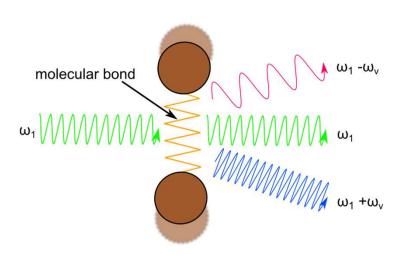
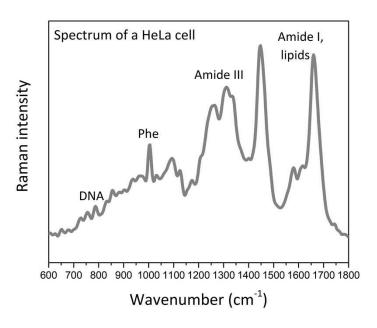
Non-supervised algorithms for Raman spectral decomposition in the in-vitro study of oxide nanoparticles effects on human cells

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### Raman spectroscopy

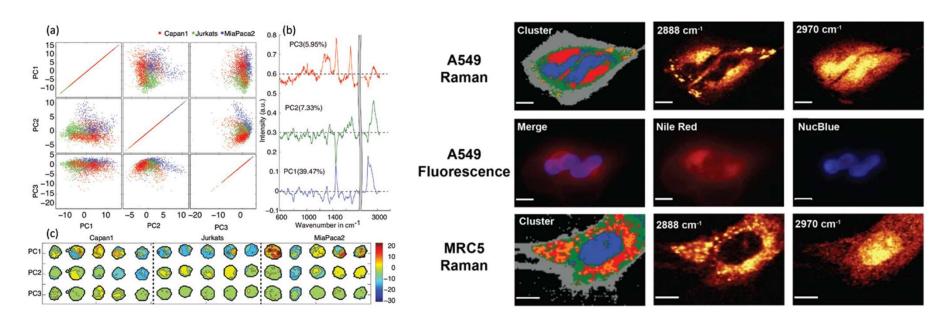
- Raman spectroscopy is a inelastic light scattering technique that can be used to probe molecular vibrations, electronic excitations, magnons etc.
- Molecular vibrations have specific energies (wavenumbers) that sometimes enable identification of molecules within a material





### Raman spectroscopy modalities

- Spontaneous, CARS, SRS, enhanced-SERS
- Common applications:
- ✓ Separating healthy and cancer cells/tissues based on vibrational spectra.
- ✓ Raman imaging obtaining Raman maps based on the vibrational wavenumbers belonging to specific biomolecules



Popp et al, Spectrosc. Suppl. 34 (2019)

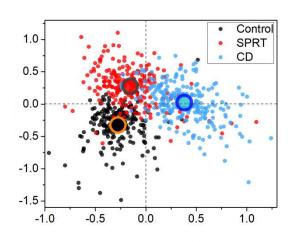
Surmacki et al, Sci. Rep. 8 (2018)

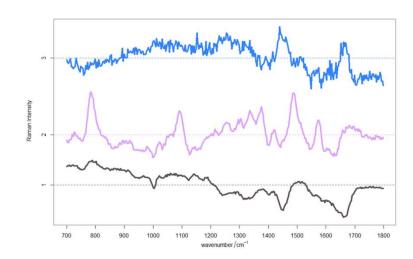
### Non-supervised algorithms

- Non-supervised multivariate methods:
  - Exploratory data analysis
  - Data dimension reduction
  - Clustering according to the components that present spectral variance
- Principal component analysis (PCA):

Loadings (basis vectors) are orthogonal and represent the directions of maximum variance between the spectra in the set

Scores – projections of original dataset onto loading vectors





### Non-supervised algorithms

### PCA:

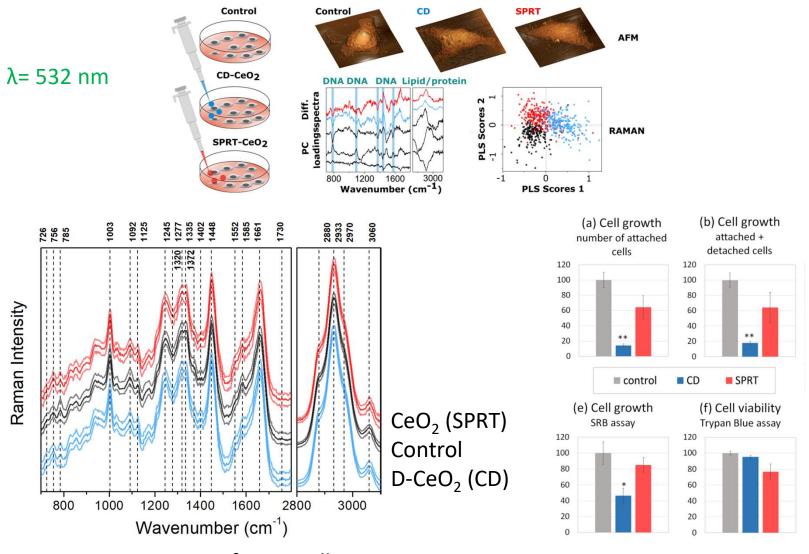
orthogonality of loading vectors, unique solution of equation, low sparseness, troublesome interpretability

- nnPCA PCA with constraint of non-negative loading vectors: non-orthogonality of loading vectors, higher sparseness, non-negative values, good interpretability
- NMF non-negative matrix factorization:
   non-orthogonality of loading vectors, non-unique solution of equation, non-negative values, good interpretability

$$X_{nxp} = W_{nxk} * H_{kxp} + R$$

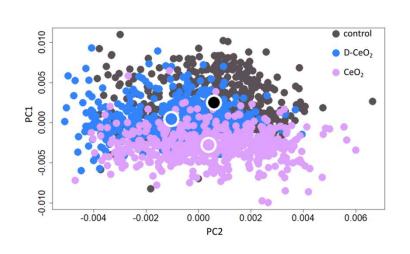
k – reduced dimension, n – number of samples, p-number of features

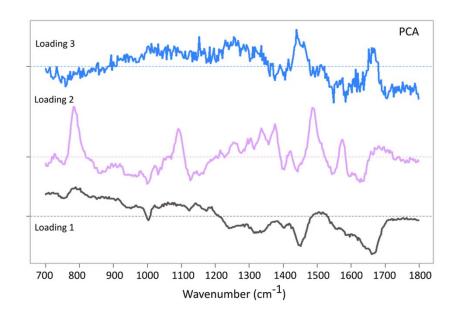
# Raman spectra of CeO<sub>2</sub> NP<sub>s</sub>-treated cells

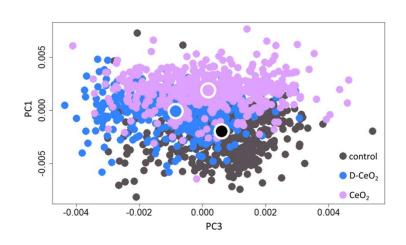


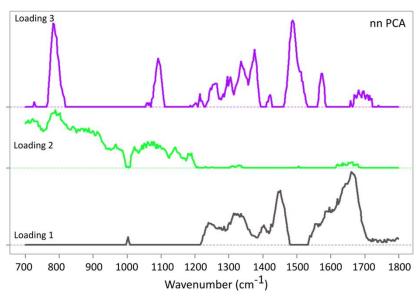
Mean spectra of HeLa cells +/- standard deviation

# Raman spectra of CeO<sub>2</sub>-treated cells



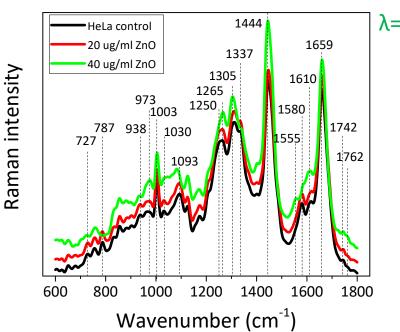


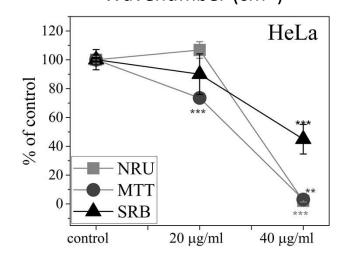




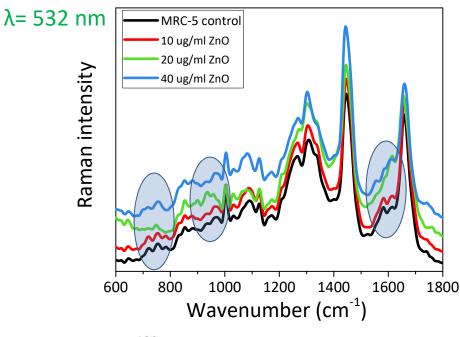
# Raman spectra of ZnO NP<sub>s</sub>-treated cells

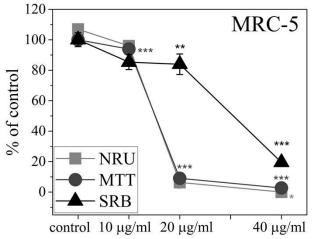






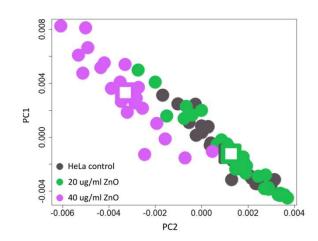
### Mean spectra of MRC-5 cells

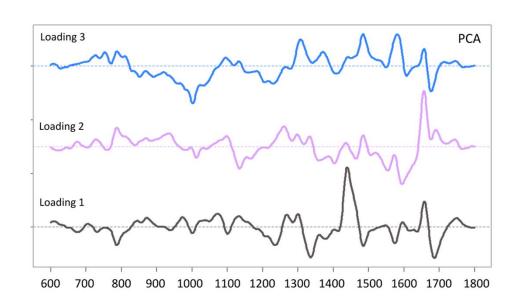


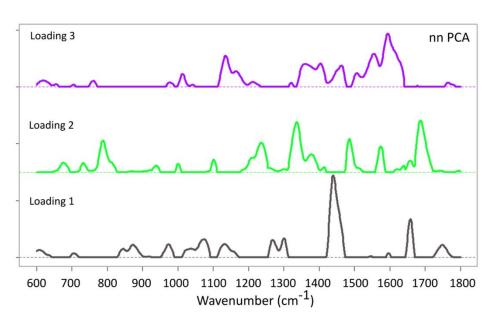


# Raman spectra of ZnO NP<sub>s</sub>-treated cells

- HeLa cells
- nnPCA loading vectors resemble phosphatydilserine and DNA spectra
- With the increased ZnO concentration, cells show:
- ✓ Higher phosphatidylserine content
- ✓ Lower DNA content

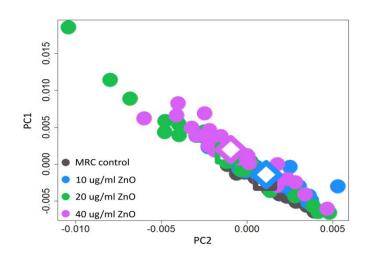


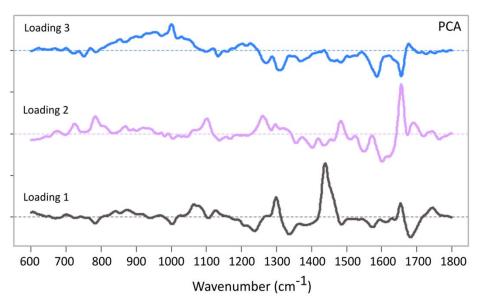


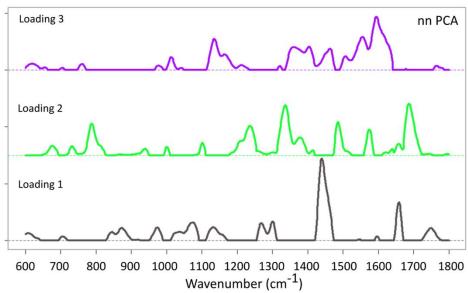


# Raman spectra of ZnO NP<sub>S</sub>-treated cells

- MRC-5
- nnPCA loading vectors resemble unsaturated fatty acid and DNA spectra
- With the increased ZnO concentration, cells show:
- ✓ Higher unsaturated lipids content
- ✓ Lower DNA content

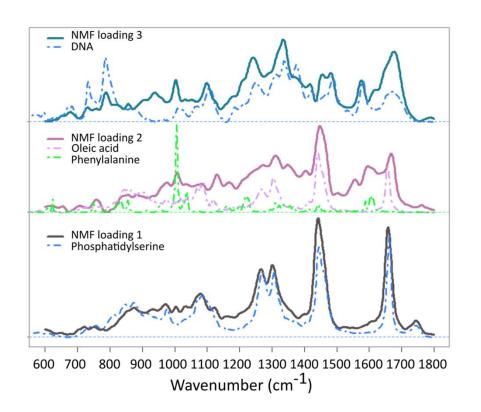


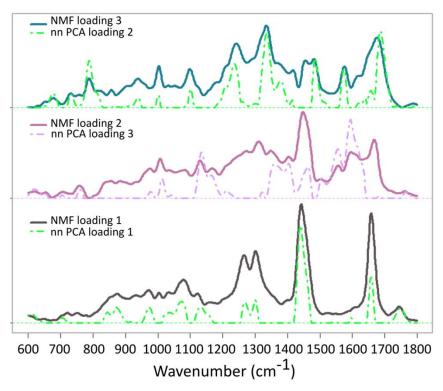




# Raman spectra of ZnO NP<sub>s</sub>-treated cells

- NMF: ns-NMF algorithm, dimension reduction to k=3
- 2 NMF loadings match the spectra of DNA and phosphathidylserine
- when number of components is higher than 3 variations of the same components reappear





### Conclusions

- PCA, nn-PCA and NMF were used for spectral data analysis of cells treated with different concentrations of CeO<sub>2</sub> and ZnO NPs
- Better interpretability of the NPs effect on cells was demonstrated for nn-PCA, especially in the case of ZnO nanoparticles
- NMF provided similar conclusions as nn-PCA, but the choice of the number of latent components influences the correlation of loadings with the spectra of real biomolecules
- For NMF the best mathematical results are not best in terms of physical interpretation

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# Thank you for the attention!

