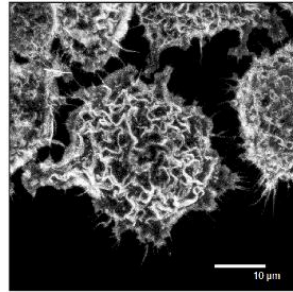
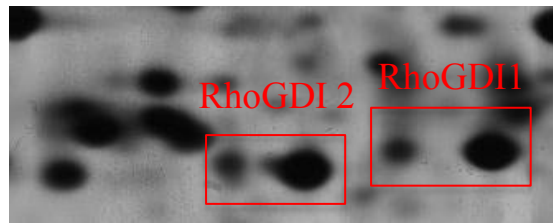
ZnO



ZrO₂

Changes in the actin cytoskeleton

Control cells



ZnO treated

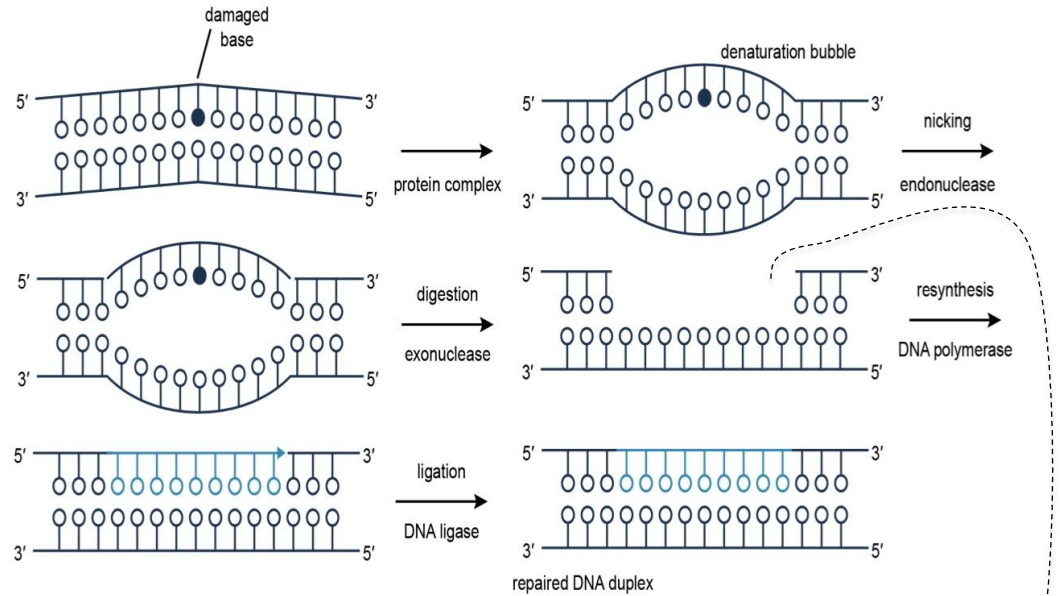
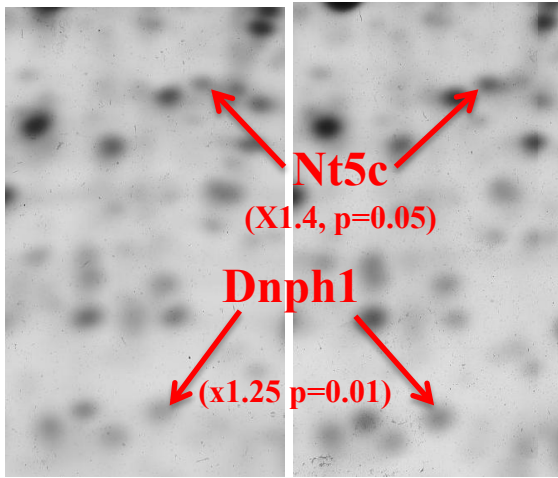
ctl

Zn⁺⁺

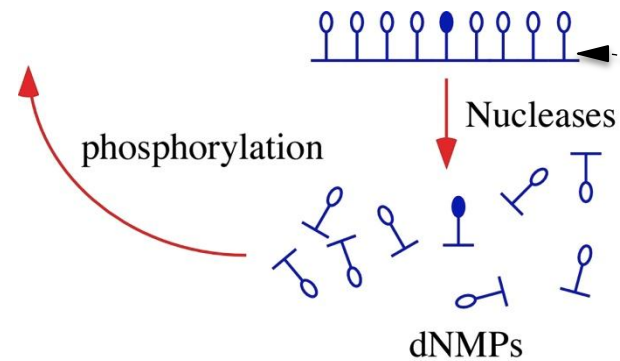
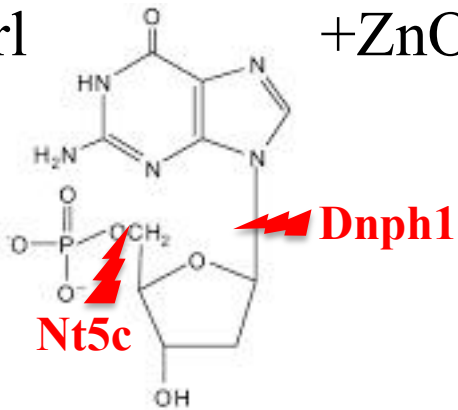
ZnO

ZrO₂

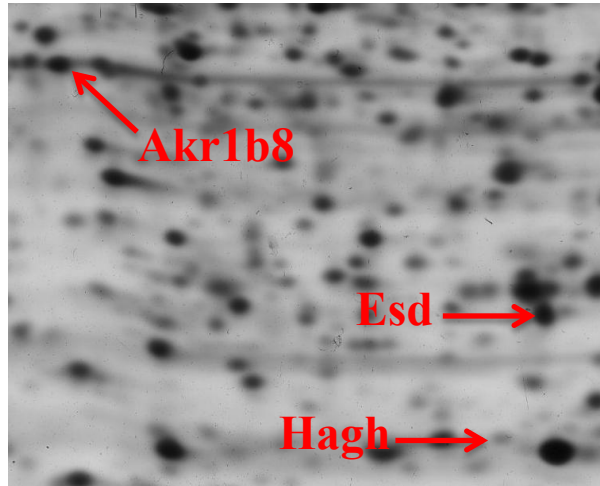
Zinc genotoxicity : the genotoxicity of a non-Fenton metal



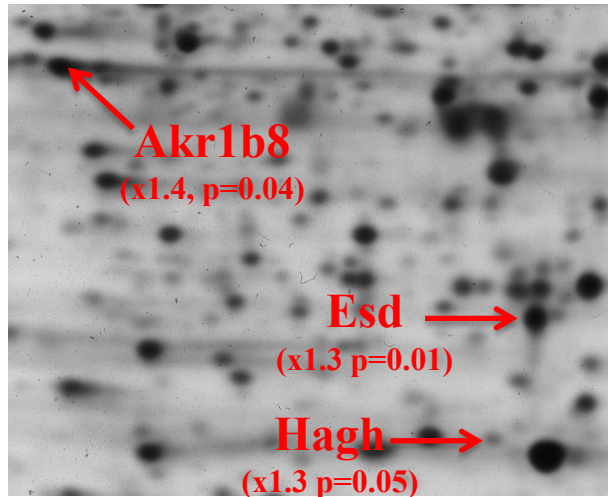
ctrl +ZnO



ctrl



+ZnO

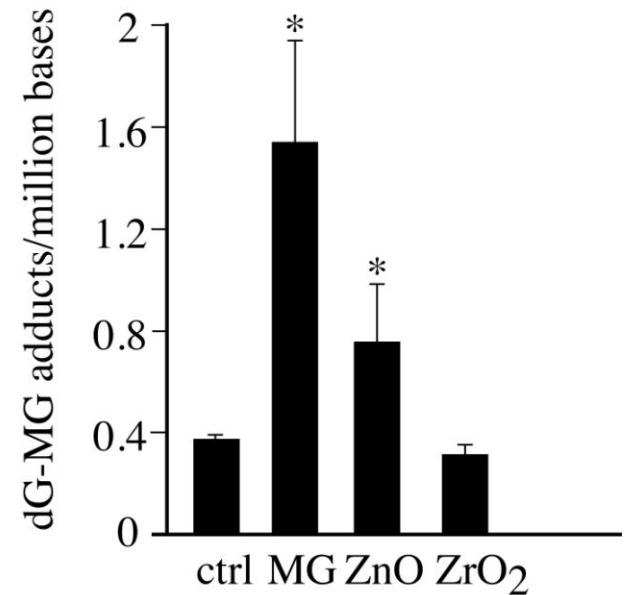
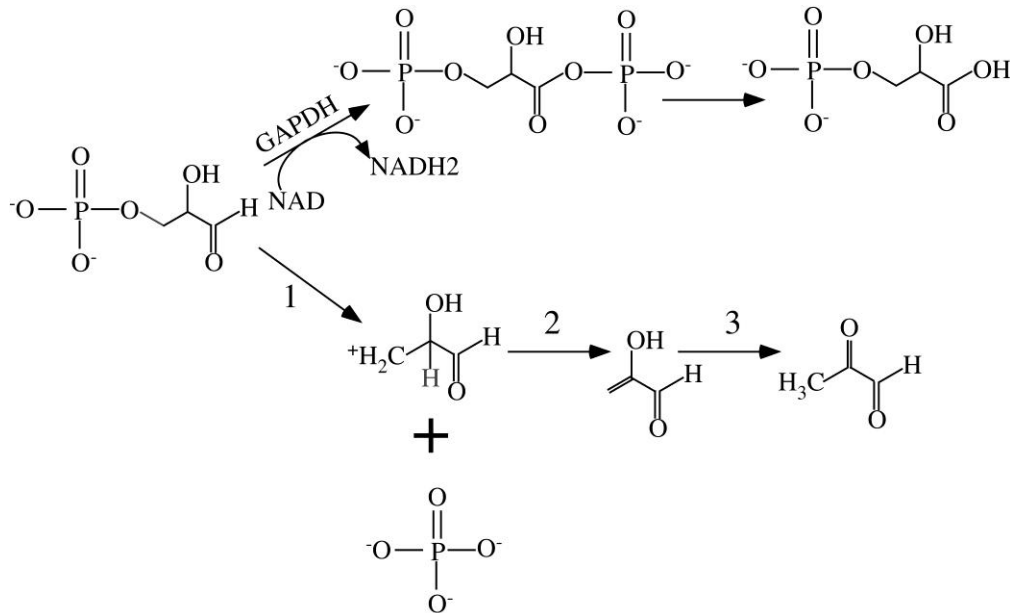


aldehydes detoxification

condition	GAPDH	condition	GAPDH
ctrl	16±2	0μM Zn	16.5±1
ZrO2	13.5±2.5	100μM Zn	11.5±0.6
ZnO	7.5±2.5	150μM Zn	7.2±0.8
		200μM Zn	4.9±1.5

The activities are expressed in units/mg protein, the unit being defined as 1μmole of substrate converted per minute

The methylglyoxal pathway in zinc toxicity



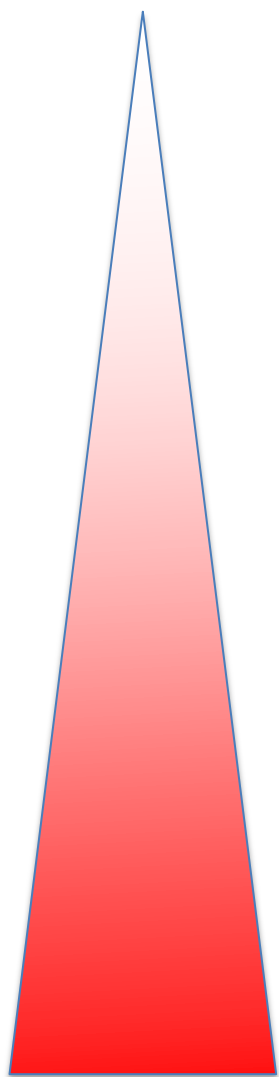
=> an indirect and composite genotoxic mechanism (DNA Pol ι and κ)

=> toward a proteomics-driven study of nanoparticles cross-toxic effects

Conclusion: proteomics can do the job

- Proteomics underscores biologically relevant responses at non toxic doses
(e.g. mitochondria, GSH biosynthesis, Hox, DOPA, methylglyoxal)
- Proteomics can sort different responses even if tox. parameters are similar
- Proteomics is able to underscore possible cross-toxicities
(e.g. Cu + rotenone, Cu+DOPA)

Full exploitation of proteomics data require functional validation



Genomics

The world of possibles

Transcriptomics

What is going to happen (maybe)

Proteomics

What is happening now

Metabolomics

What has happened

Chemical diversity
Dynamic range

Less is more, Less is bore, or mess in more ?
(Mies van der Rohe) (Robert Venturi)