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Plasma Proteins Interaction with Curcumin Nanoparticles: Implications in Cancer Therapeutics

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Abstract

Curcumin, a natural bioactive polyphenol, has been widely investigated as a conventional medicine for centuries. Over the past two decades, major pre-clinical and clinical trials have demonstrated its safe therapeutic profile but clinical translation has been hampered due to rapid degradation, poor water solubility, bioavailability and pharmaco-kinetics. To overcome such translational issues, many laboratories have focused on developing curcumin nanoformulations for cancer therapeutics. In this review, we discuss the evolution of curcumin nanomedicine in cancer therapeutics, the possible interactions between the surface of curcumin nanoparticles and plasma proteins, the role of nanoparticle-protein complex architecture parameters, and the rational design of clinically useful curcumin nanoformulations. Considering all the biologically relevant phenomena, curcumin nanoformulations can be developed as a new neutraceutical or pharmaceutical agent.

Keywords

Polyphenol; drug delivery; nanomedicine; cancer therapeutics; bioavailability; protein corona

1. CURCUMIN IN CANCER THERAPY

Cancer remains the second leading cause of death worldwide. The American Cancer Society estimated there were approximately 1,638,910 new diagnoses and 577,190 deaths in 2012 due to cancer in the United States [1]. Present treatment modalities such as surgery, chemotherapy, hormone therapy, adjuvant therapy and radiation have been successful to reduce the morbidity or mortality of cancer. Chemotherapy is a common means of treatment for all stages of cancers. Eventhough the use of chemotherapy typically results in an improved survival rate, there are numerous side effects associated with it. Since ancient days

CONFLICT OF INTEREST

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many natural products have been used to improve human health outcomes. In addition, due to low toxicity and low hypersensitive reactions, use of these natural products increases the patient's compliance and strength. Therefore, there are a growing number of scientists and clinicians interested in exploring the use of traditional or herbal medicine for preventive and therapeutic purposes to control cancer growth. Traditional medicines derived from natural sources come under preventive treatment options according to complementary and alternative medicine by the National Cancer Institute (NCI) of the National Institutes of Health (NIH, USA). In fact, three-fourths of drugs approved by the Food and Drug Administration (FDA, USA) for cancer treatment are either natural compounds or their analogues [2] which exhibit favorable therapeutic potential. Significant data among the herbal medicinal molecules demonstrates that curcumin exhibits antitumor effects against various cancer cell lines and animal models. Curcumin (CUR) is principally a bioactive polyphenol (chemical structure Fig. 1A), derived from the rhizome of Curcuma longa Linn that has proven to have a pleiotropic property. This molecule can target and modulate a number of carcinogenic intracellular signaling pathways that regulate cell growth, inflammation and apoptosis in cancer therapeutics [3–5]. Additionally, curcumin enhances the potency of chemotherapeutic agents and radiation therapies in cancer treatments [6–7]. More importantly, curcumin is nontoxic to healthy cells and therefore does not hinder health care practitioners from using it. Many clinical trials for cancer treatments have investigated the roles of curcumin both alone and in combination with standard therapeutic agents [8, 9]. Ongoing clinical and animal studies have demonstrated the therapeutic significance of curcumin. Curcumin could also emerge as a potential therapeutic and chemopreventive agent in cancer therapeutics, by increasing its solubility, absorption, bioavailability, pharmaco-kinetics and specificity to improve plasma and tissue levels. Recent investigations focusing on developing new approaches based on nanotechnology such as selfassemblies, polymer complexes, curcumin loaded nanoparticles (NPs) or layers of micelles, aim to increase the systemic availability for efficient therapeutics at the tumor site to (1) control and prevent the cancer from progression, (2) detect and image simultaneously, (3) provide adjuvant therapy in addition to conventional therapies, and (4) have multi-functional therapeutic purposes [10] (Fig. 1B).

2. CURCUMIN-HUMAN SERUM ALBUMIN IN DRUG DELIVERY

Orally administered curcumin was found in the liver, kidney and blood in rats due to albumin, an effective carrier [11]. Subsequent investigation further confirmed that curcumin interacts with human serum albumin (HSA) at two distinct binding sites [12]. A Leung and Kee study [13] delineates that among plasma proteins, curcumin binds to HSA and fibrinogen with a binding constant of $1.22 \pm 0.35 \times 105$ M⁻¹ and $5.99 \pm 1.75 \times 104$ M⁻¹, respectively. Such binding phenomena reduce hydrolytic degradation of curcumin by 95% compared to curcumin in PBS. Similarly, a significant reduction of curcumin degradation was also observed in the presence of a 6–8% serum protein in a buffer solution [14]. Curcumin has shown to bind the sub-domain IIA site of HSA [15]. Curcumin-HSA interactions in synchronous fluorescence spectral analysis are 2 fold higher for tryptophan than tyrosine residue in HSA, and proves that curcumin-HSA binding interaction mainly takes place at tryptophan residue in HSA [16]. The curcumin-HSA binding interaction occurs at 2.8 nm

förster critical distance, 3.1 nm binding distance to the tryptophan residue of protein with 2.209×10^{-14} cm³ m⁻¹ overlap integral and 0.35 energy transfer efficiency [17]. Upon curcumin binding, it alters the content of the secondary structure of HSA, i.e., a-helix structure increased while the random coil percent decreased. All investigations and literature proposed that HSA can act as an efficient transporter for the curcumin molecule [18]. The physiological role of HSA is widely known and it can bind to numerous lipophilic molecules that may lead to an increase in its pharmacology profile [19]. In a review article, Xiao et al. [20] concluded that the effect of the polyphenol-HSA interaction on biological activity is not equivocal and depends entirely on the structure-affinity relationship. Although attempts have been made to improve the solubility, stability, and degradation of curcumin in vitro, delivering it in vivo in the presence of organic solvent/surfactant (needed to dissolve curcumin) which are may be toxic to humans. Additionally, achieving optimal therapeutic concentration of curcumin at the tumor site is highly difficult using the HSA-binding or HSA-complex transportation method because no specific target mechanism is involved. Alternatively, nanoparticle (NP) carriers have been chosen for effective delivery of curcumin [10]. However, a number of studies showed that relatively small fractions of NPs reached the bloodstream due to hazardous agglomeration or cluster formations [21]. In fact, many curcumin and anti-cancer therapeutic nanoformulations have a bioavailability less than 5% injected dose volume (IDV) at tumor sites. Therefore, bioavailability of curcumin NPs is strongly dependent on the extension of adsorption of proteins in blood. The Centre for BioNano Interaction at University College Dublin, Belfield (Dublin, Ireland) has initiated studies which examine the interaction of NPs with various plasma proteins [22]. Based on these observations, a detailed interaction of curcumin NPs with plasma proteins and implications for their use in cancer therapeutics will be discussed in forthcoming sections.

3. NANOPARTICLES-PLASMA PROTEINS BINDING: GENERAL VIEW

3.1. Nanoparticles-plasma Proteins Interaction in Drug Delivery Applications

The use of NPs has tremendously increased for drug delivery applications. However, it is a highly challenging task to design a NP system that exhibits superior *in vivo* behavior. Many of the conventional nanoformulations may be compromised by the reticuloendothelial system (RES) prior to reaching the tumor tissue or the blood brain barrier. For improving such biological phenomenon, several methods have been implemented to control particle size and increase the stealthiness of nanoformulations. Curcumin NPs are considered to be a novel alternative therapeutic option in cancer therapeutics [20, 23, 24]. Studies have confirmed that NPs are robustly bound with plasma proteins, cells or tissues and are never found in the form of parent particles [21]. After adsorption of plasma proteins, NPs may either lead to beneficial or harmful biological actions. Therefore, in order to develop effective translational therapeutic outcomes for curcumin NPs, it is essential to learn its complex role in the biological plasma system.

In general, NPs contain a larger active surface area than their origin bulk material. Active surfaces of NPs are first exposed and begin to interact with human plasma proteins when they are in blood circulation [25, 26]. During this process, a selective exchange and

interface of the protein corona on NPs are constructed either with "short" or "long" lived structures which can be determined by the state of exchange of proteins. For superior drug delivery applications, NPs must be generated to allow the formation of hard corona which will likely improve the interface between NPs, formulations and biological milieu.

3.2. Evaluation of Nanoparticles-plasma Proteins Interaction

Dell'Orco *et al.* [30] proposed a model after considering the time evolution of the NPprotein corona formation in body fluids (HSA, high density lipoprotein and fibrinogen) with 70 nm polymer NPs. This model provides binding data fitted by a simple binding model of inter-protein interaction phenomena and the dynamics of the NP -protein interactions of a reversible biochemical equation (Equation 1). The overall biochemical system differential equation and the time evolution of individual proteins and NP complexation was assessed by numerical kinetic models carried out by SBTOOLBOX2 (Matlab, http:// www.sbtoolbox2.org) (Equation 2). This modeling technique can further extend to evaluate the specific molecular interactions of nanoparticle-cell/tissues. NP, PR, and t represent concentration of nanoparticles, proteins and time of measurement, respectively. The association and dissociation of NPPR complexes were represented as concentrations, individual values of the rate constants κ^{on} , κ^{off} , and the stoichiometry in a single layer.

$$NP + PR \leftrightarrow NP.PR$$
 (1)

$$\frac{d[NP.PR]}{dt} = nPR\kappa PR^{on}[NP][PR] - \kappa PR^{off}[NP.PR] \quad (2)$$

The charge/density of NPs and plasma protein coronas may also have a major biological impact for *in vivo* applications [31]. Fibrinogen showed higher association with NPs at a higher membrane charge density while apolipoproteins and C4b-binding proteins bound NPs showed less membrane charge density. Charge density can be calculated from the relative amount of charge on the protein corona by multiplying the total charge on the protein κ ($q\kappa$), probability threshold (Φ) by protein abundance, molecular weight of normalized scaffold for protein κ (MWNSC κ):

$$Q(\Phi) = \sum_{k=1}^{n} \mathbf{q}\kappa \times \mathrm{MWNSC}\kappa(\Phi)$$
 (3)

Recently, quantitative proteomics analysis has gained much attention as a means to evaluate the adsorbed plasma proteins on NPs [32–34]. Zhang *et al.* [32] have studied different surface modifications and particle sizes of NP interactions with plasma proteins. All NPs showed significant binding with 42 plasma proteins. Only one out of three clusters from these 42 proteins exhibited a different protein binding pattern with different surface modifications. Two clusters of proteins presented opposite patterns of relative abundance

As expected, when NPs make contact with biological fluids, the surfaces are coated with proteins which change their surface charge and properties. However, high NP protein binding is not always disadvantageous. In some cases, these interactions may lead to the delivery of NPs to a specific organ or tissue. For instance, a paclitaxel nanoformulation [(albumin-coated paclitaxel nanoformulation (nab-PTX) (Abraxane®)] showed specific interactions with secreted acidic and rich extratumoral proteins help with targeted delivery of paclitaxel [36, 37]. Further, such interactions efficiently modulate the transportation of paclitaxel across endothelial barriers through gp60 receptor and caveolin-1 mediated pathways [38].

to identify new therapeutic targets in cancer via proteins bound to NPs [35].

Dell'Orco *et al.* proposed that theoretical systems-level analysis of the protein corona of plasma protein formations on the engineered and functionalized surface of NPs which have specific biological targets is often neglected [39]. Their suggestion for effective targeted delivery of drug(s), in addition to the biophysical quantities of early/hard/dynamic corona, is that systems-level mathematical approach generated stoichiometry and the kinetics of the relative interactions must be evaluated with a particular tissue(s). Global sensitivity analyses of selected dynamic conditions of the network models will improve the targeted drug delivery success rate through nanomedicine. We believe such global sensitive models can also be implemented for effective delivery of curcumin to the tumor site for an enhanced therapeutic potential.

3.3. Techniques Used to Investigate Nanoparticles-plasma Proteins Interaction

The molecular structure and affinity relation of curcumin with serum proteins, observed as non-covalent binding, can be investigated by the binding constants obtained from spectral, dynamic light scattering, microscopic analysis and other analytic methods [22, 40–43]. In the case of curcumin nanoparticles, bound plasma proteins were separated by normal or gradient centrifugation, and then pellets containing curcumin nanoparticle-plasma protein corona (Fig. 2) are subjected to various analytical methods for accurate evaluation. Gebauer and co-workers demonstrated that the protein corona of HSA around citrate-functionalized silver NPs stabilized the particles without inducing agglomeration and protein structure changes [44]. CD spectral analysis can provide signals of proteins arising from electronic transitions in secondary structural elements such as α -helix, β -helix, turn and random coils. The quantitative analysis of these data provides insight into the alterations which occurred in the original protein structure by the interaction. Atomic force microscopy is a relatively new technique that can provide additional information on currently used methods. This technique delineates the amount of proteins present on NPs and the force of adhesion and surface free energy on the particles before and after modification with proteins [45]. Hydrophobicity in NP composition is also highly favored for extensive binding of plasma proteins [46]. Lindman et al., [46] demonstrated that a fully covered HSA layer was achieved by hydrophobic NPs but at a lower curvature. These results were confirmed experimentally by isothermal titration calorimetry, yielding stoichiometry, affinity, and enthalpy changes from

the NP and HSA binding. Most of the NP and plasma protein interaction studies were evaluated by fluorescence quenching or the decay of fluorescence of proteins. These measurements can be achieved by steady-state absorption, synchronous fluorescence, threedimensional (3D) fluorescence and circular dichroism (CD) spectroscopic techniques [43]. In addition, one and two-dimensional polyacrylamide gel electrophoresis (1D and 2D-PAGE), liquid chromatography mass spectrometry (LC-MS) and liquid chromatography mass spectrometry/mass spectrometry (LCMS/MS) can also help to quantify NPs and

plasma proteins interaction. Table 1 illustrates various analytical techniques used to evaluate nanoparticles and plasma proteins interactions.

4. CURCUMIN NANOPARTICLES-PLASMA PROTEIN CORONA IN CANCER THERAPEUTICS

Deeper penetration of curcumin NPs into cancer cells can be achieved through proper protein corona formation on the surface of NPs. Based on the plasma protein interaction with curcumin NPs, their *in vivo* biologically relevance can be determined (Fig. 3). Table 2 illustrates how curcumin NPs can significantly show increased bioavailability *in vivo* [10, 49–63] due to their favorable plasma protein corona formation i.e., less number of proteins are required to increase the NPs in blood circulation [47, 48]. The NP-protein corona is highly associated with cellular uptake in cancer cells [35]. Modulation of protein corona around NPs is a promising approach and an effective therapeutic target. Serum proteins have ensured a role in the feeding process which permits cellular entry through contacting bare NP surfaces. Such types of internalization include the uptake of NPs or drug loaded NPs in living cells followed by (1) passive diffusion, (2) endocytosis, and/or (3) phagocytosis. Phagocytosis occurs when NP-plasma protein interactions make particles larger in size, typically several hundred nanometers to microns, due to aggregation. Endocytosis of NPs is possible through an internalization mechanism in which NPs are coated slightly with human serum proteins.

Lundqvist *et al.* [28] suggested a possible NP trafficking scenario in which the role of protein corona formation is highly significant. The adsorbed-protein complexes on NPs are dynamically changed from one biological environment into another by replacing proteins from the new biological fluid. In the end, the NP and adsorbed protein may control signals that can be explained from the protein corona. In contrast, Jiang *et al.* [64] revealed that in the presence of $100 \,\mu$ M transferrin or HSA coating on 1 nM NPs, the amount of NPs on the cell surface, as well as in the intracellular binding, was substantially reduced. This behavior was further verified through confocal analysis of a group of 15 cells from two to three independent experiments. Interestingly, lower intracellular fluorescence was observed from the NPs than the membraneassociated particles. The media used for *in vitro* experiments has a major role in internalization of curcumin NPs in cancer cells. Gold NPs suspended in DMEM media display a red shift and increased band intensity over time [65]. Such significant changes were not noticed in RPMI, which indicated DMEM endorsed abundant protein corona on the NPs. This in turn influenced the uptake and safe internalization in U937 and HeLa cells [65].

Recent findings have reported on the roles of particle aggregation and plasma protein coating on cell uptake by the iron content estimation [66]. According to this study, at low plasma concentrations NPs form clusters by proteins like fibrinogen, whereas at high plasma concentrations NPs stabilized by apolipoprotein coating. However, magnetic fractionation was useful to separate clusters and stabilize NPs which have a specific protein fingerprint. Overall, these results indicate that administration of a single population of NPs (protein-NP complexes) may be helpful to increase the availability in bloodstream. A drug delivery system with an entrapped cargo with transferrin corona spontaneously bound to a transferrin receptor (TfR) and displayed a superior cellular uptake and delivered payload to the cytosol [67]. Mulik et al. [68] demonstrated a significant increase in the uptake of curcumin NPs covered with transferrin protein corona in MCF-7 breast cancer cells. This condition greatly promoted curcumin activity by inducing increased loss of mitochondrial membrane potential and generation of excessive reactive oxygen species leading to superior apoptotic characteristics. Another formulation, namely superparamagnetic iron oxide NPs with transferrin composition, showed efficient targeting of curcumin against K562 myeloid leukemia cancer cells [69]. Further, curcumin loaded fibrinogen nanocarriers showed a preferential internalization and stimulated the apoptosis pathway in MCF-7 cancer cells compared to L929 fibroblast cells [70]. Similarly, a number of curcumin NPs decorated with plasma proteins have been documented as possessing improved therapeutic potential in cancer cells [10].

One of our studies has delineated the efficient internalization of PLGA-CUR NPs in cancer cells [71]. The serum corona drives curcumin NPs in a time and cell type dependant internalization. This implies that internalized NPs may release curcumin in its native active form for therapeutic purpose. Additionally, the NPs designed by PLGA with a polyvinyl alcohol stabilizer have the capability to escape from late endosomes and deliver a larger amount of drug to the cytosol [72]. Among five curcumin nanoformulations (Fig. 4A) tested for HSA protein binding, the PLGA based nanoformulation has the lowest protein binding which suggests it is a good carrier for delivering curcumin to target site(s) over an extended period (Fig. 4B) [73]. Higher HSA protein binding was noticed with dendrimer curcumin formulations which can lead to aggregation of the formulations over time and probably elimination from the body. However, the adsorption of different plasma proteins onto curcumin NPs varies, and in all types of formulations fibrinogen and immunoglobulin G adsorption was found to be very low. Overall, the degree of protein adsorption to curcumin NPs is as followed: HSA > transferrin > immunoglobulin G > fibrinogen. Furthermore, our un-published data on HSA binding on PLGA NPs stabilized by PVA in combination with other polymers, F127, F68, dextran, hydroxyl cellulose, chitosan, and gum acacia, did not show significant differences (Fig. 4C).

Jedlovszky-Hajdú *et al.* [74] investigated the influence of composition and structure of magnetic NP interactions with human plasma. The citric acid (CA), poly(acrylic acid) (PAA), and oleic acid double layer (OAOA) coated magnetic NPs have shown increased NP size in soft (10% serum) and hard corona (55% serum) determined by Dynamic Light Scattering (DLS), NP Tracking Analysis (NTA), and Differential Centrifugal Sedimentation (DCS) methods [74]. More importantly, OAOA and CA have clearly shown a variation in

protein coronas associated with lipid carrier proteins, complement factors and immunoglobulin after exposure with 10% and 55% plasma [74]. From this it can be considered that oleic acid layered magnetic NPs have enriched lipoproteins and albumin while complement and immunoglobulin proteins likely escape the immune system and RES. With this in mind, we have generated oleic acid/cyclodextrin layers with Pluronic polymer stabilizers for delivery of curcumin/doxorubicin [75, 76].

It was demonstrated that the structure of proteins was significantly altered upon incubation with polystyrene NPs [77]. No charge and negatively charged NPs encouraged formation of the helical structure in apoAI or HDL (negatively charged) while positively charged NPs decreased the amount of helical structure. Similarly, the secondary structure is affected by NPs carrying an opposite surface charge relative to the protein. This indicates that surface charge of NPs and proteins are critical parameters for predicting protein adsorption and corona formation. In our comparative plasma protein binding, the dendrimer curcumin nanoformulation significantly alters secondary structures while other formulations did not have considerable influence (Fig. 5). This severe binding potential is achieved because of the positive nature of dendrimer curcumin formulations.

5. FACTORS INFLUENCE CURCUMIN NANOPARTICLEPLASMA PROTEIN CORONA

Design and engineering of the NP–protein interface itself is a broad field in drug delivery. In our opinion, protein corona formation on curcumin NPs is dependent on the physicochemical properties of NPs (Fig. 6). A recent review [78] presents that the protein corona formation on nanoparticles depends on (1) the characteristics of the nanomaterial, such as nanoparticle size, shape, and composition, (2) the physiological environment, such as blood, interstitial fluid, cell cytoplasm, etc., and (3) the physiological response, including kind of transport, signaling mechanism, pharmacokinetic, accumulation process, and toxicity. This review also discussed in detail approximately 125 plasma proteins that have higher association with various nanoparticles.

The bound plasma protein corona on curcumin NPs is strictly a highly complex phenomenon, as such, the corona contained bound protein cannot be presumed based on profusion in the plasma and particle size. Commonly, PEGylated NPs protect from over plasma binding and avoid clearance by the RES mechanism. Kodiyan *et al.* [79] successfully designed NPs coated with cysteine-functionalized alginate based polymers which exist up to 4 weeks in circulation of peripheral blood in mice. Such plasma protein repellent stabilizers can be employed as an alternative to overcome the existence of limitations of PEG coating or PEGylation in therapeutic applications. A recent report of NP–histodine complexes demonstrated that the orientation, geometry and stoichiometry and its specificity can be regulated by the size and binding sites on the NP's surface [80].

In some instances, plasma proteins adsorbed on the surfaces of NPs enhance the interaction with partially unfolded proteins that are directed toward cluster formation [81, 82]. These radical transformations favor extensive development of fibrils that promote pathological conditions ranging from neurodegenerative disorders to systemic amyloidoses [83]. Our

studies have demonstrated that curcumin nanoformulations made with PLGA NPs, cyclodextrin assembly, cellulose NPs, magnetic NPs, and nanogel do not show any signs of fibril formation in a transmission electron microscopy evaluation [73] (Fig. 7). However, the dendrimer curcumin formulation exhibited linear filament formation but not fibrils. In addition, chemical surface modification of NPs with small molecules that can target proteins of the associated corona in biological milieu can stimulate the protein misfolding phenomenon [84]. This misfolding phenomenon allows access for cell specific internalization.

PLGA NPs have exhibited the adsorption of three major plasma proteins: fibrinogen > albumin > globulin (γ -globulin) [85]. Thiol modified PLGA NPs (57.16 ± 6 8.5 µg –SH groups/100 mg of particles) resulted in maximum adsorption reduction with globulin (up to 20% reduction) while only 10% reduced adsorption was noticed with albumin and fibrinogen. PLGA-CUR NPs with smaller sizes had a significantly higher uptake (2–6 fold) in A2780CP and MDA-MD-231 cancer cells which is responsible for the superior anticancer potential [71]. A recent study suggested that proteins can adopt different orientations depending on the size of the NPs [86]. Different orientations of adsorbed proteins affect the level of biological activity of the formulation. Small particle size formulations were found not to be precipitated or bound to protein A, which illustrates the principle that the size of NPs can strongly influence the binding activity of adsorbed proteins. An extensive Pathway and Network Analysis Software of Genespring GX and the Ingenuity Study revealed that not only does the particle size of formulations determine its protein binding capacity but also its various functional components involved in the biological system such as immune response, lipid and cholesterol metabolism, activation, coagulation and acute response [34].

We have shown in our recent publication that surface chemistry plays a significant role in the formation of protein corona on NPs [41]. These structural variations lead to different uptake profiles in MDA-MB-231, HPAF-II and SKBR-3 cancer cells. Wiogo *et al.* [87]. validated that surface chemistry assisted protein corona formation. Branched polyethylene glycol coated magnetic NPs binds ~ 95% serum protein while polymethacrylic acid coated magnetic NPs binds ~ 10% serum protein. Linear polyethylene glycol coating also helps to reduce serum binding to 40% compared to un-coated NPs. The higher serum bound protein corona on the magnetic NPs can also cause a reduction to their MRI characteristics. Additionally, these inherent properties are also dependent on the type of corona formed on NPs. Casals *et al.* [42] examined the hard and soft corona density in terms of thickness by the dynamic light scattering method. They concluded that soft corona has more thickness than hard corona because soft corona densities were correlated with particle size.

Hemolysis is a measure of an acute toxicity assay that indicates the extent of damage to red blood cells *via* release of hemoglobin. This assay is widely used to evaluate the hemocompatibility of the nanoformulation. To assess the curcumin nanoformulations for human use, the formulations must be tested for hemolytic compatibility. Curcumin nanoformulations demonstrated almost no hemolysis existed at the concentrations tested compared to the positive controls (up to 1 mg/mL) mL) [70, 76, 88]. The noted hemolytic ratio of curcumin formulations was < 3–4% (below 5%) which is considered a critical safe

ratio for nanomaterials or biomaterials according to ASTM F756 and ISO/TR 7406. This suggests that the risk of hemolysis by curcumin nanoformulations is completely absent. Blood-NP interactions by TEM and SEM studies revealed that NPs can interact with platelets, red blood cells (RBCs), and white blood cells (WBCs) [89]. A study showed that gold NPs not only attached to these cell surfaces but were engulfed in small channels in the cell membranes regardless of their surface properties [89]. However, RBC membrane damage may lead to increased spleen uptake due to the damaged RBCs recycling process. Therefore, design of curcumin NPs that are highly associated with RBCs without damage would be highly suitable for cancer therapeutics. PLGA NPs are widely employed for curcumin delivery. PLGA NPs have shown a lower aggregation tendency in blood cell fractions and are almost equivalent to the control RBCs in terms of membrane integrity [85].

Targeted motif mediated delivery of curcumin NPs is reportedly limited at this time and their plasma protein binding is unknown. However, a combination of inert fresh anti-fouling surfactants and bioactive moieties (*via* ligand, aptamer, penetrating peptide or protein, and antibody mediation) of particular delivery mechanisms will increase cancer therapeutic potential [89]. Our preliminary data on curcumin loaded magnetic particles showed improved tumor uptake that could be due to optimum plasma protein corona formation in the HPAF-II xenograft mouse model (Fig. 8). The targeting can be further improved by antibody or aptamer conjugation to NPs for treatment of pancreatic cancer.

6. FUTURE CONSIDERATIONS FOR CANCER THERAPEUTICS

Understanding interactions of curcumin NPs and plasma proteins in the pharmaceutical fields of research is important. Structural, functionality, configuration and morphological alterations occur during the formation process of plasma protein coronas which involves a number of biological phenomena that determines the *in vivo* fate of curcumin NPs. Some of these interactions can protect the integrity of curcumin NPs and their ability to reach the intended target organs. However, to determine such phenomena by any one specific method is difficult because NPs are made with different chemical substances and surfactants. The secondary interactions may be dominant in some cases. Studying complex curcumin NP and plasma protein biological interactions should account for: (i) physicochemical properties of curcumin NPs, (ii) thermodynamic feasibilities (enthalpy and entropy driven process where H < 0, S > 0 and Gibbs free energy G < 0), and (iii) electrostatic interactions [43]. Further, it is important to develop unique rational methods to evaluate NP and protein biointerfaces. These studies may become landmark modules to evaluate curcumin NP-plasma protein interactions which can be extrapolated for their in vivo utility and for cancer therapeutics. Investigation of complement system recognition protein (C1_a, ficolins, mannose binding lactin) interactions with nanocurcumin particles will determine its opsonise characteristics. This information will enhance the specificity towards tumors, provided the NP avoids interactions with complement system recognition proteins.

There are 200 types of cells with distinguished membranes responsible for various cellular uptake. Plasma protein corona on NPs recognized by the cell may depend on the extent of internalization by the "sees-cell observer" mechanism [90]. In this way, uptake of encapsulated drugs in NPs is robustly connected to their transport through the cellular

membrane. However, no known specific incubation time of NPs and plasma proteins have been reported for predicting such internalization in cancer cells.

Both legislative policies and the pharmaceutical industry need to encourage academic and translational research that is focused on the NP mediated delivery of curcumin. These policies can expedite novel findings related to the use of curcumin nanoformulations for cancer therapeutics. Further, these policies can also help to increase the utilization of various polyphenols, such as resveratrol, quercetin, epigallocatechin-3-gallate, and genistein. The McNeil group [91] proposed an overview of the number of *in vitro* assays in addition to protein binding NPs that needed to be tested for NP compatibility with the immune system, including hemolysis, platelet aggregation, plasma coagulation, complement activation, phagocytosis, leukocyte proliferation, nitric oxide production, and chemotaxis CFU-GM test. The next important consideration is to develop a curcumin nanoformulation utilizing existing and approved technology, such as Abraxane or Doxil, which follows good laboratory and manufacturing (cGLP and GMP) practices. This process will help to obtain the FDA's approval for curcumin NPs as a New Drug Application (NDA), provided there is enough clinical and preclinical significance for the treatment of cancer(s).

CONCLUSION

Interaction of curcumin nanoformulations with serum plasma proteins can provide a new platform for the interpretation of NP-interfaces from the biological perspective and to estimate the formulation's fate *in vivo*. The method, polymer, and surfactants used to prepare curcumin nanoformulations, as well as the particle size, aggregation phenomenon and surface charge of particles can influence the binding of plasma proteins. It is necessary to develop a standard analytical procedure and optimization for evaluating nanoformulation and plasma protein binding. The discussed curcumin nanoformulations and plasma protein binding are foundational to establish a rational basis for developing improved curcumin nanoformulations for cancer therapeutics.

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Fig. (2).

Schematic representation of sample preparation from nanoparticles-plasma proteins corona for estimation of bound plasma proteins. Incubation time, centrifugation time, and number of washing for obtaining nanoparticle-plasma protein corona varies depend on the specific protocol.

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Fig. (3).

In vivo fate of curcumin nanoparticles after interacting with human serum or human plasma proteins.



Fig. (4).

Influence of plasma proteins interaction with curcumin nanoformulations. (**A**) Structurally varied curcumin nanoformulations generated by different methods. Curcumin nanoformulations (20 µM) were incubated in 100 µg human plasma proteins and after 2 hours of incubation, adsorbed proteins were separated by ultra centrifugation and sodium dodecyl sulfate-polyacrylamide gel electrophoresis was run at 150 V for 60 minutes and resultant stained with Coomassie® G-250 stain. (**B**) Fibrinogen, immunoglobulin G (IgG), transferrin, and serum albumin bound to curcumin nanoformulations. (**C**) HSA associated

with PLGA nanoformulations which are prepared in the presence of various stabilizers. CD: β -cyclodextrin; CUR: curcumin; PLGA: poly(lactide-*co*-glycolide); Nanogel: poly(N-isopropyl acrylamide); Dendrimer: polyaminoamide (4 generation) based curcumin NPs. F127 and F68 are Pluronic polymers and HC indicates hydroxy cellulose. Figure 3A–B has been reprinted with permission from Dove Medical Press (Copyright Clearance Center, Order License No: 3059490805921) See Ref. 73.

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Fig. (5).

Curcumin NPs and serum albumin complex formation in PBS solution. Serum albumin (12.5 μ M) was incubated with curcumin nanoformulations (0–50 μ M) and the circular dichroism spectra were obtained at room temperature. CD: β -cyclodextrin; CUR: curcumin; PLGA: poly(lactide-*co*-glycolide); MNP: magnetic NPs; Nanogel: poly(N-isopropyl acrylamide); Dendrimer: polyaminoamide (4 generation) based curcumin NPs. Note: Large variation in circular dichroism spectra of dendrimer curcumin formulation indicates significant structural changes occurred in serum albumin.



Fig. (6).

Schematic representation of the influence of various structural, chemical and biological parameters on protein plasma corona formation on curcumin NPs.

200 nm



Fig. (7).

Human serum interaction leads to change in the morphology of curcumin nanoformulations. Curcumin nanoformulations (1 mg/mL) were incubated with and without human serum albumin (100 μ g) for 2 hrs and after centrifugation and washing with water, curcumin NPs were viewed under a transmission electron microscopy. CD: β -cyclodextrin; CUR: curcumin; PLGA, poly(lactide-*co*-glycolide); MNP: magnetic NP; Nanogel: poly(N-isopropyl acrylamide); Dendrimer: polyaminoamide (4 generation) based curcumin NPs. Filament formation was observed only in dendrimer curcumin formulation (bar arrows represent filaments). Reprinted with permission from Dove Medical Press (Copyright Clearance Center, Order License No: 3059490805921) See Ref. 73.



Fig. (8).

Curcumin loaded magnetic nanoparticle (MNP-CUR) formulation targets tumor cells in HPAF-II xenograft mouse model. (A) Schematic representation of possible targeting behavior of curcumin NPs to tumor cells. (B) MNP-CUR internalization in tumor cells *in vivo*. MNP-CUR (20 µg per mice) was administered *via* intraperitoneal injection and internalization in tumor cells was detected with Prussian blue staining. Black arrows indicate MNP-CUR NPs internalization in tumor cells.

Table 1

Various Analytical Techniques Used to Evaluate Nanoparticles and Plasma Proteins Interactions

Analytic Method	Type of Evaluation and Comments	
1D-polyacrylamide gel electrophoresis (PAGE) and 2D PAGE	Identification of proteins in corona on nanoparticles. The order of protein adsorbed onto the surface of nanoparticles can be determined. With the help of densitometry, the half-max protein adsorption and hill slope for protein. Highly laborious and not helpful/useful for quick kinetic adsorption evaluations.	
BSA Quantification	Half-max protein adsorption. Hill slope for protein adsorption.	
UV-Vis extinction spectroscopy	Half-max protein adsorption. Hill slope for protein adsorption. Apparent thickness, orientation and refractive index of the adsorbed proteins. Simultaneous protein adsorption measurements.	
Gel permeation chromatography	Determination of molecular weight of plasma proteins bound to nanoparticles. Comparative evaluation of bound proteins on nanoparticles. Requires larger volumes of samples for analysis.	
LC-MS/MS	Efficient separation of bound plasma proteins. Identification of protein/peptide sequence. Accurate molecular weight distribution. Requires larger volumes of samples for analysis.	
Capillary electrophoresis	Analysis of protein–nanoparticle interaction with high degree of resolution. Efficient separation of plasma proteins and very small sample volumes are required for analysis. Determination of the fate of nanoparticles <i>in vivo</i> including delivery to tumors due to their interactions with specific plasma proteins. Stable nanoparticle-protein complexes and transient complexes were resolved by capillary zone electrophoresis and affinity capillary electrophoresis. This technique is also used in combination with other analytical methods to evaluate each type and percentage of various bound proteins on nanoparticles.	
Atomic force microscope	Measure of change of indention before and after protein corona <i>in situ</i> modification. Determined the work of adhesion. Evaluation of surface free energy during protein corona formation. Force of adhesion involved in protein corona formation.	
Fluorescence correlation spectroscopy	Used to quantitatively monitor protein binding or adsorption onto fluorescent nanoparticles. Correlation analysis of the fluorescence emission time traces yields a characteristic time scale of diffusion from the hydrodynamic radius. Size of the nanoparticles after protein deposition on their surfaces can be monitored. Nanomolar concentrations of the nanoparticles in microlitre-sized volumes required.	
Particle Size and Distribution (TEM and DLS)	Plasma protein corona structure examination. Particle's dispersibility before and after protein corona formation on nanoparticles. Particle's zeta potential before and after protein corona formation on nanoparticles.	

Table 2

Improved Bioavailability of Curcumin with Curcumin Nanoformulations

Formulation	Study Outcome	
	Dose and Type of Administration	Improved Bioavailability
Curcumin loaded poly(lactide- <i>co</i> -glycolide) (PLGA) NPs [49]	250 mg/kg (curcumin), 250 mg/kg (curcumin) + 10 mg/ kg(piperine), and 100 mg/kg (PLGA NPs) body weight of rat Oral	The <i>in vivo</i> oral bioavailability was increased up to 9- fold compared to curcumin+piperine composition.
PLGA and a stabilizer polyethylene glycol (PEG)-5000 [50]	2.5 mg/kg body weight of mice Intravenous	Curcumin NPs or curcumin were maintained at 430– 100 ng/ml and 275–25 ng/ml in serum for 1–24 hrs. All time points tested curcumin NPs showed at least 2 fold more bioavailability.
Nanocrystal solid dispersion (CSD-Cur), amorphous solid dispersion (ASD-Cur), and nanoemulsion (NE-Cur) [51]	100 mg/kg body weight of rat for curcumin and 20 mg/kg body weight of rat for CSD-CUR, ASD-CUR, and NE-CUR Oral	Oral bioavailability was increased from 0.9 (cur- cumin) to 14.3, 10.7, and 7.9% for CSD-CUR, ASD- CUR, and NE-CUR formulations, respectively.
Curcumin nanosuspension with D-a - tocopheryl polyethylene glycol 1000 succinate (TPGS) [52]	15 mg/kg body weight of rabbit Intravenous	Approximately 3.8-fold greater bioavailability $(145.42 \pm 9.29 \ \mu g/mL \ min)$ was achieved with a mean residence time of $194.57 \pm 32.18 \ (11.2$ -fold longer vs. curcumin).
Glycerol monooleate-poly(vinyl alco-hol)- pluronic (F127) polymer NPs [53]	30 mg/kg body weight of mice Intravenous	At 1–24 hrs time points, curcumin NPs exhibited 17.5-4 μg/ml curcumin in serum while native cur- cumin was barely detected (0.2 μg/ml serum).
Nanocur™ [54]	25 mg/kg body weight of mice Intraperitoneal	> 8 µg/ml plasma levels of Nanocur TM was present between 1 to 8 hrs whereas free curcumin in corn oil was barely detectable.
Solid lipid NPs C-SLNs: very high (VH); high (H); medium (M); and small (S) and free curcumin (C-S) [55]	50, 50, 25, 12.5, and 1 mg/kg body weight of rat for C-S, VH, H, M, and S, respectively Oral	Approximately, 0.292, 14.29, 8.00, 7.87, and 1.00 μg/ ml of maximum of curcumin was present in serum by C-S, VH, H, M and S administrations, respectively.
Organogel-based nanoemulsion [56]	240 mg/kg body weight of mice Oral	8.5–9.8 fold increase in bioavailability.
PLGA-CUR LMw and HMw nanofor- mulations [57]	50 mg/kg body weight of rat for LMw and HMw nanocurcumin formulations and 1000 mg/kg body weight of rat for free curcumin Oral	No significant difference in bioavailability between LMw and HMw curcumin nanoformulations but 1.67 and 40-fold higher bioavailability compared to free curcumin
Solid lipid NPs [58]	400 mg/kg body weight of mice Intraperitoneal	20 micro molar curcumin was noticed at 30 min.
Poly(e–caprolactone)-b-poly(ethylene glycol)- b-poly(e–caprolactone) triblock copolymer NPs [59]	15 mg/kg body weight of rat Intravenous	Increased mean residence time from 0.169 to 40.148 hours and the area under the concentration- time curve increased 4.178-fold.
PLGA-CUR [60]	7.5 mg/kg body weight of rat Intravenous	PLGA-CUR was found to be 6.139 mg/L h.
PLGA and PLGA-poly(ethylene glycol) (PEG) [61]	50 mg/kg body weight of rat Oral	PLGA and PLGA-PEG NPs increased the curcu-min bioavailability by 15.6- and 55.4-fold, respectively.
Silica-coated flexible liposomes as a nanohybrid [62]	50 mg/kg body weight of rat Oral	The bioavailability of CUR-SLs and CUR-FLs was 7.76- and 2.35-fold higher, respectively.
Lipid based oral formulation [63]	250 mg/kg rat Oral	Improved C_{max} and AUC _{0-∞} by 11.6 and 35.8 fold, respectively, over control.