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1 Size-Dependent Protein-Nanoparticle Interactions in Citrate-Stabilized 2 Gold Nanoparticles: The Emergence of the Protein Corona

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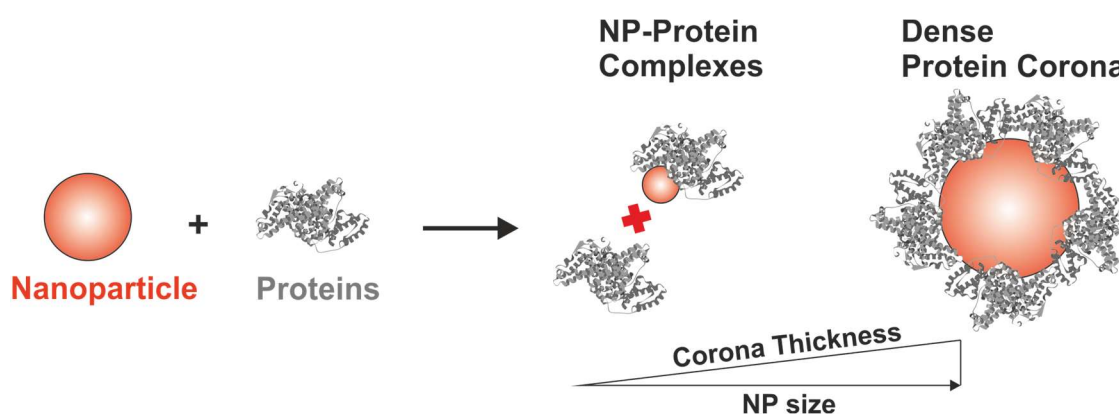
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10 Abstract

11 Surface modifications of highly monodisperse citrate-stabilized gold nanoparticles (AuNPs) with
12 sizes ranging from 3.5 to 150 nm after their exposure to cell culture media supplemented with fetal
13 bovine serum were studied and characterized by the combined use of UV-vis spectroscopy, dynamic
14 light scattering, and zeta potential measurements. In all the tested AuNPs, a dynamic process of
15 protein adsorption was observed, evolving toward the formation of an irreversible hard protein
16 coating known as the Protein Corona. Interestingly, the thickness and density of this protein coating
17 were strongly dependent on the particle size, making it possible to identify different transition regimes
18 as the size of the particles increased: (i) NP-protein complexes (or incomplete corona), (ii) the
19 formation of a near-single dense protein corona layer, and (iii) the formation of a multilayer corona.
20 In addition, the different temporal patterns in the evolution of the protein coating come about more
21 quickly for small particles than for the larger ones, further revealing the significant role that size plays
22 in the kinetics of this process. Since the biological identity of the NPs is ultimately determined by the
23 protein corona and different NP-biological interactions take place at different time scales, these results
24 are relevant to biological and toxicological studies.

25 TOC



27 Introduction

28 Noble metal nanoparticles (NPs) are among the most widely used nanomaterials in biomedicine
29 because of their outstanding optical, electromagnetic, and photothermal properties. These unique
30 properties, together with the high reactivity of the NPs and their affinity for binding many different
31 (bio)molecules, make them attractive candidates in a wide variety of uses and applications including
32 delivery, diagnostics, and therapy.^{1, 2} However, despite the rapid development of strategies for particle
33 design and functionalization, relatively little is known about their *in vitro* and *in vivo* behavior in
34 complex biological systems.³

35 The ability to integrate NPs in biological systems ultimately relies on the fundamental understanding
36 of the interaction of these inorganic NPs with biological fluids.⁴⁻⁶ A large number of biological fluids
37 (plasma serum or otherwise) are complex aqueous media composed of electrolytes, proteins, lipids,
38 and metabolites, able to adsorb (by electrostatic, hydrophobic, van der Waals, and dispersive forces)
39 onto the surfaces of the NPs forming a protein dense coating known as Protein Corona (PC),⁷⁻¹¹ a
40 phenomena extensively studied previously in implants.¹² The PC shields the original surface
41 properties of the NPs acting as a “complex” surfactant, and alters their size and composition providing
42 the NPs with a new biological identity.^{10, 13, 14} This corona is what is ultimately “seen” by cells, and
43 critically determines the physiological response and interaction of the NPs with living systems,
44 including cellular uptake, circulation lifetime, signaling, biodistribution, therapeutic effects, and
45 toxicity.¹⁵⁻²¹

46 The composition, structure and kinetics of the PC formation depends on (i) the specific characteristics
47 of the biological environments in which NPs are dispersed, especially protein composition²² and
48 concentration;^{16, 23} (ii) the physicochemical features of the dispersed material, such as NP chemical
49 composition²⁴, morphology,^{7, 25, 26} surface charge (functional groups),^{16, 27} and hydrophobicity,^{7, 28} and
50 (iii) the exposure time, which directly correlates with the relative abundance of proteins and the
51 different protein-NP binding constants (Vroman’s effect).^{24, 27, 29} Among these parameters, the
52 morphology of the NPs relating to size and shape has been suggested to play an important role. Due to
53 surface curvature effects,³⁰ molecules tend to adopt different conformations on top of NPs as
54 compared to flat surfaces indicating different protein-binding affinities, giving rise variation of the
55 structure of PC according to the size of the NPs.³¹ Pioneering studies were performed by Cerdervall et
56 al.^{7, 32} who investigated the role of surface curvature in copolymer NPs of 70 and 200 nm after their
57 dispersion in plasma, and further studies were later extended to other sizes and compositions
58 including AuNPs,^{22, 33-35} AgNPs,²⁰ polystyrene NPs,³⁶⁻³⁸ SiO₂,^{14, 15, 25, 36} metal oxide NPs,^{24, 39, 40} and
59 polymer-coated FePt and CdSe/ZnS.⁴¹ Although all of the authors reported consistent results, it was
60 often difficult to separate the effect of NP size from other physicochemical aspects, such as particle
61 composition and surface chemistry. Some of them found that particle size and surface curvature
62 influences the amount of bound protein and (to a lesser extent) the identity of these proteins.^{15, 25, 32} In

63 contrast, other studies reported significant qualitative size-dependent changes in protein adsorption.⁴⁰
64 So far, the underlying mechanism explaining size-dependent protein-particle specific interaction is not
65 yet resolved. Moreover, in all these studies the diameter of the tested NPs was above that of most
66 abundant serum proteins (6-12 nm). As a result, changes in the PC of NPs of a similar or even smaller
67 size than proteins were not addressed. In this regard, only a few studies have provided data on this
68 size regime, including AuNPs (2 nm, 15 nm),^{16, 22} CdSe/ZnS (9-11 nm),⁴² and FePt (10 nm).⁴¹

69 Little is known about the PC formation of NPs when they are smaller than most of the proteins
70 themselves, with subsequent lack of knowledge of their biological consequences. Smaller NPs present
71 some features that make them very attractive for biomedical applications: they are less prone to
72 opsonization and have substantially longer lifetimes in the bloodstream, in such a way that a particle
73 as small as 10 nm would not significantly activate the immune system.^{43, 44} They also have faster
74 biodegradation (e.g., dissolution) together with excretion profiles which may avoid unwanted
75 bioaccumulation.⁴⁵ Moreover, the small size of the NP may affect the PC formation process. Thus,
76 when NPs have sizes similar to (or even smaller than) those of proteins it may be difficult to form a
77 compact corona since the number of proteins at the surface of the particle is too small for a crowding
78 cooperative effect to be achieved.^{24, 46} Conversely, as the size of the particles becomes larger, it may
79 be possible for a proper full density protein corona to develop.

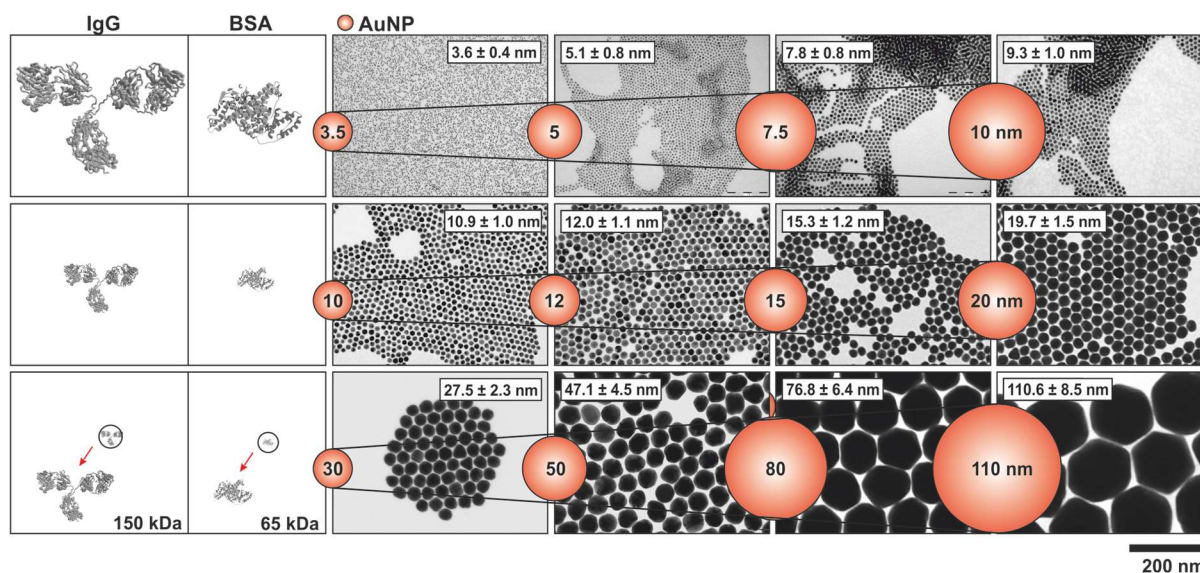
80 Taking advantage of the recent improvements in the synthesis of sub-10 nm AuNPs,^{47, 48} we have
81 studied the formation of PC on highly monodisperse citrate-stabilized AuNPs of different sizes (from
82 3.5 to 150 nm) after their exposure to cell culture media (CCM) supplemented with fetal bovine serum
83 (FBS). By the combined use of UV-vis spectroscopy, dynamic light scattering, and zeta potential, a
84 dynamic process of protein adsorption has been identified and characterized for all the tested particles,
85 leading to the formation of an irreversible coating that mediates the colloidal stability of the AuNPs in
86 the physiological medium. Interestingly, the thickness (and compactness) of this protein coating was
87 strongly determined by the size of the NPs being possible to identify a transition regime from AuNP-
88 complexes (or sparse corona) in which the thickness of the corona was smaller than that expected for a
89 single protein layer (AuNPs \leq 12 nm), to the formation of a full corona composed by a dense protein
90 monolayer for AuNPs sizes from 12 to 80 nm. Finally, for very large particles ($>$ 80 nm), an
91 additional regime could be identified in which a multilayered corona composed of 2-3 protein layers,
92 similar to that reported in flat surfaces, seemed to be favored.⁴⁹

93 **Results and Discussion**

94 The protein adsorption process was studied by incubating citrate-stabilized AuNPs for 48 h at 37⁰C in
95 complete cell culture media (cCCM); composed of Dulbecco's Modified Eagle Medium (DMEM)
96 supplemented with 10% of fetal bovine serum (FBS). Highly monodisperse citrate-stabilized AuNPs
97 with sizes ranging from 3.5 to 150 nm were synthesized following recently reported kinetically-

98 controlled seeded-growth approaches.^{47, 48} Citrate is a biocompatible reducer commonly used in the
 99 synthesis of AuNPs which provides a negative electrostatic stabilization once adsorbed on the NP
 100 surface. Besides, different from other surfactants that hinder the formation of the protein corona (PC),
 101 i.e., polyethylene glycol (PEG),^{26, 50, 51} the citrate layer is easily replaced, allowing the spontaneous
 102 adsorption of proteins under normal conditions of *in vitro* and *in vivo* exposure.^{50, 52} Moreover, FBS
 103 is a biological fluid that contains more than 3700 different proteins, with 12 orders of magnitude
 104 difference in their relative concentrations.⁷ Among all proteins in serum, albumin represents the most
 105 abundant fraction and accounts for ~60% in mass of the total proteins in solution (1 mM), with
 106 globulins the second most abundant (~40%) and all the rest representing less than 1%.⁵² With this in
 107 mind, in Figure 1 the relative size of the particles used in this study is compared to proteins, using
 108 bovine serum albumin (BSA) and Immunoglobulin G (IgG) as model proteins.

109 For the study of the PC formation, it is important to previously select the proper exposure condition of
 110 the AuNPs in cCCM. cCCM is a high ionic strength medium in which the stability of citrate-coated
 111 AuNPs is strongly compromised due to the screening of the electrostatic repulsion between particles
 112 by the high concentration of charges in solution. This process is evidenced when performing exposure
 113 experiments in DMEM media (**Fig. 2A**), where the immediate aggregation of 10 nm AuNPs is clearly
 114 observed by UV-vis spectroscopy.⁵³ Thus, in the absence of serum proteins, the localized surface
 115 plasmon resonance (LSPR) band of the particles rapidly vanishes, red-shifts and broadens, and a new
 116 band between 600 and 800 nm emerges, which can be univocally ascribed to the coupling of the
 117 plasmon modes of individual NPs when they come into contact.^{52, 54}

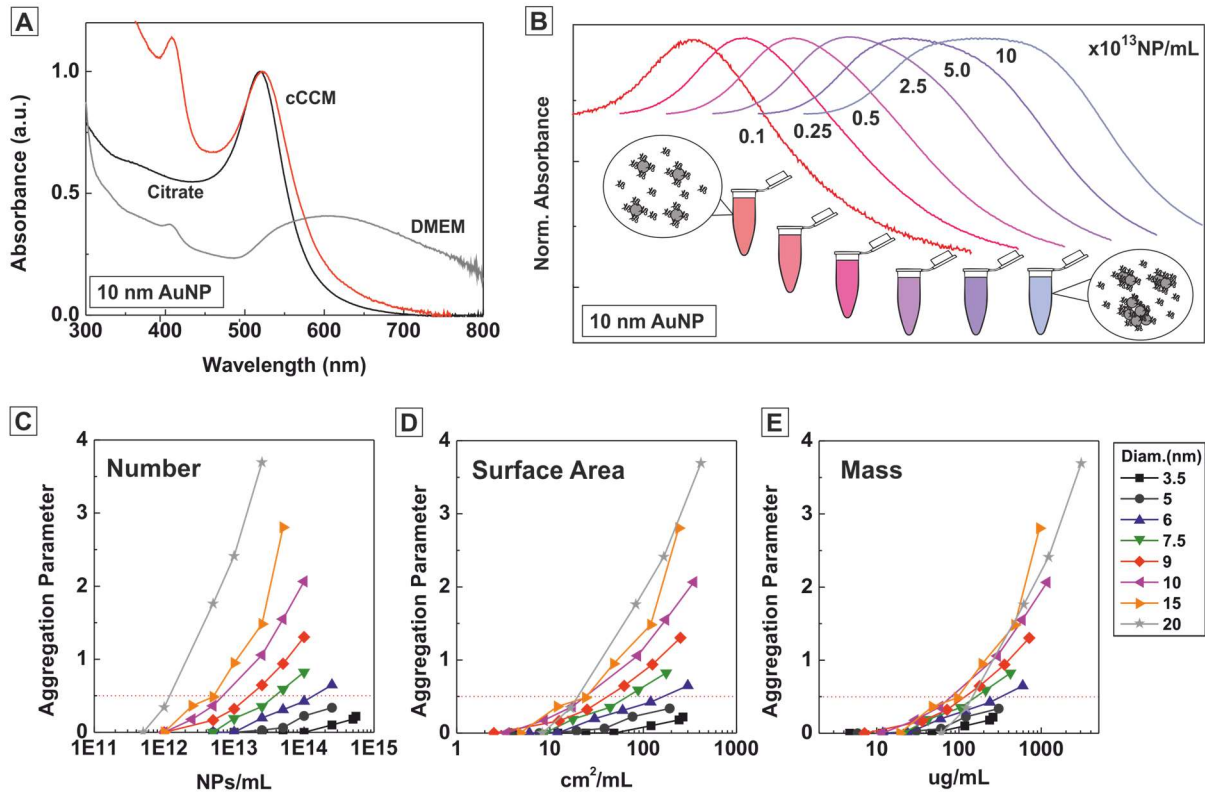


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 119 **Figure 1. Representative TEM images of citrate-stabilized AuNPs of selected sizes used in this study.** The
 120 relative difference in size between AuNPs and proteins has been compared by choosing BSA (65 kDa) and IgG
 121 (150 kDa), the two most abundant protein in serum, as model proteins. The inset spheres represent the size of
 122 the AuNPs scaled to the model proteins schematically drawn in the column on the left. The smallest AuNPs (3.6
 123 nm) cannot accommodate more than 1-2 proteins while AuNPs larger than 30 nm can accommodate hundreds of
 124 them on their surfaces.

125 In the presence of proteins (fixed at the level of standard *in vitro* studies), particle concentration is the
126 main driver for aggregation. Stability conditions of 10 nm AuNPs in cCCM were initially studied by
127 exposing the same volume (1 mL) of particles of adjusted concentrations (from 10^{12} NP/mL to 10^{15}
128 NP/mL) to an equal volume of cCCM (9 mL), thus keeping the concentration of proteins constant
129 (**Fig. 2B**). The extent of aggregation was systematically quantified from UV-vis spectra, by
130 calculating the aggregation parameter (AP) according to Lévy *et al.*⁵⁴ after 1 h of exposure. AP values
131 higher than 0.5 indicated significant aggregation. It is important to note that the LSPR peak and the
132 calculation of this parameter were not affected by the presence of free protein in solution, since
133 proteins absorb at wavelengths far lower than the characteristic AuNPs band. For the selected NP-
134 protein exposure conditions, AuNPs were stable in cCCM at a low concentration of $2.5 \cdot 10^{12}$ NPs/mL
135 while they formed aggregates when the concentration was increased above $2.5 \cdot 10^{13}$ NPs/mL. Since the
136 concentration of proteins in the media does not represent a limiting factor (in all conditions the
137 amount of proteins in solution is largely above that needed to fully cover all available NP surface), the
138 aggregation is likely due to kinetic factors when mixing both solutions. That is, NPs destabilize and
139 aggregate before proteins succeed in protecting their surfaces.

140 When the NP and the protein solutions are mixed, two different kinetic processes are competing at the
141 same time: (i) the destabilization (and further aggregation) of the NPs promoted by the high ionic
142 strength of the media in which they are dispersed and (ii) the stabilization of the NP surface against
143 aggregation via protein adsorption. Thus, if the formation of a protective protein layer is faster than
144 the characteristic time of destabilization (that is the time needed by two individual “bare” particles to
145 bind to each other), then the resultant colloidal solution will remain stable. Conversely, the sample
146 will destabilize and (partially or totally) agglomerate, forming aggregates immediately coated by
147 proteins that will progressively sediment.⁵⁵ Based on these results, it becomes clear that these two
148 processes take place in the same time scale. For completeness, stability conditions were additionally
149 evaluated for particles of different sizes (3.5 to 20 nm) and the results were normalized to: (1) the total
150 number of NPs in solution (**Fig. 2C**), (2) the total surface area exposed by the NPs (**Fig. 2D**) and (3)
151 the total gold mass in each sample (**Fig. 2E**), with it being possible to observe that curves hardly
152 converge in any case. Only when the results were normalized to the total gold mass in each sample
153 did they become comparable, which suggest that, in excess of free proteins, the aggregation of the
154 particles is not exclusively determined by the probability of collisions of two individual particles but
155 may be affected by additional parameters. Interestingly, small particles appear to be slightly more
156 stable than large ones and thus are able to reach higher concentrations. This could be related to faster
157 protein coating or a faster dispersion of the NP due to increased Brownian motion as size decreases.
158 Consequently, stability conditions depend on both the characteristics of the AuNPs and the ratio
159 between the NPs and protein concentration in the media.

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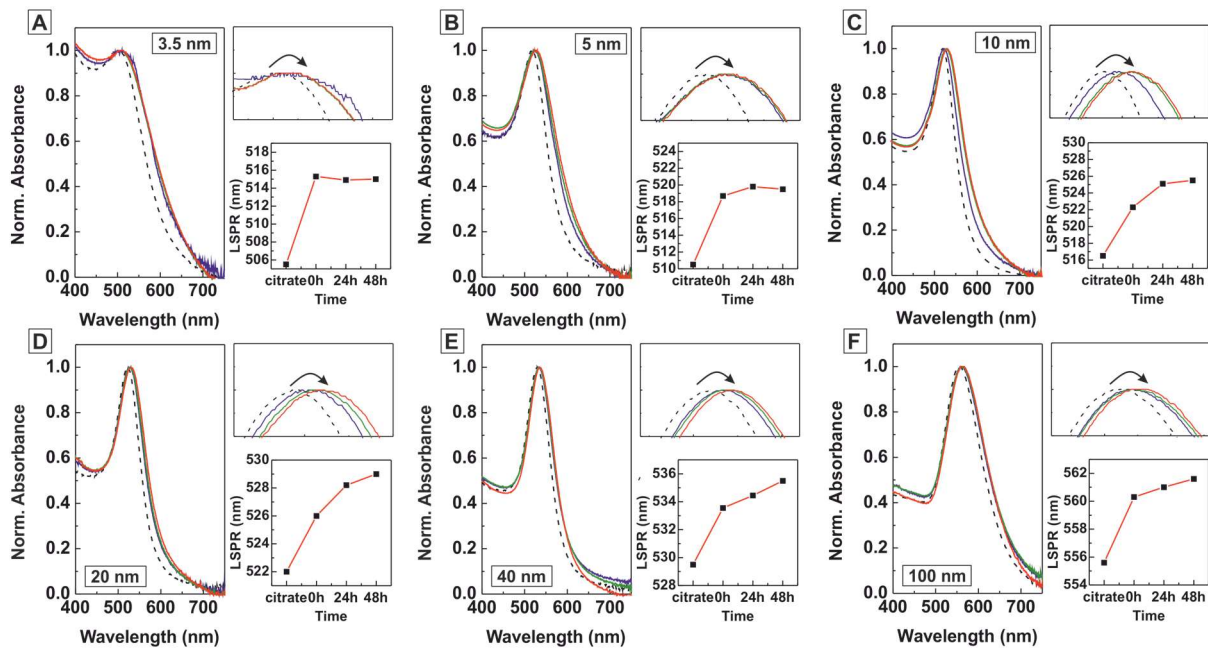
Figure 2. Optimization of exposure conditions. (A) UV-vis spectra of AuNPs in different media. AuNPs aggregate in DMEM, even at low particle concentration, while they remain stable in cCCM. (B) Normalized UV-vis spectra of AuNPs in cCCM at different concentrations. The changes in the LSPR band indicate that AuNPs aggregate in cCCM when they were exposed to the medium at high concentration while their stability was not compromised at low concentration. (C-E) Experimentally measured aggregation parameter of AuNPs of different sizes after their exposure to cCCM at increasing NP concentrations. Results are normalized by particle number (NPs/mL) (C), total particle surface area (cm^2/mL) (D) and total mass of Au atoms ($\mu\text{g}/\text{mL}$) (E). In all cases 1 mL of a solution of AuNPs 10x concentrated with respect to the desired concentration was added to 9 mL of the medium and rapidly mixed for a couple of minutes before being left undisturbed for 1 h under mild stirring.

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Considering these results, we adjusted the concentration of AuNPs, from $\sim 10^{13}$ NPs/mL for the smallest particles (3.5 nm) to $\sim 10^9$ NP/mL for the largest ones (150 nm), to be low enough to promote the formation of a protein layer before the characteristic time of destabilization. It is worth mentioning that these studies show how the exposure of NP to physiological media is concentration dependent and how dispersion of highly concentrated NP samples in these media (as i.v. injection or high NP dose toxicology studies) may be challenging.

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Protein adsorption profile is a kinetic process in which a soft protein corona composed of loosely bound proteins in equilibrium with the free proteins in solution initially forms and evolves slowly toward an irreversible hard protein corona consisting of tightly bound proteins that do not readily desorb.²⁴ In this process, the proteins adsorb/desorb and rearrange at the surface of the particle, leading to the hardening of the corona, a process that determines the final identity of the particle. In this stable (hard) conformation of the protein corona, crowding/cooperative effects between attached proteins are a key factor: once the protein layer is formed, the incorporation of a new protein implies



185

186 **Figure 3. UV-vis characterization of purified AuNPs of selected sizes after their exposure to cCCM for**
 187 **different times: (A) 3.5 nm, (B) 5 nm, (C) 10 nm, (D) 20 nm, (E) 40 nm, (F) 100 nm.** UV-Vis spectra before
 188 (dotted-line) and after their exposure at different times (blue $t < 30$ min, green $t = 24$ h, red $t = 48$ h).

189 the rearrangement of many, which is translated into a highly stable PC. This process takes about 48 h
 190 in the case of AuNPs that end up having a permanent PC that does not detach from the surface when
 191 NPs are redispersed in protein-free media.²⁷ To obtain a PC hardening profile, the NP has to be
 192 purified from the exposure media to remove the unbound and loosely bound proteins, which is indeed
 193 a critical step. Purification of the AuNPs was performed by centrifugation followed by subsequent
 194 dispersion in protein-free aqueous solution after a proper optimization of the process. Additionally,
 195 the number of purification steps was determined to be at least 2. Otherwise the presence of unbounded
 196 free proteins was directly translated into bimodal size distributions in dynamic light scattering (DLS)
 197 measurements, in particular for the small nanoparticles (more details can be found in the SI).

198 The size-dependent evolution of the hardening of the protein corona was analyzed by UV-vis
 199 spectroscopy, DLS, and zeta potential measurements of purified samples. This combination of
 200 characterization techniques provides a remarkably robust analysis of the interaction between the
 201 proteins and the NPs in the colloidal state,²⁷ bypassing complex postprocessing steps and/or avoiding
 202 chemical labeling strategies, such as luminescent or radioactive labelling, usually needed for protein
 203 detection. As observed in **Table 1**, important changes in the physicochemical properties of the
 204 purified AuNPs after exposure to cCCM are identified. The LSPR band of the AuNPs systematically
 205 red-shifts which can be ascribed to the change of the refractive index at the vicinity of the particles⁵⁶,
 206 and it is consistent with the spontaneous binding of media proteins.^{27, 51} The shift is clearly observed
 207 within minutes and it takes 24-48 h to reach a stationary value, indicating a dynamic process of
 208 hardening of the protein corona, reaching a constant value faster for small particles than for the large

209 ones (**Fig. 3**). The extent of the red-shift also depends on particle size being $\Delta\lambda \approx 9$ nm for the
 210 smallest AuNPs (3.5 nm) and $\Delta\lambda \approx 6$ nm for the largest ones (150 nm). However, this cannot be
 211 directly correlated with the thickness and density of the protein layer since particles of different sizes
 212 present different sensitivity toward similar modifications of the surrounding environment (thickness,
 213 refractive index and chemical anchoring).⁵⁶ The absence of absorbance peaks at longer wavelengths is
 214 further evidence that under optimized conditions the AuNPs remain stable and well-dispersed in
 215 cCCM for all the sizes under study.⁵⁴

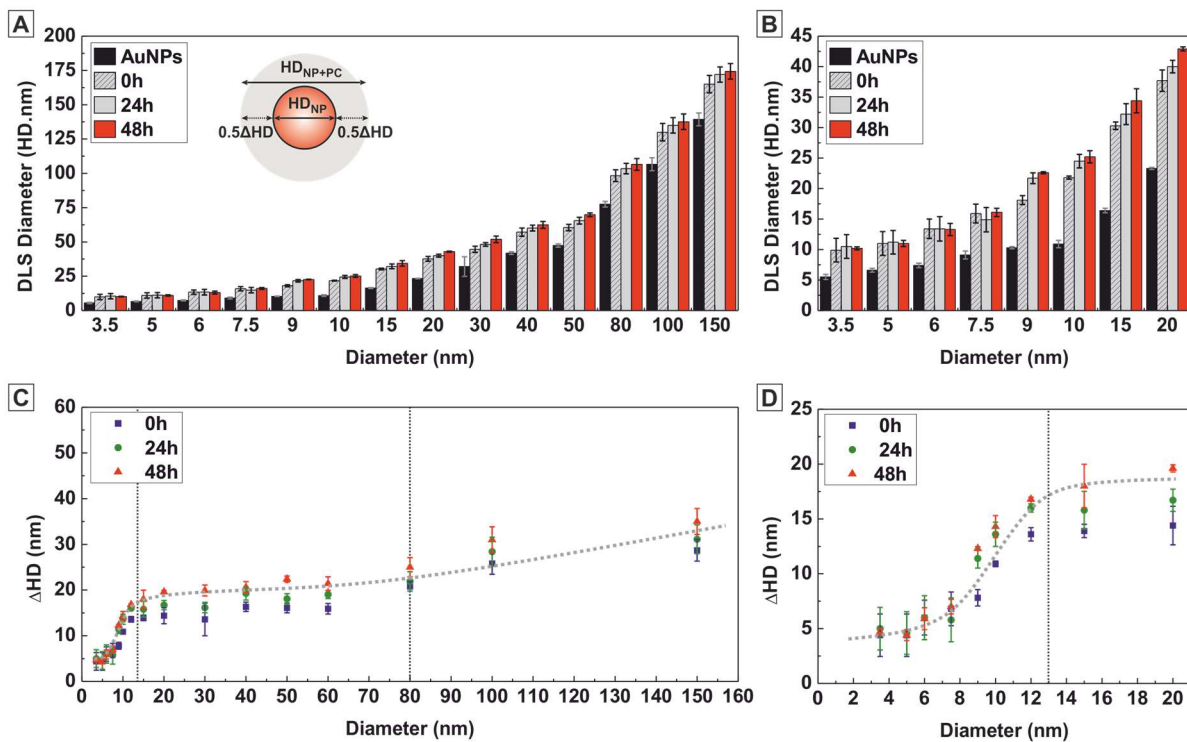
216 **Table 1. Physicochemical Characterization of AuNPs before and after Exposure to cCCM for 48 h and**
 217 **Further Purified.**

target Size (nm)	citrate				cCCM		
	diameter TEM (nm)	diameter DLS (nm)	ζ -Potential (mV)	LSPR maximum (nm)	diameter DLS (nm)	ζ -Potential (mV)	LSPR maximum (nm)
3.5	3.6 ± 0.4	5.5 ± 1.2	-36.8 ± 8.7	505.5	10.2 ± 3.9	-32.8 ± 5.1	515.0
5	5.1 ± 0.8	6.6 ± 1.4	-37.2 ± 14.3	510.5	11.0 ± 3.6	-31.0 ± 7.8	519.5
6	6.2 ± 0.6	7.4 ± 1.6	-39.6 ± 11.1	513.0	13.3 ± 3.2	-26.1 ± 4.3	522.5
7.5	7.8 ± 0.8	9.1 ± 1.9	-42.6 ± 8.6	515.0	16.1 ± 4.3	-24.5 ± 6.2	523.5
9	9.3 ± 1.0	10.3 ± 2.2	-43.6 ± 19.8	516.0	22.6 ± 4.1	-22.5 ± 5.1	524.0
10	10.9 ± 1.0	10.9 ± 3.4	-45.8 ± 14.6	516.5	25.2 ± 5.3	-16.7 ± 5.3	525.5
12	12.0 ± 1.1	12.9 ± 2.7	-42.9 ± 16.4	516.5	29.7 ± 6.9	-16.5 ± 4.8	524.0
15	15.3 ± 1.2	16.4 ± 3.5	-42.9 ± 17.3	518.5	34.4 ± 8.3	-17.1 ± 4.6	527.0
20	19.7 ± 1.5	23.3 ± 4.8	-46.9 ± 15.6	522.0	42.9 ± 9.3	-18.9 ± 6.5	529.0
30	27.5 ± 2.3	32.0 ± 6.2	-48.3 ± 22.8	526.0	51.9 ± 11.9	-20.7 ± 5.9	532.0
40	42.0 ± 3.7	41.8 ± 8.2	-45.7 ± 23.4	529.5	62.4 ± 13.0	-20.9 ± 6.8	535.5
50	47.1 ± 4.5	47.4 ± 11.8	-45.7 ± 17.8	533.0	69.8 ± 15.2	-19.9 ± 7.4	539.0
60	58.3 ± 4.6	55.8 ± 14.6	-42.8 ± 15.6	536.5	77.2 ± 19.3	-25.1 ± 6.6	542.5
80	76.8 ± 6.4	77.5 ± 16.6	-50.2 ± 22.8	546.5	106.5 ± 29.1	-24.2 ± 6.6	553.0
100	110.6 ± 8.5	106.6 ± 34.1	-44.2 ± 14.5	556.5	137.6 ± 43.4	-20.5 ± 8.2	561.3
150	149.1 ± 11.9	139.3 ± 40.8	-44.1 ± 15.2	586.0	174.3 ± 57.5	-21.4 ± 7.4	597.5

218 DLS was used for an in-depth study of the PC and to determine variations in the hydrodynamic
 219 diameter (HD) of the samples. Obtained results are summarized in **Table 1** and **Figure 4**, where it can
 220 be seen how the HD of the AuNPs systematically increases after their exposure to cCCM, which is
 221 associated with the direct adsorption of the proteins onto the surface of the NPs (**Fig. 4-A, B**). This
 222 increase strongly depends on the NP size and can be directly correlated with the thickness of the PC
 223 (thickness = 1/2 Δ HD) (**Fig. 4-C, D**). Thus, for small 3.5 nm AuNPs a protein layer of ~2.5 nm in
 224 thickness is observed (Δ HD ~5 nm). As the size of the particles progressively increases the protein
 225 coating becomes thicker, reaching a rather constant value of 8-10 nm (Δ HD from 16 to 20 nm) for
 226 AuNPs from 12 to 80 nm. Finally, for particles larger than 80 nm, the thickness of the corona seems
 227 to be even larger, obtaining an increase of 15-16 nm in thickness (Δ HD from 30 to 32 nm).

228 Considering the results obtained earlier and assuming that albumin (65 kDa, HD ~6-7 nm) and other
 229 high molecular mass (>65 kDa) proteins¹³ have been extensively found as the major constituents of
 230 the PC, it is reasonable to identify three different regimes: (i) small size regime, where the particles

231 are smaller than mentioned proteins, and the thickness of the PC is thinner than that of a single layer
 232 of proteins, (ii) medium size regime, where the size of the particles starts to be significantly larger
 233 than of the proteins, and where the obtained thickness of the PC supports the overall formation of a
 234 single layer of proteins and (iii) large size regime, where the size of the particles is much larger than
 235 that of proteins, approaching a flat surface, and where the obtained PC thickness suggests the
 236 formation of a multilayer of proteins. It is worth noting that despite the PC not being completely
 237 developed in the small AuNP regime; it is robust and provides enough stability to prevent them from
 238 aggregation.

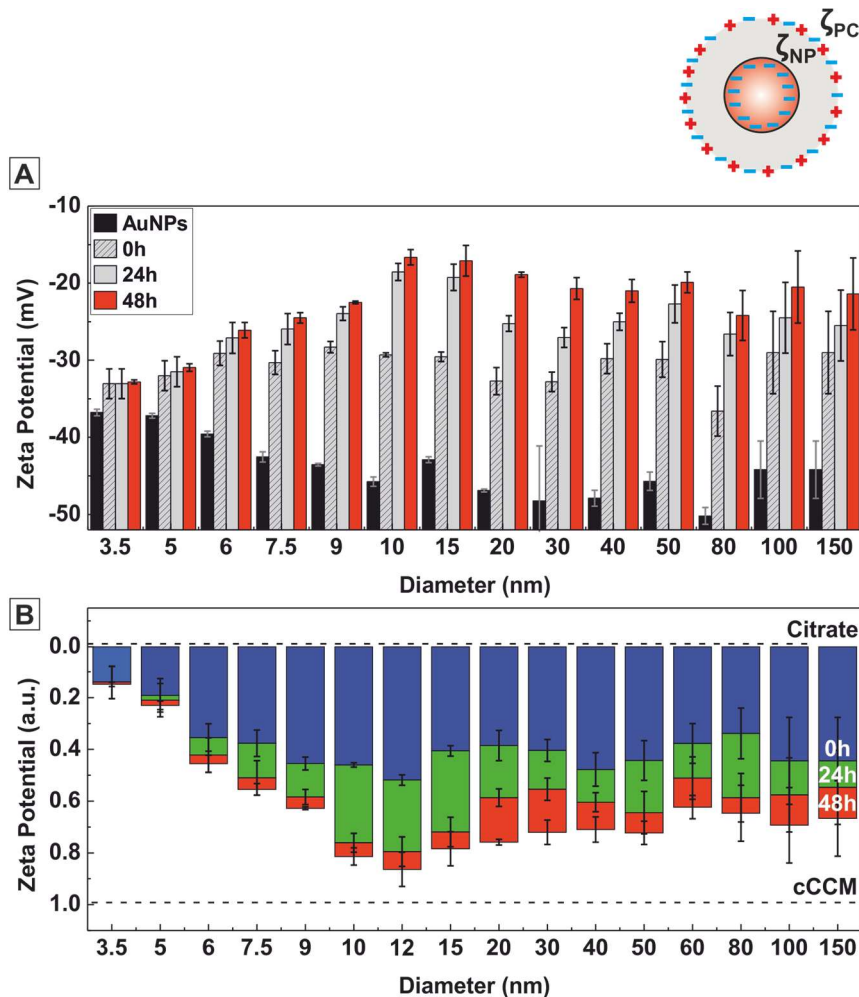


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 240 **Figure 4. Hydrodynamic diameter (HD) measured by dynamic light scattering of purified AuNPs of**
 241 **different sizes after their exposure to cCCM for different times.** (A) HD obtained by DLS for particles
 242 between 3.5 and 150 nm and (B) detailed for particles between 3.5 and 20 nm. (C, D) Respective increases in the
 243 HD showing doubling of the corona thickness. Straight lines indicate different tendencies in the resultant protein
 244 corona.

245 Similar evidence was obtained by zeta potential measurements used to assess the changes in particle
 246 surface charge after exposure to cCCM. Thus, concomitant with the observed red-shift of the LSPR
 247 position and the increase of the NP size measured by DLS, a drop in absolute surface charge of the
 248 particles toward the average value of the medium proteins was systematically observed. It accounts
 249 for the progressive screening of the negatively charged citrate-metal surface by the coating protein
 250 layer, which provides a long-term colloidal stability to the AuNPs in biological media. Interestingly,
 251 here again, zeta potential results depend on AuNP size (Fig. 5A), changing from -40 mV (citrate) to -
 252 30 mV for the smallest particles, and to -(15-20) mV for larger particles after protein coating. These
 253 results supports the previous idea that proteins are unable to form a dense layer on the surface of very

254 small AuNPs, while a proper protein corona is formed in the large ones, either by a single dense
 255 protein layer or a multilayered coating. Since the surface charge itself is not high enough to prevent
 256 NPs from electrostatic aggregation (an absolute value of 30 mV is usually considered a limit for
 257 particle stability), these results are further evidence that the stability of the NPs is mediated by steric
 258 repulsion provided by proteins (which stay in their native state²⁷) rather than via the former
 259 electrostatic repulsion provided by citrate.

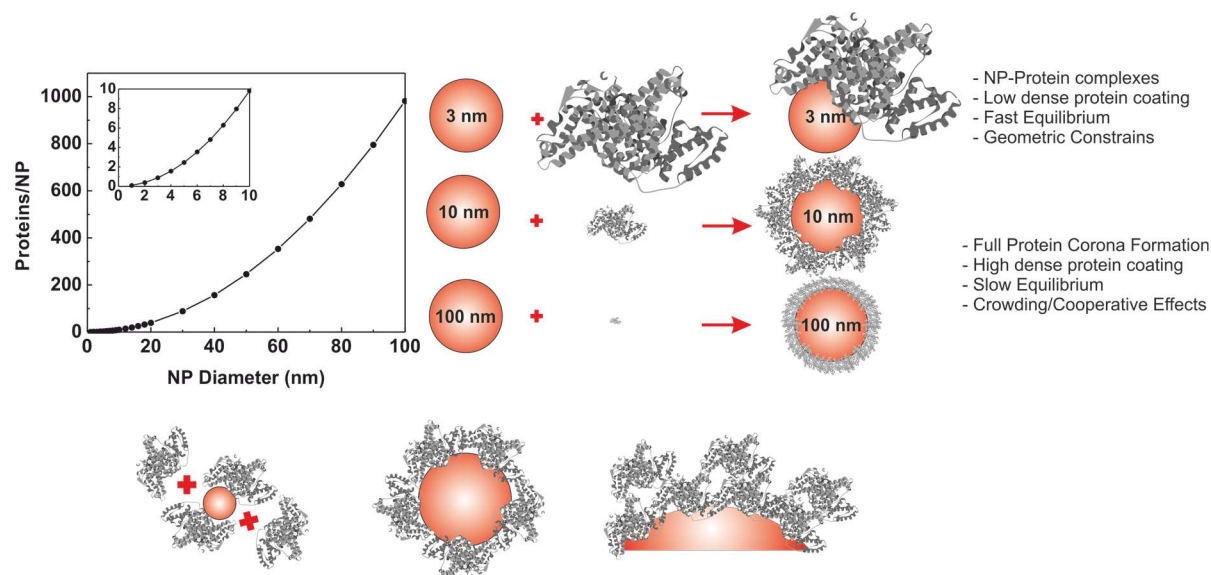
260 As previously observed by UV-vis spectroscopy, time-dependent DLS and zeta potential results
 261 confirm the kinetic evolution of a hard protein corona and its dependence on particle size (**Fig. 4** and
 262 **Fig. 5**), being faster (24 h) for smaller sizes than for larger ones (48 h). These results suggest that
 263 particle size plays a critical role in both the kinetic and the thickness of the protein layer: the smaller
 264 the particle the faster the kinetics evolution and the thinner (incomplete) the protein layer.



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 266 **Figure 5. Zeta potential of purified AuNPs of different sizes after their exposure to cCCM for different**
 267 **times.** (A) Absolute values of zeta potential measurements and (B) relative values in which 0 represents the
 268 surface charge of the citrate coated particles and 1 the charge of the proteins. The size-dependent protein corona
 269 evolution is clearly observed; where small particles present the most negative charge after exposure as a
 270 consequence of not forming a dense layer. For large particles, zeta potential is closer to that of the proteins
 271 indicating the formation of a denser layer.

272 Based on the obtained results, there seems to be a size limit below which a complete PC cannot fully
 273 develop. Presumably, this can be related to geometric constraints or other effects (crowding,
 274 cooperative effects) that could be a subject of further studies. As described by Nygren and
 275 Alaeddine,⁵⁷ and contrary to what might be expected, proteins do not distribute on surfaces randomly.
 276 Instead, once the first proteins are attached they serve as nucleation centers for the deposition on new
 277 ones, thereby stabilizing them, and this mechanism is repeated until the entire surface is filled. These
 278 and other attempts to explain the irreversibility of this process seem to have in common that the initial
 279 attachment of a protein to a surface is followed by a series of movements and/or rearrangements to
 280 make the interaction more stable, ultimately resulting in an irreversible attachment. These processes
 281 appear to be restricted when the particles become very small and their surface is limited to very few
 282 proteins per NP (**Fig. 6**). On the other hand, the formation of a hard protein corona composed of more
 283 than one layer of proteins is a plausible explanation for larger particles. In this case, when
 284 approaching flat surfaces, proteins are subjected to larger conformational changes upon adsorption
 285 leading to multiple protein adsorptions. Thus, the increase in the corona thickness of the larger
 286 particles with respect to the smaller particles could be related to this effect along with the appearance
 287 of defects on the first protein layer docking other proteins (**Fig. 6**). However, it remains unknown
 288 whether particle size determines the nature of the PC only quantitatively (the number of proteins
 289 attached) or also in a qualitatively way (the composition of proteins).¹⁵

290 In order to see whether and how obtained results can be extrapolated to other systems we studied the
 291 PC formation in the presence of pure albumin. Considering the high abundance of albumin in serum



292
 293 **Figure 6. Representative scheme of the formation process of a hard protein corona in citrate-stabilized**
 294 **AuNPs of different sizes.** (i) NP-protein complexes (or incomplete coating), (ii) the formation of a single dense
 295 protein corona layer and (iii) the formation of a multilayer corona. Graphic on the left details the number of
 296 proteins that can attach on the surface of the NP as a single monolayer. Calculations has been performed based
 297 on the size of BSA chosen as model protein (see SI).

298 (represents ~60% of the total mass serum proteins), its high affinity for many different surfaces, and
 299 low specificity, albumin is known to be one of the main components of the protein corona when
 300 AuNPs, and other NPs, are exposed to serum or blood plasma.^{15, 25} Indeed, this high affinity of
 301 albumin for gold surfaces has been widely proven, resulting in the spontaneous formation of a dense
 302 and tightly bound albumin-rich coating in a process usually known as albuminization.^{41, 49, 51, 58, 59}
 303 With the aim to elucidate whether the albuminization of the particles showed similar size dependent
 304 effects to those observed in the PC formation by serum proteins, analogous experiments were
 305 performed by exposing the particles to BSA in phosphate buffer solution. The obtained results are
 306 summarized in **Table 2**, confirming the formation of an irreversible BSA coating layer onto all the
 307 citrate-stabilized AuNPs after 48 h of exposure. BSA adsorption did not induce colloidal aggregation
 308 of the NPs, comparable to that given by FBS serum proteins in cCCM. This further indicates that the
 309 BSA preserves their native state and do not undergo denaturalization. The obtained results are in
 310 general agreement with those observed in the case of cCCM suggesting that BSA is unable to form a
 311 proper corona for the very small particles while a single layer or multilayer is formed on larger ones.
 312 Slight differences observed between FBS and BSA may account for the presence of other proteins in
 313 combination with BSA. Briefly, the differences in the shift of the LSPR position (smaller in the case
 314 of BSA than FBS) along with the variations of the HD observed by DLS (smaller in the case of BSA
 315 than FBS) suggest that the presence of pure BSA may lead to the formation of a more structured, and
 316 consequently thinner, layer.

317 **Table 2. Physicochemical Characterization of AuNPs before and after Exposure to BSA for 48 h and**
 318 **Further Purified.**

Target Size (nm)	BSA		
	Diameter	ζ -Potential (mV)	LSPR
	DLS (nm)		maximum (nm)
3.5	9.6 ± 1.6	-30.7 ± 7.3	513.5
5	10.6 ± 1.7	-29.5 ± 6.8	518.0
6	12.6 ± 2.5	-27.7 ± 4.5	520.0
7.5	14.9 ± 3.4	-23.5 ± 6.1	522.5
9	19.8 ± 4.4	-23.5 ± 4.7	522.5
10	22.2 ± 5.1	-18.5 ± 4.4	523.0
12	22.3 ± 5.2	-20.8 ± 5.2	522.5
15	29.6 ± 6.7	-21.9 ± 5.1	524.5
20	34.8 ± 7.8	-19.7 ± 6.8	527.5
30	44.4 ± 10.8	-21.8 ± 7.4	531.0
40	56.1 ± 11.0	-25.1 ± 6.6	534.0
50	60.9 ± 13.2	-22.2 ± 5.3	536.5
60	69.5 ± 19.3	-23.4 ± 5.9	540.5
80	98.3 ± 19.1	-25.2 ± 6.1	550.5
100	135.6 ± 35.4	-22.5 ± 7.7	560.5
150	159.1 ± 53.2	-24.6 ± 7.6	594.0

319

320 **Conclusions**

321 The optimization of exposure conditions (stability and purification) allowed us the study of the
322 protein corona in citrate-stabilized AuNPs of different sizes. The proper combination of UV-vis
323 spectroscopy, DLS, and zeta potential measurements has been proven as a potential tool to describe
324 qualitatively the protein corona formation on the surface of AuNPs. These techniques need to be
325 discussed together in order to avoid misleading interpretations. For example, the largest red-shift in
326 the LSPR, measured by UV-vis is observed for the smallest particles, while DLS measurements show
327 that these small particles form the thinnest corona. On the other hand, UV-vis spectroscopy and zeta
328 potential do not predict the formation of a multilayer protein coating while DLS allows
329 characterization of this process. Finally, all three techniques are effective in monitoring kinetic protein
330 adsorption profiles. As a result we observed how both the kinetics of protein corona hardening and the
331 characteristics of the protein corona are particle size-dependent, despite the question of debate
332 remaining whether particle size determines the PC not only quantitatively but also in a qualitatively
333 way.

334 The obtained results present relevant biological consequences as different NP–biological interactions
335 take place at different time scales. Thus, tightly associated proteins may stay adherent to the particle
336 when the particle is endocytosed from the extracellular fluid to an intracellular location, whereas
337 proteins with a fast exchange rate are replaced by intracellular proteins during or after such a transfer.
338 As a result, the same NP can give different biological responses depending on the portal of entry,
339 history and preincubation in serum among others, thereby illustrating the importance of the study of
340 size-dependences of the protein corona in a biological environment.

341 **Experimental**

342 **Chemicals.** Sodium citrate tribasic dihydrate ($\geq 99\%$) and gold (III) chloride trihydrate $\text{HAuCl}_4 \cdot$
343 $3\text{H}_2\text{O}$ (99.9% purity) were purchased from Sigma-Aldrich. Fetal Bovine Serum, FBS (research grade,
344 sterile filtered) and Dulbecco's Modified Eagle Medium, DMEM (with 1000 mg/mL glucose and
345 sodium bicarbonate, whiteout L-glutamine, sodium pyruvate, and phenol red, liquid, sterile-filtered),
346 were purchased from Sigma. All reagents were used as received without further purification and all
347 glass material was sterilized and dehydrogenated in an oven prior to use. Milli-Q water was used in
348 the preparation of all solutions.

349 **Nanoparticle Synthesis.** Aqueous solutions of citrate-stabilized AuNPs with different sizes (3.5-150
350 nm) were synthesized according to the previously developed seeded-growth method in our group.
351 Detailed synthetic procedure and full characterization of the resultant solutions can be found in the
352 respective articles.^{47, 48} Briefly, AuNPs with increasing sizes were obtained from an initial AuNP
353 solution after different sequential steps of growth; consisting of sample dilution plus further addition
354 of gold precursor. This means that the number of particles in the solution decreases as the size

355 increases. Consequently, the concentration of smallest particles was $\sim 10^{14}$, which decreased down to
356 $\sim 10^{10}$ for the largest 150 nm particles. All the particles were used within 20 days after synthesis. The
357 use of aged particles may result in increased experimental variability, since after a large period of time
358 the stability of the particles may be compromised by the degradation of the surfactant.

359 **Exposure to Cell Culture Media.** Complete cell culture medium (cCCM) consisted of DMEM
360 supplemented with 10% of FBS. AuNPs and cCMM were mixed (1:10 by volume) and placed in an
361 incubator at 37°C for different incubation times. As a result, the final concentration of AuNPs was
362 that of the synthesis divided by the dilution factor: $\sim 10^{13}$ NP/mL for the smallest particles and $\sim 10^9$
363 for the largest particles (approximately 0.05 mM of Au). At these working concentrations, AuNPs
364 were stable throughout the experiments and the number of proteins in larger excess with respect to the
365 available particle surface area. After incubation, the samples were centrifuged between 2000 and
366 35000 g for 10-20 min. (the larger the size, the lower the speed) to remove the excess of unbound or
367 loosely bound proteins on the NP surfaces followed by resuspension in the particle original medium
368 (sodium citrate 2.2 mM). Any possible aggregated fraction resulting from the centrifugation process
369 was removed by filtering the samples with a 200 nm-pore-size cellulose acetate membrane. In the
370 case in which the AuNPs solution needed to be concentrated with respect to the synthesis
371 concentration, such as for stability tests, a regenerated cellulose centrifugal filter (Millipore 10KDa)
372 was used.

373 **Physicochemical Characterization of the NPs and NP-Protein Corona.** AuNPs and the time
374 evolution of their coating by proteins were characterized before and after exposure to cCCM using
375 different techniques. The proper combination of these techniques has been used in other similar
376 studies by our group²⁷ and proven to be reliable when performed with adequate controls. *TEM:*
377 Diameter of the synthesized particles were obtained from analysis of TEM images acquired with a
378 JEOL 10101 electron microscope operating at an accelerating voltage of 80 kV. Samples were
379 prepared by drop-casting 10 microliter of the sample on a carbon-coated copper TEM grid and left to
380 dry at room temperature. At least 500 particles from different regions of the grid were counted. In
381 order to avoid aggregation of the particles during TEM grid preparation they were previously
382 conjugated with 11-mercaptoundecanoic acid. *UV-Visible Spectroscopy:* UV-Visible spectra were
383 acquired with a Shimadzu UV-2400 spectrophotometer. 1 mL of sample was placed in a plastic
384 cuvette, and spectral analysis was performed in the 300 to 800 nm range at room temperature. AuNPs
385 exhibit a characteristic absorbance peak in this region, the so-called localized surface plasmon
386 resonance (LSPR) band. The LSPR position is sensitive to the surrounding of the NPs at the
387 molecular level, and therefore the changes in the close environment of the NPs (such as the protein
388 adsorption) can be investigated using this technique. Water was taken as the reference for all samples.
389 *DLS and zeta potentia measurements:* The hydrodynamic size and surface charge of the AuNPs
390 before and after incubation in cCCM were determined by Dynamic Light Scattering (number mean)

391 and Laser Doppler Anemometry respectively, using a Zetasizer Nano ZS instrument equipped with a
 392 light source wavelength of 532 nm and a fixed scattering angle of 173°. Aliquots (0.8 mL) of the
 393 colloidal NP solutions were placed into specific plastic cuvette and the software was arranged with the
 394 parameters of refractive index and adsorption coefficient of gold, and solvent viscosity of water at 25
 395 °C. Each value was the average of at least 3 independent measurements.

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404 **Supporting Information**

405 Full characterization of the cell culture media by UV-vis spectroscopy, dynamic light scattering and
 406 zeta potential, verification of the excess of proteins compared to the surface area of the nanoparticles
 407 and detailed purification steps of the particles.

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Supporting information for:

Size-dependent Protein-Nanoparticle Interactions in Citrate-Stabilized Gold Nanoparticles: The Emergence of the Protein Corona.

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1. Characterization of complete cell culture media (cCCM)

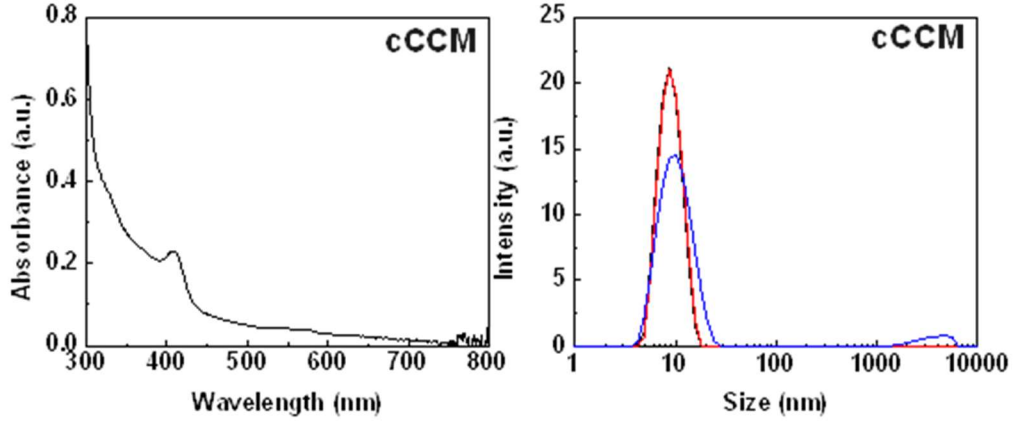


Figure S1. Characterization of complete cell culture media (cCCM): DMEM supplemented with 10% of FBS. UV-vis spectrum (left) shows a strong absorbance in the region below 500 nm, while almost no absorbance is observed above this value, where the Localized Surface Plasmon Resonance peak characteristic of the AuNPs is usually located. Dynamic light scattering plot (DLS) (right) of the cCCM displaying an intense peak in the region between 5 and 15 nm attributed to the proteins. This peak may overlap with the AuNPs peak if they have not been previously purified. Finally, the zeta potential of the cCCM results in -11.2 ± 0.9 mV which is characteristic of the slightly negative charge of most of the proteins.

2. Excess of proteins in cCCM compared to the total surface area of AuNPs

For a specific colloidal solution of AuNPs, the average NP diameter (\bar{D}), average NP surface area (\bar{S}) and average NP mass (\bar{M}) can be calculated from the size distribution obtained by TEM analysis:

$$\bar{D} \left(\frac{cm^2}{NP} \right) = \frac{\sum D_n}{n} \quad (1)$$

$$\bar{S} \left(\frac{cm^2}{NP} \right) = \frac{\pi \sum D_n^2}{4n} \quad (2)$$

$$\bar{M} \left(\frac{mols}{NP} \right) = \bar{V}_{NP} \delta M_W = \frac{4}{3} \frac{\sum (D_n)^3}{n} \delta M_W \quad (3)$$

where n and D_n are the number and diameter of the particles that has been counted by TEM (at least 500 NPs are needed for a representative population), δ is the density of gold, 19.3 g/mL,

and M_w the molecular weight, MW 197. Notice that a common error is to calculate \bar{S} and \bar{M} from \bar{D} (the polydispersity degree of the particles is erroneously omitted in this case). Finally, the concentration of gold (C_{Au}), obtained either from the concentration of precursor used in the synthesis of the particles and the yield of the reaction or by ICP-MS, can be translated to the number of NPs and the surface area in solution as follow:

$$number\ of\ NPs\ \left(\frac{NP}{L}\right) = concentration\ of\ Au\ (M) \cdot \frac{1}{M} \left(\frac{mol}{NP}\right)^{-1} \quad (4)$$

$$surface\ area\ \left(\frac{cm^2}{L}\right) = number\ of\ NP\ \left(\frac{NP}{L}\right) \cdot \bar{S}\ \left(\frac{cm^2}{NP}\right) \quad (5)$$

The total surface area in a standard synthesis procedure range from 2 to 30 cm^2/mL , which corresponds to $\sim 10^9$ NPs/mL for the larger particles (150 nm) and to $\sim 10^{13}$ for the smallest ones (3.5 nm).

On the other hand, the total number of proteins in cCCM is at least three orders of magnitude larger than that of the particles in solution. The total surface area they can cover upon adsorption on a surface is the result of multiplying their cross section (CS) for their concentration (C):

$$area_{proteins}\ \left(\frac{cm^2}{L}\right) = \sum(C_{protein} \cdot CS_{protein}) \quad (6)$$

For example, the triangular CS of the BSA in the folded conformation has been previously estimated to be $32\ nm^2 = 32 \cdot 10^{-14}\ cm^2$ while their concentration in the cCMM is around $6 \cdot 10^{16}$ (0.1 mM). This represents enough BSA in the cCCM to cover a total area of 19300 (cm^2/mL); at least more than $5 \cdot 10^2$ times higher than that needed to fully surround the AuNPs surface in solution. Thus, it is reasonable to consider the number of proteins in the cCCM largely in excess with respect to the AuNPs, even if more than one layer of proteins is adsorbed on the surface of the particles, and the fraction depleted from the medium upon adsorption insignificant.

3. Number of washing steps for a proper DLS characterization

cCCM free of NPs gives a background peak in the range of 5-15 nm by DLS (see section 1), which is comparable to the hydrodynamic size of the small AuNPs used in the experiments here presented. Consequently, interferences from the unbounded proteins in solution when measuring the hydrodynamic size of the AuNPs by DLS is something that should not be ruled out if the particles has not been previously washed and dispersed in a solution free of proteins.

The interferences that free proteins in solution may induce to the DLS measurements¹ can be estimated from their respective abilities to scatter light assuming Rayleigh scattering properties, in which scattering intensity of a specific molecule is proportional to its number in solution (N) and its size raised to the six power (D^6):

$$I \propto \varepsilon N D^6 \quad (7)$$

where ε is the refractive index for each molecule. The relative contributions of the unbounded proteins with respect to the AuNPs can be then quantified by defining a dimensionless parameter (γ) as follow:

$$\gamma = \frac{I_{NP}}{I_p} = \frac{\varepsilon_{NP} N_{NP} D_{NP}^6}{\sum \varepsilon_P N_P D_P^6} = \frac{0.47 N_{NP} D_{NP}^6}{\sum \varepsilon_P N_P D_P^6} \quad (8)$$

where the refractive index for AuNP is $\varepsilon_{NP} = 0.47$, while the ε_P is ~ 1.5 for proteins. Previous studies with BSA have demonstrated that γ should be at least > 0.1 to avoid the contributions of free proteins in the DLS measurements.¹

In this regard, rough calculations based on the approximate BSA concentration in cCCM ($6 \cdot 10^{16}$ protein/mL, HD 7 nm and $\varepsilon_P = 1.447$), demonstrated that $\gamma \ll 0.1$ for AuNPs of 3.6 nm in the range of concentration here working (10^{13} NP/mL) and at least 2 cycles of washing were needed to avoid the contribution of unbound proteins (note that each washing step consist on replacing 980 uL of supernatant from a 1000 uL of solution decreasing the concentration of proteins in

solution by a factor of 50x). Indeed, based on the previous approximation the concentration of these particles should be in the range of $1 \cdot 10^{18}$ to be properly measured in cCCM without requiring any washing step. On the other hand, γ already approach to that of 0.1 for AuNP of 60 nm at any concentration larger than $4 \cdot 10^{10}$, which is in the range of working concentration in our experiments. In agreement with that, DLS plot showed monomodal curves even in the absence of any washing steps for large AuNPs (Figure S2-S3).

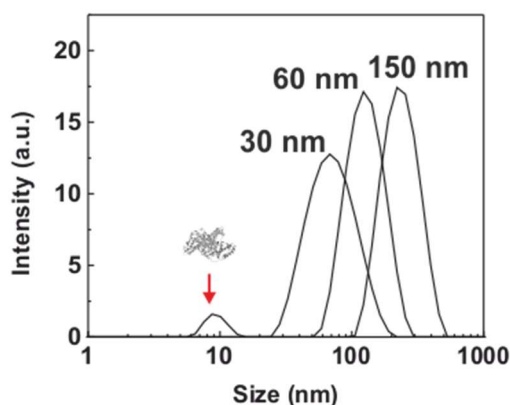


Figure S2. DLS plot of AuNPs measured in cCCM without purification of the particles. Particles of 60 nm and 150 nm show a monomodal curve, while particle of 30 nm, and below, present a bimodal curve due to interferences from unbound proteins in the medium.

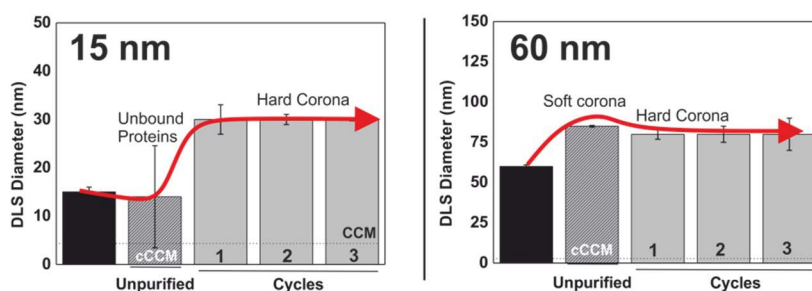


Figure S3. Hydrodynamic diameter measured by DLS. For 15 nm particles the measured diameter decreases after incubation in cCCM as a result of contributions from unbound proteins. For 60 nm particles unbound proteins in cCCM does not significantly contribute to the measured values.

4. Maximum number of proteins per NP considering a monolayer coating

The theoretical available number of binding sites (NBS) for a monolayer coverage of proteins (considering BSA as a model protein) on a NP with a diameter (D) is estimated by dividing the surface area of the NP (πD^2) by the triangular cross section of the folded BSA (32 nm^2):²

$$\frac{BSA}{NP} = \frac{CS_{BSA}}{\pi D^2} = \frac{32}{\pi D^2} NBS = \frac{\pi D^2}{CS_{BSA}} = \frac{\pi D^2}{32} \quad (9)$$

References

- (1) Tsai, D.-H., DelRio, F. W., Keene, A. M., Tyner, K. M., MacCuspie, R. I., Cho, T. J., Zachariah, M. R., and Hackley, V. A. (2011) Adsorption and Conformation of Serum Albumin Protein on Gold Nanoparticles Investigated Using Dimensional Measurements and in Situ Spectroscopic Methods. *Langmuir* 27, 2464-2477.
- (2) Dominguez-Medina, S., McDonough, S., Swanglap, P., Landes, C. F., and Link, S. (2012) In Situ Measurement of Bovine Serum Albumin Interaction with Gold Nanospheres. *Langmuir* 28, 9131-9139.