

Supporting Information

Effect of Surface-engineered AuNPs on gene expression, bacterial interaction, protein denaturation, and toxicology assay: An *in vitro* and *in vivo* model

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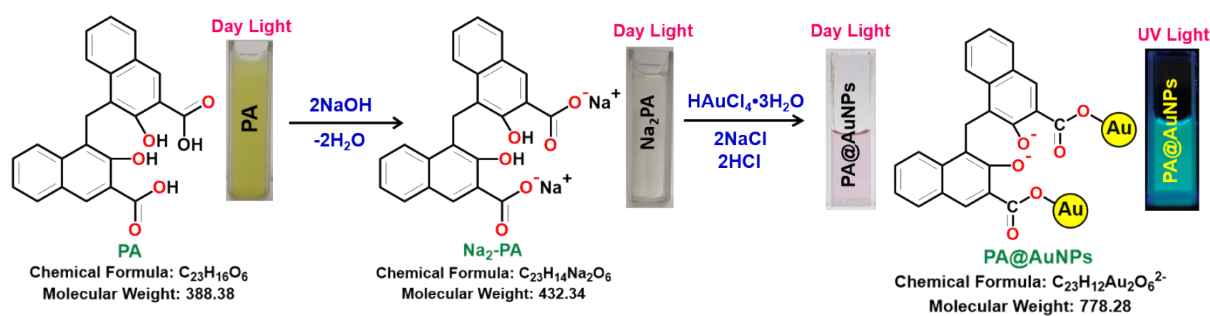
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1. Synthesis of PA@AuNPs



Scheme SI1. Synthesis of PA@AuNPs (inset images: corresponding reaction solutions under daylight and UV light).

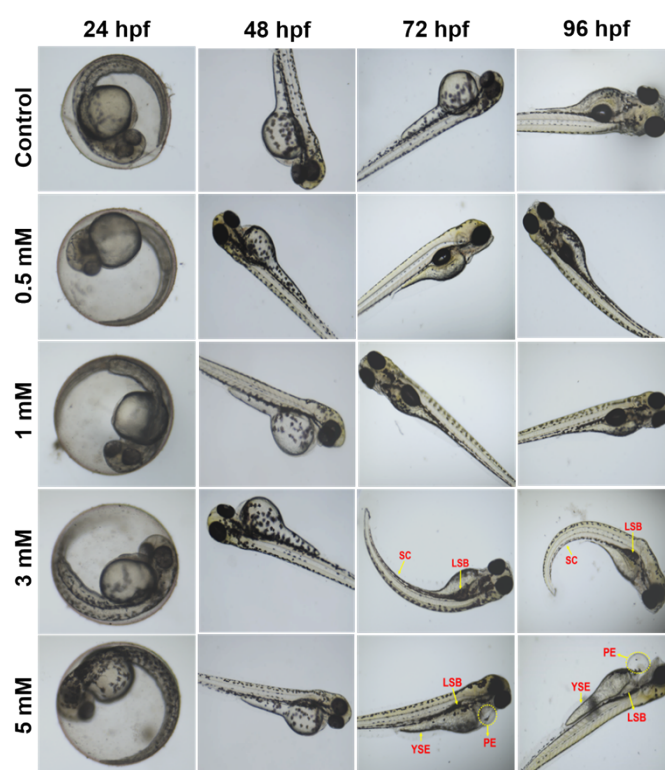


Figure SI1. Morphological changes in zebrafish embryos affected by various concentrations of PA@AuNPs (0.5, 1, 3, and 5 mM). The control showed normal development of an embryo, and the PA@AuNPs exposed to the embryo showed deformities like SC-spinal curvature, LSB-lack of the swim bladder, YSE-yolk solk edema, and PE-pericardial edema.

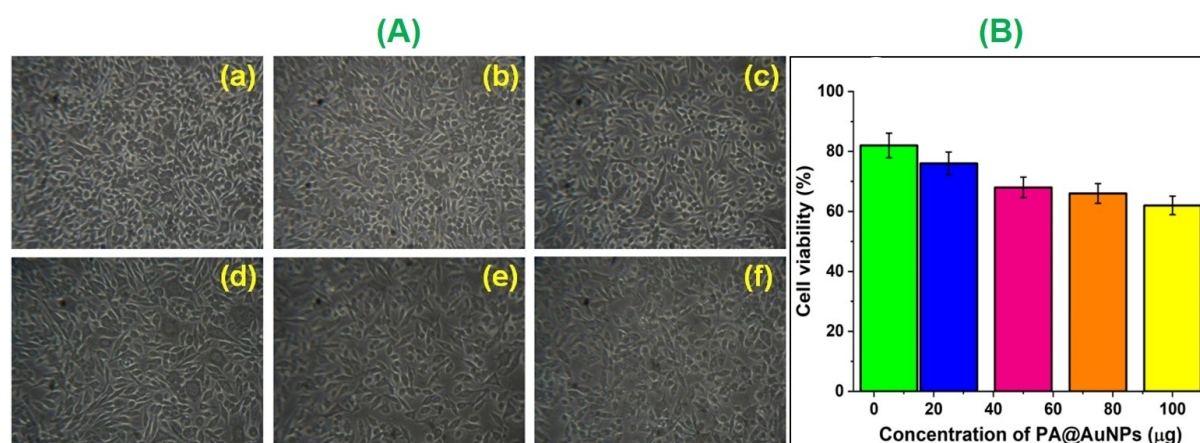


Figure SI2. (A) The cytotoxic effect of PA@AuNPs (0, 5, 25, 50, 75, and 100 μL) with L929 fibroblast cells by the standard MTT viability assay. L929 fibroblast cells were incubated with PA@AuNPs for 24 h (images (a) untreated L929 fibroblast cells, (b) L929 fibroblast cells + 5 μL PA@AuNPs, (c) L929 fibroblast cells + 25 μL PA@AuNPs, (d) L929 fibroblast cells + 50 μL PA@AuNPs; (e) L929 fibroblast cells + 75 μL PA@AuNPs, and L929

fibroblast cells + 100 μ L PA@AuNPs, respectively). (B) Evaluation of cell viability of PA@AuNPs (0, 5, 25, 50, 75, and 100 μ g) by the standard MTT assay.

2. Method

2.1. Cell culture

The cell line was cultured in a 25 cm² tissue culture flask with DMEM supplemented with 10% FBS, L-glutamine, sodium bicarbonate, and an antibiotic solution containing, Penicillin (100U/ml), Streptomycin (100 μ g/ml), and Amphotericin B (2.5 μ g/ml). Cultured cell lines were kept at 37 °C in a humidified 5% CO₂ incubator. The viability of cells was evaluated by direct observation of cells by an inverted phase contrast microscope and followed by the MTT assay method.

2.2 Cells seeding in 96 well plates

A two-day-old confluent monolayer of cells was trypsinized and the cells were suspended in 10% growth medium, 100 μ L cell suspension (5×10^3 cells/well) was seeded in 96 well tissue culture plate and incubated at 37 °C in a humidified 5% CO₂ incubator.

2.3. Isolation of total RNA by TRIzol Method

A total RNA sample was isolated following the manufacturer's instructions. By adding TRIzol solution, cells are disrupted and release RNA. As a result of adding isopropanol, RNA precipitates as a white pellet around the tube's sides and bottom. After adding 1 ml of TRIzol reagent, the culture well plate was incubated for 5 min. This mixture was transferred to a new, sterile Eppendorf tube, and 200 μ L of chloroform was added. It was then forcefully shaken for 15 sec and allowed to incubate for two to three minutes at ambient temperature. Finally, it was centrifuged for fifteen minutes at 4 °C at 14000 rpm. After gathering the obtained aqueous layer, 500 μ L of 100% isopropanol was added, and the mixture was incubated for 10 minutes at ambient temperature before being centrifuged for 15 min at 4 °C at 14000 rpm. The pellet was cleaned with 200 μ L of 75% ethanol after the supernatant was removed, and it was then centrifuged for five minutes at 4 °C at 14000 rpm in a cooling centrifuge. After drying the RNA pellet was suspended in TE buffer.

2.4. cDNA Synthesis

With the TRIzol reagent, total RNA was isolated, and then total RNA's concentration and purity were assessed. With the help of a cDNA preparation kit, the template

complementary DNA was synthesized. About 5 μ L of RT easy mix, 0.5 μ L of oligodT, and an RNase-free tube were filled with 2 L of RNA template (0.5 g total RNA). Then the total reaction solution was made up to 10 μ L with the addition of sterile distilled water, and then the solution was mixed gently. For, cDNA synthesis by using a thermal cycler. The cycling conditions were applied for 20 min at 42 °C and 5 min at 85 °C (Table SI1).

Table SI1. Temperature and time variations in cycling conditions.

Step	Temperature °C	Time (minutes)	No. of cycles
Synthesis of cDNA	42	20	1
Inactivation	85	5	1

3. Characterization

Table SI2. Statistical analysis of anticancer assay by MTT Method

Cell line - MDA MB 231									
SAMPLE CODE									
	OD1	OD2	OD3	% viability 1	% viability 2	% viability 3	Average	Stdev	Std error
Control	0.72	0.72	0.71	100	100	100	100	0	0
6.25	0.61	0.62	0.62	84.52	85.50	87.44	85.82	1.48	0.49
12.5	0.51	0.51	0.51	70.03	70.93	71.13	70.70	0.58	0.19
25	0.41	0.40	0.40	57.55	54.94	56.15	56.21	1.30	0.43
50	0.34	0.33	0.33	47.50	45.47	46.25	46.41	1.02	0.34
100	0.23	0.22	0.22	32.80	31.34	31.92	32.02	0.73	0.24

Table SI3. Statistical analysis of Gene expression study

	Experimental Well 1	Experimental Well 2	Experimental Well 3	Control Well 1	Control Well 2	Control Well 3	Average Experimental Ct Value	Average Experimental Ct Value	Average Control Ct Value	Average Control Ct Value	Δ Ct Value (Experimental)	Δ Ct Value (Control)	Delta Delta Ct Value	Expression Fold Change
	Raw Ct Value	Raw Ct Value	Raw Ct Value	Raw Ct Value	Raw Ct Value	Raw Ct Value	TE	HE	TC	HC	Δ CTE	Δ CTC	$\Delta\Delta$ Ct	$2^{-\Delta\Delta$ Ct
Housekeeping Gene	20.43	20.16	20.38	22.29	22.08	22.24	-	20.32	-	22.20	7.25	6.96	0.29	0.816014485
Gene being Tested	27.88	27.59	27.26	29.35	29.06	29.08	27.58	-	29.16	-				

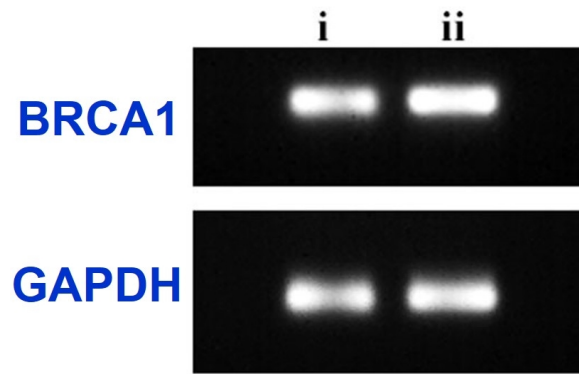


Figure SI3. Agarose gel electrophoresis image of GAPDH and BRCA1.

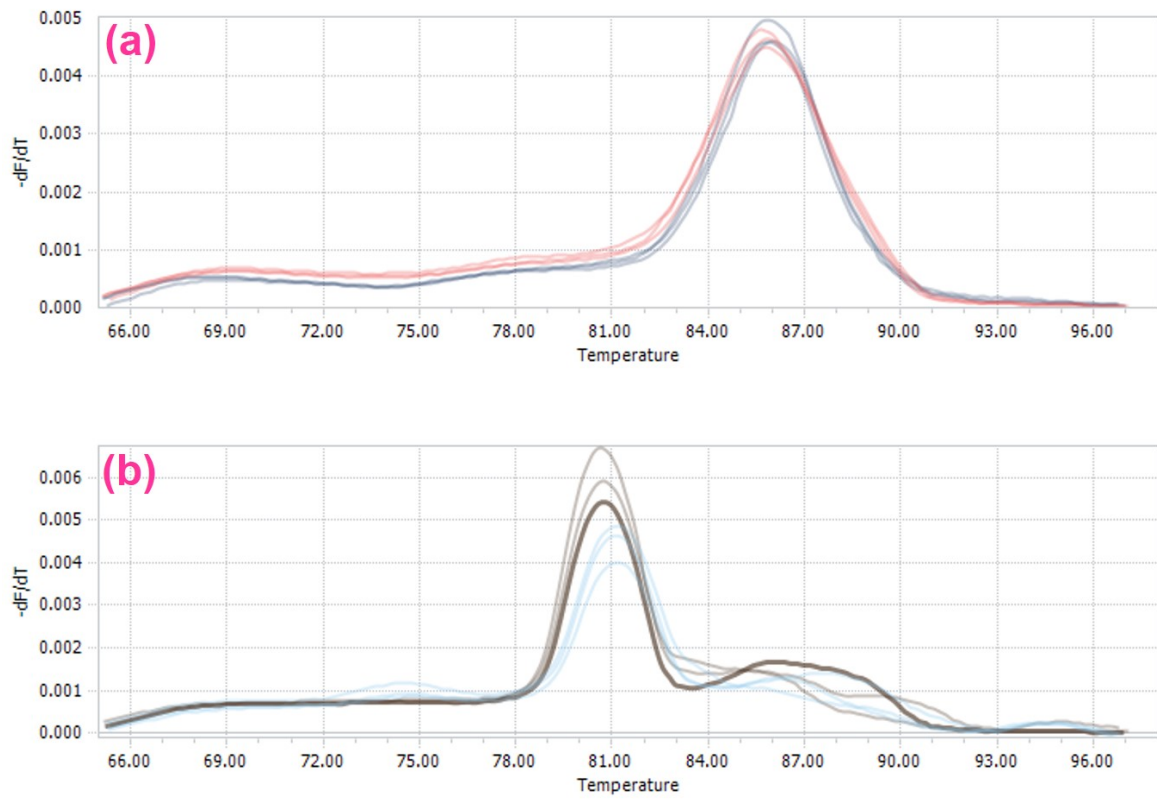


Figure SI4. Real-time melting peaks of PA@AuNPs, (a) GAPDH and (b) BRCA – 1.

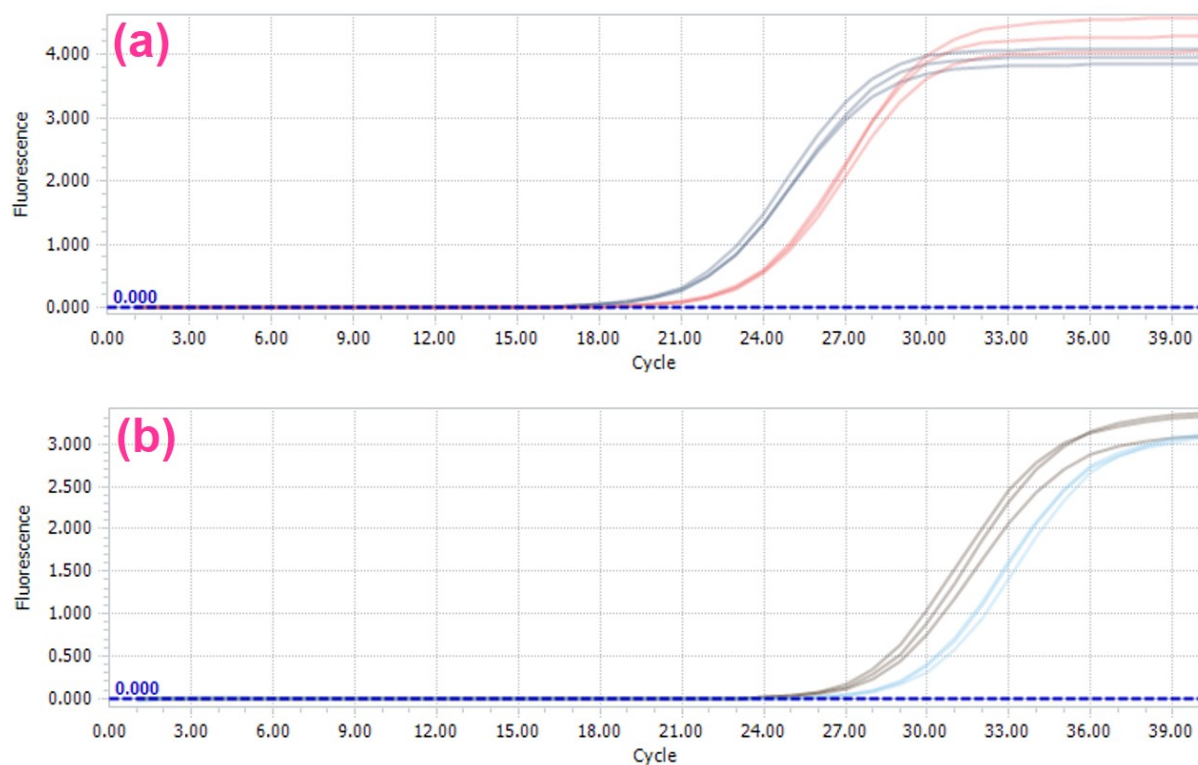


Figure SI5. Real-time amplification curves of PA@AuNPs, (a) GAPDH and (b) BRCA – 1.

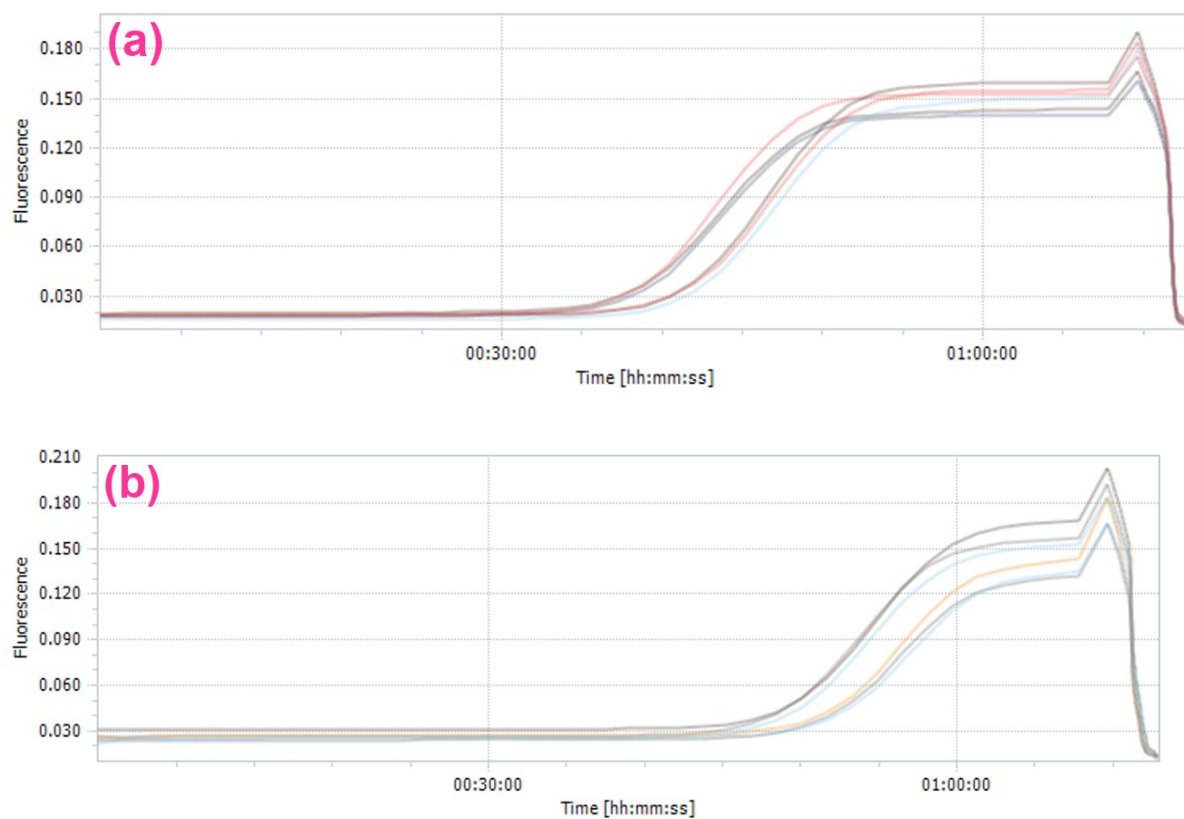


Figure SI6. Real-time fluorescence curves of PA@AuNPs, (a) GAPDH and (b) BRCA – 1.

Table SI4. The various steps involved, the time taken, and the temperature of each step.

Steps	Time required	Temperature
Initial activation step	2 min	95 °C
3 step cycling		
Denaturation	10 sec	95 °C
Annealing	1 min	58 °C
Extension	1 min/kb	72 °C
Number of cycles	40 cycles	68 °C
End of PCR cycling	Indefinite	4 °C

Table SI5. Gross morphological changes in vital organs after acute toxicity testing of PA@AuNPs in Sprague-Dawley rats

Parameters	Groups					
	Control			PA@AuNPs		
	Heart	Liver	Kidney	Heart	Liver	Kidney
Organ Weight (gms)	1.06 ± 0.24	11.07 ± 1.02	1.08 ± 0.12	1.04 ± 0.12	11.69 ± 1.19	1.08 ± 0.04
Discoloration	Nil	Nil	Nil	Nil	Nil	Nil
Lesion	Nil	Nil	Nil	Nil	Nil	Nil
Structural change	Nil	Nil	Nil	Nil	Nil	Nil

Table SI6. Histological scoring of stained organ sections after acute toxicity testing

Sl. No.	Group	Organs		
		Heart	Liver	Kidney
1	Control	0	0	0
2	PA@AuNPs	0	0	0

Table SI7. Parameters under consideration during acute toxicity study.

Sl. No.	Parameters	Control (Out of 5)	PA@AuNPs (Out of 5)
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		animals)	animals)
1	Mortality	0	0
2	Convulsions	0	0
3	Weight loss ($\geq 2g$)	0	0
4	Salivation	0	0
5	Piloerection	0	0
6	Abnormal Reflexes	0	0
7	Gastro-intestinal abnormality	0	0
8	Muscular tone	0	0
9	Motor Control	0	0
10	Analgesia	0	0

Table SI8. Gross morphology surrounding PA@AuNPs after subcutaneous implantation.

Sl. No.	Parameter	PA@AuNPs	
		Week 2	Week 4
1	Edema	Nil	Nil
2	Hemorrhage	Nil	Nil
3	Pus or discharge	Nil	Nil
4	Lesion or ulceration	Nil	Nil
5	Necrosis of adjacent tissue	Nil	Nil
6	Vascularization	moderate	excellent

Table SI9. Comparative table of AuNPs for *in vitro* and *in vivo* analysis.

S. No.	Materials used	<i>In vitro</i> and <i>in vivo</i> models	Ref.
1	AuNPs	TNBC – MDA-MB-231	1
2	AuNPs	TNBC – MDA-MB-231	2
3	Polyethylene Modified Au-based SiRNA	TNBC – MDA-MB-231	3

4	Thioglucose-bound AuNPs	TNBC – MDA-MB-231	4
5	AuNPs	BC – MCF-7	5
6	AuNPs	TNBC – MDA-MB-231	6
7	Citrate-capped-AuNPs	TNBC – MDA-MB-231	7
8	Au-Curcumin (Au-C), Au-Paclitaxel (Au-P), Au-Curcumin-Paclitaxel (Au-CP)	TNBC – MDA-MB-231	8
9	AuNPs	foodborne bacterial pathogens	9
10	A-AuNPs and M-AuNPs	<i>E. coli</i> , MDR <i>E. coli</i> , <i>P. a</i> , MDR <i>P. a</i> , MDR <i>K. p</i>	10
11	Polyelectrolyte-wrapped AuNPs	<i>Shewanella oneidensis</i> and <i>Bacillus subtilis</i>	11
12	AuNPs (1000 mg /kg)	Mice and Rats	12
13	Pectin-mediated-AuNPs 500 – 1000 µg	Sprague Dawley Rats	13
14	AuNPs 0.9, 9, 90 µg	Sprague Dawley Rats	14
15	PA@AuNPs	TNBC – MDA-MB-231, Sprague Dawley Rats, Bacterial species, and Protein denaturation	Current work

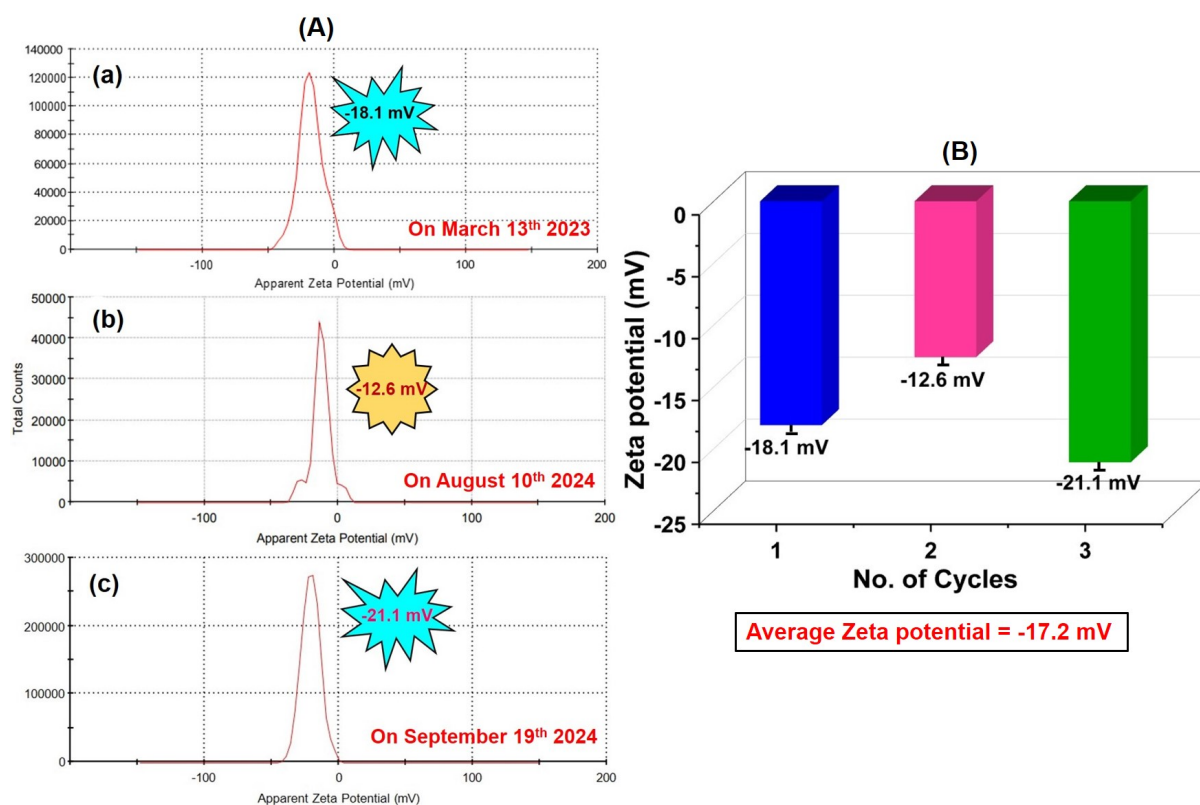


Figure SI7. (A) Different time interval measurements of zeta potential of PA@AuNPs (Average zeta potential = -17.2 mV), (B) Corresponding bar graph (n = 3%).

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