Therapeutic Effects of Plant-Extract Synthesized Gold Nanoparticles against Alcohol-Induced Inflammatory Activity - A Proof of Concept

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Abstract

The present study describes potential beneficial and adverse effects of plant-extract synthesized gold nanoparticles (AuNPs) on ethanol toxicity in SH-SY5Y cells. Although Kudzu root extract (K), edible-gum extract (G), alone or in combination (KG), reduced Au$^{3+}$ into AuNPs, the extract’s composition and the reaction temperature determined their size (AuNP$_{KG}$ ($90 < 50 < 37$) $<_{<}$ AuNP$_{K}$ ($90 < 50 < 37$) $<_{<}$ AuNP$_{G}$ ($90 < 100$)), the subscript KG, K or G is extract identification and numerical are reaction temperature in Celsius) and biological properties (AuNP$_{KG}$ ($90 < 50 > 37$) $<_{<}$ AuNP$_{K}$ ($90 < 50 > 37$) $<_{<}$ AuNP$_{G}$ ($90 < 50$)). The surface of each AuNP contained the extract’s active ingredients that were analyzed and confirmed using LDI and low-matrix MALDI. AuNP$_{KG-50}$ was (i) Least toxic to SH-SY5Y cells, but and most effective in suppressing the adverse effects of ethanol on SH-SY5Y cells and (ii) More effective than a combination of free kudzu and gum extracts. The beneficial and adverse effects of AuNPs may have been modified by the formation of proteins corona. This study provides a proof of concept for possible application of plant-extract synthesized AuNPs in mitigating alcoholism and ethanol toxicity.

Keywords

Ethanol, Gold nanoparticles, Plant extracts, Kudzu root, Edible gum, Cell viability, Apoptosis, Necrosis, Lactate dehydrogenase, Lipid peroxidase

Introduction

Alcoholism, a complex illness of the Central Nervous System (CNS), impacts the lives of millions around the world. Untreated alcoholism devastates families, depresses economic vitality, and burdens the country’s health care systems, costing the U.S. over $249 billion annually [1-4]. Although possible mechanisms underlying the etiology of alcoholism are not fully understood, a critical role of opiate, Gamma Amino Butyric Acid (GABA), serotonin and NMDA/AMPA/Kainate receptors has been implicated [5-7]. Consequently, FDA has approved four drugs for treatment of alcoholism: naloxone (an opiate antagonist), acamprosate (GABA receptors), disulfiram (acetaldehyde dehydrogenase inhibitor), and topiramate (AMPA/Kainate receptor), that can be used alone or in combination with psycho-behavioral therapies and counseling [8]. Many other receptor-based drugs are being developed for future use [9]. The current pharmacotherapy, although partially effective, has certain disadvantages. First, pharmacotherapy is a one-size-fits-all approach of addiction treatment that has not been largely successful, possibly because alcoholics constitute varying subtypes with differing biological and psychosocial contributions to the disease [10]. Second, alcoholism is a multifaceted disorder associated with neuro chemical heterogeneity and behavioral complexities, but current and prospective anti-alcoholism drugs selectively target specific receptors or transporters [8,9]. Therefore, patients may have to take many pills daily for comprehensive treatment/protection of alcoholism. This may be partially responsible for the patients’ non-adherence to pharmacotherapy regimens. Third, the current

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therapeutic drugs exhibit poor bioavailability, serious side effects and probability of development of addiction (for benzodiazepines).

Currently, the following alternative treatment approaches may constitute novel broad-spectrum pharmacotherapies for alcoholism and addiction.

- **Herbal medications**: Although many Asian countries have used medicinal plants such as *Tabernanthe iboga*, *Panax ginseng*, *Salvia miltiorrhiza*, *Hypericum perforatum*, *Pelargonium graveolens*, *Lippia citriodora*, *Punica granatum*, *Morinda citrifolia*, *Mirabilis jalapa*, *Aloysia triphylla* etc. to treat addiction for centuries, only recently has the West begun to understand their pharmacological possibilities and clinical applications in alcoholism treatment [9-11]. Despite large volume of supporting research data, the clinical application of medicinal plants are hindered by their poor solubility and bioavailability of the active ingredients, inability to cross the blood-brain barrier, and high variability [12].

- **Traditionally synthesized nanoparticles**: Development of Nanoparticle (NP)-based pharmacotherapy is a promising advancement in diagnosing and designing personalized treatment of addiction and other diseases. Studies have used colloidal gold and silver NPs, functionalized with multiple pharmaceuticals and other active ligands such as a blood-brain barrier permeant peptide, in treatment of alcoholism [13-15]. NPs, because of their unique properties, may circumvent the disadvantages of current pharmacotherapy by improving bioavailability and therapeutic efficacy of the drugs, loading multiple drugs in a single nanocarrier, improving compliance because patients will not have to take multiple pills, and facilitating on-demand drug release via external (UV or NIR beam or ultrasound) or internal (pH or selected enzymes) cues [16,17]. However, the traditionally synthesized gold and silver NPs have some disadvantages: they require stabilization to prevent rapid aggregation, difficult to functionalize with certain ligands, and undergo de-functionalization, releasing toxic NPs. Since the NPs contain the FDA approved drugs listed above, thus exhibit the same limitations listed above for pharmaceutical preparation.

- **Plant-extract synthesized nanoparticles**: Earlier studies have described synthesis of metal NPs using plant extracts (also known as ‘green’ synthesis) that are environment friendly, cost effective, easily scaled up for large scale syntheses, and do not require stabilizers such as polyethylene glycols [18-32]. Most importantly, the ‘green’ NPs retain the therapeutic potency of the plant and the unique properties of NPs. However, the key problems associated with the ‘green’ NPs are lack of methodology to identify the surface ligands, poor dose-response studies and lack of information regarding therapeutic doses.

Based on the observations discussed above, it is hypothesized that AuNPs synthesized with plant-extracts, especially using a combinations of kudzu root and gum extract because of their relevance to alcohol disorders [33-35], provides an effective and safe therapy for alcoholism and addiction. However, for broader acceptance, there is an urgent need to develop remedies for the disadvantages associated with plant-extract synthesized NPs. The overall goal of this study was to synthesize and characterize Gold Nanoparticles (AuNPs) using aquatic extract of kudzu root with or without edible gum. The specific aims were to (1) Characterize the AuNPs; (2) Identify and quantify the surface ligands; (3) Determine the AuNPs uptake into the SH-SY5Y cells and AuMP-protein interactions; (4) Assess the beneficial and adverse effects of AuNPs in SH-SY5Y cells. The methodologies were improved by including (i) An internal standard (4d-daidzein) in the extract used for synthesis of AuNPs preparations that allowed extract standardization and (ii) LDI and low-matrix MALDI for analysis of AuNP surface ligands. The preliminary studies indicated that inclusion of ‘gum’ in the reaction medium improved the AuNP's synthesis by kudzu root extract.

**Materials and Methods**

**Preparation of plant extracts**

Commercial, pre-calibrated, kudzu root extract powder (20 g, Sigma Chemicals) and 10 g Gum Arabic (Sigma Chemicals) were mixed in 100 ml Double Distilled Water (DDW) and incubated under stirring 50 °C for 4 h. The mixture was filtered through a 0.45 μm filter (Millipex™ poly vinylidene difluoride; EMD Millipore). The composition of the filtrate was analyzed as described in the supplementary section. The filtrate was freeze dried and stored at -80 °C for later use.

**Nanoparticle synthesis and characterization**

The traditional synthesis of AuNP using a reducing agent was performed as described by Singh, et al. [15]. For AuNP synthesis using plant extract, in a 250 ml flask, 1 mM AuCl₃ was mixed with 50 mL of filtered extracts (kudzu, gum, or mixture of the two) and 0.1 mL (1 mg/ml stock) of the internal standard (4d-daidzein) and heated to 37 °C, 50 °C and 90 °C that was maintained for 4 h (concentrations, temperature and time was determined in a preliminary study). The UV-Vis spectrum was measured at excitation band ranging from 500 to 600 nm to assess NP formation. After incubation (the preliminary study showed that 4 h of incubation at 50 °C or 2.5 h at 90 °C was sufficient for conversion of AuCl₃ to AuNPs, respectively)
5 to 10 nm), the mixture was centrifuged and the AuNP pellets were collected and washed 3 x with distilled water. Nanoparticles were characterized by measuring the following indices:

**UV-VIS spectrum:** Synthesized AuNPs were confirmed by sampling the aqueous component at different time intervals and the absorption maxima was scanned by UV-VIS spectrophotometer at the wavelength range of 300-800 nm on Perkin-Elmer Lambda 25 spectrophotometer.

**FT-IR spectrum:** AuNP samples were dried and grinded with KBr pellets for FTIR measurements. The spectrum was recorded in the range of 4000-400 cm⁻¹ using Varian FTS 1000 FT-IR Spectrometer in the diffuse reflectance mode at resolution of 4 cm⁻¹.

**Surface topology by TEM:** Morphology and size of the silver nanoparticles were investigated by TEM images using Hitachi model HF-2000 field emission transmission electron microscope with a resolution of 0.1 nm. A thin film of the sample was prepared on a carbon coated copper grid by just dripping a very small amount of the sample on the grid and drying under lamp.

**Identification of AuNPs and the surface functional groups:** Laser Desorption Ionization (LDI) without matrix and low-matrix-assisted LDI-TOF-MS (Bruker Autoflex III Smartbeam mass spectrometer that is equipped with a 355 nm Nd:YAG laser) were used to characterization of AuNP and functionalized groups. The nanoparticle preparations (1 µl) were mixed with 1 µL of (i) 50% acetonitrile (without matrix) and 0.1% Tri Fluoroacetic Acid (TFA) and (ii) 3 mg (for low-matrix) and 20 mg (for full matrix) of α-Cyano-4-Hydroxy Cinnamic Acid (CHCA) solution. The mixtures were crystallized onto the LDI-disc. MS experiments were done in positive mode on a reflectron-type (20.92 kV) time-of-flight (to acquire a single spectrum, 200 laser shots were fired at a frequency of 100 Hz at 35-40% of the full laser power). In all cases, multiple spectra were averaged to obtain the reported data.

**Beneficial efficacy and adverse effects of AuNPs**

Beneficial and adverse effects of the AuNPs were investigated using SH SY5Y neuronal cell-lines. For culture, the vials containing SH SY5Y cells were thawed rapidly in a 37 °C water bath and the contents were transferred in a centrifuge tube containing 9.0 ml complete culture medium (1:1 mixture of Eagle's Minimum Essential Medium and F12 Medium mixed with fetal bovine serum, final concentration 10%). The cells were centrifuged at 125 × g for 7 minutes and the pellets were suspended with the complete medium (pH 7.0 to 7.6). The cells were incubated at 37 °C in 5% CO₂ in air atmosphere for 3 days. Then the cells were plated at a density of 10⁷ cells in Cyto One filter cap flasks and cultured for 3 days. Then, the media was replaced with the one containing either different concentrations of AuNP preparations (AuNP-positive cells) or the matrix alone.

**Figure 1:** Experimental protocol. SH-SY5Y cells were expanded in growth media for 3 days. On day-4, 3 × 10⁷ cells were transferred into each well and incubated for 4 days in media containing different doses of AuNPs (AuNP positive (A⁺) cells) or matrix alone (AuNP negative (A⁻) cells). At day-8, cells were washed and incubated in media containing 54 mM (final concentration) ethanol (A⁺ ethanol exposed (A⁺E⁺) or A⁻ ethanol exposed (A⁻E⁻) cells) or the matrix alone (AuNP and ethanol negative (A⁺E⁻) or A⁻ ethanol negative (A⁻E⁻) cells) for 24 hours. Then cells were washed and incubated in media alone. At different time intervals after ethanol exposure, cells were harvested and analyzed for AuNP internalization, AuNPs’ adverse effects and modulation of ethanol toxicity.
(AuNP-negative) cells that continued for 4 days (day-4 to day-7). On day-8, cells were exposed for 24 h to either 50 mM ethanol (ethanol-positive) or saline (ethanol-negative) cells as matrix.

This yielded the following cell groups: (1) AuNP-negative/ethanol-negative (An+/En); (2) AuNP-negative/ethanol-positive (An+/Ep); (3) AuNP-positive/ethanol-negative (An+/En); and (4) AuNP-positive/ethanol-positive (An+/Ep) cells (Figure 1). At day-9, the cells were incubated in AuNP and ethanol free culture media for another 16 days. At predetermined times (day-9 to day-25 (i.e. day-0 days post ethanol), cells were harvested and processed for the following:

**Determination of internalized AuNPs by measuring intracellular Au concentrations (iAuCs):** The Atomic Absorption Spectrophotometric method described by Pedersen and Graaebek [36] and Benhabib, et al. [33,34] was used for analysis of Au in tissue samples. In brief, the cells were harvested, treated with trypsin, etched with Potassium Iodide (KI) and Iodine (I₂) at 1:6 ratio (0.34 mM I₂) to remove the surface AuNPs [37], washed with plasma buffered saline and mixed with 5 mL of aqua regia and digested for 24 h. The digest was diluted 1:1 with a solution of 2 ppm yttrium in dilute nitric acid serving as an internal standard. The instrument was calibrated using a solution of 1 ppm Au and 1 ppm yttrium in 50% aqua regia. The samples were introduced in a segment flow mode of a flameless atomic absorption spectrophotometer (Beckman model 485). Au concentration was determined using a calibration curve. The brain Au concentrations were normalized to total tissue weight.

**Lactate dehydrogenase release assay:** A 50 μl of clear cell media was mixed with the reconstituted 2 × LDH assay buffer (223 mg 2-p-iodophenyl-3-p-nitropheryl-5-phenyl tetrazolium chloride, 57 mg N-methylphenazonium methyl sulfate, 575 mg NAD, and 3.2 g lactic acid in 480 ml 200 mM Tris buffer solution, pH 8.0). The mixture was gently shaken for 30 seconds and incubated for 20 min at 37 °C for different time intervals. The Carboxy-tetramethyl-Rhodamine (TMR)-labeled oligonucleotide (0.5 ng) was then added to the reaction mixture and incubated for 20 min at room temperature [39].

**Cell viability:** The MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide) tetrazolium reduction assay kit from Millipore (Chemicon, CT02) was used to assess cell viability.

**Analysis of cells undergoing apoptosis and necrosis:** Cells were exposed to the AuNPs followed by ethanol as shown in Figure 1. At different time intervals after cessation ethanol, the cells were harvested and double stained with Annexin V-FITC (An) and Propidium Iodide (PI) as described previously [15,38] and were subjected to FACS analysis using on a FAC Scan flow cytometer (Becton Dickinson, San Jose, CA). All the experiments were performed in triplicate. Cells were classified as An+PI or An+PI (normal cells), An+PI (necrotic cells), An+PI (early apoptosis stage), An+PI (late apoptosis stage).

**NFkB activation by Electrophoretic Mobility Shift Assay (EMSA):** Cell cultures from in vitro studies or finely chopped tissue samples from in vivo studies were suspended in HEPES hypotonic buffer A (10 mM HEPES, pH 7.9, 10 mM KCl, 0.1 mM EDTA, 0.1 mM EGTA, 1 mM Dm, 0.5 mM phenylmethysulfonyl fluoride, and 10 pg/ml leupeptin, antipain, aprotinin, and pepstatin) for 15 min on ice. Samples were vortexed for 10 s with 0.6% Nonidet P-40 and centrifuged at 12,000 × g for 60 s. The pellets containing Nuclei were resuspended in nucleus buffer (20 mM HEPES, pH 7.9, 25% glycerol, 0.4 M NaCl, 1 mM EDTA, 1 mM EGTA, 1 mM DTT, 0.5 mM phenylmethysulfonyl fluoride, and 10 µg/ml leupeptin, antipain, aprotinin, and pepstatin) and briefly sonicated on ice. For EMSA nuclear extracts (10 pg of protein) were incubated in 25 µl of total reaction volume containing 20 mM HEPES, pH 7.9, 50 mM NaCl, 0.1 mM EDTA, 1 mM DTT, 5% glycerol, 200 pg/ml bovine serum albumin, and 2.5 pg of poly(d1-dC) for 15 min at 4 °C. The Carboxy-tetramethyl-Rhodamine (TMR)-labeled oligonucleotide (0.5 ng) was then added to the reaction mixture and incubated for 20 min at room temperature [39].

**Reference probe:** TMR-AGT TGA GGG GAC TTT CCC AGG CAA-3

**ReLa-p50 probe:** TMR-AGT TGA GGG GAC TTT CCC AGG CAA-3

**Gels were scanned on the Hitachi FMBIO II using 585 channels for the isotope probe.**

**Characterization of AuNP-Protein Interaction**

To investigate the AuNP-protein interaction, different AuNP preparations were incubated with fresh serum (1:10 by volume) at 37 °C for different time intervals. Thereafter, an equal volume of sucrose solution (0.75 M) was carefully layered and the samples were centrifuged at 10,000 g for 1 hour to separate unbound proteins (remained in water phase) from the AuNPs (settled at the bottom). The pellets were collected and washed three times with isotonic saline to remove soft corona. The
hard corona proteins were desorbed using CHAPS buffer (DNase 1 (100 U/mL), RNAse A (10 U/mL), CaCl₂ (20 mM), MgCl₂ (50 mM), Tris HCl pH 7 (100 mM), SDS 0.02%, ampholine (1%), CHAPS (0.8%), β-mercaptoethanol (1%) and urea (8 mM)). The mixture was centrifuged (15,000 rpm, 3 min) to separate the AuNPs (in the pellets) and the desorbed proteins (in the supernatant). The desorbed proteins were concentrated, mixed with Bromophenol blue (0.5 µL) and run on 1-DE gels. The separated proteins were staining with Coomassie dye or viewed for fluorescence under UV light [40].

For protein identification, the surface-adsorbed proteins were reduced with 10 mM Dithiothreitol (DTT) at 37 °C for 1 h and alkylated with 25 mM Iodoacetamide (IAA) at room temperature for 50 min. For the digestion, the samples were incubated with porcine trypsin (Promega) for 8 h at 37 °C. The digested peptides were concentrated and purified in reverse phase C18 µ column (Promega) for 8 h at 37 °C. The digested peptides were reduced with 10 mM Dithiotreitol (DTT) at 37 °C for 1 h and alkylated with 25 mM Iodoacetamide (IAA) at room temperature for 50 min. For the digestion, the samples were incubated with porcine trypsin (Promega) for 8 h at 37 °C. The digested peptides were concentrated and purified in reverse phase C18 µ column (Promega) for 8 h at 37 °C. The digested peptides were reduced with 10 mM Dithiotreitol (DTT) at 37 °C for 1 h and alkylated with 25 mM Iodoacetamide (IAA) at room temperature for 50 min. For the digestion, the samples were incubated with porcine trypsin (Promega) for 8 h at 37 °C. The digested peptides were concentrated and purified in reverse phase C18 µ column (Promega) for 8 h at 37 °C.

For immune staining, the separated proteins were transfected onto a PVDF membrane, blocked and incubated with rabbit anti-mammalian-IgG selective antibodies (reactivity to rat proteins were determined), followed by Horse Radish Peroxidase (HRP)-conjugated goat anti-rabbit IgG antibodies. For detection, an HRP-conjugated secondary antibody enhanced chemiluminescence kits from Abcam were used.

### Statistical analysis

Data were recorded as mean ± SD using Microsoft Excel. Statistical analysis was performed using ANOVA followed by Tukey’s Multiple-Comparison Test or t-test at < 0.05 significance level.

### Results

#### Nanoparticle synthesis and characterization

Physicochemical characterization: 

- **Physicochemical characterization**: 

  - AuCl₄⁻ (a yellow colored aqueous solution) exhibited strong absorption at 217 nm and a weak absorption at 287 nm. Exposure of AuCl₄⁻ to reducing agents immediately changed its color from yellow to red/brown characterized by a strong absorption from 500-665 nm range, depending upon the nanoparticle size and synthesis procedure (Figure 2a). A combination of Kudzu and Gum (KG) extract at 90 °C, 50 °C and 37 °C yielded AuNPs with average diameter of 8 nm, 12 nm and 20 nm, respectively (Figure 2b). The Kudzu (K) extract alone at 90 °C, 50 °C and 37 °C formed relatively larger AuNPs (diameter 30 to > 50 nm). The Gum solution (G) alone yielded particle > 50 nm in diameter. The FTIR spectra of K extract and different nanoparticles are shown in Figure 3. The extract exhibited strong absorption at 2928, 1609, 1078 and 1069 cm⁻¹, and weak absorption at 250-500 cm⁻¹. The AuNPs (i: AuNP KG-50; ii: AuNP KG-95 and iii: AuNP KG) exhibited strong absorption at 3420, 1609 and 1078 nm, and weak absorption at 2920, 1456, 1156, 981 and < 500 cm⁻¹ (Figure 2 and Figure 3).

The surface characterization: Low-matrix MALDI detected Au⁺ (m/z 193), Au³⁺ (m/z 396) and Au³⁺ (m/z 792). Table 1:

#### Table 1: List of fragment ions monitored for protein identification

| Protein                | Test: IS or MS: sequence | m/z (+1) | m/z (+2) |
|------------------------|--------------------------|----------|----------|
| Bin1 protein           | MS: AEEELIK              | 832.2    | 418.1    |
| Casein Kinase 1 y2     | MS: MEYVHTK              | 908.0    | 456.3    |
| cAMP/cGMP phosphodiesterase | MS: VOYHNWK             | 944.4    | -        |
| Glutaryl-Co-A dehydrogenase | MS: CEDNCIR            | 853.3    | 439.1    |
| Arginine N-methyltransferase | MS: QYKDYK             | 845.5    | 424.3    |
| Lactate dehydrogenase  | MS: MVSGESR              | 766.52   | 384.3    |
| MAPK kinase 7          | MS: LCGFISGR             | 968.7    | -        |
| Parvalbumin            | MS: FFQMVGLK             | 970.5    | 486.4    |
| Protein Tyr Phosphatase | MS: MPVIVSR            | 803.1    | -        |
| Transferrin            | MS: EDQTFYY              | 815.5    | 209.3    |
| Triose phosphate isomerase | MS: AISDNYK           | 731.4    | -        |
The dose-internalization curves for different AuNPs (measured as iAuC) were linear from 5 to 100 mg/L AuNP dose (Figure 5). The $r^2$ for Ap/En and Ap/Ep cells ranged from 0.97 to 0.99 and 0.78 to 0.95, respectively, indicating greater variability in Ap/Ep cells. As shown below, the iAuCs were highest in cells exposed to AuNP

onto the surface (Figure 4).

**AuNPs internalization in cells**

The dose-internalization curves for different AuNPs (measured as iAuC) were linear from 5 to 100 mg/L AuNP dose (Figure 5). The $r^2$ for Ap/En and Ap/Ep cells ranged from 0.97 to 0.99 and 0.78 to 0.95, respectively, indicating greater variability in Ap/Ep cells. As shown below, the iAuCs were highest in cells exposed to AuN-
P_{KG-50} and lowest in cells exposed to AuNP_{G}. The slopes of the curve ranged from 0.06 to 0.1 (Figure 5).

Figure 4: Mass spectral identification of surface ligands. A: Low-level matrix-LDI that detects Au ions; B: Typical LDI (matric free) analysis of AuNP_{KG-50} synthesized by a mixture of kudzu root and gum; C: Typical LDI analysis of a mixture of kudzu and gum solutions; D: Typical LDI (matric free) analysis of AuNP_{G-50} synthesized by a gum solution alone; E: Typical LDI (matric free) analysis of AuNP_{K-50} synthesized by a kudzu extract alone; and F: Analysis of a blank solution.

Figure 5: The dose (mg/L of AuNPs) versus response (intracellular Au concentration-iAuC) curves for internalization of different AuNPs at day-16 of cell incubation. A: A/P/E^α cells and B: A/P/E^β cells. Values are mean ± SD.

Ap/En cells: AuNP_{KG-50} > AuNP_{KG-90, KG-37, K-37} > AuNP_{K-37, K-90, K-50, PEG + KG} > AuMP_{G-50}
A\textsuperscript{p}/E\textsuperscript{n} cells: AuNP\textsubscript{KG-50} > AuNP\textsubscript{KG-90, KG-37, K-37} > AuNP\textsubscript{K-30, K-90} > AuNP\textsubscript{PEG+K+G} > AuNP\textsubscript{G-50}

Figure 6 shows the time-course of iAuC changes in cells harvested at (i) Culture days 4 to 7 (days 1 to 4 of AuNP exposure), (ii) Culture day 8 (24 h ethanol exposure), and (iii) Culture day 9, 10, 14, 18 and 26 (16 days free of ethanol and nanoparticles) as shown in Figure 1. From day-4 to day-7, iAuCs increased gradually and, at day-4, the iAuCs showed the following pattern: AuNP\textsubscript{KG-90} > AuNP\textsubscript{KG-50} and AuNP\textsubscript{KG-37} > AuNP\textsubscript{K-90} > AuNP\textsubscript{K-37} > AuNP\textsubscript{PEG+K+G} and AuNP\textsubscript{G-50}. In A\textsuperscript{p}/E\textsuperscript{n} cells, a linear decrease in iAuCs occurred. The values at day-24 (16 days after discontinuation of ethanol) ranged from 0 to 4 ng/well. In A\textsuperscript{p}/E\textsuperscript{n} cells, the post-ethanol iAuCs were significantly lower than those in corresponding A\textsuperscript{p}/E\textsuperscript{n} cells (Figure 6). The day-24 (16 days after discontinuation of ethanol) values for A\textsuperscript{p}/E\textsuperscript{n} and A\textsuperscript{p}/E\textsuperscript{n} cells did not differ significantly (Figure 6).

Characterization of surface ligands in AuNPs collected from exposed cells

Figure 7 shows a typical time-course of change in surface ligands of intracellular AuNP\textsubscript{KG-50} at different time intervals in A\textsuperscript{p}/E\textsuperscript{n} cells (Figure 7A - fAuNP\textsubscript{i}, fAuNP\textsubscript{j}, fAuNP\textsubscript{k}, fAuNP\textsubscript{l}, fAuNP\textsubscript{m}, fAuNP\textsubscript{n}, fAuNP\textsubscript{o}, fAuNP\textsubscript{p} represent 1, 2, 4, 8 and 16 days after ethanol exposure). Figure 7B shows time-course of change in total number of peaks observed onto the AuNP\textsubscript{KG-50} surface. The peak numbers remained unchanged for up to 8 days, but decreased significantly at day-16 after ethanol exposure. The Np/Ep values were significantly higher than the Ap/Ep values. This may be due to desorption and degradation of the ligands. Figure 7C shows a linear relationship between the AuNP\textsubscript{KG-50} dose and the IS-peak-area/ligand-peak-area ratio (Figure 7).

Beneficial and adverse effects of AuNPs

To evaluate the adverse effect of AuNPs alone, A\textsuperscript{p}/E\textsuperscript{n} cells were exposed to different concentrations of individual AuNPs for 4 days, the saline (matrix for ethanol) for 24 h. The cells were assayed for (i) Viability using MTT metabolism assay, (ii) Apoptotic and necrotic cells, and (iii) Lipid peroxidase and lactate dehydrogenase activities. The beneficial effects were studied by incubating A\textsuperscript{p}/E\textsuperscript{n} cells with different doses of AuNPs for 4 days, followed by 50 mM ethanol, and then assayed as above.

Cell viability:

Dose-response: In A\textsuperscript{p}/E\textsuperscript{n} cells, a 5, 10, 20, 40 or 60 mg/L dose of different AuNP preparations did not significantly affect, while a 100 mg/L dose of AuNPs caused
Figure 7: Identification of surface ligands on AuNPs collected from A^+/E^- and A^+/E^+ cells. A: the plant component peaks at 1 (fAuNP LD1), 2 (fAuNP LD2), 3 (fAuNP LD3), 4 (fAuNP LD4), 8 (fAuNP LD8), and 16 (fAuNP LD16) days after ethanol exposure; B: Time-course of change in total number of peaks adsorbed onto the AuNP surface. Total number of peaks decreased significantly at day-16. The time-course of change in intracellular AuNPs as measured by iAuCs. A: Intracellular AuNP_{KG-50}; B: Intracellular AuNP_{K-50}; C: Intracellular AuNP_{PEG} in A^+/E^- cells (filled circle) and A^+/E^+ cells (open circle). C: AuNP dose versus (ligand peak/IS peak) ratio plots for (i) PUM, KAM, DI, DE, AR, GAM, FR in AuNP_{KG-50} isolated from A^+/E^- cells, and (ii) PUE, KAE, DI,E, DE,E, AR,E, GAE, FR in AuNP_{KG-50} isolated from A^+/E^+ cells. A linear response between AuNP dose vs. ligand/IS ratio suggest that a dose-response could established by using the ligand concentration as the dose. Values are mean ± SD.

Figure 8: (A) Plots of dose (AuNP mg/L) versus viable A^+/E^- cells (% of control) at day-16. A 5 to 60 mg/L dose of all AuNP exhibited comparable response. Only 100 mg/L dose caused significant decrease in viable cell enumeration. Values are mean ± SD; (B) Plots of dose (AuNP mg/L) versus viable A^+/E^+ cells (% of control) at day-16. Different AuNPs yielded variable response, but maximum protection was provided by AuNP_{KG-50} at 20 mg/L dose. Values are mean ± SD.; p < 0.05 when compared with A^+/E^- response and # or @: p < 0.05, when viability of A^+/E^- cells exposed to a given dose of AuNP compared with viability of A^+/E^- exposed to the same dose of other AuNPs; (C) The time-course of change in viability of A^+/E^- (broken line) and A^+/E^+ (solid line) cells pre-exposed to a 20 mg/L dose of AuNP_{KG-50}*: p < 0.05 when exposed to day-0 values and x: p < 0.05 when A^+/E^- and A^+/E^+ values were compared at each time interval.
30% to 60% decrease in viable cells (Figure 8A, hour-16 data). Ethanol exposure of A/Ep or A/Ep cells significantly decreased cell viability (Figure 8B). A 5, 10, 20, 40 or 60 mg/L dose of different AuNP preparations protected (10% to 40%) against the ethanol-induced decrease in cell viability (AuNP KG-50 or KG-90 >> AuNP KG-50 or KG-90 > AuNP K-90 or K-50 > AuNP KG or KG-100, > AuNP KG-50 or KG-50 or AuNP PEG). A 100 mg/L dose of all AuNP preparations either did not change or augmented ethanol toxicity. This suggests that the 20 and 40 mg/L dose of AuNP KG-50 and AuNP KG-90 were most effective against ethanol toxicity in SH-SY5Y cells.

**Time-course:** At 1 and 2 days after cessation of ethanol exposure in AuNP KG-50-pretreated (A/Ep) or AuNP KG-50-negative (A/Ep) cells, about 70% of total cells were viable and the values did not differ significantly between the two cell groups. Thereafter, the AuNP KG-50-pretreated values were significantly greater than AuNP KG-50-ve cells at all time intervals (Figure 8).

**Lipid peroxidase and lactate dehydrogenase activities:**

**Dose response:** In A/Ep cells, a 5, 10, 20, or 40 mg/L dose of all AuNP preparations did not increase, a 60 mg/L dose slightly increased, and a 100 mg/L dose significantly increased the enzyme activities (Figure 9A for lipid peroxidase and Figure 10A for LDH). The enzyme activities (Figure 9A for lipid peroxidase and Figure 10A for LDH) showed the following patterns in A/Ep cells: AuNP G-50 and AuNP K-90 >> AuNP PEG > AuNP PEG+K+G, AuNP KG-50 and AuNP KG-50 and AuNP KG-90. In A/Ep cells or A/Ep cells, exposure to AuNP G-50, AuNP K-90, AuNP KG-50, AuNP KG-37, AuNP PEG and AuNP PEG+K+G caused 125% to 155% increase in the enzyme activities (Figure 9B for lipid peroxidase and Figure 10B for LDH). In A/Ep cells, exposure to AuNP KG-50 or AuNP KG-90 (dose: 20, 40 and 60 mg/L) ethanol exposure did not significantly increased the enzyme activities (Figure 9B for lipid peroxidase and Figure 10B for LDH). However, ethanol exposure of AuNP KG-50-pretreated (dose: 100 mg/L) cells significantly increased the enzyme activities. This suggests that a 20 to 60 mg/L dose of AuNP KG-50 or AuNP KG-90 protected A/Ep cells against ethanol toxicity (Figure 9 and Figure 10).

**Time-course (Figure 9C for lipid peroxidase and Figure 10C for LDH):** The two enzymes exhibited different time-course of change in response to ethanol exposure. Lipid peroxidase activities were significantly higher at day-1 and day-2, and then decreased gradually to 130% of control at day-16. The ethanol-induced increase in peroxidase activity in A/Ep cells were significantly greater than the activity in A/Ep (AuNP KG-37, -50 or AuNP KG-50-pretreated, 20 mg/L) cells, indicating neutralization of ethanol-induced oxidative stress by AuNP KG. Other AuNPs were variably effective, providing < 20% protection. Unlike lipid peroxidase activities, LDH activities increased gradually and peaked at day-4. Then a gradual decrease occurred. LDH activities in A/Ep cells and A/Ep (AuNP KG-pretreated) cells did not differ significantly.
at days 1, 8 or 16, but differed significantly at days 2 and 4. Other AuNPs were variably effective, showing 5% to 30% increase (Figure 9 and Figure 10).

Enumeration of apoptotic and necrotic cells:

**Dose-response (16-day data):** Figure 11 shows distribution of different cell phenotypes in $A^+/E^+$, $A^+/E^+$ (different AuNP dose) and $A^+/E^+$ cells. $A^+/E^+$ and $A^+/E^+$ (5 to 40 mg/L AuNP dose) cells exhibited mostly An PI- phenotypes. $A^+/E^+$ (60 mg/L AuNP dose) cells exhibited An PI- and An +++PI- phenotypes, while $A^+/E^+$ (100 mg/L AuNP dose) exhibited An PI+, An +++PI- and An PI++ phenotypes. This suggests that 5 to 40 mg/L doses of AuNP KG-50 did not induce apoptosis or necrosis in cell culture. The cell phenotypes in $A^+/E^+$ cells exposed to other AuNPs are shown in Table 2.

$A^+/E^+$ cells exhibited An PI+, An ++PI- and An PI++ phenotypes. The $A^+/E^+$ (AuNP KG dose 5 and 10 mg/L) cells exhibited An PI- and An +++PI- phenotypes. $A^+/E^+$

**Figure 10:** (A) Plots of dose (AuNP mg/L) versus Lactate Dehydrogenase (LDH) activity in $A^+/E^+$ cells (% of control) at day-16. A 5 to 60 mg/L dose of all AuNP exhibited comparable response. Only 100 mg/L dose caused significant increase in enzyme activity. Values are mean ± SD. •: p < 0.05 when $A^+/E^+$ or $A^+/E^+$ values were compared with $A^+/E^+$ values, x: p < 0.05, when enzyme activity of $A^+/E^+$ cells exposed to a given dose of and AuNP compared with enzyme activity of $A^+/E^+$ cells exposed to the same dose of other AuNPs; (C) The time-course of change in LDH activity of $A^+/E^+$ (broken line) and $A^+/E^+$ (solid line) cells pre-exposed to a 20 mg/L dose of AuNP KG-50: •: p < 0.05 when exposed to day-0 values and x: p < 0.05 when $A^+/E^+$ and $A^+/E^+$ values were compared at each time interval.

**Figure 11:** Effects of different doses of AuNPs on incorporation of annexin V and propidium iodide (PI) in $A^+/E^+$ (bottom 0 mg/L), $A^+/E^+$ (bottom 5 to 100 mg/L), $A^+/E^+$ (top 5 to 100 mg/L AuNP) and $A^+/E^+$ (top 0 mg/L) cells.
was confirmed by competition experiments in which the formation of protein-DNA complex was specifically competed by non-labeled specific probe and by different amounts of nuclear proteins. Figure 13B shows fluorescence intensity of NFκB-DNA band. In the absence of ethanol, a 5, 10, 20 or 40 mg/L dose of AuNPKG-50 did not modulate, while 60 mg/L and 100 mg/L dose significantly activated p65-p50. Ethanol exposure, in the absence of AuNPs, gradually increased NFκB activation that peaked at day-8. At day-16, the values either did not change or decreased slightly. The 5 and 10 mg/L dose of AuNPKG-50 did not, while the 20 mg/L and 40 mg/L doses significantly attenuated NFκB activation by ethanol, thus protecting the cells against the ethanol's adverse effects. Higher AuNP doses augmented NFκB activation and ensuing toxicity by ethanol (Figure 13).

Formation of intracellular protein-corona: Figure 14A shows electrophoretic separation of cytosolic and nuclear proteins. Formation of intracellular protein-corona: Figure 14A shows electrophoretic separation of cytosolic and nuclear proteins.

| Table 2: Effects of AuNP dose on sh-sy5y cell phenotype. |

| NP (mg/L) | 0  | 5  | 10 | 20 | 40 | 60 | 100 |
|-----------|----|----|----|----|----|----|-----|
| A+/E- cells (no AuNPs or ethanol) |   | An-PI |   | An-PI |   | An-PI |   | An-PI |
| A+/E+ cells (AuNPs positive but ethanol negative) |
| AuNP<KG-90, 50, 37 | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI |
| AuNP<KG-60, 50, 37 | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI |
| AuNP<KG-50, 90 | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI |
| AuNP<PEG, PEG+K+G | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI | An-PI |
| A+/E+ cells (AuNPs negative, but ethanol positive) |
| AuNP<KG-90, 50, 37 | An-PI | An++PI | An++PI | An++PI | An++PI | An++PI | An++PI |
| AuNP<KG-60, 50, 37 | An-PI | An++PI | An++PI | An++PI | An++PI | An++PI | An++PI |
| AuNP<KG-50, 90 | An-PI | An++PI | An++PI | An++PI | An++PI | An++PI | An++PI |
| AuNP<PEG, PEG+K+G | An-PI | An++PI | An++PI | An++PI | An++PI | An++PI | An++PI |

(AuNP<kg dose 20 mg/L) cells exhibited An++PI phenotypes, while A+/E- (AuNP<kg 60 and 100 mg/L) cells exhibited An++PI, An++PI, An++PI and An PI++ phenotypes. The cell phenotypes in A+/E+ cells exposed to other AuNPs are shown in (Table 2).

Time-course (20 mg/L data): Distribution of cell phenotypes at different time intervals after ethanol exposure in A+/E+ and A+/E- (AuNP<kg dose 20 mg/L) cells are shown in Figure 12. At day-16, A+/E+ cells exhibited An++PI, An++PI, An++PI and An PI++ phenotypes, while A+/E- (AuNP<kg dose 20 mg/L) cells exhibited An++PI phenotypes, suggesting that a 20 mg/L dose of AuNP<kg protected against ethanol toxicity. The data for other NPs are not shown since they provided poor protection against ethanol toxicity (Figure 11, Figure 12 and Table 2).

NFκB activation: Figure 13A shows typical gel images of NFκB’s (p65-p50)-DNA complex by the fluorescent EMSA. The specificity of the DNA-binding protein was confirmed by competition experiments in which the formation of protein-DNA complex was specifically competed by non-labeled specific probe and by different amounts of nuclear proteins. Figure 13B shows fluorescence intensity of NFκB-DNA band. In the absence of ethanol, a 5, 10, 20 or 40 mg/L dose of AuNP<kg did not modulate, while 60 mg/L and 100 mg/L dose significantly activated p65-p50. Ethanol exposure, in the absence of AuNPs, gradually increased NFκB activation that peaked at day-8. At day-16, the values either did not change or decreased slightly. The 5 and 10 mg/L dose of AuNP<kg did not, while the 20 mg/L and 40 mg/L doses significantly attenuated NFκB activation by ethanol, thus protecting the cells against the ethanol’s adverse effects. Higher AuNP doses augmented NFκB activation and ensuing toxicity by ethanol (Figure 13).

Formation of intracellular protein-corona: Figure 14A shows electrophoretic separation of cytosolic and nuclear proteins.
Figure 12: The time-course of incorporation of annexin V and propidium iodide in A/V/E° cells (C), A/V/E° cells (bottom, 1 to 16 days, fixed AuNP dose of 20 mg/L), A/V/E° cells (top, 1 to 16 days, fixed AuNP dose of 20 mg/L) and A/V/E° cells (top day-1) cells.

Figure 13: A: Typical electrophoretic separation of NFkB dimer, p65-p50, in A/V/E° (matrix), A/V/E° (E+matrix), A/V/E° (E + 5 to 100 mg/L, data for AuNP KG-50 is shown) and A/V/E° (saline + 5 to 100 mg/L, data for AuNP KG-50 is shown) cells. [a] matrix exposed control, A/V/E° cells, [b] AuNP negative but ethanol positive A/V/E° cells, [c] AuNP and ethanol negative A/V/E° cells, [d] A/V/E° cells receiving 5 mg/L AuNP KG-50, [e] A/V/E° cells receiving 5 mg/L AuNP KG-50 [f] A/V/E° cells receiving 10 mg/L AuNP KG-50, [g] A/V/E° cells receiving 10 mg/L AuNP KG-50, [h] A/V/E° cells receiving 20 mg/L AuNP KG-50, and [i] A/V/E° cells receiving 10 mg/L AuNP KG-50, 20 mg/L AuNP KG-50, and 10 mg/L AuNP KG-50, respectively. B: Scan of the p65-p50 fluorescence - dose-response and time-course studies. [0 - Pre E] A/V/E° cells were exposed to different doses of AuNP KG-50 [1 to 16 days post ethanol exposure]: A/V/E° cells were exposed to different doses of AuNP KG-50 for 4 days then ethanol for 24 hours. Cells were analyzed at, day-1, day-2, day-4, day-8 and day-16 post ethanol. Values are mean ± SD. x: p < 0.05 when values for cells exposed to 60 mg/L were compared with cells exposed to lower AuNP doses. #: p < 0.05 when values for cells exposed to 200 mg/L were compared with cells exposed to lower AuNP doses. ¥: p < 0.05 when values for cells exposed to lower AuNP doses. %: p < 0.05 when values for cells exposed to lower AuNP doses. @: p < 0.05 when values for same AuNP dose collected at different time intervals were compared. Values are mean ± SD, n = 5.
AuNP-corona proteins from cells exposed to different AuNP preparations. The cytosol proteins ranged from 240 kDa to < 10 kDa, while the AuNP corona proteins ranged from 100 kDa to < 10 kDa in both control and ethanol-exposed cells. The protein composition exhibited the following differences (Figure 14):

i. Out of 35 proteins detected in control (A^+/) cells, 9 cytosolic proteins were confirmed. Tyr phosphor- ylase, Arg N-methyl transferase and MAPK kinase-7 yielded stronger bands (scan density 2.2 to 2.6) than parvalbumin, ADH, Glu-CoA dehydrogenase, cAMP/gAMP phosphodiesterase, transferrin and casein kinase 1y2 (scan density 0.3 to 1.4). Day-1 to day-16 values did not differ significantly.

ii. The AuNP corona protein/cytosol protein ratio for Tyr phosphorylase, Arg N-methyl transferase and MAPK kinase-7 increased from < 10 at day-1 to > 20 at day-16. The ratio for other proteins either did not change or increased slightly.

iii. Ethanol exposure did not significantly alter the protein scan values, but significantly suppressed the AuNP corona protein/cytosol protein ratio. Ethanol specifically altered binding of MAPK kinase-7, casein kinase, Tyr phosphorylase and cAMP/gAMP phosphodiesterase to AuNPs, possibly by altering their concentrations in the cytosol. This suggests that ethanol exposure may have directly affected the AuNP-protein interaction.

iv. The composition of protein corona changed temporally. AuNPs extracted from day-1 and day-2 A^+/E^a or A^+/E^E^ cells interacted with proteins ranging from < 37 kDa to 150 kDa. The band intensity increases with increasing incubation time (Figure 14).

Discussion

The present study showed that aqueous extracts of kudzu root and edible gum provided a simple one-step synthesis of AuNPs that had many advantages over the traditional synthesis method. The AuNPs were well dispersed and retained the advantages, while minimizing the disadvantages of the medicinal plant extract. The following sections describe the characteristics and medicinal properties and of AuNPs synthesized by Kudzu root and gum extracts.

Synthesis and characterization of AuNPs

AuNP size, UV absorption and FTIR spectroscopy: The UV-spectra of AuNPs indicated sharp absorption maxima at around 526 to 586 nm, a characteristic absorbance for AuNPs arising possibly due to an interaction of electromagnetic radiations with surface Plasmon of AuNPs. Like the observations of Amendola and Mene-ghetti [41], the present study also showed that AuNPs synthesized in this study exhibited considerable differences around 200-586 nm, but no differences after the Surface Plasmon Resonance (SPR) region around 500 to 600 nm [42].

In general, absorption maxima around 526 nm are determined by the AuNP size and incubation time. AuNPs of 22, 48 and 99 nm diameter, respectively, exhibited absorption maxima at 520, 540, 533 and 575 nm. Temperature affected UV absorption of AuNPs in a size dependent manner: smaller nanoparticles (< 20 nm diameter) were relatively resistant to temperature changes, while an increase in size is asso-
associated with an increase in temperature sensitivity [43]. The present study indicated that AuNPs synthesized at different temperatures exhibited differences in the absorption maxima. More research is needed to explain this divergence.

Figure 3 show the FTIR spectra for gum extract, kudzu extract and AuNP KG-50. Gum extract exhibit the absorption bands at (i) 3246 cm⁻¹ possibly due to the O-H stretching band; (ii) 2901 cm⁻¹ due to aliphatic C-H stretching; (iii) 1442, 1374 and 1339 cm⁻¹ due to C-H bending vibrations; (iv) 1220 to 998 cm⁻¹ due to C-H and O-H deformation and (v) 1145 to 554 cm⁻¹ due to the C-O and C-C groups’ vibration modes [44]. These bands characterize the carbohydrates present in gum solution. Kudzu extract exhibited broad band between 3600 and 3000 cm⁻¹ that is assigned to O-H stretching and to H-bonding involving the -OH groups. The band at 2920 cm⁻¹ is assigned to CH₂ symmetrical stretching vibrations. The band at 1639 cm⁻¹ is attributed to the scissoring of two O-H bonds, while the bands at 750 cm⁻¹ are due to skeletal stretching vibrations. The FTIR spectra for AuNP KG-50 (synthesized by kudzu-gum mixture at 50 °C) exhibited bands at 3306 cm⁻¹ assigned to O-H stretching vibration of alcohols and phenols, 1630 cm⁻¹ assigned to C=O stretching vibration of tertiary amides, and 1044 cm⁻¹ assigned to C=O stretching of primary alcohols [43]. From these observations, it is clear the poly-phenols present in the extract are responsible for the reduction and stabilization (Figure 3). The observation that FTIR spectra for AuNP KG-50 contains bands that are also present in kudzu and gum extracts suggests that the many of the kudzu and gum ingredients are adsorbed onto the AuNP surface (Table 3).

Characterization of surface ligands: An important observation of this study was that the two MS methods, LDI-TOF-MS and low matrix MALDI-TOF-MS, detected different sets of surfaces and ligands: low-matrix MALDI detected the AuNPs surface charge, while LDI-TOF-MS detected the desorbed organic ligands including an internal standard added to the plant extract. The identities of the ligands were confirmed by analyzing a mixture of authentic standards by the same method. As shown in Figure 4, AuNP KG-50 contained the components of kudzu extract (daidzein, puerarin and kakkasaponin III), gum (arabinose, galactose and fructose) and the internal standard (d4-daidzein). The intensity of natural and d4 daidzein did not significantly change during 16 days of cell maintenance. The internal standard allowed semi-quantitative determination of ligand concentrations onto the AuNP surface. This allowed determination of ligand dose as well as the extent of ligand depletion. Earlier studies have shown that puerarin, a naturally occurring isoflavone C-glycosides found in the roots of Pueraria lobata and P. thomsonii are widely used in China and India for the treatment of alcoholism, cerebrovascular and cardiovascular diseases [45]. Dietary gum is exhibits anti-inflammatory and anti-oxidative properties [46]. Thus, the AuNP’s surface ligands may be responsible for beneficial effects reported for the phytochemical-synthesized AuNPs.

![Image of FTIR spectra](https://example.com/ftir_spectra.png)

**Figure 15: Characterization of protein corona on AuNP surface (A) Effects of ethanol exposure on composition of corona proteins.**

**Abbreviation:** 1: AuNP KG-50; 2: AuNP KG-90; 3: AuNP KG-50; 4: AuNP KG-50 and 5: AuNP KG-50. Protein identification: 1: Alcohol dehydrogenase; 2: cAMP/cGMP phosphodiesterase; 3: Tyr phosphatase; 4: Transferring; 5: Glutamate CoA phosphodiesterase; 6: Arginine N-methyltransferase; 7: Casein kinase; 8: MAPK kinase; 9 and 10: Parvalbumin.

(B) Time-course of change in individual protein concentrations in cytosol and desorbed from AuNP KG-50.

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**Table 3: Description of FTIR peaks.**

| Peak cm⁻¹ | Functional group/vibration mode |
|-----------|--------------------------------|
| 750       | Aromatic protons               |
| 927       | C-O-C vibration modes of α-1,4 glycosidic linkage |
| 1040/1162 | Carboxyl (C=O) group, C-O stretching alcohols (primary, secondary and tertiary), carboxylic acids, esters and ethers |
| 1020/1079 | Carbonyl (C=O) group, C-O stretching alcohols (primary, secondary and tertiary), carboxylic acids, esters and ethers |
| 1116/1156 | C-O/C stretching in exo/endo-cyclic bonds and C-O/H/C deformation modes of oligo/polysaccharides and absorption bands for carbohydrates |
| 1383/1456 | The spectral bands for C-O/H stretching in pyranose and C-H deformation in esters |
| 1621/1637 | Aromatic C=C/C-OH stretching in exocyclic bond |
| 2923      | Glucoside ring due to the stretching vibration of O-H |
| 3420      | O-H, as also the H-bonded alcohols and phenols, carboxylic acid |
Cellular uptake and extrusion of AuNPs

Figure 15 shows uptake, fate and extrusion of AuNP in cells. NPs, upon encountering cell membranes, interact with (i) Caveolin-receptors targeted by small AuNPs (5 nm > 20 nm >> 50 nm), (ii) Clathrin-receptors for non-phagocytic uptake targeted by relatively larger AuNPs (50 nm) in the absence of serum, and (iii) Scavenger-receptor that mediated phagocytic uptake targeted by relatively larger AuNPs (50 nm) in the absence of serum [46-50] (Figure 15).

Once internalized, NPs either enter the cytosol directly and interact with cytosol-proteins [49] or trapped in endosome-lysosomes complex for processing (NP de-functionalization, protein hydrolysis and surface modification) before being transported to the cytosol and the cell’s periphery for extrusion via exocytosis [47,51-53]. In general, the endosomal NPs are extruded within 30 to 60 minutes [54], NPs accumulated in multi-vascular bodies’ exits cells within 6 days [55], and microtubule accumulated NPs are extruded constitutively via Golgi apparatus [56]. Internalization of nanoparticles can be modified by surface modification [57,58]. PEG coupling of nanoparticles surface significantly reduced the cytotoxic effects of nanoparticles in vitro and in vivo [59,60].

The present study showed that AuNP synthesized using different plant extracts and incubation temperature exhibited differences in size (AuNP<sub>KG</sub> > AuNP<sub>PEG</sub> > AuNP<sub>PEG/KG</sub>) and composition of surface ligand. Time-dependent decreases in density of certain ligands were observed (Figure 7). In Ethanol-negative cells, AuNP intake maxed within 3 days and the intracellular NPs concentrations exhibited the following patterns: AuNP<sub>KG</sub> > AuNP<sub>PEG/KG</sub> > AuNP<sub>PEG</sub> > AuNP<sub> KG,90 and 50</sub>. After discontinuation of NPs, the iAuC decreased gradually and returned to basal level within 16 days. Earlier studies have shown that AuNPs and other NPs are internalized by cells in a size- and surface-dependent manner through (i) Caveolin-mediated uptake targeted by small AuNPs (5 nm > 20 nm >> 50 nm), (ii) Clathrin-mediated non-phagocytic uptake targeted by relatively larger AuNPs (50 nm) in the absence of serum, and (iii) Scavenger-receptor mediated phagocytic uptake targeted by relatively larger AuNPs (50 nm) in the absence of serum [46-50]. Since SH SY-5Y are non-phagocytic, they may be internalized via caveolin-mediated and clathrin-mediated endocytosis. This suggests that, unlike the AuNPs uptake that varied according to the synthesis procedure, their excretion from the cells did not differ significantly.

In ethanol-positive cells, harvested at 1, 2, 4 and 8 days after ethanol discontinuation, iAuCs were either significantly lower than (as for AuNP<sub>KG</sub> and AuNP<sub>PEG</sub> pretreated cells) or comparable to (as for AuNP<sub>K</sub> or AuNP<sub>G</sub> pretreated cells) the corresponding values in ethanol-negative cells. At day-16, the two cell groups exhibited comparable values. This suggests that ethanol may differentially affect internalization of AuNPs that may be associated with unique properties of ethanol. Earlier studies have shown that ethanol, due to their polar and nonpolar characteristics, can (i) Leak through the cell membrane and form hydrogen-bonds with phospholipids head-group that disrupts the hydrophobic chain orientation and (ii) Increases the membrane fluidity by denaturing proteins and affecting the lipids, resulting in formation of small holes in the membrane [50,61,62]. Ethanol, by forming H-bonds may orient the phospholipids outwards, thus increasing their movement and the membrane fluidity increases and affecting the integrity of the membrane receptors [63]. Therefore, AuNP<sub>KGG</sub> being relatively smaller nanoparticles, may escape the cells via exiting through the ethanol-induced membrane holes.

Toxicity of plant-extract synthesized AuNPs

Although surface functionalized AuNPs, because of their unique properties, have emerged as promising candidates for biomedical applications such as sensing, imaging, and drug delivery [64], their application in clinical setting has not materialized due to uncertainties in their adverse effects. Since the NP toxicity is impacted by their size, shape, charge and surface functionalization, toxicity of each individual NP preparation must be evaluated before testing the beneficial effects. The present study showed that a 5 to 20 mg/L dose of AuNP preparations did not significantly affect (i) Viable cell enumeration; (ii) LDH and lipid peroxidase activities; (iii) NFkB activation and (iv) Apoptotic/necrotic cell enumerations. However, 40 to 100 mg/L dose caused a dose-dependent increase in these indices. This suggests that relatively higher doses (> 60 mg/L) of AuNPs may cause different degrees of cytotoxicity in SH SY5Y cells. These results are consistent with previous investigations performed with MCF-7, PC-3 and dermal fibroblasts [65,66] which demonstrated that the AuNPs, at higher doses (> 100 mg/L), impaired the proliferation of dermal fibroblasts and induced an abnormal formation of actin filaments, causing therefore a reduced cellular motility and influencing the cell morphology. Vijayakumar, Ganesan [67] reported that citrated and biotinylated 18 nm gold nanoparticles did not induce toxicity, while smaller particles were much more toxic in leukemic K562 cell lines. Similar results were reported by Cho, et al. [68] in adenocarcinomic human alveolar basal epithelial cell line, A549. These observations suggest that a 5 to 20 mg/L dose of AuNPs can be used safely in this experimental setup. However, toxicity must be reevaluated when the NPs physicochemical properties or the experimental conditions change.
Therapeutic potency of plant-extract synthesized AuNPs against ethanol toxicity

Ethanol cytotoxicity: It is well established that acute or chronic ethanol exposure (1) Up-regulated oxidative stress, dysregulated inflammation and cell signaling [20]; (2) Up-regulated expression of cytokines, increased amounts of TNF-α and depleted both cytosolic and mitochondrial GSH that may contribute to an increase in oxidative stress [69]; (3) Significantly inhibited activities of p450 enzymes CYP1A1, CYP2B6, CYP2C19, and CYP2D6 that may alter xenobiotic metabolism and drug pharmacokinetics [70], and (4) Significantly reduced total number of embryonic stem cells, inducing the formation of cardiomyocytes [62]. The present study demonstrated that a 54 mM concentration of ethanol exposure exerted immediate cytotoxicity in SH SY5Y cells possibly via the mechanisms listed above. The cytotoxic effects of ethanol persisted for up to 16 days after the cells were treated with ethanol, suggesting that 24 h ethanol exposure may have long-term cytotoxicity, possibly dysregulating oxidative stress and inflammation (Figure 16).

Possible mechanisms for protection against ethanol: The protective effects of plant-extract-synthesized AuNPs may be attributed to the adsorption of bioactive compounds onto the AuNPs surface. In the present study, the following ligands have been identified from the AuNPs surface: (1) Up-regulated NF-κB (a pro-inflammatory dimer of p50-p50) and suppression of p50-p65 (an anti-inflammatory dimer of p50-p65), (2) Up-regulated expression of cytokines, increased amounts of TNF-α and depleted both cytosolic and mitochondrial GSH that may contribute to an increase in oxidative stress [69], (3) Significantly inhibited activities of p450 enzymes CYP1A1, CYP2B6, CYP2C19, and CYP2D6 that may alter xenobiotic metabolism and drug pharmacokinetics [70], and (4) Significantly reduced total number of embryonic stem cells, inducing the formation of cardiomyocytes [62]. The present study demonstrated that a 54 mM concentration of ethanol exposure exerted immediate cytotoxicity in SH SY5Y cells possibly via the mechanisms listed above. The cytotoxic effects of ethanol persisted for up to 16 days after the cells were treated with ethanol, suggesting that 24 h ethanol exposure may have long-term cytotoxicity, possibly dysregulating oxidative stress and inflammation (Figure 16).

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matory dimer of NFκB), resulting in dysregulation of inflammation and activation of pro-apoptosis signaling [32]. The addictive effects of ethanol are mediated by the brain addiction pathways including dopamine, GABA, NMDA, and serotonin neurons [5,98,99]. Earlier studies have shown that kudzu root extract the brain addiction circuits, especially the dopamine, GABA, NMDA and Glu neurons/receptors in protecting the brain against the adverse effects of ethanol and suppressing the severity of the withdrawal symptoms [100-102].

Taken together, these observations suggest that pretreatment of cells with the AuNP-adsorbed plant-components are more potent in protecting the cells against ethanol toxicity than pretreatment of cells with crude plant extracts their purified components.

**AuNP-Protein Interaction:** It is well established that NPs in biological and environmental media (blood, intracellular fluid, plant extracts or environmental samples) interact with the media components such as soluble proteins, lipids and other ingredients, and form a surface corona that may modulate the NPs’ physicochemical and biological (cellular/tissue responses such as cellular uptake, kinetics, signaling, accumulation, transport, and toxicity) properties [103-108]. Formation of protein corona in blood or intracellular fluids may depend upon the composition and organization of surface proteins [109]. In general, the corona may be divided into two components: the “hard” corona with long exchange time and “soft” coronas with short exchange times [109]. In a biological fluid, the surface corona’s protein composition, amount and presentation on the surface, modify their beneficial, therapeutic or adverse effects [110-115]. In the present study, 9 proteins, ranging from <10 to 200 kDa, have been identified in cytosol and hard-corona of AuNPs synthetized using different plant extracts and incubation temperature (Figure 14A). AuNP<KG>50 (average diameter - 8 nm) AuNP<K>G50 (average diameter - 9 nm) and AuNP<K>G50 (average diameter - 41) interacted with protein <37 kDa, 50 to 37 kDa and >75 kDa, respectively. This suggests that the size and surface ligand composition may determine the composition of proteins corona. As shown in Figure 14B, the time-course of change in individual corona proteins paralleled the change in corresponding cytosol proteins, suggesting the possibility of rapid exchange between corona and cytosol. It is hypothesized that poor protein-binding capacity of AuNP<KG>50 may contribute to its high therapeutic activity and relatively lower toxicity.

**Conclusion**

Although kudzu extract, edible gum extract and a combination of both (AuNP<KG>37, -50, -90) incubated at 37 °C, 50 °C and 75 °C, reduced Au<sup>3+</sup> into colloidal gold nanoparticles (kudzu synthesized: AuNP<KG>50, -90, -37°, gum synthetized: AuNP<sup>G</sup>G<sup>KG</sup>50, G<sup>KG</sup>G<sup>G</sup>50 and kudzu+gum synthesized: AuNP<sup>KG</sup>G<sup>KG</sup>37, -50, -37°), they differed in size (AuNP<KG><sub>K</sub> << AuNP<sub>K</sub> < AuNP<sub>G</sub> < AuNP<sub>KG</sub>) and physicochemical properties, including their interaction with cytosol proteins (AuNP<sub>KG</sub> << AuNP<KG>). The surface of each AuNP contained the extract’s active ingredients that were analyzed and confirmed using LDI and low-matrix MALDI. AuNP<sub>KG</sub> was least toxic to SH-SY5Y cells and most effective in suppressing the adverse effects of ethanol on SH-SY5Y cells. The beneficial and adverse effects of AuNPs may have been modified by the formation of proteins corona.

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