

ESI 5: PCA scores plots of omics data

Principal components analysis (PCA) scores plots of transcriptomic, lipidomic and metabolomic data derived from exposure of A549 cells to silver, zinc and CeO₂ nanoparticles, microparticles and/or ions.

For silver, control samples are shown in black, Ag NP-NM300K treated in red, Ag MP in cyan, Ag⁺ in blue.

For zinc, control samples are shown in black, ZnO NP-NM110 in red, ZnO NP-NM111 in cyan, Zn²⁺ in blue and ZnO MP in pink.

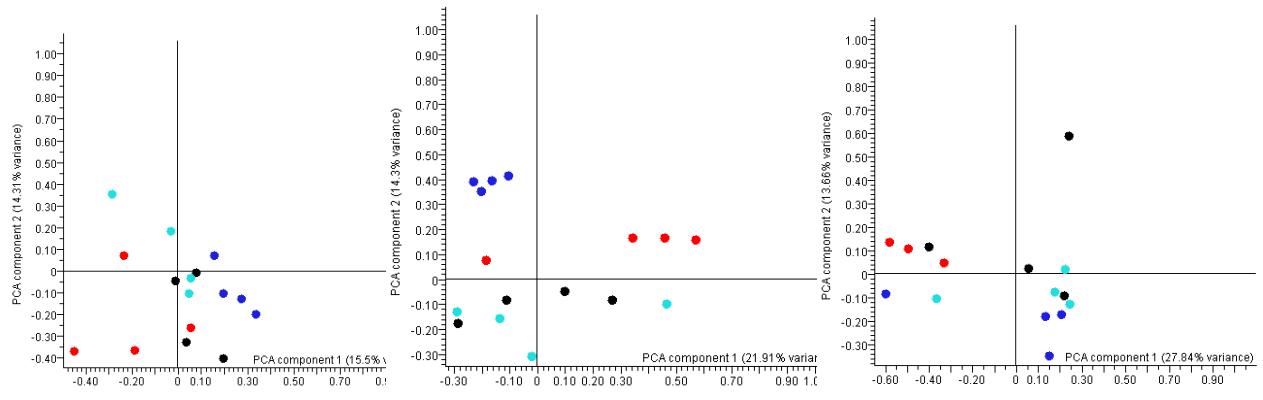
For CeO₂, control samples are shown in black, CeO₂ NP-A treated in red, CeO₂ NP-C in cyan, CeO₂ NP-E in blue and CeO₂ NP-NM212 in pink.

Supplementary material of:

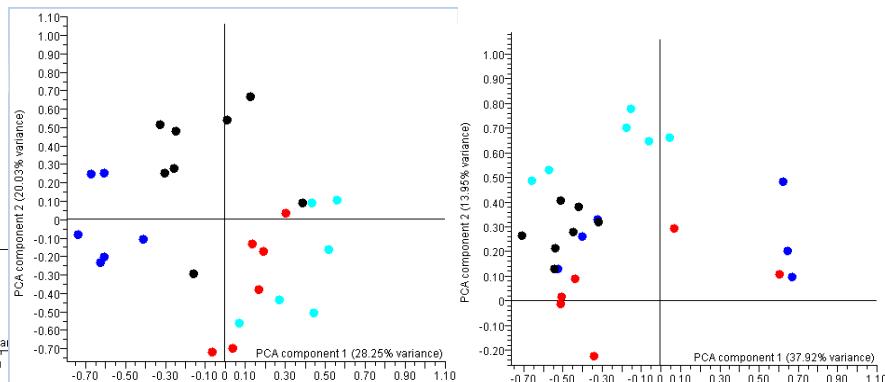
Susan Dekkers^{*1}, Tim D. Williams^{*2}, Jinkang Zhang², Jiarui (Albert) Zhou³, Rob J. Vandebriel¹, Liset J.J. De La Fonteyne¹, Eric R. Gremmer¹, Shan He³, Emily J. Guggenheim⁴, Iseult Lynch⁴, Flemming R. Cassee^{1,5}, Wim H. De Jong¹, Mark R. Viant². Multi-omics approaches confirm metal ions mediate the main toxicological pathways of metal-bearing nanoparticles in lung epithelial A549 cells. Environmental Science Nano. Corresponding author: susan.dekkers@rivm.nl.

Transcriptomics

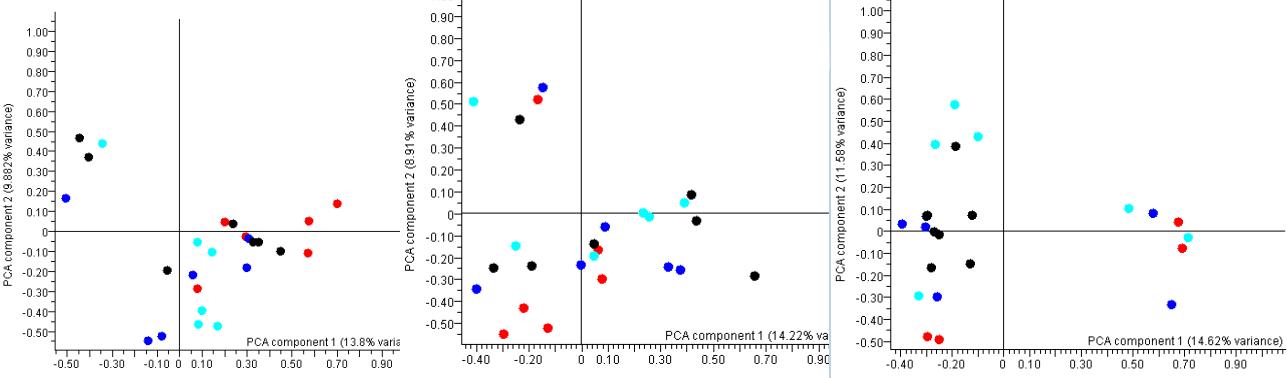
Silver 1 hour 6 hours 24 hours



Polar metabolomics

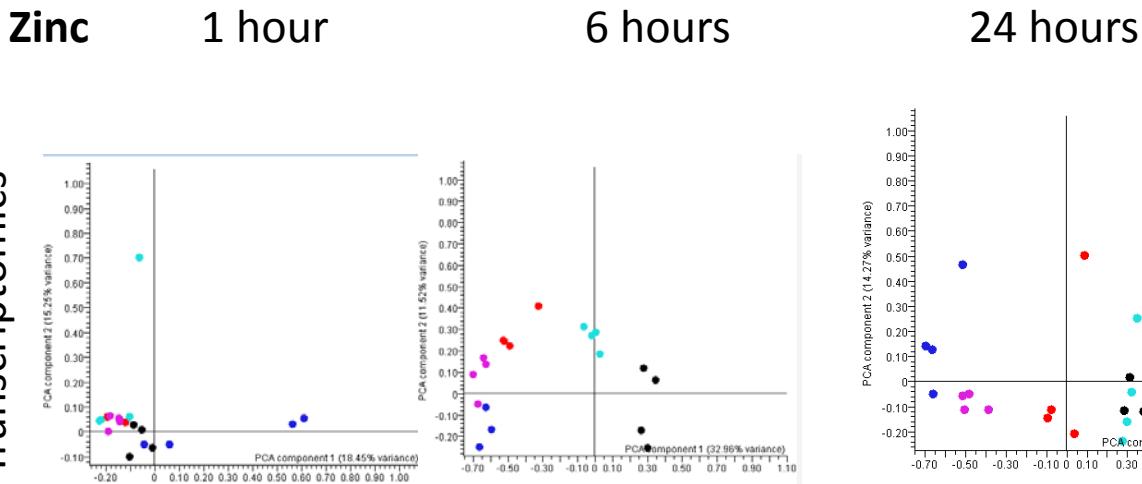


Lipidomics

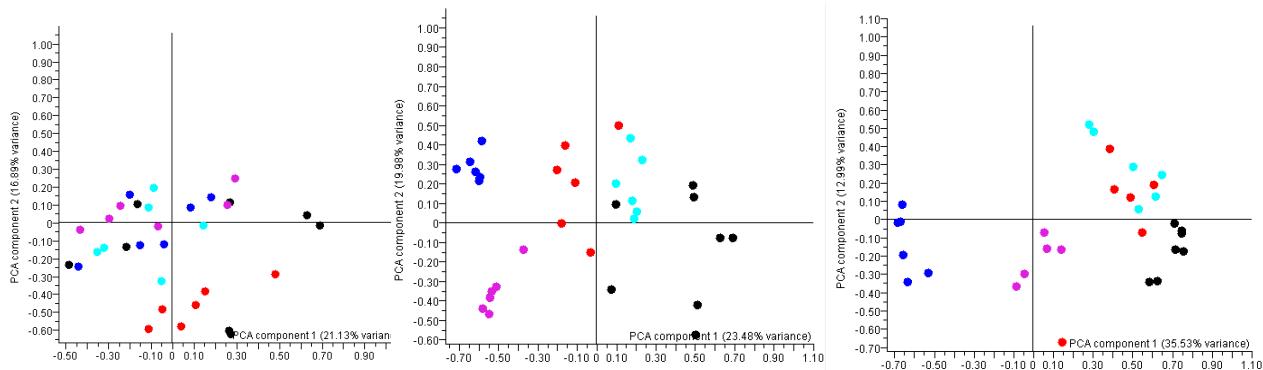


control samples are shown in black,
 Ag NP-NM300K treated in red,
 Ag MP in cyan,
 Ag⁺ in blue.

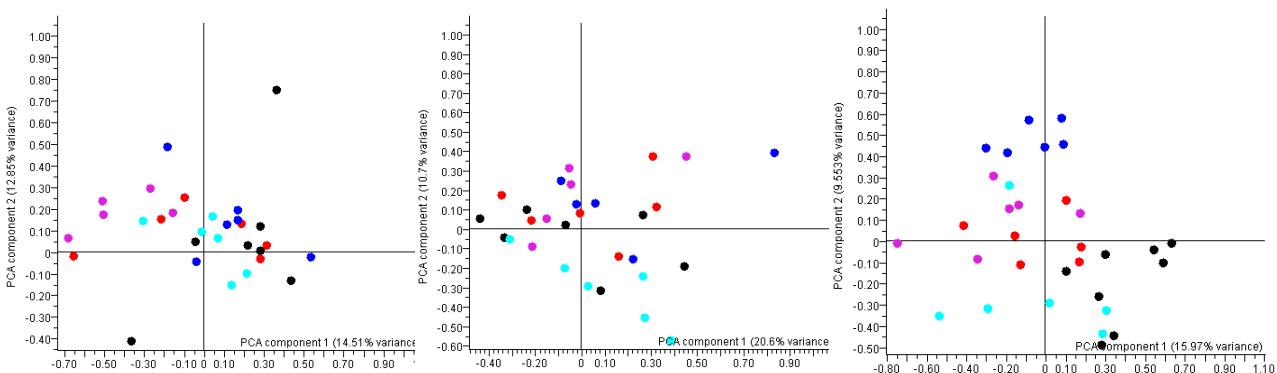
Transcriptomics



Polar metabolomics



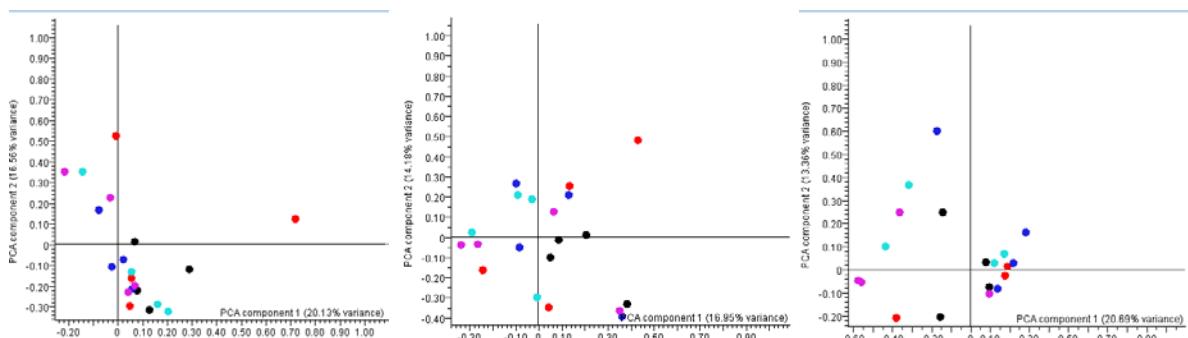
Lipidomics



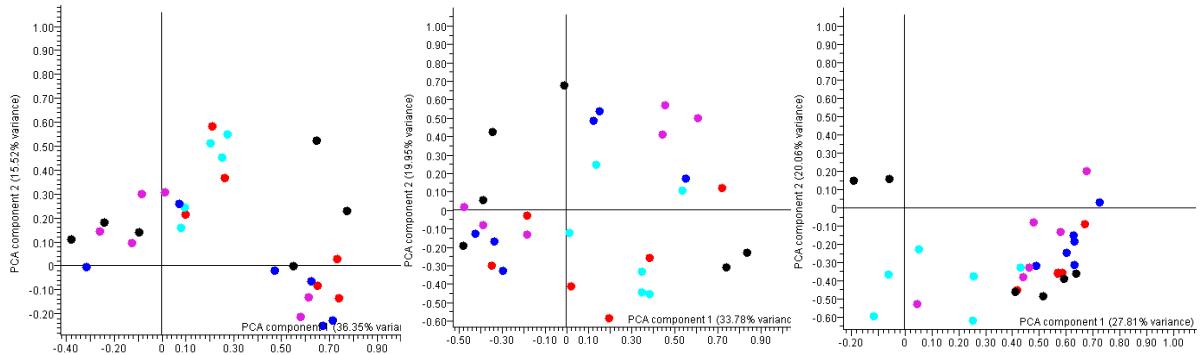
control samples are shown in black,
 ZnO NP-NM110 treated in red,
 ZnO NP-NM111 in cyan,
 Zn₂₊ in blue
 ZnO MP in pink.

Transcriptomics

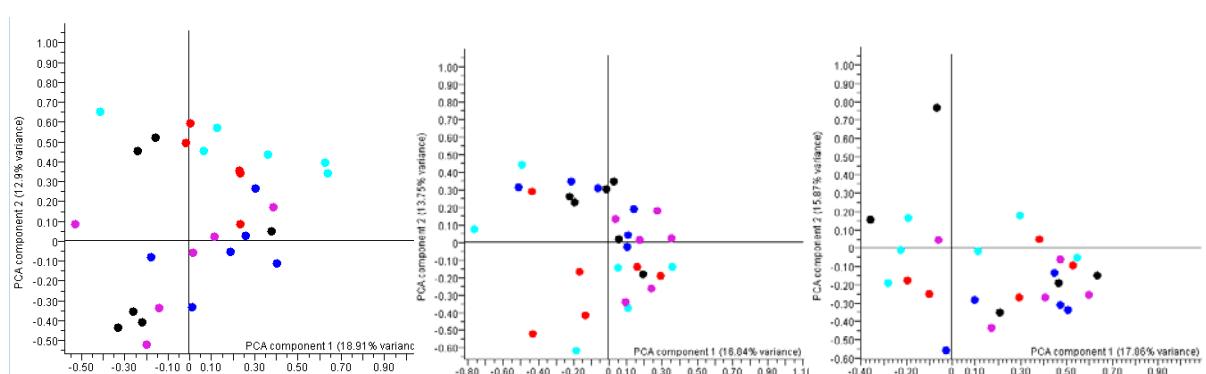
CeO₂ 1 hour 6 hours 24 hours



Polar metabolomics



Lipidomics



control samples are shown in black,
CeO₂ NP-A treated in red,
CeO₂ NP-C in cyan,
CeO₂ NP-E in blue
CeO₂ NP-NM212 in pink.