



# Cell based biosensor approach to characterize interactions between nanoparticles and cell surface receptors

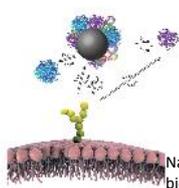
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## Background

- When nanomaterial come into contact with complex biological fluids, they rapidly get covered by a selected groups of biomolecules and this layer act as the interface between nanomaterial and the environment<sup>1</sup>.
- In particular protein molecules get adsorbed onto the surface of nanomaterial and form a coating around the nanomaterial, which refers as 'protein corona'.



The protein corona signifies the biological identity of the nanomaterial thus effect their applications

Despite the rapid increase of the understanding of the bio-nano-interface *in vitro*, its behaviour in biologically relevant environment remains obscure

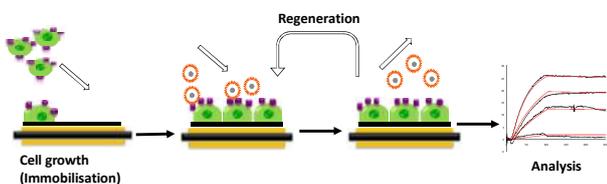
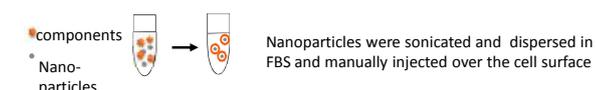
Nanoparticles-corona complex in a biological environment<sup>1</sup>

- The work presented is aimed developing a QCM based platform for the characterisation of interactions between bio-nano interface and its cell binding partners using Attana label free cell based biosensor system<sup>2</sup>
- This new biosensor system is based on the quartz crystal microbalance technology (QCM) and utilised adherent cells directly grown on the surface of a sensor in contrast to the conventional biosensor system where the bio receptors are often the purified biomolecules

## Methods

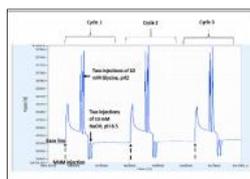
### Assay set up

- Adenocarcinoma human alveolar basal epithelial cells (A549) and adenocarcinoma human colon cell line (Caco-2) were selected as model cell lines.
- A549 cells were selected to model the cell exposure to NPs after inhalation and Caco-2 cells were selected as cell exposure via ingestion.
- TiO<sub>2</sub> and SiO<sub>2</sub> nano-particles (50 nm) were used for the assay development



### Assay workflow

- Experiments were performed using Attana cell™ 200 biosensor (Attana AB) at a flow rate of 20 ul /min at 22 °C.
- Binding of nanoparticles was studied by sequential injections (105 seconds) of the nanoparticle solutions followed by dissociation up to 200 seconds and the surface was regenerated using 10mM Glycine, pH2 followed by 10mM NaOH at pH 8.5
- The frequency changes during binding experiments were recorded using Attestar software (Attana AB) and analysed using Evaluation software (Attana AB).



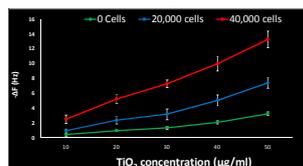
Three sequential injections of TiO<sub>2</sub> nano-particles and regeneration

## References

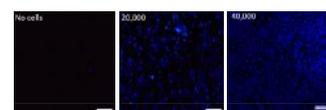
1. Marco P. Monopoli, M.P., Åberg, C., Anna Salvati, A., Dawson, K.A. Nature Nanotechnology. 2012; 7; 779-786
2. Peiris, D., Markiv, A., Curley, G. P., Dwek, M. V. Biosensors and Bioelectronics. 2012; 35:160-166
3. Diluka Peiris, Daniel Wallinder, Teodor Aastrup. Eurolab. 2015 (1) 63-6

## Results

### Effect of seeding density on NPs interactions



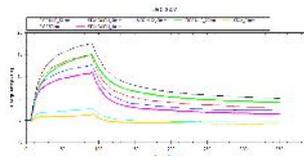
Maximum frequency response of TiO<sub>2</sub> (PROM-TiO<sub>2</sub> uncoated) as a function of seeding density. ( Error bars SD for triplicate injections on three different sensor chips with corresponding seeding density)



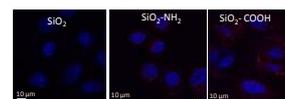
To-Pro3 (1mM) stained A549 cells on QCM sensor surfaces prepared using increasing seeding density. Scale bars 200 µm.

- The maximum frequency response increased with increasing seeding density for all TiO<sub>2</sub> concentrations

### Effect of surface modifications on NPs binding

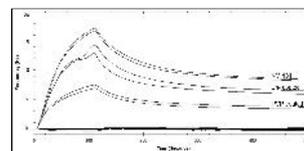


Sensograms for SiO<sub>2</sub> (JRC) binding to A549 cell surface. Amine or carboxyl modified SiO<sub>2</sub> particles showed binding to A549 cell surfaces while unmodified SiO<sub>2</sub> showed no binding at the same conditions.



Confocal images showing Ru (BFPy)<sub>3</sub> labelled SiO<sub>2</sub> binding to A549 cells.

- Biosensor data is in agreement with the immunofluorescence data, where unmodified Amine or carboxyl modified SiO<sub>2</sub> particles showed binding to A549 cell surfaces

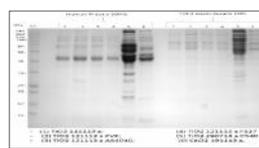


Sensograms for TiO<sub>2</sub> binding to A549 cell surface. Uncoated, PVP coated and rutile hydrophilic (NM104) TiO<sub>2</sub> particles showed binding to cell surface while TiO<sub>2</sub> (NIST), Pluronic F127 coated, Dispex AA4040 coated showed no binding

Particle	kd1	kd2	Off rates of TiO <sub>2</sub> particle dissociation
TiO <sub>2</sub> -uncoated	6.3 E-3	4.98 E-4	
TiO <sub>2</sub> (PVP coated)	5.7 E-3	5.31 E-4	
TiO <sub>2</sub> (rutile, hydrophilic)	5.1 E-3	5.03 E-4	

- There were no significant difference between dissociation rate constants of three bound particles. This might imply involvement of same protein from protein corona in the cell surface binding. However this need further investigation using proteomics approach.

### Characterization of protein corona



SDS-PAGE analysis of nanoparticles incubated in plasma and FBS

Protein Name	Accession ID	100 BSA	100 PVP	100 TiO <sub>2</sub>	100 FBS
Albumin	P02768	1.00	1.00	1.00	1.00
Immunoglobulin heavy chain	A01974	0.00	0.00	0.00	0.00
Immunoglobulin light chain	A01973	0.00	0.00	0.00	0.00
Transferrin	P02747	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin	P02746	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein	P02745	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02744	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02743	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02742	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02741	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02740	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02739	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02738	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02737	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02736	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02735	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02734	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02733	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02732	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02731	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02730	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02729	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02728	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02727	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02726	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02725	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02724	0.00	0.00	0.00	0.00
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Alpha-1-acid glycoprotein chain	P02719	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02718	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02717	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02716	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02715	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02714	0.00	0.00	0.00	0.00
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Alpha-2-macroglobulin chain	P02712	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02711	0.00	0.00	0.00	0.00
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Alpha-1-acid glycoprotein chain	P02635	0.00	0.00	0.00	0.00
Alpha-2-macroglobulin chain	P02634	0.00	0.00	0.00	0.00
Alpha-1-acid glycoprotein chain	P02633	0.00	0.00	0.00	0.00
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