The corona of protein - gold nanoparticle systems:

role of ionic strength

Cristina Cantarutti,^{a‡} Yamanappa Hunashal,^{b,a‡} Carmelo La Rosa,^c Marcello Condorelli,^c Sofia Giorgetti,^d Vittorio Bellotti,^{d,e} Federico Fogolari^{f,g} and Gennaro Esposito *^{b,g}

^a DAME, Università di Udine – 33100 Udine, Italy.

^b Science Division, New York University Abu Dhabi – Abu Dhabi, UAE.

^c Dip. Scienze Chimiche, Università di Catania – 95125 Catania, Italy.

^d Dipartimento di Medicina Molecolare, Università di Pavia – 27100 Pavia, Italy.

^e Centre for Amyloidosis and Acute Phase Proteins, Division of Medicine, University College London – London NW3 2PF, UK.

^f DMIF, Università di Udine – 33100 Udine, Italy.

^gINBB, Viale Medaglie d'Oro 305 – 00136 Roma, Italy.

[†] These authors contributed equally.

*Corresponding author

rino.esposito@nyu.edu (Gennaro Esposito)

ELECTRONIC SUPPLEMENTARY INFORMATION

Determination of AuNP surface occupancy by β2m

 β 2m is a globular protein whose three-dimensional structure can be assimilated to a cylindroid with longitudinal and transverse axes of 4.3-3.8 and 2.5-2.0 nm, respectively.

Through simple geometrical calculations, it is possible to estimate the number of proteins that can be adsorbed on the surface of a single nanoparticle. If the ratio between the volumes is considered as proposed by Calzolai *et al.*,¹ it is possible to obtain the number of proteins (N) from this equation

$$N = 0.65 \times \frac{(R_{complex}^3 - R_{NP}^3)}{R_{protein}^3} \qquad \qquad \text{Eq. S1}$$

where $R_{complex}$ is given by the sum of the NP radius and the diameter of $\beta 2m$, R_{NP} is the radius of the nanoparticle and $R_{protein}$ is the radius of $\beta 2m$. Considering the crystallographic cylindrical shape of $\beta 2m$, the previous equation (Eq. S1) becomes

$$N = 0.65 \times \frac{\frac{4}{3}(R_{complex}^3 - R_{NP}^3)}{h_{cyl} \times r_{cyl}^2}$$
 Eq. S2

where $R_{complex}$ is given by the sum of the NP radius and the height of the β 2m cylindroid, R_{NP} is the radius of the nanoparticle and h_{cyl} and r_{cyl} are the height and the base radius of β 2m cylindroid. Following this equation, a spherical Cit-AuNP with average diameter of 7.5 nm can accommodate 55-95 ß2m monomers. This estimation uses the maximum packing density factor possible for spherical proteins. Since the β2m cylindroid base dimension is a half with respect to height, considering symmetric tetramers instead of cylindroids can improve the geometrical model, leading to 15-30 tetramers per NP that means 60-120 monomers. Both cylindrical and tetrameric volume-based packing densities, however, appear unrealistic. Instead, the number of protein monomers per NP is reduced to 45-56 if the ratio of Cit-AuNP surface area and protein cylindroid base is considered.² This last calculation was experimentally confirmed by UV-Vis experiments in which the shift of the AuNP surface plasmon band (SPB) was followed by increasing β2m concentration (Figure S1). The UV-Vis titration experiment showed that a plateau in the redshift started when $\beta 2m/AuNP$ ratio was around 50. Based on this experimental assessment done with 7.5 nm Cit-AuNPs, we determined the occupancy of the 5 nm Cit-AuNPs used in this work by calculating the ratio of NP surface area and protein cylindroid base leading to an estimation of 14-20 proteins per NP.



Figure S1. a) UV-Vis spectra of 25 nM Cit-AuNPs (7.5 nm diameter) after the progressive addition of β 2m (0-20 μ M). b) Plot of SPB shift as a function of [β 2m]/[Cit-AuNP] ratio. The experiment was done in triplicate and the error is the standard deviation



Figure S2. SEM image of Cit-AuNP monolayer used as an active SERS substrate.



Figure S3. SERS spectra of 0 nM (black), 200 nM (red), 400 nM (green), 800 nM (blue) and 1600 nM (magenta) β 2m dissolved in 50 mM HEPES (pH 7) and deposited on a layer of Cit-AuNP solution. The main bands observed in the highest concentration trace can be tentatively assigned to β -sheet Amide II (1550-1600 cm⁻¹), random coil (~1250 cm⁻¹), helix-like (from turns and loops) Amide I and III (~1620 and ~1300 cm⁻¹, respectively) and aromatic side chains (Trp, at ~750 and ~1400 cm⁻¹, and Phe, around 950 cm⁻¹ and 1150 cm⁻¹). An analogous ensemble of bands, albeit with a different intensity pattern and an additional β -sheet Amide I (~1700 cm⁻¹) band, can be recognized also in the spectrum reported in Figure 1 of the main text and relative to 10nM β 2m dissolved in 0.12 mM HEPES.



Figure S4. Chemical shift invariance of β 2m NMR spectra with Cit-AuNPs at high and low ionic strength. a) Overlay of ¹⁵N-¹H HSQC maps of 8µM β 2m in 0.1 mM HEPES without (black contours) and with (red contours) 60 nM Cit-AuNPs. b) Overlay of ¹⁵N-¹H HSQC maps of 8µM β 2m in 50 mM HEPES without (black contours) and with (blue contours) 60 nM Cit-AuNPs. a) Overlay of ¹⁵N-¹H HSQC maps of 8µM β 2m with 60 nM Cit-AuNPs at 50 mM (blue contours) and 0.1 mM (red contours) HEPES. The chemical shift invariance indicates that the secondary and tertiary structure of the protein is preserved in the presence of NPs at low and high ionic strength.



Figure S5. Time course of the QCMD Δm changes for Cit-AuNPs, D76N β 2m without and with Cit-AuNPs at 50 mM (a) and 0.1 mM (b) HEPES. The Δm values were calculated using the Sauerbrey equation (see main text) from the total Δf values measured over 4,000 s flow.

Figure S6. Time courses of QCMD Δf and ΔD obtained from 0.5 mM D76N β 2m with 25 nM Cit-AuNPs at low ionic strength. The sampled experimental points are shown without decimation. The enlargement highlights the coincidence of the initial deposition phase with the single dissipation step.

Figure S7. Zeta potential for citrate-coated AuNPs with 5 nm diameter. The determination was carried out on a \sim 5 nM NP sample in 0.12 mM, HEPES at 25°C and pH 7 (the same condition as used in SERS spectra), using a Horiba Scientific nanoPartica SZ100 instrument.

References

- 1 L. Calzolai, F. Franchini, D. Gilliland and F. Rossi, Nano Lett., 2010, 10, 3101–3105.
- 2 Y. Teshima and T. Ogawa, Forma, 2001, 15, 347–364.