



HAL
open science

Protein corona characterization through analytical quality-by-design

Jeanne Deleforterie, Thierry Bastogne

► **To cite this version:**

Jeanne Deleforterie, Thierry Bastogne. Protein corona characterization through analytical quality-by-design. 8èmes Rencontres Annuelles en Nanometrologie, Jun 2019, Paris, France. hal-02632035

HAL Id: hal-02632035

<https://hal.science/hal-02632035v1>

Submitted on 27 May 2020

HAL is a multi-disciplinary open access archive for the deposit and dissemination of scientific research documents, whether they are published or not. The documents may come from teaching and research institutions in France or abroad, or from public or private research centers.

L'archive ouverte pluridisciplinaire **HAL**, est destinée au dépôt et à la diffusion de documents scientifiques de niveau recherche, publiés ou non, émanant des établissements d'enseignement et de recherche français ou étrangers, des laboratoires publics ou privés.

Protein Corona Characterization through Analytical Quality-by-Design

J. Deleforterie¹, T. Bastogne^{1,2,3}

¹CYBERnano, ²CRAN UL-CNRS UMR 7039, ³INRIA BIGS

Background. After administration of nanoparticles (NP) into biological fluids, the NP may interact with surrounding biomolecules like proteins [1]. Those interactions can then cause protein corona (PC) that affect the physicochemical and biological properties of NP. Numerous studies have been carried out to characterize PC effects in simple and complex biological media but our understanding is far from being complete. Moreover, the variety of methods and tools used for this purpose prevents any robust conclusion and sometimes introduces confusions and uncertainty. Analytical Quality by Design (AQbD), introduced in 2010 [2], is the application of the QbD concept, strongly recommended by FDA, to analytical method development [3]. AQbD aims at optimizing accuracy and robustness of analysis results by identifying and controlling critical quality variables and risk factors over the complete protocol, including biological sample preparation, metrological protocol and statistical analysis.

Objectives

Our goal is to establish a safe and repeatable protocol through AQbD to characterize the Protein Corona by minimizing errors due to data misinterpretation and by reducing and controlling lab-to-lab variability in order to better predict the PC consequences.

Methods

We have analyzed 90 articles devoted to PC characterization and revisited their contributions in the AQbD paradigm. The articles were selected according to the following criteria : published within the last years (2000-2019), published in English language and relevant with the search related terms.

Results

We have firstly proposed a generic analytical target profile of the PC characterization. We have also identified the critical biological and physicochemical properties to be controlled as well as the critical risk factors among all the protocol parameters. Then, we have reported the main analytic methods used to analyze the PC within their limits and advantages [4], applied to NP. Finally, we have exposed how to best exploited data analysis derived from these methods according to AQbD requirements.

Conclusion

We have outlined which analytical techniques are the most suitable at developing efficient predictive tests to systematically identify and validate physicochemical and biochemical markers correlated to corona formation on Nanoparticles using the novel approach of Quality by Design (AQbD).

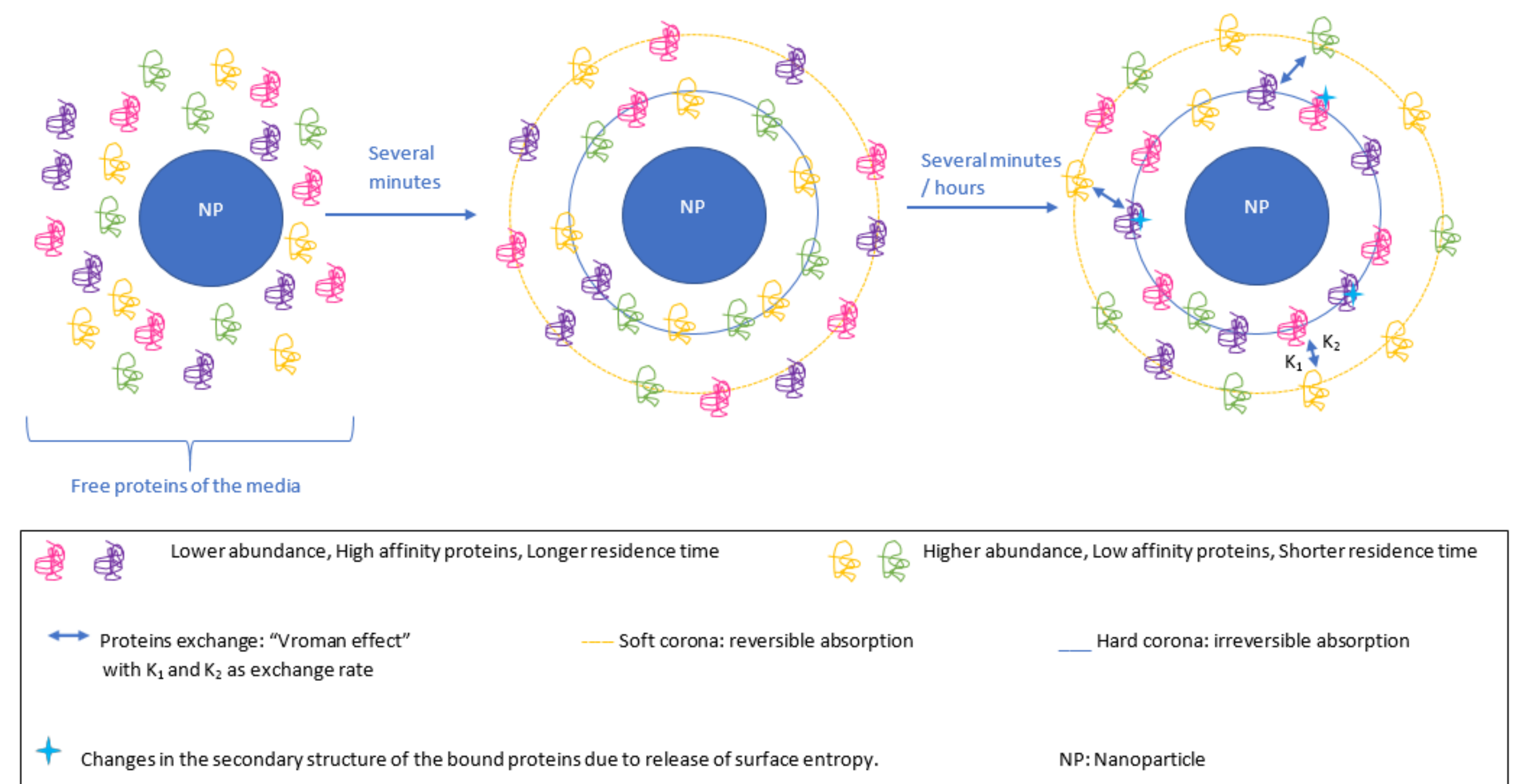


Fig.1: Schematic representation of dynamics exchanges of proteins forming the protein corona.

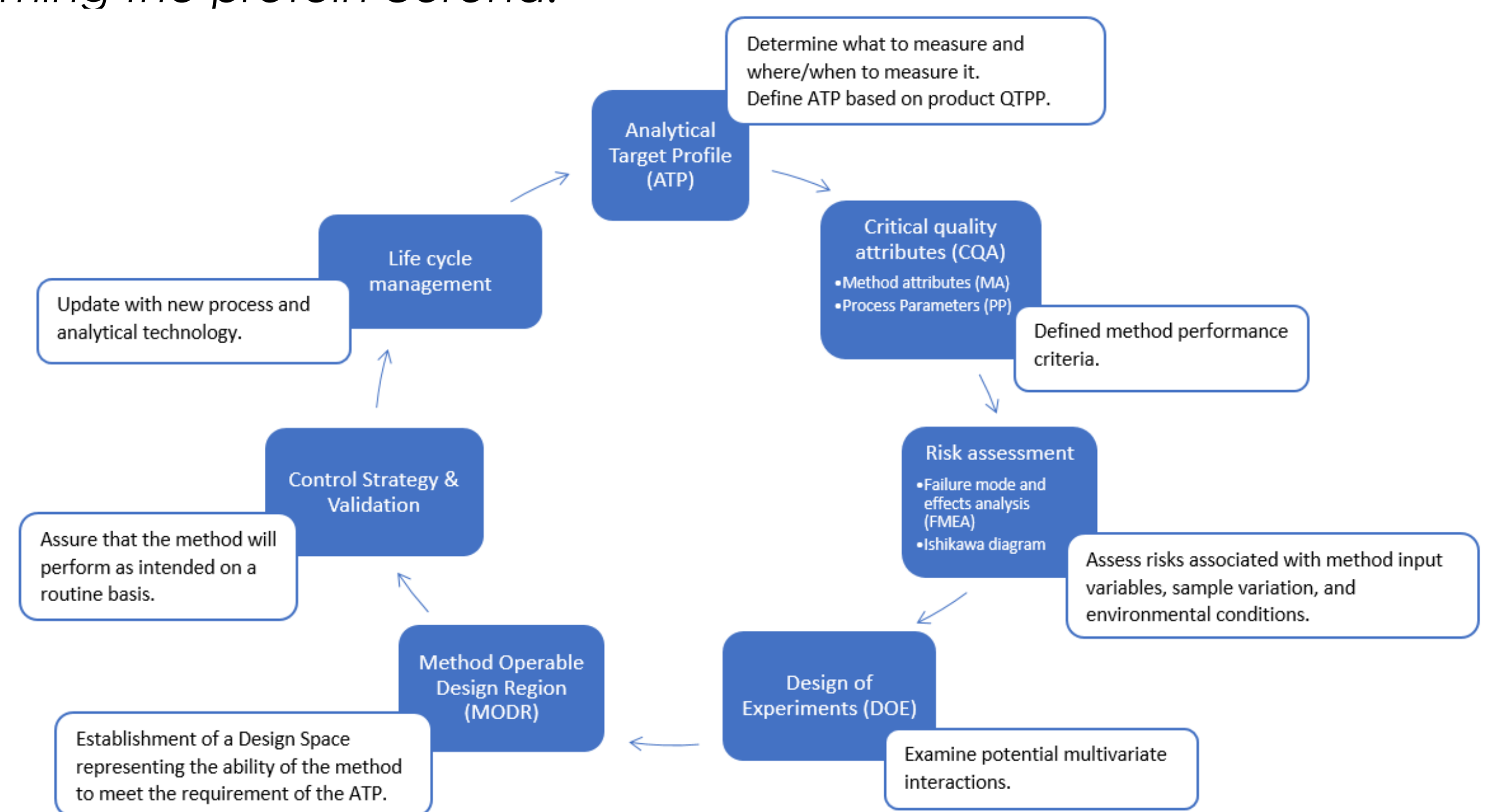


Fig.2: Analytical Quality by Design Workflow.

Main Methods	Physicochemical													Biological															
	Stability	Aggregation / Agglomeration rate	Chemical composition	Secondary structure	Shape/Dimension/ Surface	Surface area	Size/ Size distribution	Surface charge / zeta potential	NP Hydrophobicity	Surface chemistry	Protein shell thickness	Surface contamination	Redox potential	Photocatalytic activity	Crystal structure	Dustiness	Water solubility/Dispersibility	Particle physicochemical structure	Porosity	Heterogeneity	Binding thermodynamics	Binding kinetics	Drug release	Kinetic of circulation	Toxicity	Complement activation	Bio-distribution	Radical formation potential	
Differential centrifugal sedimentation (DCS)	X						X		X	X				X					X	X									
Centrifugal liquid sedimentation (CLS)		X					X			X				X															
Thermogravimetric analysis (TGA)	X								X					X															
Circular Dichroism	X		X																										X
Cyanide digestion	X																												
DTT competition	X																												
Lyophilization	X																												
Differential Thermal Analysis (DTA)	X																												
Dynamic Light Scattering (DLS)	X	X			X	X	X	X	X								X												
Small-angle X-ray scattering (SAXS)	X				X	X	X	X	X																				
Wide angle X-ray scattering (WAXS)	X																												
Surface-enhanced Raman Scattering (SERS)	X				X					X																			
Multi-angle light scattering (MALS)	X	X								X							X												
Transmission Electron Microscopy (TEM)	X	X			X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X
Scanning Electron Microscopy (SEM)	X	X			X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X	X
Cryogenic SEM	X																												
Scanning tunneling Microscopy (STM)	X				X	X	X	X	X																				
Atomic force Microscopy (AFM)	X				X	X	X	X	X																				
Confocal Microscopy	X																												
Isothermal Adsorption (BET)	X					X	X																						
Isothermal titration calorimetry (ITC)	X								X																				

Fig.3: Presentation of some of the main methods to analyze physicochemical and biological properties of Nanoparticles.

References:
 [1] Monopoli, Marco P., et al. "Physical- chemical aspects of protein corona: relevance to in vitro and in vivo biological impacts of nanoparticles." *Journal of the American Chemical Society* 133.8 (2011): 2525-2534.
 [2] Schweitzer, Mark, et al. "Implications and opportunities of applying QbD principles to analytical measurements." *Pharmaceutical Technology* 34.2 (2010): 52-59
 [3] Guideline, ICH Harmonized Tripartite. "Validation of analytical procedures: text and methodology Q2 (R1)." International conference on harmonization, Geneva, Switzerland. 2005.
 [4] Rasmussen, Kirsten, et al. "Physico-chemical properties of manufactured nanomaterials - Characterization and relevant methods. An outlook based on the OECD Testing Programme." *Regulatory Toxicology and Pharmacology* 92 (2018): 8-28.