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Effect of Patient Specific Blood Biomarkers on Nanoparticle - Cell Interactions

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EFFECT OF PATIENT SPECIFIC BLOOD BIOMARKERS ON NANOPARTICLE – CELL
INTERACTIONS

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To my family, friends, mentors, lab mates and school colleagues who supported and encouraged me throughout my whole academic career. Thank you for teaching me that there is not a one and only definition to success nor a limit that can't be reached.

EFFECT OF PATIENT SPECIFIC BLOOD BIOMARKERS ON NANOPARTICLE – CELL
INTERACTION

by

VERONICA GABRIELA CONTRERAS GUERRERO, B.S.

THESIS

Presented to the Faculty of the Graduate School of

The University of Texas at El Paso

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of the Requirements

for the Degree of

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Department of Metallurgical, Materials and Biomedical Engineering

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Abstract

Nanoparticles are currently known to be a promising material class for bio-applications in drug delivery and vaccine development. Using gold nanoparticles of varied sizes, in this case 45 and 100 nanometers as a model nanomaterial system, we investigated how patients' blood physiology and chemistry (such as solute, protein, lipid levels) affect the biological response to bionanomaterials. When nanoparticles are injected into the body, biomolecules in the blood adsorb to the nanoparticle's surface to form a biomolecular corona that is specific to the patient's unique blood composition. This biomolecular corona is important because it affects the in vivo fate and biodistribution of the administered nanoparticles. Here we show that the abundance of certain proteins and lipids in patient blood changes the biomolecular corona of nanoparticles of the varied sizes. Furthermore, the presence of these proteins and lipids affected in vitro cell uptake of nanoparticles. Our preliminary results suggest that certain blood composition biomarkers can predict patient-specific biomaterial interaction. Future experiments will follow up on these candidate protein and lipid biomarkers and investigate if they are applicable in predicting nanoparticle interactions of different physicochemical properties, such as lipid-based nanoparticles. Studying how patient-specific blood biomarkers interact with biomaterials can help us rationally design personalized nanomaterials for therapeutic applications to improve their health.

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Chapter 1: Background & Introduction

The United States is facing an increase of overweight and obese populations. As of 2024, the US Center of Disease Control has estimated that over 70% of the population of ages 20 and older may be facing obesity and with it a development of conditions that negatively affect a patient's health (Figure 1.1). As medical advancements have improved and grown overtime, better and more precise methods for drug delivery have been tested in hopes of making personalized medical advancements for patient treatments. Research is progressing to investigate the correlation between drug delivery efficiency and uptake with respect to differences in body physiology. More specifically, we are interested in studying the differences in nanoparticle uptake between healthy individuals and patients with obesity. To date, lipid-based nanoparticles (LNPs) have become the focal point of study for drug and gene delivery research due to their ability to deliver therapeutic nucleic acids. However, it has been found that LNPs face certain challenges in terms of efficacy and targeting within the body. To address this, studies are exploring new ways to enhance LNPs targeting while ensuring safe cargo delivery. Interestingly, a difference in uptake and biodistribution have been observed based on patient-specific physiological factors. This is directly related to the environment that the nanoparticles travel through in the body, and the resulting creation of their protein corona. This phenomenon is attributed to nanoparticles interacting with biological fluids, which cause different proteins to coat the nanoparticles.

For this thesis, we are developing a model that studies the relationship between protein coronas of specific blood biomarkers on gold nanoparticles (AuNPs) and how their uptake differs under each healthy and patients' physiological conditions. Although gold nanoparticles are unlikely to be clinically translatable since they are inorganic nanomaterials, they have several major advantages that make them ideal as tools to study nanomaterial-biological interactions. With the use of AuNPs, we are able to have direct control on the size of the nanoparticles synthesized, and its' uniformity throughout. AuNPs are also known to be economical to synthesize and ease of surface modification by thiol-based

conjugation. AuNPs are also amenable to characterization by Transmission Electron Microscopy (TEM) and UV-Visible Spectroscopy (UV-VIS), and quantification by inductively-coupled plasma mass spectrometry (ICP-MS). The long-term goal of this study is to enable the engineering of personalized nanoparticles for drug and gene delivery by tailoring nanoparticle design to each patient's physiology for optimized uptake and efficacy of their treatment. (Figure 1.2) [1][2][3][4] As a first step towards this goal, we used two different serum conditions to investigate differences in nanoparticle protein corona formation and subsequent cell uptake. We hypothesized that nanoparticle cell uptake will depend on patients' physiological differences as represented with the two serum conditions, because the different protein compositions in the serum will lead to the formation of different nanoparticle protein coronas.

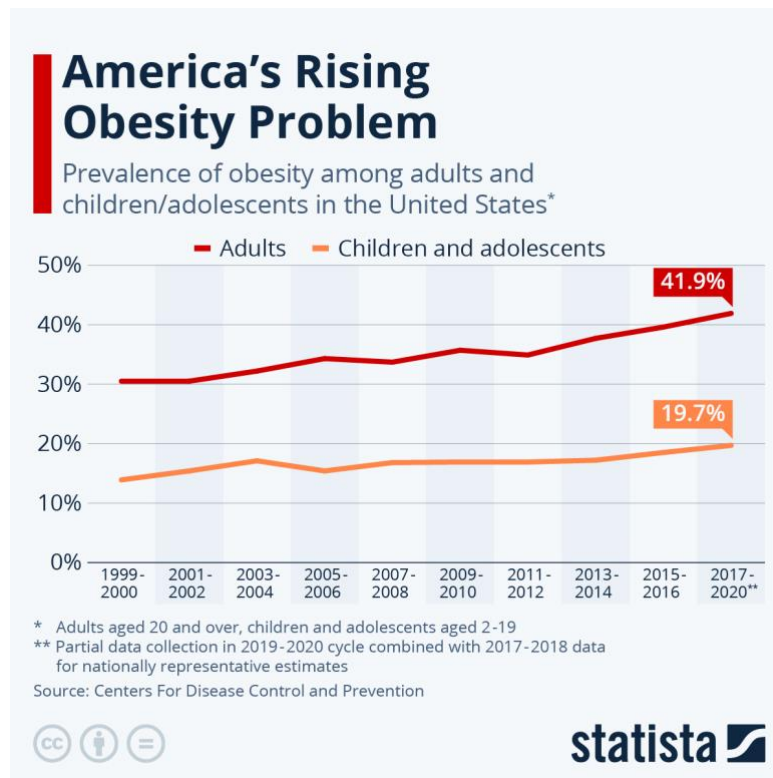


Figure 1.1: Statistics on America's obesity growth [5]

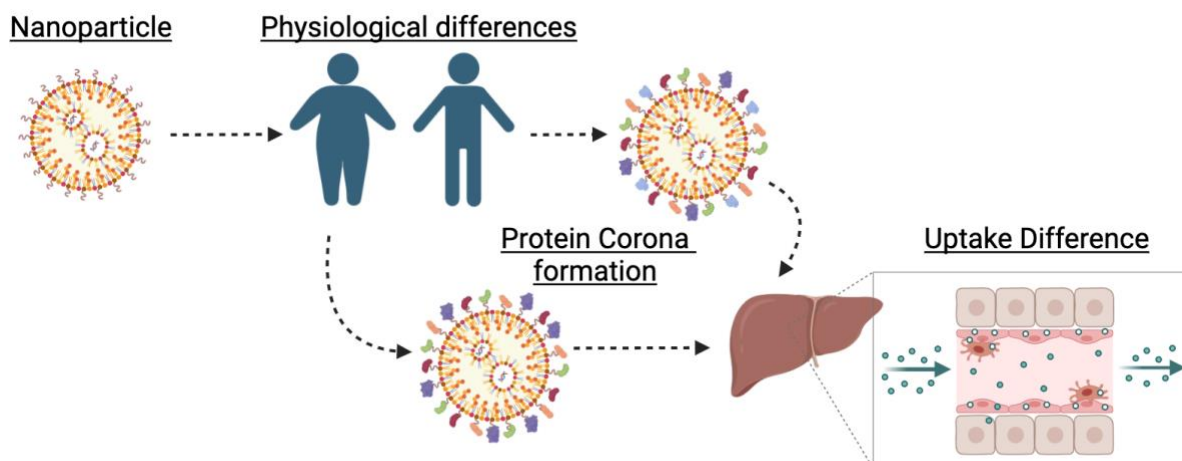


Figure 1.2: Nanoparticle bio-applications and the effects in uptake due to physiological difference (Made in Bio render)

1.1 MATERIAL SELECTION & PROPERTIES

To first understand the difference of surface treatments that can be subjected onto nanomaterials, it is important to understand the materials paradigm, known as processing-structure-properties-performance (Figure 1.3). [6] These parameters identify the goals, means with cause and effect of a material, and its systematic design. The process will be the work and conversion of raw material into its' final stage. In this case, the process of the material will be the synthesizing of the gold into AuNPs. Structure of the material relates to the association of arrangement of its fundamental components. Relating to this study, this refers to the size and surface treatment of the AuNPs. Moving forward with the properties, this refers to any specific characteristics that are given by the material that will at the end be what identifies the performance. In this case, the properties are the surface preparation of the AuNPs which will be treated with polymer chains that will give resistance to adherence to proteins, cells and prevent agglomeration. Finally, the material's performance is mainly the qualities that the material meets for its' intended design and purpose given. With this, the behavior it will have due to all the other

components together, in this case the interaction of the cell uptake with the AuNPs when the surface is treated.

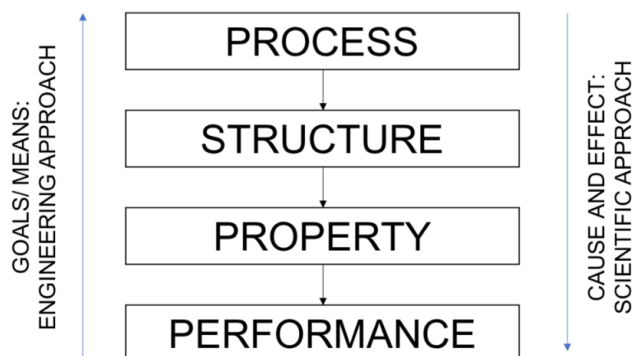


Figure 1.3: Material's paradigm schematic [7]

1.2 EXPERIMENTAL PROCEDURE

To accomplish the objectives of this project, the main approach and experimental steps are outlined in Figure 1.4. To start, the process involves the synthesis of AuNPs and determining the specific sizes, processes, and surface preparations that will be added to the material for the next steps. Following this is the material characterization of the synthesized AuNPs. Next, serum work is performed to study the protein corona formation process. Subsequently, the cell uptake of the AuNPs under various conditions is examined and correlated to all results obtained.

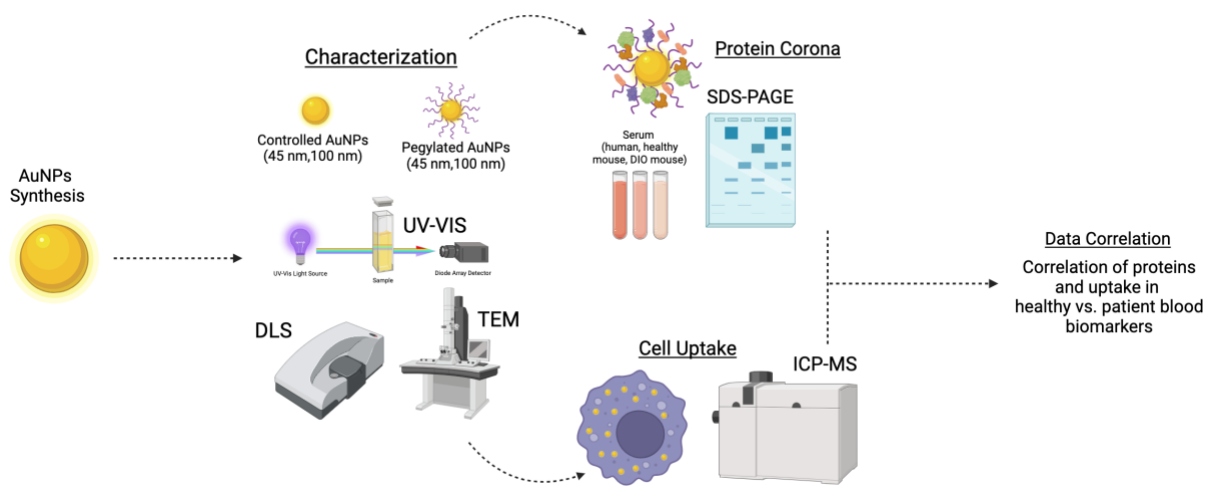


Figure 1.4: Approach to the thesis project divided into sections (Made in Bio render)

1.2.1 Gold Nanoparticles Synthesis

The synthesis of AuNPs starts by creating a gold stock. This stock is used to create AuNPs seeds, which are typically ~15nm in size. We use these seeds to grow different sizes of larger AuNPs. In this project, two standard AuNP sizes for testing were chosen, including a smaller 45nm and a larger 100nm in diameter. A difference in color of the nanoparticles in the solution can be observed when they grow from 15nm seeds to larger sized nanoparticles. This is a result of the change in how light interacts with nanoparticles of different sizes and perceived by the human eye. After AuNPs of different sizes are synthesized, their surface treatment will then be conducted. To start, it is important to understand that AuNPs can agglomerate, meaning that the nanoparticles will start to adhere to one another. This is due to high surface energy of nanoparticles, and physical processes in solution where their surfaces can contact and adhere to one another via van der Waals interactions. [8] To prevent unwanted agglomeration, a surface treatment is completed on the AuNPs, called PEGylation. This process is done by covering the surface of the AuNPs with polyethylene glycol (PEG) chains. Specifically in our case, we will cover the surface with a mixed PEG layer of methoxy PEG (mPEG), which will serve as a backfill PEG; and amino PEG (amPEG), which will be used to chemically conjugate fluorescent dye to the AuNPs for tracking in cell imaging. The purpose of PEGylation is to improve colloidal stability of the AuNPs, which will show a dependance on electrostatic interactions meaning the charge of the nanoparticle, and steric hinderance which will be creating a layer around the gold nanoparticle to prevent direct surface contact.

1.2.2 Materials Characterization

After AuNP synthesis, characterization of the resulting nanoparticles needs to be completed to confirm the correct size and polydispersity index (PDI). The PDI measures how

even the particle sizes are within the solution of nanoparticles. Dynamic Light Scattering (DLS) was used for size and PDI characterization, and UV-Visible Spectroscopy (UV-VIS) was used to determine the wavelength and plasma resonance peak shift (PRPS). PRPS is a phenomenon that occurs when there is an intersection of light with a metal surface, causing their electrons to be excited at a specific resonance wavelength that will then depend on size, shape and spacing between particles., We characterized the AuNPs in different conditions, including bare (without PEGylation surface treatment), and when surface treated after PEGylation. In addition, we utilized Transmission Electron Microscopy (TEM) for visual characterization of the morphology of the AuNPs. Our gold nanoparticles are expected to react differently due to their surface energy change when the addition of PEG to the surface of them due to these polymer chains serving as a colloidal stabilizer, meaning that visually our AuNPs may look different on the TEM images. [9]

1.2.3 Protein Corona

The next section of the study consists of the protein corona study of the tested AuNPs. For this, the AuNPs are covered with only mPEG and later combined with different serum conditions, leading to protein corona formation. For the serum conditions, we first used healthy human serum (HHS) to have a basic idea of the interaction of the human body with nanoparticles. The second set of serums consists of mouse serums: a healthy mouse serum (HMS), which serves as the control of the interaction between nanoparticles and the normal mouse physiology; and a diet-induced obese mouse serum (DIO) which serves as the model for the obese patient physiology (Figure 1.5). The DIO mouse model was created to resemble the metabolic human syndrome. This model consisted of providing a high fat diet along fructose water to normal C57BL/6 mice to allow the study of obesity and other obesity-related metabolic abnormalities. [10] Previous literature and characterization demonstrates that DIO mice serum

has elevated levels of cholesterol, high-density lipoproteins (HDL) and low-density lipoproteins (LDL). These lipoproteins are made from a phospholipid sphere that is filled with triglycerides and cholesterol having an outer shell of apolipoproteins, which might affect our AuNPs protein corona with a higher amount of those proteins involved in the DIO serum model (Figure 1.6). [11]



Figure 1.5: Mouse model comparison from healthy to DIO (Made in Bio render)

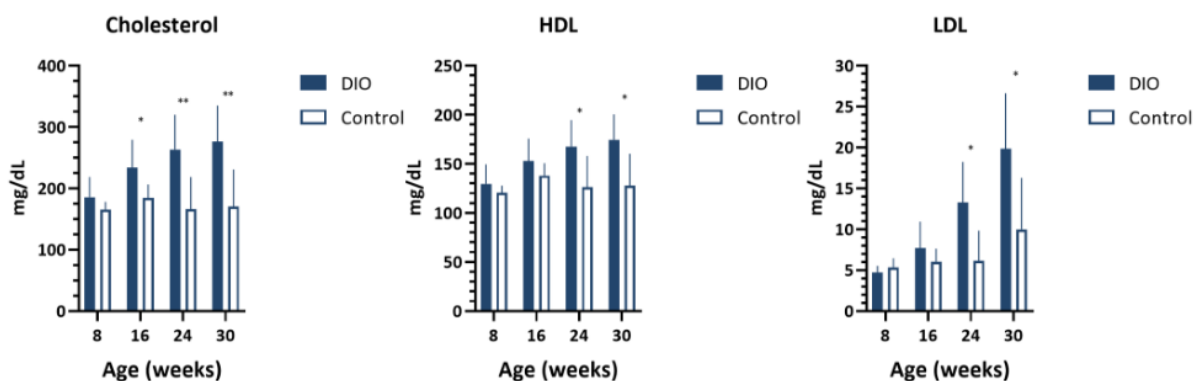


Figure 1.6: Serum Chemistry from DIO and healthy mouse serum [11]

wavelength with the UV-VIS after protein corona formation on the AuNPs. With the use of Sodium dodecyl-sulfate polyacrylamide gel electrophoresis (SDS-PAGE), we can also identify relative abundance of proteins in the nanoparticle corona through mass-based electrophoretic separation. To accomplish this, proteins are isolated from AuNPs by dithiothreitol (DTT) and centrifugation. The proteins are then denatured with DTT and SDS and separated by mass with electrophoresis through a polyacrylamide gel. The gel is then stained with Coomassie blue that labels proteins to give the final band sizes of the proteins present as a protein corona signature. [12]

1.2.4 Cell Uptake

Moving on to the next step, we sought to investigate the interaction between AuNPs and cells by using RAW264.7 mouse macrophages. We perform optical microscopy to observe any physical differences in the cell morphology and culture behavior when incubated with all the serums. This can help us understand how the serum physically affects the cell, and their interaction with AuNPs subsequently. Next, we performed fluorescence microscopy cell-imaging to observe cell uptake differences in macrophages. We fluorescently labelled PEGylated AuNPs with Alexa Fluor 488 NHS ester, which is a green-fluorescent dye for imaging. [13] AuNP localization inside cells can be tracked, and cell nuclei were counterstained with NucBlue fluorescent dye. Finally, we use Inductively Coupled Plasma – Mass Spectrometry (ICP-MS), which measures the concentration of elements in the sample by ionizing them using high-energy plasma and measuring the mass-to-charge ratio of specific ions with the mass spectrometer. [14] In this case, we will measure the amount of gold and magnesium per cell as a proxy for gold nanoparticle uptake and cell number respectively.

1.2.5 Data Correlation

After completion of the three main steps of this study, we will then analyze the data relevant to nanoparticle uptake differences and the different proteins present in the nanoparticle protein corona. This will allow us to gain understanding in the effect of patient-specific blood biomarkers on nanoparticle cell interactions, as well as devise future experiments to complement these findings.

Chapter 2: Methods

In this section, we will cover the specific procedures that were followed to perform the experiments, as well as analyze and gather all the data for our study. The methodology will be divided by the sections the project as outlined in the approach.

2.1 GOLD NANOPARTICLE SYNTHESIS

To begin gold nanoparticle synthesis, we prepared gold stock from chloroauric acid (HAuCl_4 , 99.995% trace metals basis) (Sigma-Aldrich, catalog number 254169) (100g/L for seeds), (10mg/mL for other sizes), sodium citrate tribasic (ACS reagent, > 99.0%) (Sigma-Aldrich, catalog number S4641) (30mg/mL for seeds) (4.4mg/mL for other sizes), hydroquinone (Reagent Plus, 99%) (Sigma-Aldrich, catalog number H17902) (2.75mg/mL) and a final concentration of AuNPs of 2.4nM. All buffers and reagent solutions were prepared using purified deionized water with resistivity of $18.2 \text{ M } \Omega \cdot \text{cm}$ at 25°C .

2.1.1 Seeds (15nm AuNPs)

For synthesis of the AuNPs seeds (15nm) we washed a 250mL Erlenmeyer flask with a magnetic 1.5” stir bar with aqua regia, this is a mixture of 3-parts hydrochloric acid (ACS reagent, 37%) (Sigma-Aldrich, catalog number 258148) and 1-part nitric acid (ACS reagent, 70%) (Sigma-Aldrich, catalog number 438073). This process is of high importance to ensure that any metal or gold impurities will be dissolved and not interfere with our synthesis process. Next, we added 100 g of water to the clean Erlenmeyer flask and heated it on a hot plate at 370°C with a slow stir until the water was boiling. Once the water was boiling, we added 1mL of sodium citrate solution, and 100 μL of HAuCl_4 and stirred for 10 minutes. We then stopped the reaction by placing the solution on ice and let it sit for another 10 minutes. Once completed, the solution

is ready for characterization, with an expected PDI <0.1 and hydrodynamic size range from 18-20nm. (Figure 2.1)

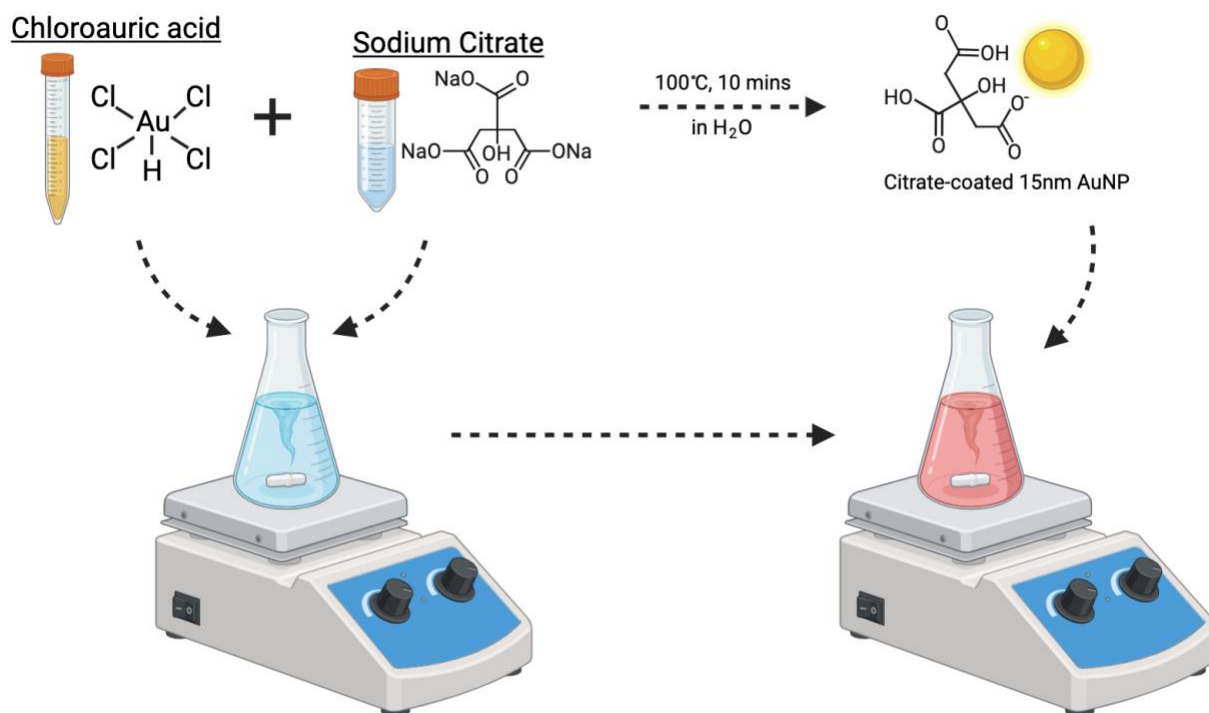


Figure 2.1: Synthesis of gold nanoparticles seeds of 15nm (Made in Bio render)

2.1.2 45nm and 100nm AuNPs

For larger AuNP synthesis, we started by cleaning a 250mL Erlenmeyer flask and a 1.5” magnetic stir bar with aqua regia as before to ensure no metal contaminants interfered with the synthesis of our AuNPs. After, we added 100g of water to the flask and stirred vigorously. Next, we added the reagents in the following order in quick succession at room temperature: AuNPs seeds, HAuCl₄, sodium citrate, and hydroquinone. The solution was left to stir continuously overnight or for about 12 hours in room temperature (Figure 2.2). The reaction was stopped with addition of 1mL of TWEEN20 (5% w/v) (Sigma-Aldrich, catalog number P2287).

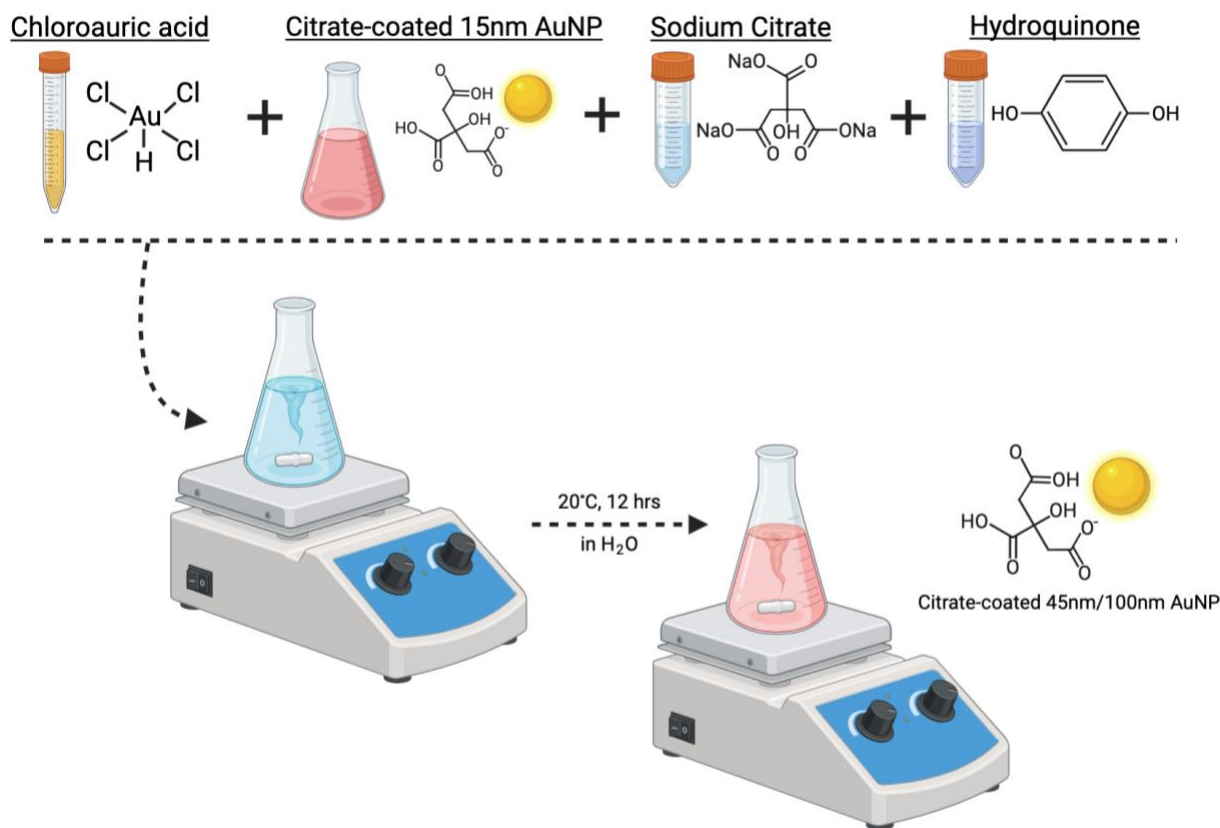


Figure 2.2: Synthesis of gold nanoparticles in 45nm and 100nm sizes (*Made in Bio render*)

2.1.3 Nanoparticle PEGylation

For the PEGylation process, we measured the amount of methoxy-5kDa-thiol PEG (Laysan Bio, custom order) and amino-10kDa-thiol PEG (Laysan Bio, custom order) needed for our different concentrations of our varied AuNPs. Once the amount of PEG is calculated, both PEGs were dissolved in water and added to AuNPs and left in a 60°C water bath for 1 hour. Once the time is completed, the mixture was optionally placed in the 4°C fridge overnight or for 12 hours. The following step consisted of washing the PEGylated AuNPs three times: the first time is to pellet down the AuNPs, second is with water, and the third and last time is 1 with 1x PBS (modified without calcium chloride and magnesium chloride) (Sigma-Aldrich, catalog

number D8537), each wash lasted 35 minutes to the recommended speed of the AuNPs at 4°C.

To finish, 500µL of 1x PBS was added to top off. (Figure 2.3)

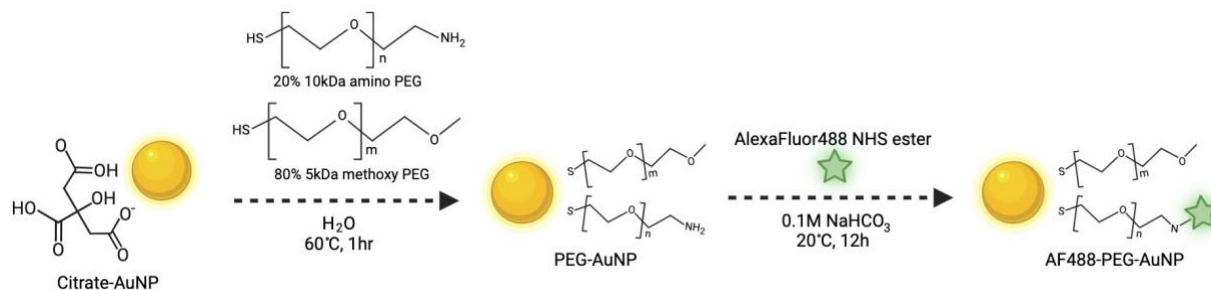


Figure 2.3: Pegylation process for amino and methoxy PEG (Made in Bio render)

2.2 MATERIALS CHARACTERIZATION

In this section, we give a basic explanation and procedure of each of the tests conducted and provide the justification for how we chose to utilize these methods for our studies.

2.2.1 Dynamic Light Scattering (DLS)

Dynamic Light Scattering (DLS) is a tool utilized for the study of diffusion of molecules in solution. to provide an analysis and measurement of size and polydispersity of the colloidal molecules within the solution. For our study, we used the Malvern Zetasizer Ultra Red to analyze each of our AuNPs in solution to measure not only the synthesized size, but how these changes with the interaction of PEG and with the change of size, as well as how PDI changes or maintains through the surface preparation of the AuNPs. [15] µLµL

2.2.2 Ultraviolet-Visible Spectroscopy (UV-VIS)

Using Ultraviolet-Visible Spectroscopy (UV-VIS), we can study how a substance interacts with electromagnetic radiation, including ultraviolet and visible regions of the spectrum. [16] In this case with the use of UV-VIS (Shimadzu UV-1900i spectrophotometer), we confirmed the AuNP size by their peak absorbance from their UV-Vis spectra, due to specific sizes absorbing characteristic wavelengths of light. [17] When we added serum to AuNPs, it is

expected that there may be agglomeration due to reduced colloidal stability, this is expected to result in a shift in the peak resonance wavelength, related to the charge differences of particles when interacting with serum. The relationship between colloidal stability with charged proteins are responsible for this peak shift. Which as explained by Mie theory - absorption and scattering efficiencies are related to the size, shape and material. [18] This is attributed to the general varied charges that the different adsorbed proteins contain; with the agglomeration of these nanoparticles, we may see a difference in the wavelength of the AuNPs because of the “color change” that is appreciated from particles absorbing and scattering light differently when adhered together. (Figure 2.4)

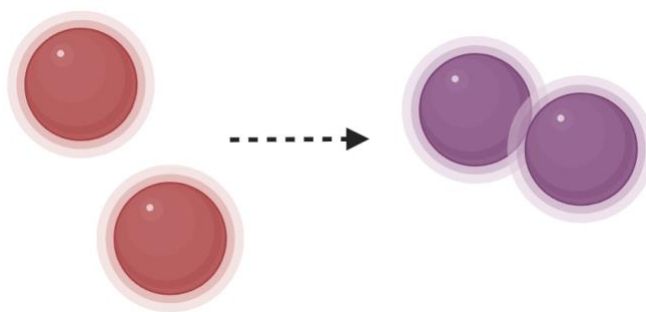


Figure 2.4: Change of visual color of nanoparticles when aggregated (Made in Bio render)

2.2.3 Transmission Electron Microscopy (TEM)

The Transmission Electron Microscopy (TEM) gives us the opportunity to take images of our nanomaterials in detail due to its' ability to “zoom” into a material past the diffraction limit of light. For sample preparation, we added AuNPs solution into copper TEM grids (Carbon Type-B, 300 mesh) (Ted Pella Inc, catalog number 01813-F). We started by plasma cleaning the grids using a Harrick Plasma - Cleaner PDC-32G (Roberts Lab, UTEP). Plasma cleaning is necessary to make the TEM grids hydrophilic, facilitating the deposition of nanoparticles when the solution is added. After plasma cleaning, we added around 5-10 μ L of sample on top of the

grid, and left it for a total of 10 minutes. Taking a filter paper, we touched the corner of the grid to absorb any excess liquid and left to air-dry for another 10 minutes.

2.3 PROTEIN CORONA

For the protein corona section of the study, it was important for us to investigate the differences in protein adsorption onto the surface of the gold nanoparticles due to the types of serum they were incubated and exposed to. The main differences we expect to see is the different protein adsorption due to the higher concentration of HDL and LDL in obese serum. These lipoproteins consist of mainly a sphere of fats covered by various apolipoproteins. The main apolipoproteins that are in HDL include Apo A-1, Apo A-II, Apo E, and Apo C; and for LDL is Apo B, meaning that these proteins have a direct correlation with the levels of concentration of lipoproteins in the obese serum. It is important to keep in mind that the measurement of 'kDa' for proteins refers to the actual molecular weight and not necessarily of the size of the proteins. These sizes can range in between 2-6nm depending on the protein three-dimensional structure, and usually has a correlation with the length of the amino acid chains and number of subunits in each protein. However, it is possible for a protein to have a relatively low molecular weight but a corresponding large diameter. Keeping in mind that the variety of proteins may depend on the actual AuNPs diameter, this is mainly because of the space available of proteins to get attached to on the surface, as well as three dimensional structural changes to the protein when it interacts with the high energy nanoparticle surface. SDS-PAGE can provide some basic insights into if there are any highly abundant possible proteins in the nanoparticle corona, or if these the serum composition itself can impede certain proteins from forming the corona. (Figure 2.5)

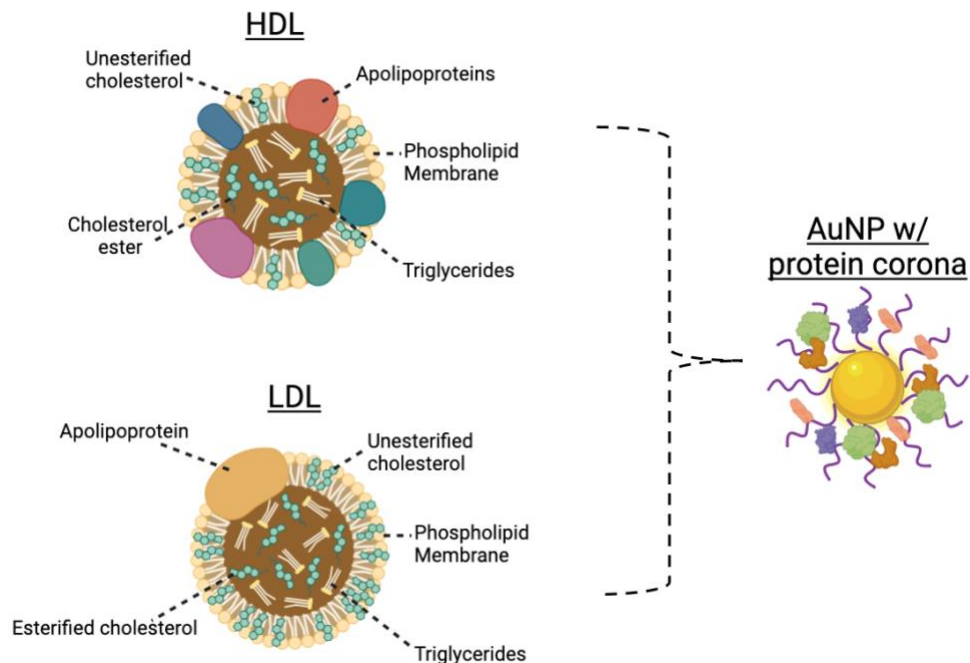


Figure 2.5: Relationship between HDL and LDL found in DIO mouse serum and protein corona (Made in Bio render)

2.3.1 Sodium dodecyl-sulfate polyacrylamide gel electrophoresis (SDS-PAGE)

We used Sodium Dodecyl-Sulfate Polyacrylamide Gel Electrophoresis (SDS-PAGE) to separate and track protein bands from the nanoparticle protein corona to identify differences in the size of proteins that are present in the protein coronas of our varied serums when interacted with all our AuNPs sizes and conditions. [19] From this, we can infer or make educated guesses to the identities of the protein bands when compared to previous protein corona studies from literature. To prepare the samples that will be analyzed by SDS-PAGE, we subjected the AuNPs to all the different serum conditions. This was done by first calculating the total surface area of both of AuNP's sizes to normalize the same amount of nanoparticle surface was being tested with the serum conditions. We then combined a total of 500 μ L of serums (HHS, HMS, DIO separately) and 500 μ L of phosphate buffered saline (PBS). In the case of AuNPs without PEG, the PBS will be exchanged for water. Healthy human serum (from human male AB plasma, USA

origin, sterile filtered) was obtained from Sigma-Aldrich, catalog number H4522, lot number SLCL9009. Healthy mouse serum (C57BL/6J mouse serum gender pooled) was obtained from BioIVT Inc, product number MSE01SRM-0101510, lot number MSE476971. DIO mouse serum (C57BL/6J DIO mouse serum male pooled) was obtained from BIOIVT, Inc, product number MSE75SRM-0119037, lot number MSE476972. Then, the samples will be left for 12 hours in 37°C. After, the samples were washed and separated from the extra serum and stored in PBS (or water in the AuNPs without PEG). To extract the protein corona from the AuNPs, a total of 20µL NuPAGE LDS Sample buffer (Invitrogen, catalog number NP0008) was added along 50µL of 2mM 1,4-Dithiothreitol (DTT) (Roche, 97% Ellman's reagent, catalog number 10197777001) and incubated for 30 minutes in a 60°C water bath. Then, we centrifuged samples for 15 minutes at 18,000G and saved the supernatant to run on the SDS-PAGE gel. After, we followed the recommended procedure on the Thermo-Fisher website for the use of the SDS-PAGE Mini Gel tank and the NuPAGE 4% to 12%, Bis-Tris Plus mini gels (Invitrogen, catalog number NW0412A). We ran samples in duplicate a: a protein ladder (Page Ruler Plus Prestained Protein Ladder, 10 to 250 kDa, Invitrogen, catalog number 26619), a control lane of the serum only, two samples for protein corona in duplicate for the different conditions: 45nm AuNPs without PEG, 45nm AuNPs with PEG, 100nm AuNPs without PEG, and 100nm AuNPs with PEG.

2.4 CELL UPTAKE

To study the physical changes of RAW 264.7 cells when exposed to different serums, we started by adding the different serum conditions into RAW 264.7 cells in 6-well plates. We used fetal bovine serum (FBS) as a control since it is the standard serum supplement for most cell culture media. The treated conditions included healthy human serum (HHS), healthy mouse

serum (HMS), and diet-induced obese (DIO) mouse serum. For all conditions, we considered a low dose and high dose of serum, corresponding to 20 μ L (2% v/v serum to media), and 100 μ L (10% v/v serum to media) respectively.

2.4.1 Inductively Coupled Plasma – Mass Spectrometry (ICP-MS)

With the use of the Inductively-Coupled plasma mass spectrometry (ICP-MS), we wanted to quantify the difference of AuNP uptake in RAW 264.7 cells in only human serum with a variety of AuNPs sizes. To prepare cell samples for ICP-MS, we removed the media from the well plates of RAW264.7 cells treated with different serum and AuNP conditions. Then, we treated the cell samples after leaving them for a day with the new serum and the AuNPs, by digesting the samples with aqua regia that is then diluted in 2% HNO₃, with a final solution of 1-100ppb range for Au. With the sample preparation we address and make a standard curve set of samples for the cell-line tested in this case our RAW 264.7 cell-line. After this process we send our sample to Dr. Lin Ma's laboratory, where he and his students process the samples in the ICP-MS to give us our results.

2.4.2 Cell - Imaging

With fluorescence microscopy cell – imaging, our goal was to visually capture the difference in nanoparticle uptake between the different serum conditions. For this process, we plated RAW264.7 macrophages (American Type Culture Collection, catalog number TIB-71) according to manufacturer recommendations (DMEM media, supplemented with 10% fetal bovine serum and 1% penicillin-streptomycin) in 6-well tissue culture plates at 37°C and 5% CO₂. Cells were left overnight in the incubator to attach to the bottom of the plate. The next day, we starved the cells for 5-10 minutes by replacing the regular supplemented media with FBS-free supplemented media. Then, we added the different serum to the cells for an additional 12 hours,

and then added the different AuNPs labelled with AF488 dye for another 12 hours. Next, we washed out excess AuNPs with 1x PBS and fixed the cells with 4% paraformaldehyde in 1x PBS (Thermo Scientific Chemicals, catalog number AAJ19943K2) and finally added NucBlue stain (Invitrogen, catalog number R37606) for 10-15 minutes before imaging with a Zeiss Axio Observer.A1 fluorescence microscope (IMSTEL Lab, UTEP).

Chapter 3: Results

3.1 MATERIALS CHARACTERIZATION

3.1.1 Dynamic Light Scattering (DLS)

For AuNP measurement by Dynamic Light Scattering, we wanted to ensure our nanoparticles matched the size we intended for the synthesis to begin with. The two main numbers we are looking to check for quality control consisted of the z-average size and the PDI, which as mentioned in the methods section, should be <0.1 for good monodispersity. As seen in figures 3.1 and 3.2, the AuNP size increased with the addition of the PEG as expected while maintaining a $PDI < 0.1$.

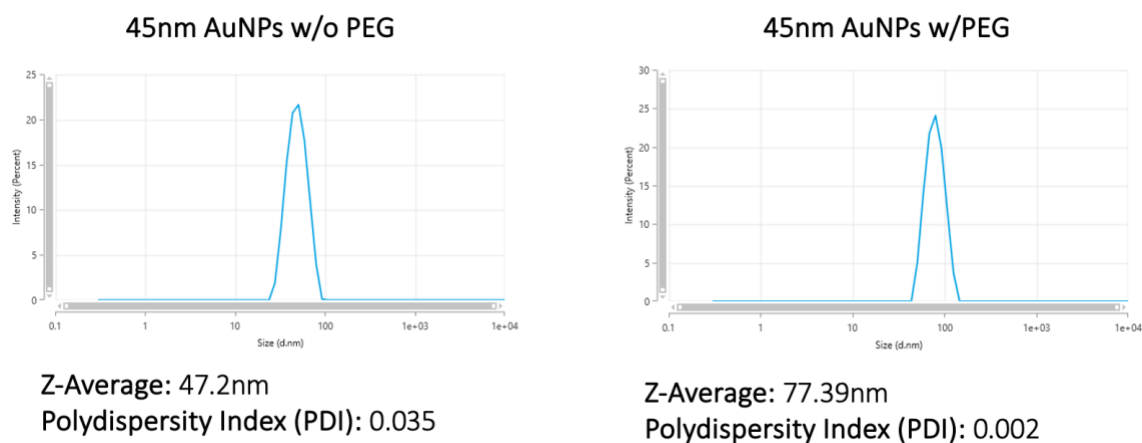


Figure 3.1: Size and Polydispersity index of both 45nm without and with PEG

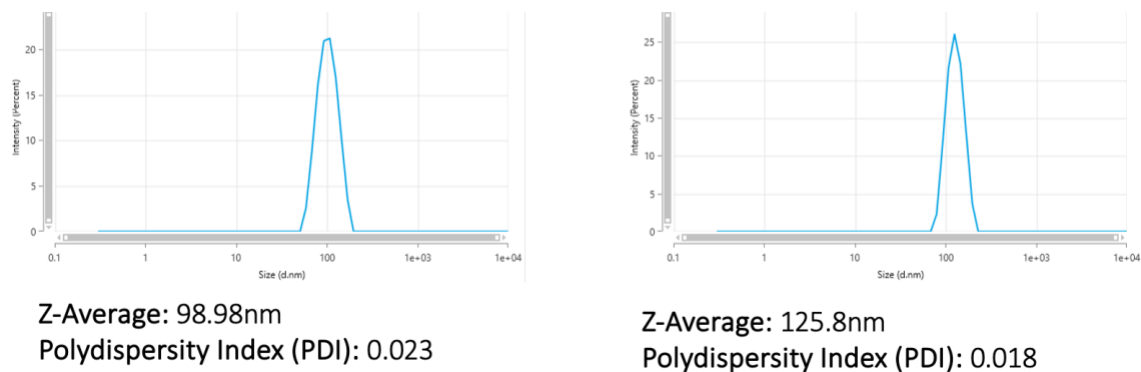


Figure 3.2: Size and Polydispersity index of both 100nm without and with PEG

Next, we wanted to compare the colloidal behavior of AuNPs with and without PEGylation when exposed to serum. From the results obtained, we can see that a change in size is mainly obtained for the AuNPs with no PEGylation. This is related to the decrease of colloidal stability due to the change of charge with the addition of serum proteins. This is due to proteins having multiple different charged domains and can exhibit zwitterionic behavior. The coated proteins therefore tend to increase adhesion between adjacent nanoparticles and induce agglomeration or aggregation. This is reflected in the DLS data as an increase in z-average and PDI. PEGylation can prevent or decrease protein adsorption onto the nanoparticle surface, as well as improves steric hinderance of AuNPs, and therefore mitigates the protein corona agglomeration effects (Figure 3.3).

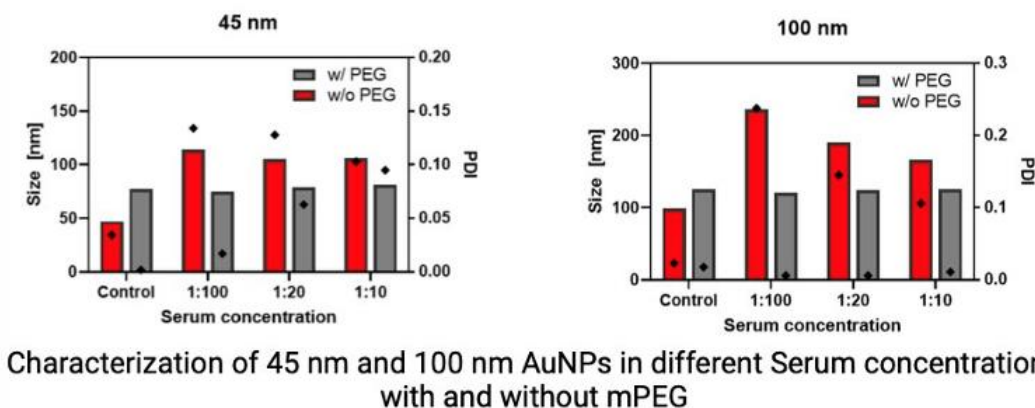


Figure 3.3: Size and Polydispersity index of both 100nm without and with PEG

To further characterize the AuNP behavior with respect to PEGylation and serum incubation, we used UV-VIS spectroscopy to determine the AuNP plasmon resonance peak shift. We expected the plasmon resonance peak in the absorbance spectra to have a peak between 500nm to 600nm, depending on the size of the AuNPs. We observed that with the addition of serum, a PRPS occurred for 100 nm AuNPs that were not surface treated with PEG. There was negligible to no PRPS for 45 nm AuNPs regardless of the presence of surface PEGylation. We

attribute this to the inherent lower colloidal stability of larger nanoparticles. This is because larger nanoparticles tend to settle and exhibit less true colloidal behavior. As such, larger nanoparticles are easier to agglomerate and aggregate. (Figure 3.4).

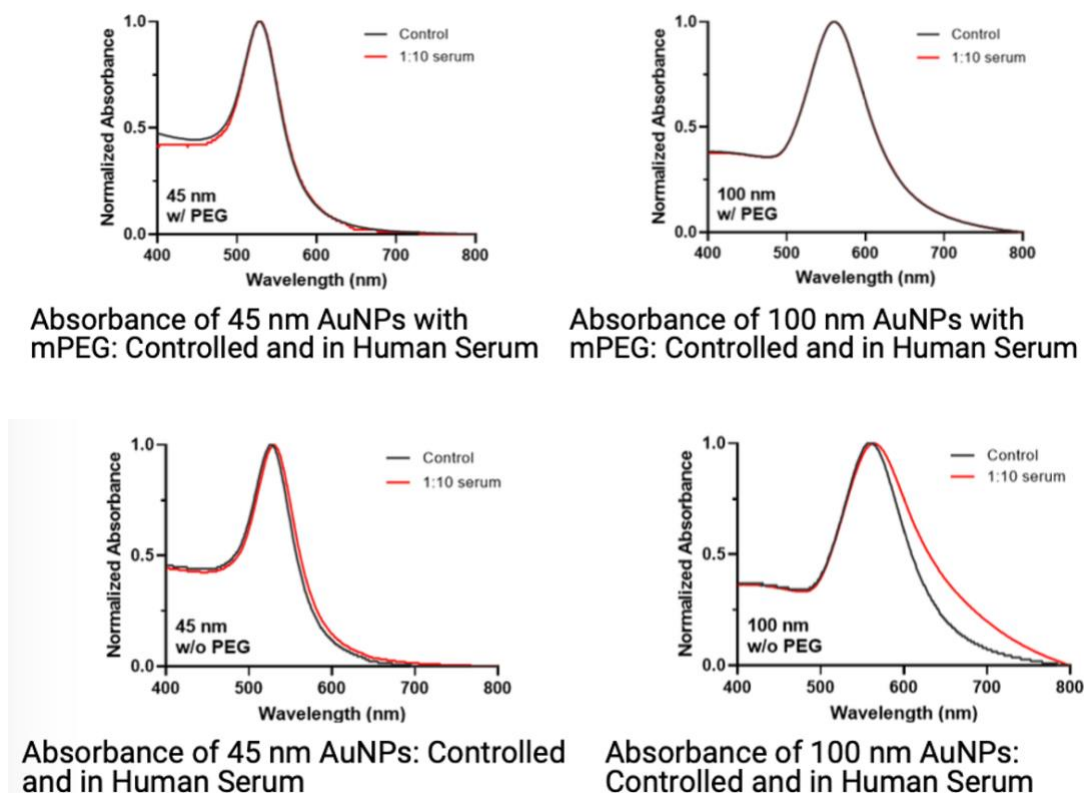
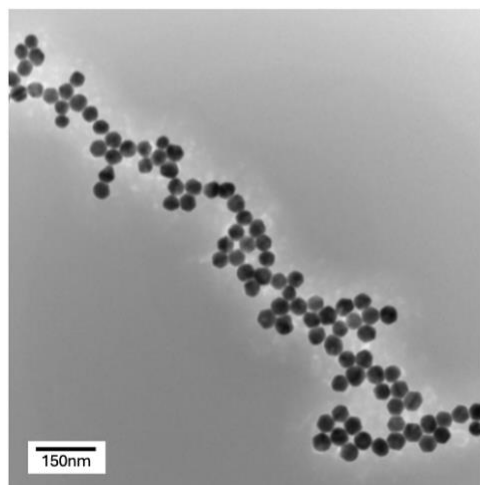


Figure 3.4: Plasmon resonance a peak shift in 45nm and 100nm AuNPs when subjected to serum

3.1.3 Transmission Electron Microscopy (TEM)

TEM results from our characterization methods guided us to visually analyze our particles morphology, not only by their size, but this process helped us understand how AuNPs interact without and with PEG surface. From our images taken, we are able to see how our particles start to sit close to almost on top of each other when not treated with PEG. This is due to the PEG functioning as a brush layer that stabilizes the AuNPs, and also physically separates them from each other. (Images 3.1, 3.2, 3.3, 3.4)

45nm AuNPs w/o PEG



45nm AuNPs w/PEG

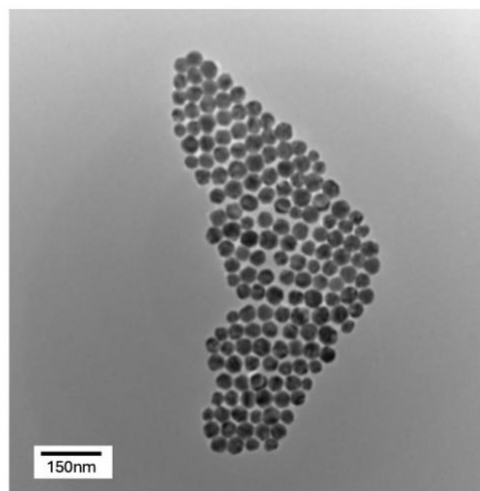
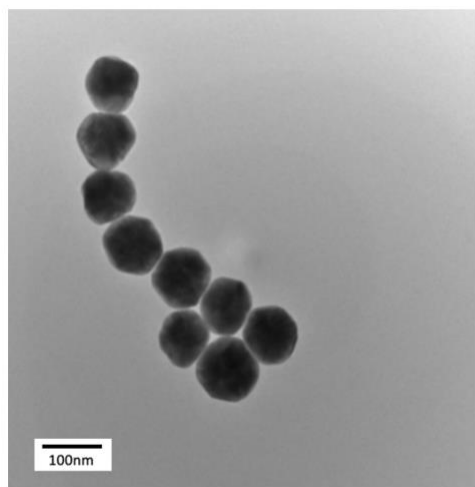


Image 3.1 & 3.2: TEM Images for 45nm with and without PEG

100nm AuNPs w/o PEG



100nm AuNPs w/PEG

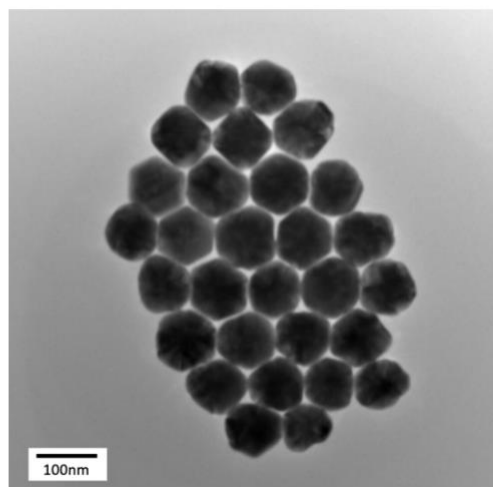


Image 3.3 & 3.4: TEM Images for 100nm with and without PEG

3.2 PROTEIN CORONA

The protein corona results obtained in our SDS-PAGE work, like mentioned before are mainly utilized to track any differences of proteins in the nanoparticle corona, without taking into consideration the specific identities of those proteins, but just the track their signature and relative abundance.

3.2.1 Sodium dodecyl-sulfate polyacrylamide gel electrophoresis (SDS-PAGE)

Given from the different serum conditions we started to see a difference from the mPEG surface coated AuNPs from the untreated ones, as seen the dark bands mean that there is a specific enrichment of proteins of that size. However, we do see a higher pattern of smearing with our AuNPs that were surface functionalized with mPEG, this can mean a result of many varieties of size proteins attached on the coronas of these condition AuNPs, decreasing the specificity of protein types.

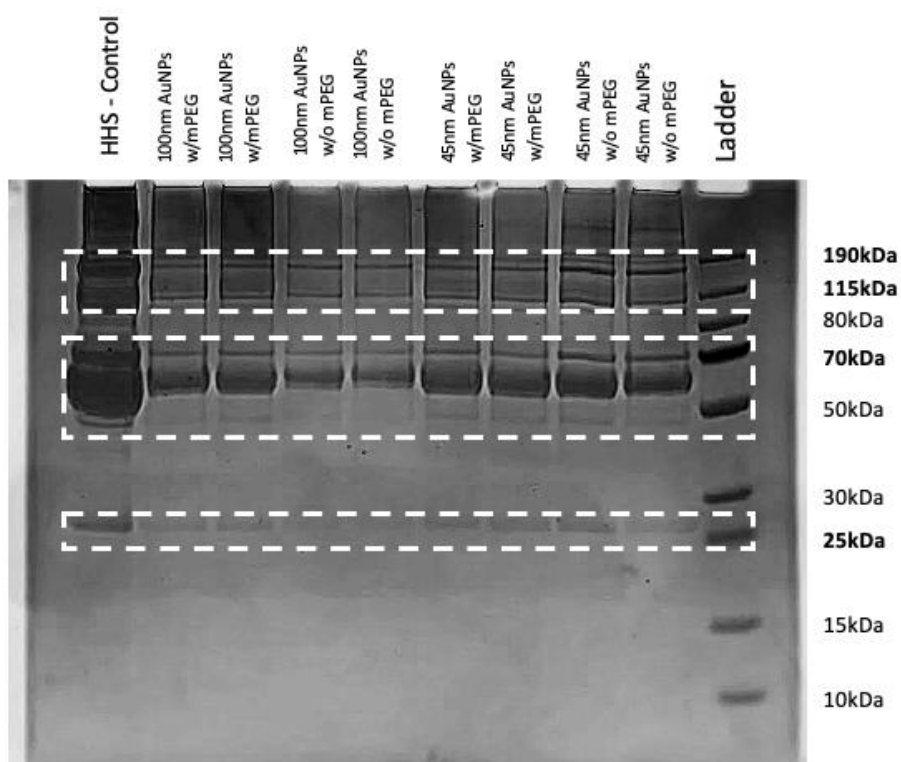


Figure 3.5: Relationship between HDL and LDL found in healthy human serum and protein corona

Table 1: Possible protein found in the HHS SDS-PAGE analysis [20]

Possible Proteins		
Type	Name	Size (kDa)
P01042	Kininogen-1	71.96 kDa
Q14624	Inter-alpha-trypsin inhibitor heavy chain H4	28.9 kDa
P01008	Serpin C1	58 kDa
P02768	Albumin	66.5 kDa
P03951	Coagulation factor <u>XIa</u> light chain	26.75 kDa

For our HHS study, we decided to reference the protein corona band signatures we obtained with the results reported in the paper, *An Analysis of the Binding Function and Structural Organization of the Protein Corona* [20], where the authors identified the main human serum proteins that attach to protein corona of AuNPs using liquid chromatography tandem mass spectrometry. (Figure 3.5) A list of potential proteins of interest in our SDS-PAGE gel that match the sizes of the most abundant listed proteins in the before mentioned paper is summarized in Table 1.

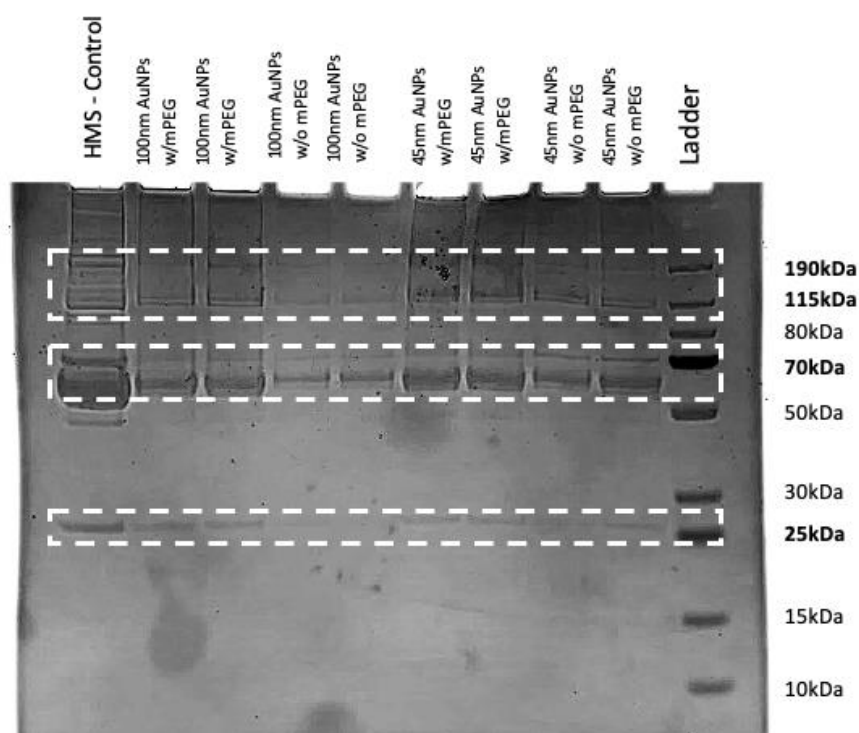


Figure 3.6: Relationship between HDL and LDL found in healthy mouse serum and protein corona

Table 2: Possible protein found in the HMS SDS-PAGE analysis [21]

Possible Proteins	
Name	Size (kDa)
Complement C3	186.5
Serotransferrin	76.7
Serum albumin	68.7
Vitronectin	54.8
Hemopexin	51.3

For our study in mouse serum conditions with both HMS and DIO mouse serums, we referenced the protein corona band signatures we obtained with the results reported in the paper, *Multomics analysis of naturally efficacious lipid nanoparticle coronas reveals high-density lipoprotein is necessary for their function*. [21] In this paper, the authors investigated the protein corona formation on lipid nanoparticles in obese and healthy rats. Although there are species differences between their rat model and our mouse model, this dataset still represents a good reference for investigating healthy versus obese protein corona in rodent models. We analyzed and compared the top 20 most significant proteins seen in protein coronas for rat serums from this paper to the protein bands from our SDS-PAGE gels for the proteins adsorbed in both HMS and DIO mouse serums on AuNPs. Upon visual inspection, for HMS there is a smearing in both no PEG and PEGylated AuNPs, however a higher intensity in the mPEG AuNPs, showing a possibility of higher variety of proteins in the corona. (Figure 3.6) From the main sizes seen in the protein bands, we compiled a table of the proteins of interest that have a size match to the ones obtained on the gel. (Table 2)

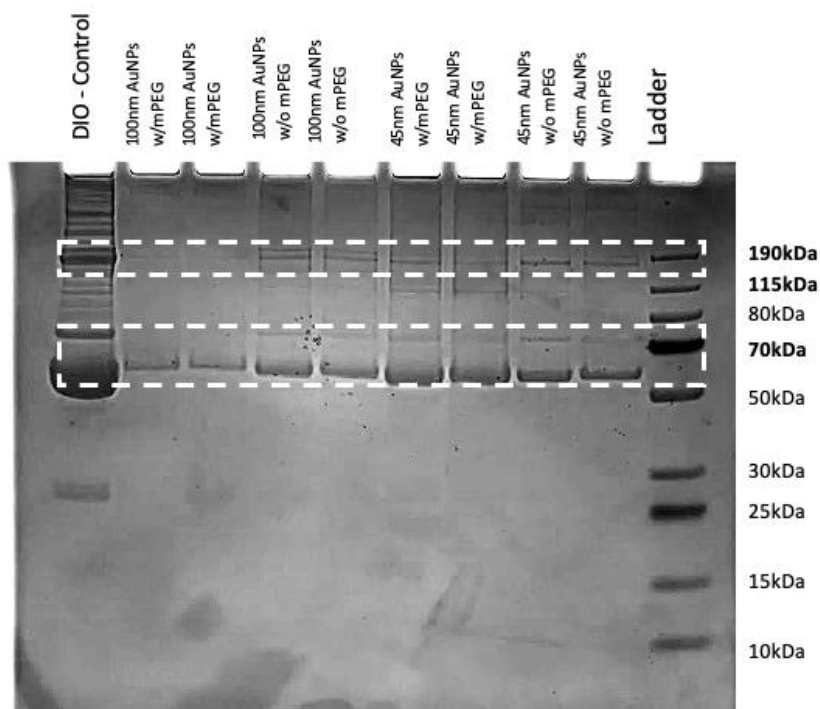


Figure 3.7: Relationship between HDL and LDL found in DIO mouse serum and protein corona

Table 3: Possible protein found in the DIO SDS-PAGE analysis [21]

Possible Proteins	
Name	Size (kDa)
Complement C3	186.5
Serotransferrin	76.7
Serum albumin	68.7
Hemopexin	51.3

Compared to our HMS gel, our DIO gel (Figure 3.7) showed a less smearing effect mainly in the 100nm AuNPs without PEG, which suggests that there is a less variety of proteins attached with this serum condition. With our next table, we did see that there was one less prospective protein of interest on this gel compared to the HMS, which may suggest that the

protein corona of this condition is relatively less diverse compared to the healthy condition.
(Table 3)

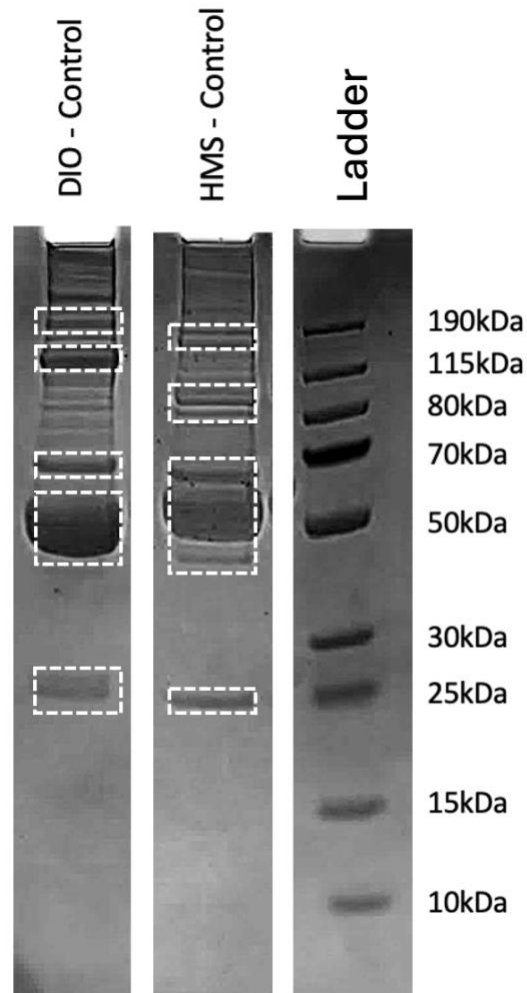


Figure 3.8: Differences between DIO and HMS mouse serum in an SDS-PAGE

When we make a direct comparison on only the serums and not necessarily the whole protein corona formed, we see a difference in protein bands present (Figure 3.8), showing an enrichment on proteins that have a 190, 115kDa in size, and a variety of proteins that can range between 25-30kDa and 45-60kDa in the DIO mouse serum. While in the HMS, we see an enrichment from 40-70kDa band sizes. From the serum biochemistry data for the DIO C57BL/6 mouse model, we expected to see the specific protein band sizes corresponding to Apo-A1,

which has a size of 28kDa, and Apo-B which is 550kDa (which might not be visible due to the gel utilized) due to the known elevated levels of HDL and LDL. From the bands obtained, in the DIO mouse serum there is a faint band seen around the size for Apo-A1, but not necessarily for the HMS.

3.3 CELL UPTAKE

With the cell uptake experiments, we first performed optical microscopy of the RAW264.7 cells incubated with different serum conditions. We observed that the macrophages exhibited extreme morphology differences when administered with FBS incubation, HHS, HMS, and DIO mouse serum conditions. With the standard FBS condition, the RAW264.7 cell maintained an even coverage of the flask culture surface, with an even size distribution and a consistent spherical morphology with elongations on some ends. (Image 3.5) When exposed to the HHS, we see the RAW264.7 cells proliferated less within 12 hours, with some agglomeration or noticeable clustering of the cells. (Image 3.6)

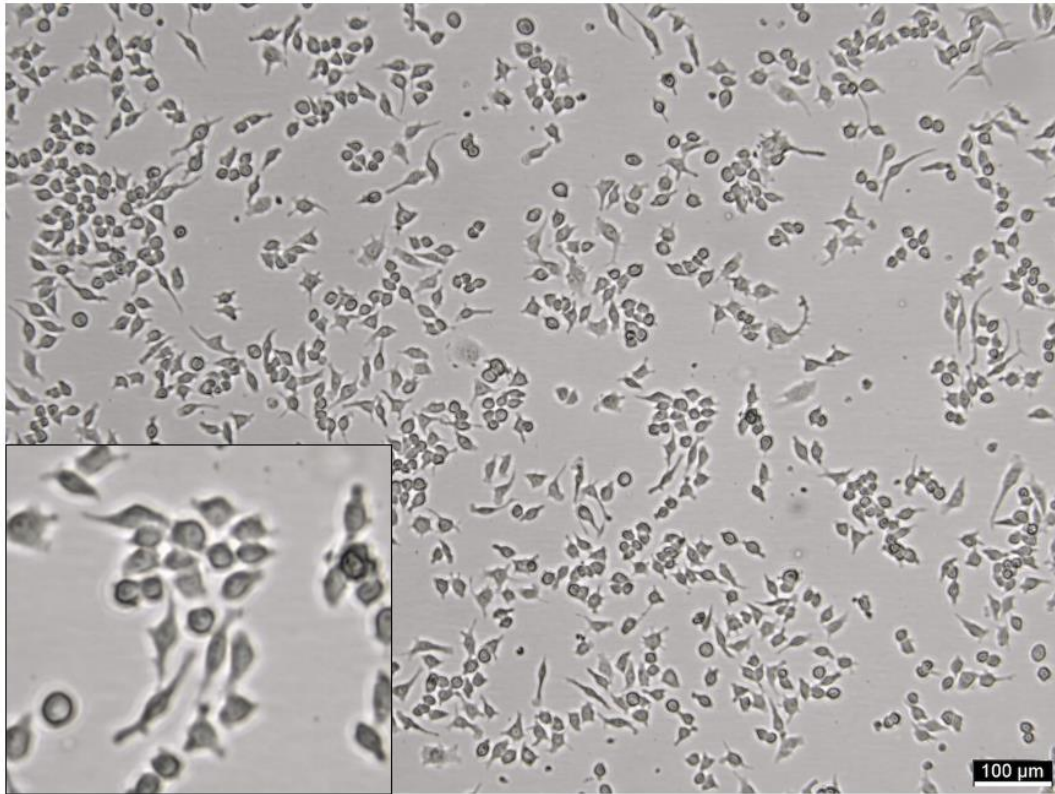


Image 3.5: RAW 264.7 Cells in FBS (Standard Incubation Condition)

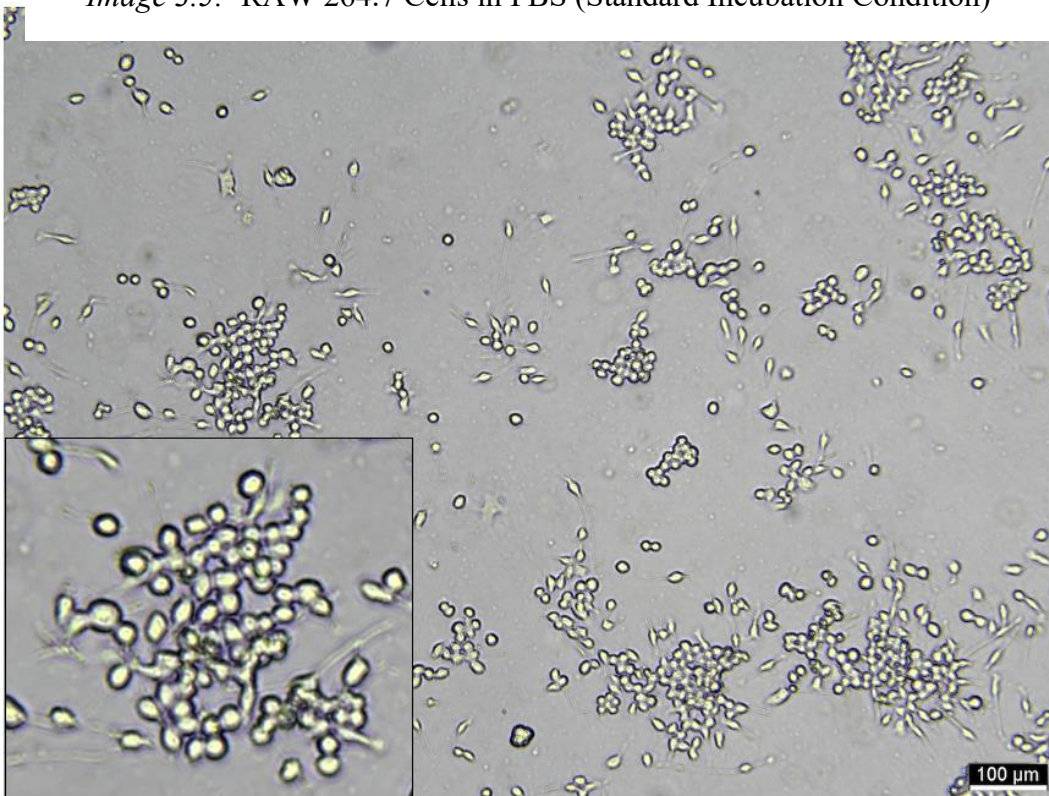


Image 3.6: RAW 264.7 Cells in HHS

However, when exposed to the mouse serum, we started to see a very different trend. Keeping in mind that RAW264.7 are mouse macrophages, we see a direct elongation of the cells when subjected to the HMS. (Image 3.8) The morphology of the cells in the healthy mouse condition helps us understand and have a close comparison to the “in vivo” scenario. When the DIO mouse serum is added, we saw a huge difference in the morphology of the cells. (Image 3.7) This condition caused the RAW264.7 cells to become chunkier and very clustered compared to the healthy condition where the cells were relatively more separated and elongated. Together, this data suggested that the RAW264.7 cells are likely responding to specific components in the different serums, and this may also have effects on their phenotypic behavior and cell state.

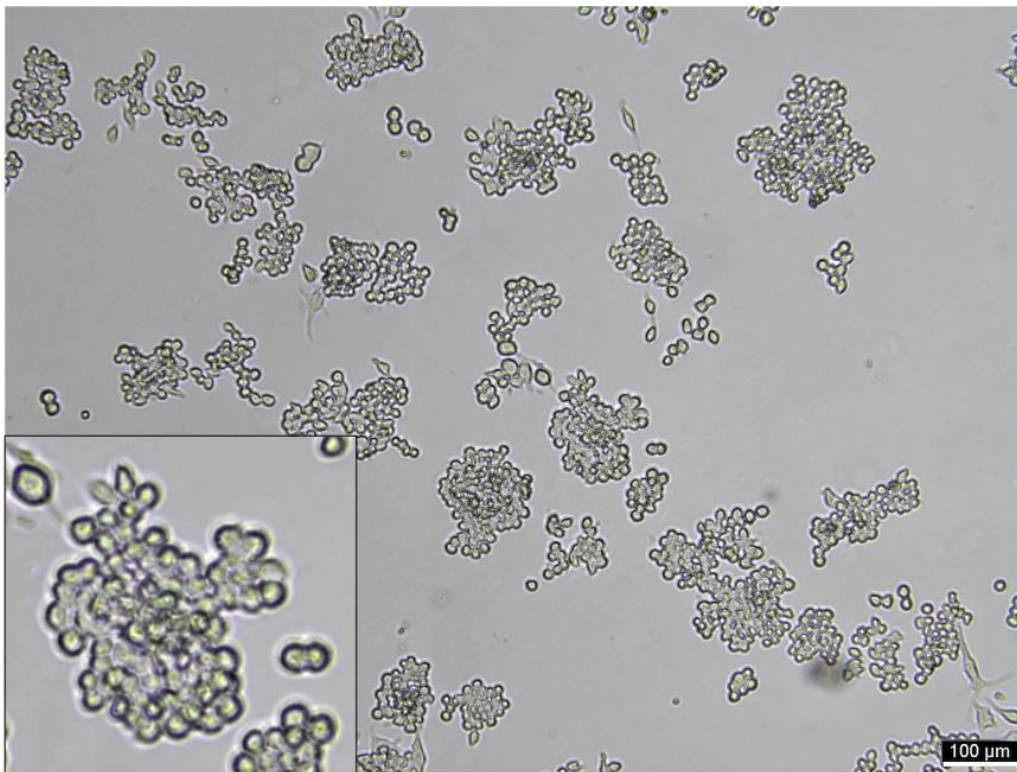


Image 3.7: RAW 264.7 Cells in DIO mouse serum

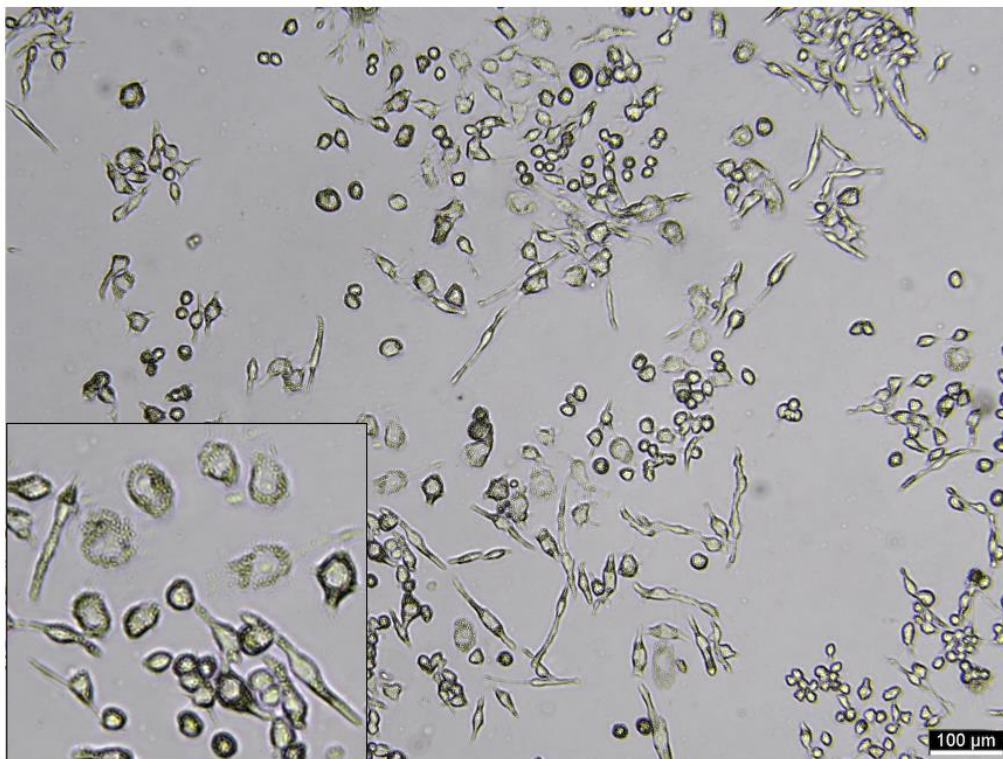
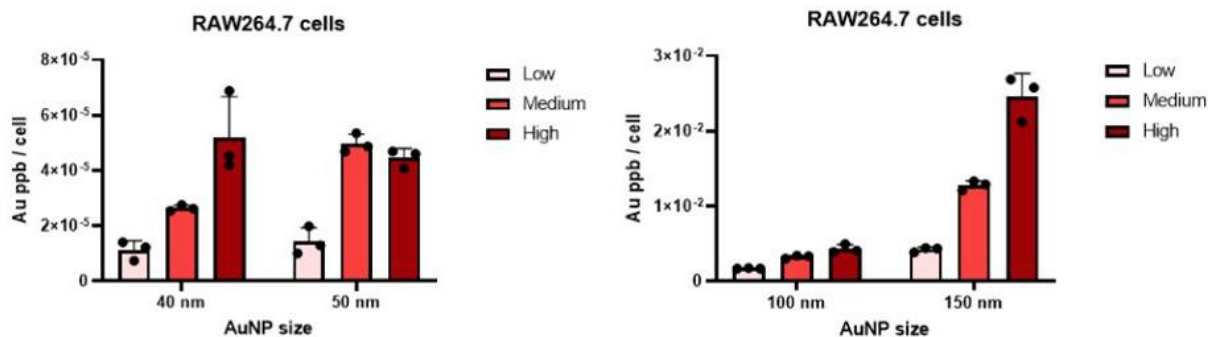


Image 3.8: RAW 264.7 Cells in HMS

3.3.1 In

From the data obtained from our ICP-MS samples, we came to the conclusion of the use of 45nm and 100nm AuNPs due to the current standard agreement for nanoparticles to be used of less than 100nm in size, and with this we decided to go close to the smallest AuNPs size tested which was the 45nm. (Figure 3.8) From this study, we were able to trial what we would be working in future experiments for the different serum conditions. There was an interesting observation of a non-linear relation of the 50nm AuNPs of the amount added with the amount of gold per cell. This can be a direct part of human error, since it is seen that the linear increment of gold stays relatively constant for the other sizes.



Gold uptake in Raw264.7 cell line with three different concentrations in 40 nm, 50 nm, 100 nm, and 150 nm

Figure 3.9: Gold uptake of AuNPs of different sizes in RAW 264.7 cells

3.3.2 Cell - imaging

With our cell-imaging, we were able to see the physical absorption of the AuNPs within each of the serum conditions and the different synthesized AuNPs sizes. The RAW264.7 cell nuclei were counterstained blue with NucBlue dye to aid in landmarking within cells. Our cells were dyed with a blue on the nucleus, meaning that for us to see an absorbance it will be around the nucleus. We are aware that auto fluorescence takes autofluorescence plays a significant part when working with green fluorescent dyes. With our results, we confirmed that this was not an issue due to our negative controls that showed no autofluorescence without the dye. For the HHS serum condition, (Figure 3.10), we observed that there was nanoparticle uptake into cells of all the different serum dose conditions and nanoparticle sizes. However, we also noted that some nanoparticle aggregates on the tissue culture plate surface, which may have led to an uneven uptake of the nanoparticles. Interestingly, we found that 45 nm AuNPs with a low dose of serum exhibited similar nanoparticle uptake into cells as the 100nm AuNPs with a high dose of serum.

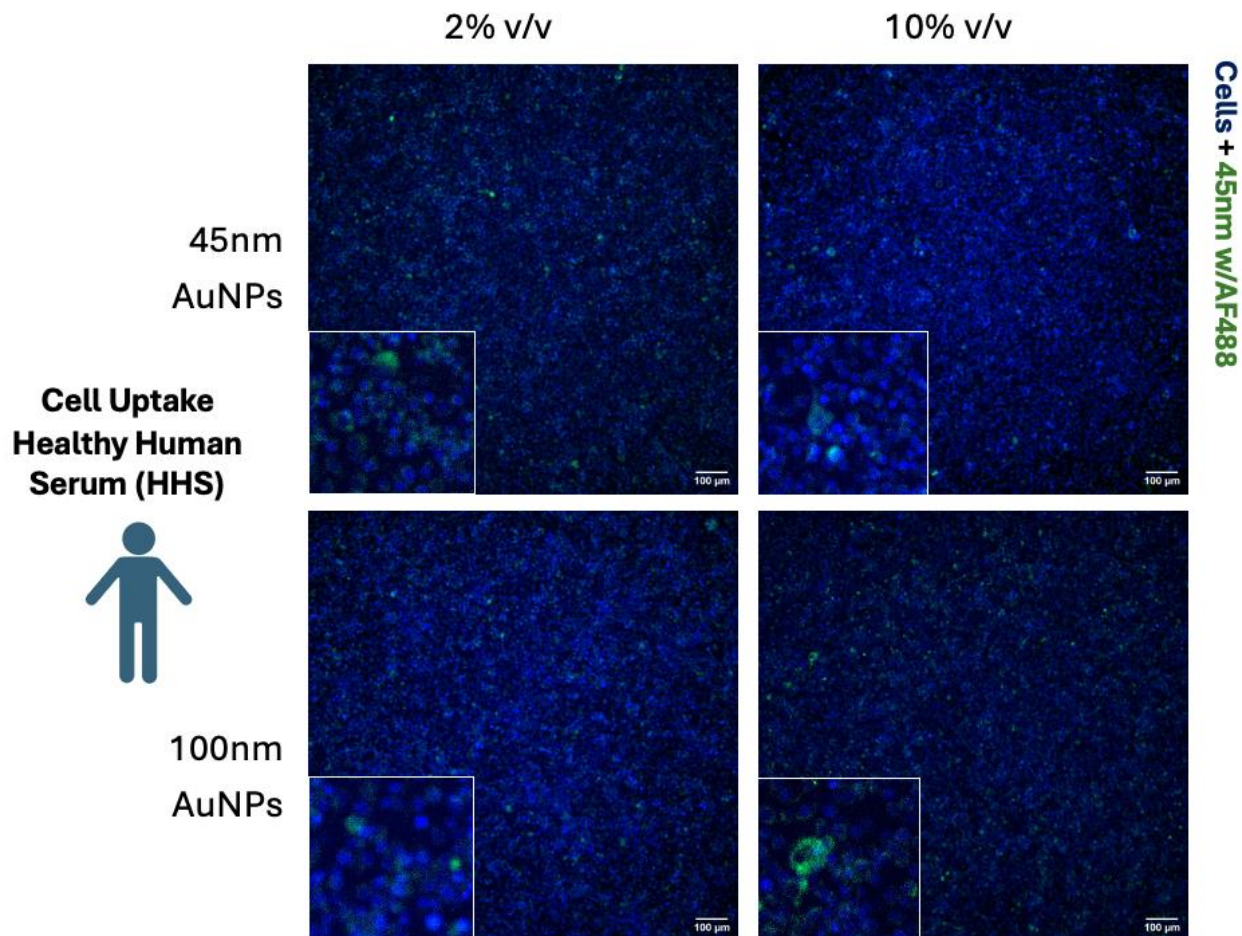


Figure 3.10: Cell uptake between AuNPs in different dose serums for HHS

For the HMS serum condition (Figure 3.11), we observed a higher uptake of the AuNPs into the RAW264.7 cells compared to the HHS serum condition. This is shown by an evenness of the green nanoparticles signal surrounding the nuclei of the cells. This suggests that the healthy mouse serum condition may be very conducive for nanoparticle cell uptake. We observed high nanoparticle cell uptake with all the different doses of serum and the different sizes of AuNPs. There were no significant observable differences in uptake between all the conditions for HMS.

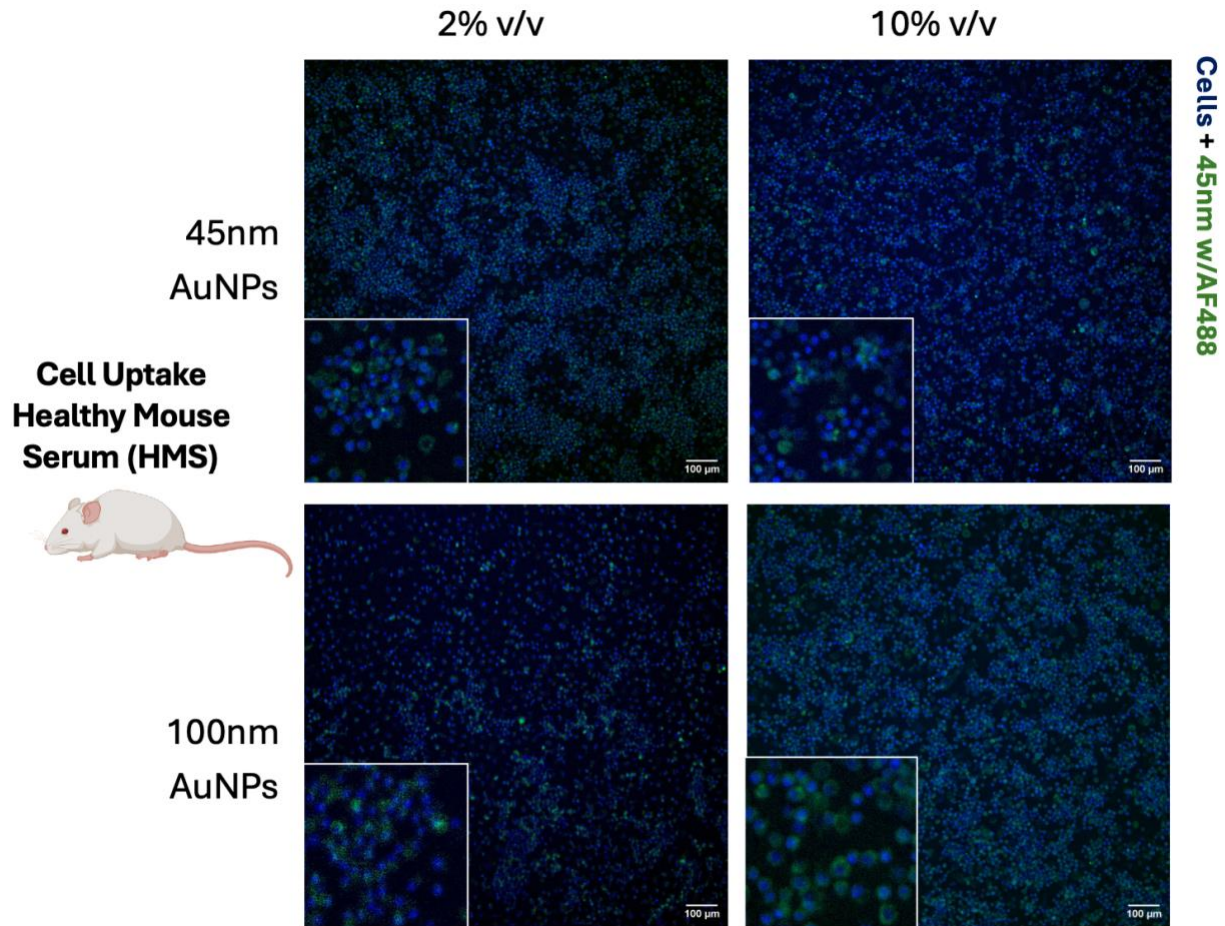


Figure 3.11: Cell uptake between AuNPs in different dose serums for HMS

Lastly with the DIO mouse serum condition (Figure 3.12), we observed that with this serum, no matter the condition given, it is seen that is very little green nanoparticle signal inside cells, especially in comparison to the HMS conditions. These images suggest that in obese serum conditions, there seems to be a negative effect on nanoparticle cell uptake.

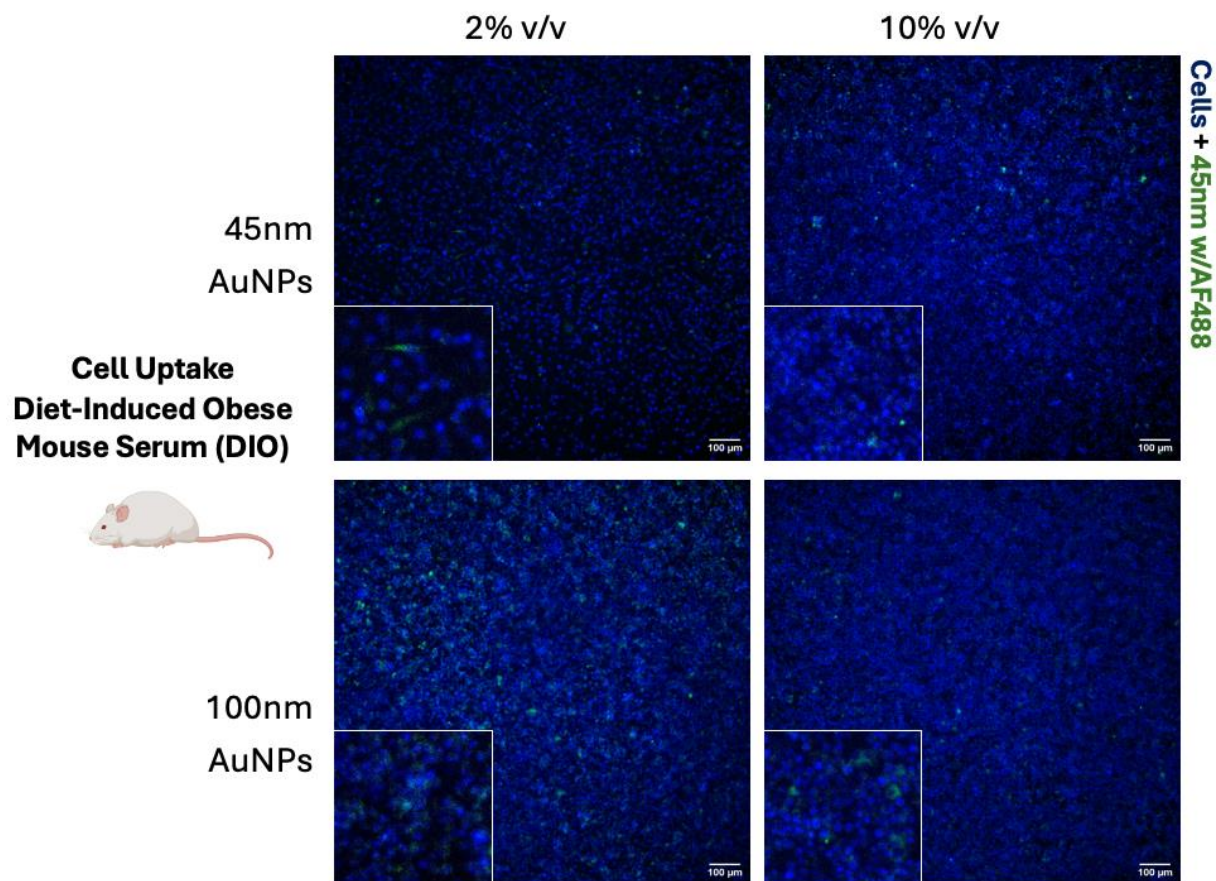


Figure 3.12: Cell uptake between AuNPs in different dose serums for DIO mouse serum

Overall, we see a difference of nanoparticle cell uptake patterns between the HMS and DIO mouse serum conditions. We hypothesize that this may be due to both the phenotypic changes from the serum itself on the RAW264.7 cells as seen with the optical microscopy without nanoparticles. A possibility of the cells having a direct relation on the mechanism of uptake of the nanoparticles due to the different proteins subjected on the DIO mouse serum, which might explain why we observed a lower amount of AuNPs inside of the cells for the obese serum.

3.4 DATA CORRELATION

From the results obtained in our study, we found a great difference in uptake and protein corona formation between the main serums tested, which were the healthy mouse serum and the

diet-induced obese mouse serum. We observed a positive uptake of both proteins in the corona and nanoparticles in the cells with the condition of the HMS. On the contrary, there was not a significant uptake compared to the healthy when nanoparticles were exposed together with the DIO mouse serum. Together, these data support the hypothesis that nanoparticle cell uptake can be different as related to body and disease physiology.

3.5 CONCLUSIONS

With the difference in both uptake and SDS-PAGE protein bands between the DIO mouse serum and the HMS, we can conclude that a lower cell-uptake and variety of proteins in protein corona formation in the DIO mouse serum is seen. From this study, we have opened a slew of new experiments to complement the results obtained, showing that there is a clear difference in uptake in different serum conditions representing body physiologies. With the initiation of this study, a future on understanding of the certain nanoparticle uptake with the differences in physiologies is needed. With additional work, it may be possible to further improve drug delivery and treatments depending on the patient-specific physiological factors.

3.6 FUTURE FORESIGHT

Future progress on this study can be obtained by completing a more thorough analysis on the specific proteins obtained in the obese and healthy protein corona, these include the work with Liquid chromatography – tandem mass spectrometry to specifically obtain the abundance and identities of those proteins. These results can be compared to literature and more sophisticated knockout studies can be performed to directly determine the contribution of different proteins and receptors on the nanoparticle cell uptake process. To improve upon the fluorescence imaging studies, it is suggested to perform more quantitative analysis, including single cell studies such as using flow cytometry. With the promising initial preliminary results on

this study, it should also be considered to use human patients' serum and with different nanoparticle types to observe the differences in a more clinically-relevant condition. It is with great hope, that one day in the near future it would be possible to make individually and personalized medical treatments to each's physiology a reality. This will undoubtedly lead to more efficacious and efficient treatment regimens to benefit patients' health.

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Glossary

AuNPs: Gold Nanoparticles

DLS: Dynamic Light Scattering

UV-VIS: UV-Visible Spectroscopy

TEM: Transmission Electron Microscopy

PRPS: Plasma Resonance Peak Shift

PEG: Polyethylene glycol

mPEG: Methoxy-PEG

amPEG: Amino-PEG

HHS: Healthy Human Serum

HMS: Healthy Mouse Serum

DIO: Diet-Induced Obese Mouse Serum

HDL: High density lipoprotein

LDL: Low density lipoprotein

FBS: Fetal bovine Serum

DTT: 1,4-Dithiothreitol

SDS-PAGE: Sodium dodecyl-sulfate polyacrylamide gel electrophoresis

ICP-MS: Inductively Coupled Plasma – Mass Spectrometry

AF488: Alexa Fluor 488 PBS: Phosphate buffered saline.

Vita

Veronica Gabriela Contreras Guerrero is a Master student who gained interest in Biomedical Engineering by going through the program in Metallurgy and Materials Science in the University of Texas at El Paso. Her interest began with the concentrations offered in the program she was enrolled in, to which biomaterials peaked an interest in her early academic career. By the end of her undergraduate degree, she started to work alongside Dr. Wilson Poon, who gave her the tools and knowledge to start working in a biomedical engineering laboratory setting. With the help of her PI and lab mates, Veronica has gained the ability to use her material knowledge and apply it to a specific field. With the help of her mentors and colleagues, Veronica wishes to pursue more experience in industry, and to later work on her doctorate degree.