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## Multi-endpoint assessments for *in vitro* nano-bio interactions and uptake of biogenic phosphorus nanomaterials using HEK293 cells†

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Phosphorus (P)-based nanomaterials (NMs) are being explored as nanofertilisers due to higher uptake in plants, enhanced nutrient use efficiency, and minimal leaching. However, their colloidal behaviour in biologically relevant media and corresponding *in vitro* nano-bio interactions have been understudied topics for the evaluation of their potential cytotoxicity. Using HEK293, in this study, we investigated the interactions and any associated cytotoxicity of four different nanohydroxyapatites (nHAPs: (i) biogenic platelet-shaped, chemically synthesized (ii) nanorods, (iii) spherical, (iv) nanoneedles) and nanophosphorus (spherical biogenic nP derived from rock phosphate). The NM's interaction with cell culture media was studied by dynamic light scattering, transmission electron microscopy, and elemental and protein corona characterisation. The investigation on acute effects such as mitochondrial and lysosomal activities, membrane disintegrity, ROS generation, and cell viability was performed by following NIH guidelines. In addition to the dose-dependent approach, the cytotoxic effects of NM on changing the initial cell seeding density were also monitored. Our results suggest that physicochemically different P-based NMs interact differently with cell culture media resulting in NM transformation, minimal P release, and protein corona formation. Lipoprotein and albumins were the major components of the hard corona. No LC<sub>50</sub> values were attained for any of the cytotoxic endpoints up to 1000 μg mL<sup>-1</sup>. Fluorescence and electron microscopy confirmed the uptake of NMs in the cells. This is the first report on the nano-bio interactions of differently shaped and sized P-based NMs using colloidal chemistry and two independent approaches-variable NM dose and variable cell densities. The findings suggest low cytotoxicity and hence the potential suitability of using the biogenic P-based NMs in agriculture as nanofertilisers.

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### Environmental significance

The use of phosphorus (P) – based nanomaterials (NMs) as fertilisers to increase agricultural productivity is being investigated. It is therefore critical to investigate the colloidal behaviour of P-based nanofertilisers in biological media relevant to human physiology while assessing their cytocompatibility. This results in dynamic changes in NM behaviour in terms of protein corona acquisition, change in size, and P release. Bridging such changes with detailed cytological effects of NMs remains an understudied topic particularly for biosynthesised NMs. This study reports the NMs interaction with cell culture media, cellular uptake, and acute effects of P-based NMs on human cells using multi-end point assays. The study provided evidence of nano-safety that supports the recommendation of environmentally relevant concentrations and dosages of P-based nanofertilisers.

## 1 Introduction

Nanomaterials (NMs) have been explored for their various applications in medicine and agriculture. Considering the depleting phosphorus (P) levels in agricultural soils across the globe, P-based NMs are proposed to be used as nanofertilisers.<sup>1</sup> Lab-scale research on P-based NMs has shown promising fertilising effects.<sup>2–6</sup> However, before agricultural applications, it is important to understand risks from the perspective of human health considering occupation safety and end-user applications. As compared to conventional fertilisers, these

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UV-vis (Shimadzu, Japan), FTIR (Nicolet 6700, Thermofisher Scientific, USA), and dynamic light scattering after dispersal in cell culture media containing 10% fetal bovine serum (FBS). The shape and size of NMs in *in vitro* cell culture media were determined using transmission electron microscopy analysis (200 kV, Tecnai G2 30-U twin microscope, FEI, USA).

The release of P from the biogenic nHAP and nP was calculated by using the previously published dialysis method.<sup>42</sup> The dialysis membranes (14k MWCO and pore size = 3.9 nm) were washed and activated. The NMs suspensions (1 mg mL<sup>-1</sup>) were prepared in DI water and serum-supplemented cell culture media (CCM) and were added to the activated dialysis membrane. These dialysis bags were added to the containers with DI water and incubated at 37 °C and 5% CO<sub>2</sub>. Samples were then taken at 0, 2, 4, 6, 8, and 24 h to measure the P-release from the NMs by measuring the P contents using flame photometry. The initial content of P was 23.22% and 0.89% in nHAP and nP, respectively.

### 2.3 Protein corona study on nHAPs and nP in serum-supplemented media

Here a comparative analysis of P-based NMs–protein interactions with the application of cell culture media supplemented with FBS was performed to investigate the formation of the hard and soft protein corona. Briefly, the NMs at 1 mg mL<sup>-1</sup> were incubated with 500 µL cell culture media + 10% FBS for 1 h at 37 °C under constant agitation. These were then separated from the supernatant by centrifugation at 15 000 rcf for 30 min. The NMs were washed with 1 × PBS in three centrifugation steps at 15 000 rcf for 30 min. The pellet obtained after copious washing was analysed for protein attachment as the hard corona. The washes were kept for soft corona assessment. Sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE), a protein quantification assay using Bradford's reagent,<sup>43</sup> and fast protein liquid chromatography (FPLC, GE AKTA, Germany) were used to analyse the protein corona.

### 2.4 Cell culture

Commercially available HEK293 was procured from National Centre for Cell Science, Pune, India. The cells were cultured with DMEM + 10% FBS, supplemented with 1% glutamine, and 1% penicillin–streptomycin. The cells were incubated at 37 °C and 5% CO<sub>2</sub>. The culture medium was changed every 3–4 days until the cells reached confluence. The adherent cells were then detached with a solution of 0.05% trypsin and 0.53 mM EDTA and the cells were then passaged.

### 2.5 Cellular metabolic activity assessment using MTT and neutral red uptake assays

A total of 1 × 10<sup>4</sup> cells were seeded in 96 well plates before the exposure to nHAP and nP. At a confluent stage of 70–80%, the cells were exposed to increasing concentrations (1.56, 3.12, 6.25, 12.5, 25, 50, 100, 200, 500, and 1000 µg mL<sup>-1</sup>) of nHAP\_B, nHAP\_C, nHAP\_Sigma, nHAP\_SRL, nP, and RP. The time of exposure was kept at 24 h to study the acute effects of NMs. The MTT assay was carried out according to the protocol described

by NCL, NIH, USA,<sup>44</sup> and a neutral red uptake assay was performed as described by Borenfreund and Puerer.<sup>45</sup> All the samples, along with positive (0.1% (v/v) Triton X), negative (no treatment, only cells), and media controls (no treatment, no cells) were run in triplicates. Absorbances were measured at 570 nm and 540 nm for MTT and neutral red assays, respectively, with a reference wavelength of 680 nm.

### 2.6 Cell membrane integrity assessment using LDH and trypan blue assays

The effects of NMs on cell membrane integrity were estimated using lactate dehydrogenase release and trypan blue dye inclusion assays. The cells were seeded and exposed to NMs as described in the section above. The LDH assay was carried out according to the protocol described by NCL, NIH, USA<sup>44</sup> and the trypan blue dye inclusion assay was performed as described by Uliasz and Hewett.<sup>46</sup> All the samples, positive (0.1% (v/v) Triton X), negative (no treatment, only cells), and media controls (no treatment, no cells) were run in triplicate. Absorbances were measured at 490 nm and 590 nm for LDH and trypan assays, respectively, with a reference wavelength of 680 nm.

### 2.7 Effect of NMs interaction on HEK293 with decreasing cell seeding density – an assessment by MTT for cellular metabolic activity

To analyze the nano-bio interactions w.r.t. variable cell densities, three different initial cell densities were chosen. A total of 1 × 10<sup>4</sup>, 1 × 10<sup>3</sup>, and 1 × 10<sup>2</sup> cells were seeded in separate 96 well plates before the exposure to nHAP and nP. The evaluation of mitochondrial activity was performed in a similar manner as stated above in Section 2.4.

### 2.8 Intracellular oxidative stress assessment using DCFH\_DA enzymatic assays

Oxidative stress due to exposure to NMs was determined using the DCFH\_DA assay. The analysis was performed by following the protocol described by NCL, NIH, USA.<sup>47,48</sup> For the DCFH\_DA assay, briefly, the cells were seeded at a density of 1 × 10<sup>5</sup> cells per well and cultured for 24 h at 37 °C in a 12-well plate. After incubation, the media was discarded, and the cells were treated with different concentrations of NMs (50 and 500 µg mL<sup>-1</sup>) for 24 h. Post-exposure, the media was discarded and 20 µM of DCFDA solution was added to the wells with fresh media. The test plate was taken at 3 h post DCFDA addition and read at ex. 485 nm and em. 530 nm. The fold change in ROS activity was reported with respect to the negative control (untreated cells) and 300 µM of H<sub>2</sub>O<sub>2</sub> was taken as the positive control.

### 2.9 Uptake of fluorescent-tagged nanohydroxyapatite particles

FITC-tagged nHAP\_B was used to see the time-dependent uptake of particles under a fluorescence microscope. The cells were seeded in a 35 mm culture dish at a seeding density of 0.3 × 10<sup>6</sup> cells. After 75–80% confluence these were exposed to 100 µg mL<sup>-1</sup> of FITC-tagged nHAP\_B. At different time points (2, 4,



6, 8, and 24 h), the exposure solution (media + NMs) was aspirated completely, and the cells were gently washed three times with  $1\times$  DPBS. Cells were fixed and after fixation imaged immediately under a fluorescence microscope (EVOS M5000 imaging system, ThermoFisher Scientific, USA) at ex. 488 nm and em. 516 nm. Untreated cells were also viewed under a fluorescence microscope to rule out autofluorescence.

### 2.10 TEM visualization of cells exposed to biogenic P-based NMs

For tracking cellular uptake and localization, TEM analysis was performed after cells were exposed to  $100\ \mu\text{g mL}^{-1}$  of biogenic nHAP\_B and nP. The cells were washed twice with  $1\times$  DPBS and detached. The cells were gently centrifuged at 800 rcf and washed with  $1\times$  DPBS. Cells were pre- and post-fixed with 2.5% glutaraldehyde solution for 30 minutes and 1% osmium tetroxide for 2 h, respectively. Post fixation, the samples were sequentially dehydrated using alcohol gradient and acetone. After dehydration, the samples were cast in blocks and infiltrated with epoxy resin. Samples blocks were sectioned using microtomy. Thin sections of 60–100 nm were placed on carbon-coated copper grids. Samples were stained, and after that, the resin sample block was trimmed and thin-sectioned at a thickness of 60–100 nm, followed by the collection on carbon-coated copper grids. The grids were stained primarily with 1% uranyl acetate and 2% lead citrate was used as a secondary stain. TEM (Tecnai G2 F20 S-TWIN, US) coupled with energy-dispersive X-ray analysis (EDX) was used to capture images and to evaluate the presence of particles inside the cells.

### 2.11 Acridine orange/ethidium bromide dual staining for cell viability

Cells seeded on a coverslip at a density of  $1\times 10^5$  cells per well were cultured for 24 h at 37 °C in a 12-well plate. After incubation, the media was discarded, and the cells were treated with different concentrations of NMs (100 and 1000  $\mu\text{g mL}^{-1}$ ) for 24 h. Untreated cells were taken as a negative control and the treatment with 300  $\mu\text{M}$  of  $\text{H}_2\text{O}_2$  was taken as a positive control. Cells were washed twice with  $1\times$  DPBS, fixed, and stained using dual fluorescent staining solution (1 : 1 mixture of 100  $\mu\text{g mL}^{-1}$  of each acridine orange and ethidium bromide) for 5 min. Excess stain was removed by washing the cells with  $1\times$  DPBS. Cells were observed under a fluorescence microscope with an automated cell counter (Cytation 5, Biotek, Agilent Technologies, USA). A total of 3715 cells as determined by the automated cell counter were analysed in each case.

### 2.12 Statistical analyses

For each of the NM (nHAPs and nP), we characterized dose-effect relationships. All statistical analyses were performed using GraphPad Prism software, Version 9 (GraphPad Software Inc., San Diego, CA, USA). Prior to statistical analysis, the normal distribution of residuals was checked with the Shapiro-Wilk test and QQ plots. Results were subjected to analysis of variance for significance followed by Dunnett *post hoc* multiple

comparison tests, to identify the treatment groups with significant differences in mean values.

## 3 Results

### 3.1 Physicochemical characterization of P-based NMs in cell culture media

The TEM micrographs (Fig. 1A) show the particles along with the media components such as proteins. The EDX results suggest that the Ca/P ratio was changed from 1.58, 1.79, 1.55, 1.62, and 3.69 to 1.6, 1.57, 1.52, 1.53, and 1.07 for nHAP\_B, nHAP\_C, nHAP\_Sigma, nHAP\_SRL, and nP, respectively. The primary sizes of NMs at a concentration of 1 mg mL<sup>-1</sup> in CCM from TEM were  $37.86 \pm 8.8$  nm for nHAP\_B; L:  $87.69 \pm 14.5$  nm and W:  $25.97 \pm 5.7$  nm for nHAP\_C;  $43.12 \pm 13.02$  nm for nHAP\_Sigma; L:  $84.04 \pm 16.76$  nm and W:  $18.42 \pm 3.76$  nm for nHAP\_SRL;  $18.41 \pm 9.69$  nm for nP. The histograms for the primary sizes obtained from TEM are presented in ESI Fig. 1.† As reported previously,<sup>30</sup> the hydrodynamic sizes of the pristine NMs in de-ionised water were  $325 \pm 37.1$  nm,  $756.2 \pm 28.8$  nm,  $874.3 \pm 51.5$  nm,  $892.8 \pm 21.1$  nm, and  $798.5 \pm 23.4$  nm, respectively, for nHAP\_B, nHAP\_C, nHAP\_Sigma, nHAP\_SRL, and nP. Upon interaction with 10% serum-supplemented CCM (Fig. 1B), the sizes were changed to  $899.7 \pm 77.27$  nm,  $842.4 \pm 86.84$  nm,  $411.5 \pm 6.56$  nm,  $530.4 \pm 31.57$  nm,  $989.9 \pm 52.11$  nm for nHAP\_B, nHAP\_C, nHAP\_Sigma, nHAP\_SRL, and nP, respectively. In contrast, when the NMs were dispersed only in the CCM without serum, the hydrodynamic sizes were found to be  $959.5 \pm 115.47$  nm,  $1122.8 \pm 373.05$  nm,  $441.4 \pm 92.09$  nm,  $794.83 \pm 231$  nm, and  $1201.5 \pm 256.34$  nm for nHAP\_B, nHAP\_C, nHAP\_Sigma, nHAP\_SRL, and nP, respectively. Correspondingly, the zeta potential values were also changed.

It is hypothesized that once these NMs disperse within the CCM, they will interact with the cells either *via* colloidal dispersion or by sedimentation in the cell culture vessel. For each of the NM tested, a theoretical analysis of the concentration-dependent particle numbers corresponding to either colloidal dispersion or sedimentation was carried out and is presented in ESI Tables 1–3.† It can be observed that for P-based NMs, the number of particles interacting with cells *via* colloidal dispersion is more than by sedimentation at any tested concentration from 1.5–1000  $\mu\text{g mL}^{-1}$ .

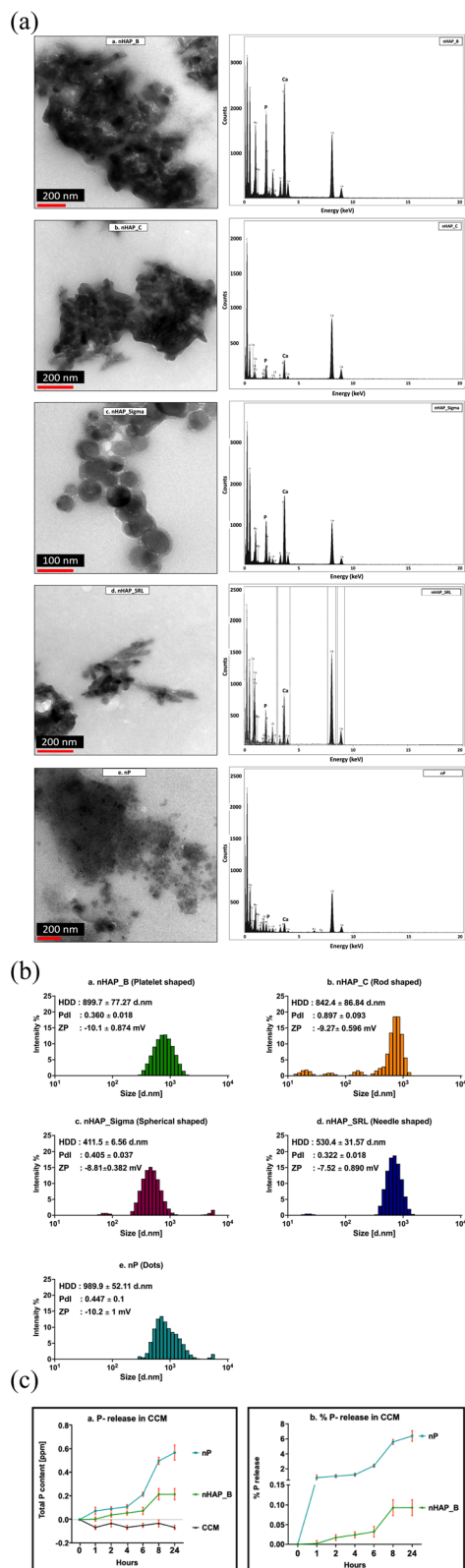
Further, on examining the release kinetics of P (Fig. 1C) from biologically synthesized NMs dispersed in serum-supplemented cell culture media at 1000  $\mu\text{g mL}^{-1}$ , it was found that for nHAP\_B, the maximum release was  $0.093 \pm 0.035\%$  at 24 h and for nP, it was  $6.38 \pm 1.24\%$  at 24 h.

### 3.2 Evaluation of protein corona formation on P-based NMs

The protein content in the corona was determined in different washes (W) and in hard corona (HC) by Bradford's method. After 4 washes (W1, W2, W3, and W4), the protein content in HC was found in the order of nHAP\_Sigma > nP > nHAP\_B > nHAP\_C (Fig. 2(I)). The SDS-PAGE results show distinct bands of HC for NM samples (Fig. 2(II)) and the quantified band intensities (Fig. 2(III)) corroborated with the protein







**Fig. 1** (A) TEM micrographs and EDX spectra for (a) nHAP\_B, (b) nHAP\_C, (c) nHAP\_Sigma, (d) nHAP\_SRL and (e) nP. (B) DLS intensity vs. size distribution for (a) nHAP\_B, (b) nHAP\_C, (c) nHAP\_Sigma, (d) nHAP\_SRL and (e) nP [HDD: hydrodynamic diameter, PdI: polydispersity index and ZP: zeta potential]. (C) Phosphorus release from NMs (nHAP, nP – 1000  $\mu\text{g mL}^{-1}$ ) dispersed in cell culture media (CCM) and control (cell culture media only) with respect to time. (a) shows

concentration trend observed by Bradford's method. Complete protein profiles are shown by FPLC graphs for the HC samples (Fig. 2(IV)). Peaks for lipoprotein, IgG, IgA, BSA, and  $\beta$ -lactoglobulin were found in 10% FBS. Of these, lipoprotein was present in HC of all the NMs, IgG was present in HC of nHAP\_B, IgA was present in HC of nHAP\_Sigma, BSA, and  $\beta$ -lactoglobulin was present in HC of nHAP\_B, nHAP\_Sigma, and nP.

### 3.3 Effect of NMs on cellular metabolic activities

A dose-dependent decrease in cellular mitochondrial activity was seen for all NMs (Fig. 3A). The corresponding linear regression graphs and equations are presented in ESI Fig. 2A.† Nearly 20% decrease in mitochondrial activity was observed for different treatments for the environmentally relevant concentration of 100  $\mu\text{g mL}^{-1}$ . The minimum cellular mitochondrial activities were noted at the highest tested concentration of 1000  $\mu\text{g mL}^{-1}$ . A neutral red uptake assay was performed to assess the lysosomal activity on exposure to NMs and a significant ( $p < 0.001$ ) dose-dependent decrease was seen (Fig. 3B) with a minimum activity at the highest tested concentration of 1000  $\mu\text{g mL}^{-1}$ . The linear regression graphs and equations for the neutral red uptake assay are presented in ESI Fig. 2B.† Similar to the mitochondrial activity, the lysosomal activity was also decreased by  $\sim 20\%$  on exposure to an environmentally relevant concentration of 100  $\mu\text{g mL}^{-1}$ . The rod-shaped nHAP\_C caused a maximum decrease in cellular metabolic activities followed by the needle-shaped nHAP\_SRL, platelet-shaped nHAP\_B, and spherical nHAP\_Sigma. For both assays, none of the NMs had  $\text{LC}_{50}$  at any tested concentrations. On the other hand, for bulk RP,  $\text{LC}_{50}$  was attained at much lower tested concentrations.

### 3.4 Effect of NMs on cellular membrane integrity

The membrane integrity was evaluated at a range of concentrations from 1.5 to 1000  $\mu\text{g mL}^{-1}$ . A dose-dependent increase in LDH leakage (Fig. 4A) was seen for all NMs. The maximum values were noted at the highest tested concentration of 1000  $\mu\text{g mL}^{-1}$ . The percentage of LDH leakage was  $< 20\%$  in all cases of exposure. The corresponding linear regression graphs and equations are presented in ESI Fig. 3A.† Trypan blue dye inclusion assay was performed to assess the membrane dis-integrity on exposure to NMs and a dose-dependent increase was seen (Fig. 4B). On comparing with 0.1% triton-positive control (100% membrane dis-integrity), it was found that the highest exposure concentration of 1000  $\mu\text{g mL}^{-1}$  resulted in maximum %membrane dis-integrity. The membrane dis-integrity was  $< 20\%$  for all tested concentrations of NMs except at the highest concentrations for rod-shaped nHAP\_C and nP. For both assays, none of the NMs had  $\text{LC}_{50}$  at any tested concentrations. The linear regression graphs and equations for the trypan blue assay are presented in ESI Fig. 3B.†

total P content in samples drawn for nHAP\_B, nP, and CCM at different time points and (b) shows % P release derived from total P-content for nHAP\_B and nP at different time points.



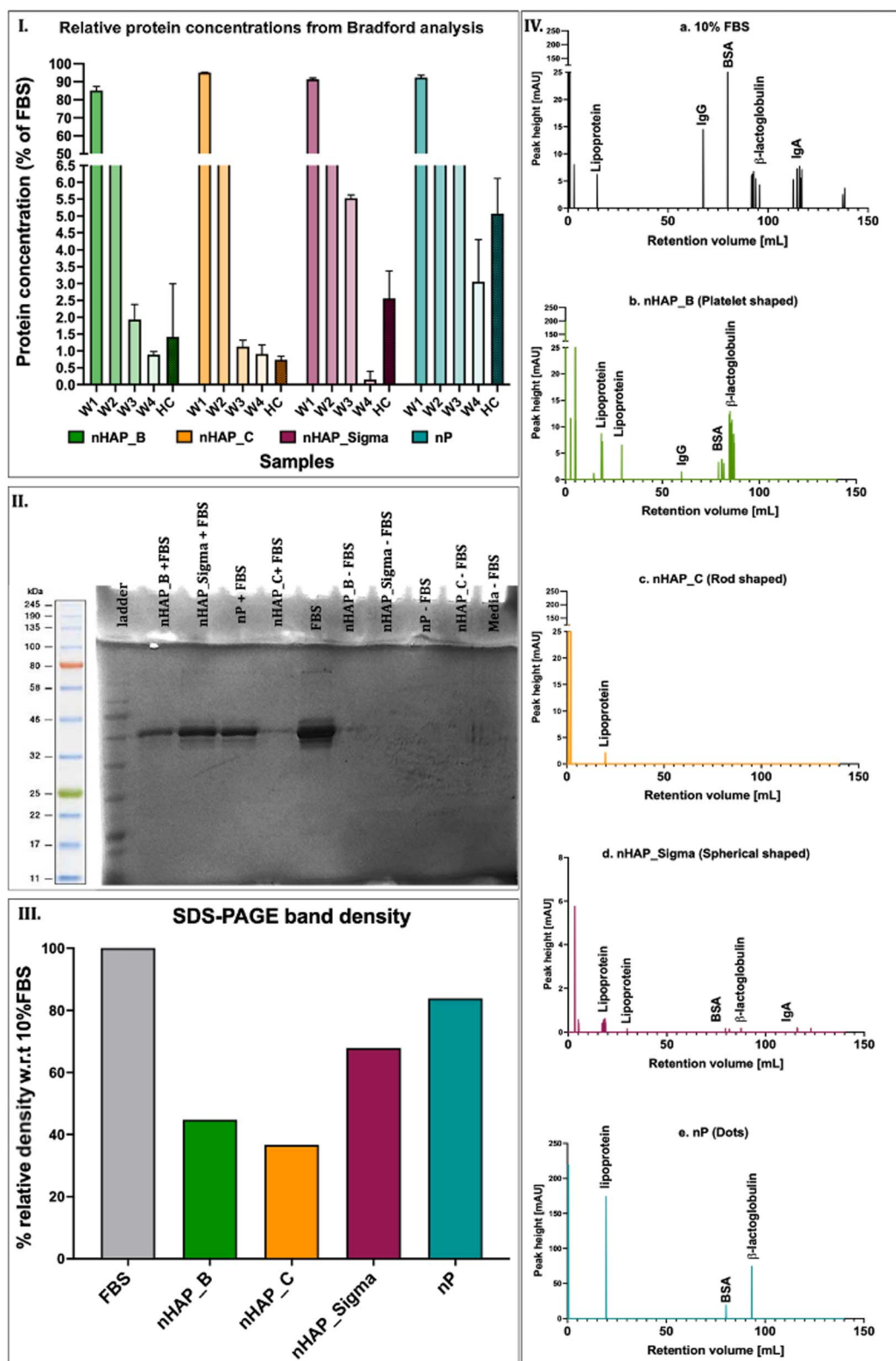


Fig. 2 (I) Protein concentration as determined by the Bradford method – W1, W2, W3, and W4 denote subsequent washes, and HC denotes hard corona. (II) SDS page gel image for hard corona over NMs. (III) Relative band intensity w.r.t. 10% FBS from SDS PAGE. (IV) Different FPLC profiles for NMs with a hard corona in comparison with FBS.



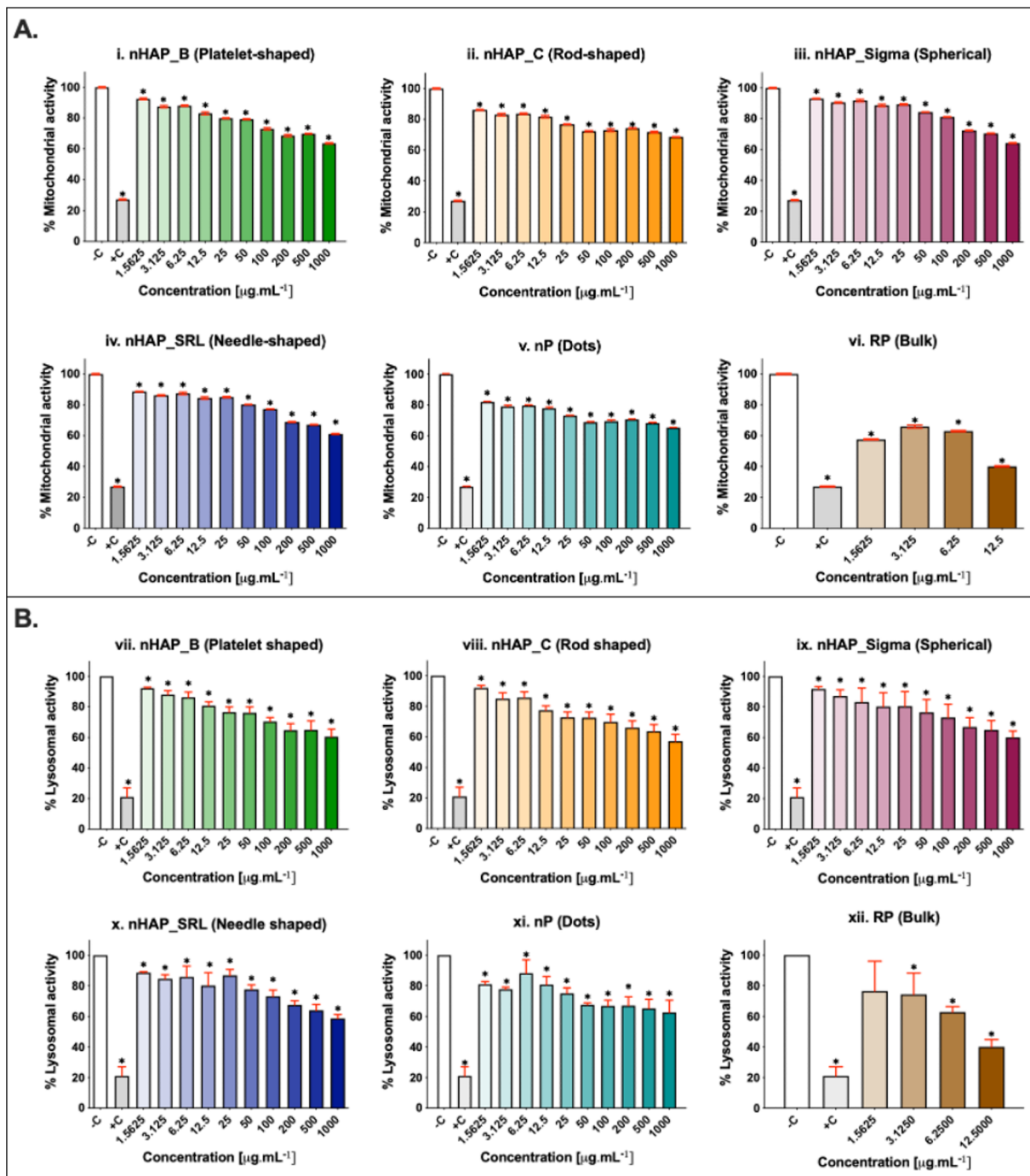


Fig. 3 Percentage cellular metabolic activity using (A) MTT assay for mitochondrial activity in case of (i) nHAP\_B, (ii) nHAP\_C, (iii) nHAP\_Sigma, (iv) nHAP\_SRL, (v) nP and (vi) RP. (B) Neutral red assay for lysosomal activity in case of (vii) nHAP\_B, (viii) nHAP\_C, (ix) nHAP\_Sigma, (x) nHAP\_SRL, (xi) nP and (xii) RP. \* show statistical significance at  $p < 0.005$  when different test concentrations are compared with the untreated control using one-way ANOVA followed by the Tukey *post-hoc* test.

### 3.5 Effects of NMs exposure on cellular activity for different initial seeding densities

Initial cell seeding densities of  $10^4$ ,  $10^3$ , and  $10^2$  were used for testing the concentration range from 1.5 to  $1000 \mu\text{g mL}^{-1}$ . For nHAP\_B, nHAP\_C, and nP, minimum mitochondrial activity was seen in the case of  $10^3$  cells and the highest was in the case of  $10^2$  cells (Fig. 5). Similar observations were made for

nHAP\_Sigma and nHAP\_SRL with some overlaps between the activity of  $10^2$  and  $10^4$  cells. However, in neither case,  $\text{LC}_{50}$  was seen for any of the NMs at any of concentrations. The minimum % metabolic activity, in any case, was found at the highest tested dose of  $1000 \mu\text{g mL}^{-1}$ . These values were statistically significant ( $p < 0.001$ ) as compared to the untreated control cells.



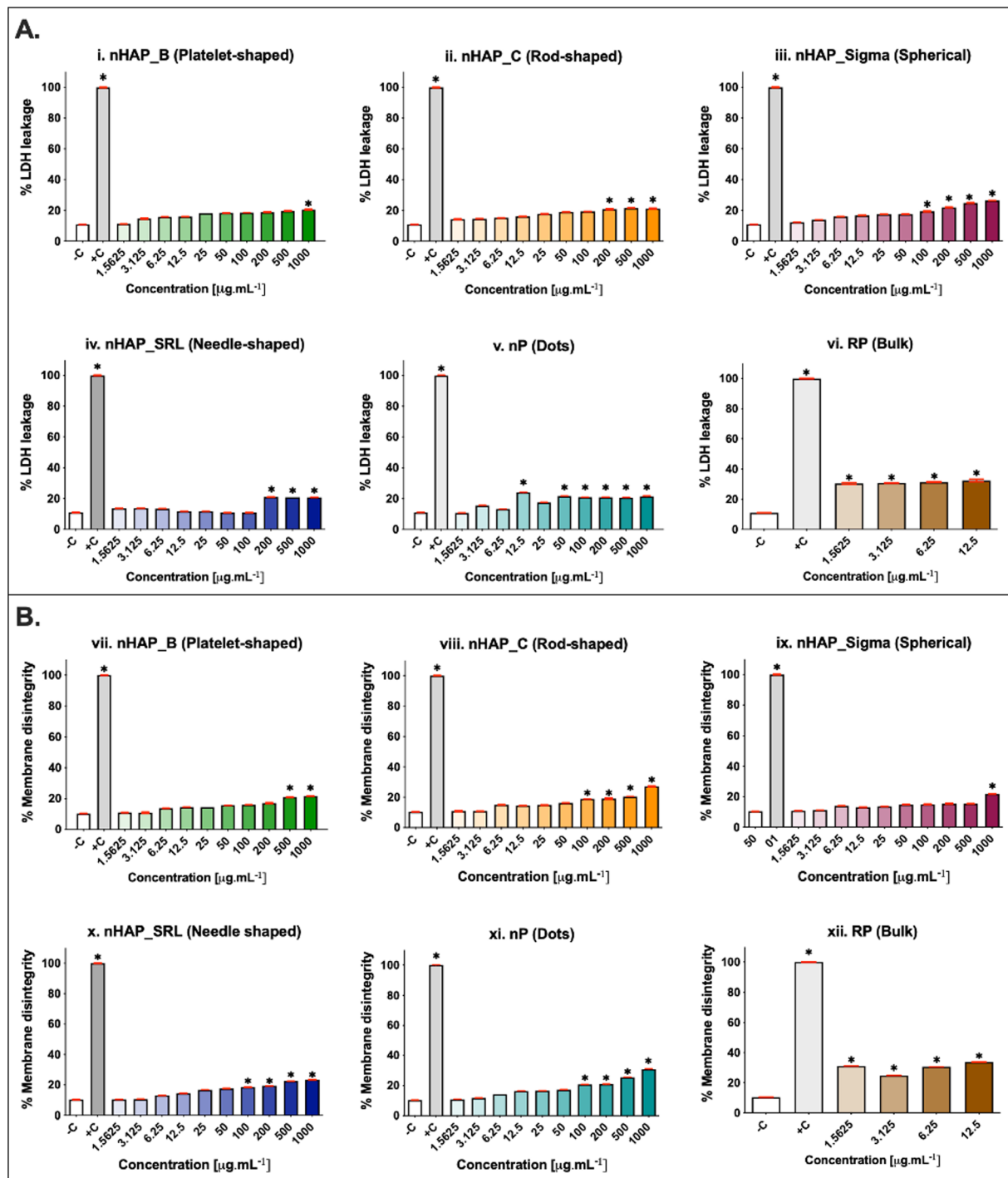


Fig. 4 (A) Percentage membrane dis-integrity using the LDH assay for (i) nHAP\_B, (ii) nHAP\_C, (iii) nHAP\_Sigma, (iv) nHAP\_SRL, (v) nP, and (vi) RP. (B) Percentage membrane dis-integrity using trypan blue assay for (vii) nHAP\_B, (viii) nHAP\_C, (ix) nHAP\_Sigma, (x) nHAP\_SRL, (xi) nP, and (xii) RP. \* show statistical significance at  $p < 0.005$  when different test concentrations are compared with the untreated control using one-way ANOVA followed by the Tukey *post-hoc* test.

### 3.6 ROS generation monitored on exposure to NMs

Intracellular ROS levels were monitored after the cells were exposed to 50 and 500  $\mu\text{g mL}^{-1}$  of NMs from 2',7'-dichlorofluorescein diacetate (DCFH-DA) oxidation. It was found that after 3 h of monitoring, results were significant for 500  $\mu\text{g mL}^{-1}$

of NMs, as compared to the untreated control. The fold changes w.r.t. untreated control were 1.5 for nHAPs and 1.9 for nP at 500  $\mu\text{g mL}^{-1}$  and at 3 h estimation (Fig. 6). These fold changes were significant ( $p < 0.001$ ) for nP from evaluation for both 50 and 500  $\mu\text{g mL}^{-1}$ . Exposure to both concentrations of RP has significant ( $p < 0.001$ ) DCF levels. This suggests that on getting







Fig. 5 Percentage cell viability for variable seeding density ( $10^4$ ,  $10^3$ , and  $10^2$  cells) using MTT for (a) nHAP\_B, (b) nHAP\_C, (c) nHAP\_Sigma, (d) nHAP\_SRL and (e) nP. Significances denoted by: (i) \* for  $p < 0.05$  when the different treatments were compared with the untreated control, (ii) (a, b, and c) for  $p < 0.05$  when effects of a particular concentration on different cell densities are compared in each case with NM treatment. Different letters show a significant difference. Statistical comparisons were performed using one-way ANOVA followed by the Tukey *post-hoc* test.

converted to nanoform from RP to nP, the oxidative stress due to the exposure decreases.

### 3.7 Time-dependent uptake of FITC-tagged nHAP\_B

Exposures to FITC\_nHAP ( $100 \mu\text{g mL}^{-1}$ ) in HEK293 were analysed in a time-dependent manner, and the data are shown in Fig. 7. Fig. 7(i) shows the control untreated cells. Fig. 7(ii)–(vii) show the time-dependent uptake at 2 h, 4 h, 6 h, 8 h, and 24 h. It can be suggested that the apparent uptake of P-based NMs starts after 4 h of treatment.

### 3.8 TEM analysis of cellular uptake and localization

TEM analysis was performed for the untreated control cells (Fig. 8(i)) and for cells after exposure to nHAP\_B and nP ( $100 \mu\text{g mL}^{-1}$ ) by processing samples immediately after NMs exposure (Fig. 8(iii) and (iv)) and after 24 h of exposure (Fig. 8(ii) and (v)–(xii)). No significant cellular internalization of NMs was seen for samples processed immediately after the exposure. NM internalization was seen in the case of both nHAP\_B and nP after 24 h of exposure, contrary to the untreated control. NMs were seen interacting with prominent cell organelles such as





Fig. 6 ROS generation estimated by DCF at 3 h of DCFH-DA monitoring after 24 h of HEK293 treatment with 50 and 500  $\mu\text{g mL}^{-1}$  of nHAPs, nP, and RP, as compared with untreated cells as negative control and H<sub>2</sub>O<sub>2</sub> (300  $\mu\text{M}$ ) as the positive control. Significance is denoted by \* for  $p < 0.05$ .

cytoplasm, nucleus, mitochondria, endoplasmic reticulum, endosomes, lysosomes, and the membranes of different cellular organelles. nHAP\_B aggregates were seen in the nucleus of the cell, while particles were also present in subcellular organelles. nP was more in the aggregated form inside the cells as compared to nHAP\_B. Fig. 8(xiii) and (xiv) show EDX for nHAP\_B and nP, respectively, indicating the presence of P and Ca.

### 3.9 Live/dead cell assessment using acridine orange/ethidium bromide dual staining

To investigate cell viability and apoptosis, dual staining using acridine orange and ethidium bromide was completed and 3715 cells in each case were auto-analysed using an automated cell counter (built-in function of Cytation 5, Biotek, Agilent Technologies, USA). In the untreated control, 100% of the HEK 293 cells appeared healthy (Fig. 9A(i) and B). Exposure of the cells to the positive control, H<sub>2</sub>O<sub>2</sub>, resulted in an increased number of apoptotic ( $61 \pm 1.73\%$ ) and necrotic cells ( $32.33 \pm 2.52\%$ ; Fig. 9A(ii) and B). Further, as shown in Fig. 9A(iii)–(viii) and B, healthy cells were observed for nHAPs. The exposure of cells to nP also caused an increase in apoptotic cells ( $36 \pm 2\%$ ) compared to the control (Fig. 9A(vii) and B). Exposure of the cells to the bulk control, RP, was also observed to cause an increase in both apoptotic ( $69 \pm 1\%$ ) and necrotic cells ( $21.67 \pm 3.51\%$ ). All these values were statistically significant ( $p < 0.01$ ) w.r.t. to untreated control cells.

## 4 Discussion

The P-based NMs are proposed for agricultural applications as P-supplementing nanofertilisers. For occupational health and

safety in response to the nanofertiliser exposure, it is critical to assess the potentially toxic effects of NMs on humans using the human cell lines or other model systems at their predicted application concentrations as fertilisers (up to  $25 \mu\text{g mL}^{-1}$ )<sup>49,50</sup> and higher concentrations ( $25\text{--}1000 \mu\text{g mL}^{-1}$ ). Farmers and field workers are most likely to be exposed to nanofertilisers when applying it either *via* the soil or foliar spray in a direct exposure concentration range of  $10\text{--}1000 \mu\text{g mL}^{-1}$ .<sup>6,8,10–12</sup> Dermal, inhalation, and ocular routes may be the primary ways of NM internalisation in the human body.<sup>51</sup> After internalisation, renal clearance of the NMs that have reached the systemic circulation is an expected and possible elimination route from the human body.<sup>38</sup> For this reason, kidneys may be one of the most important secondary target organs, and NM-induced renal toxicity should be evaluated. Further, excess Ca from HAP crystallites on the surface of renal epithelial cells is known to induce the formation of Randall plaques and even kidney stones.<sup>39,40</sup> Due to the diversity of P-based NMs, especially their physicochemical properties that influence their characteristics and behaviour, many nano-bio interactions are yet to be investigated. Therefore, further studies are required to provide a comprehensive and critical evaluation of the toxicity of NMs in the renal system. In addition to providing insights into the effects of P-based NMs on nephrotoxicity, HEK293 cells, being epithelial cells, can also be used to assess the toxicity of P-based NMs on epithelial cells. Previously, the effects of P-based NMs, especially nHAP have been tested on mammalian cell lines and some results have shown toxicity<sup>20–23</sup> and inflammatory<sup>22–26</sup> responses of rod- and needle-shaped nHAPs. Recently, a study on rod-shaped mesoporous silica-coated nHAP reported that the NM was toxic to MC3T3-E1 cells on doses above  $10 \mu\text{g mL}^{-1}$ .<sup>52</sup> However, a thorough investigation of the physicochemical modifications and protein corona formation remains a gap to understand the cytological behaviour of differently shaped P-based NMs.

In this study, we investigated the *in vitro* nano-bio interactions of multiple variants of nHAPs and nP, which differed in physicochemical properties and origin as described previously.<sup>35,41</sup> Four differently shaped nHAPs namely, nHAP\_B (platelet-shaped), nHAP\_C (rod-shaped), nHAP\_Sigma (spherical), and nHAP\_SRL (needle-shaped), were used in this study. Another P-based NM used in this study, nP (dots) was derived from rock phosphate. A systematic and thorough impact of these NMs was carried out in HEK293. In the serum-supplemented CCM, the hydrodynamic behaviour, physicochemical properties, P release, and protein absorption on the NM surface were investigated. Cell viability, cellular mitochondrial and lysosomal activity, membrane disintegrity, oxidative stress, uptake, and localization of these NMs in the cell and cellular organelles were studied. The effects of these NMs were also tested on the cellular metabolic activity at varying seeding densities.

The addition of the P-based NMs to serum-supplemented media resulted in increased hydrodynamic diameters for all nHAPs. The Z-average size of nHAP\_C was the highest followed by nHAP\_SRL, nHAP\_B, and nHAP\_Sigma. It is worth noting here that nHAP\_C and nHAP\_SRL are rods- and needle-shaped,



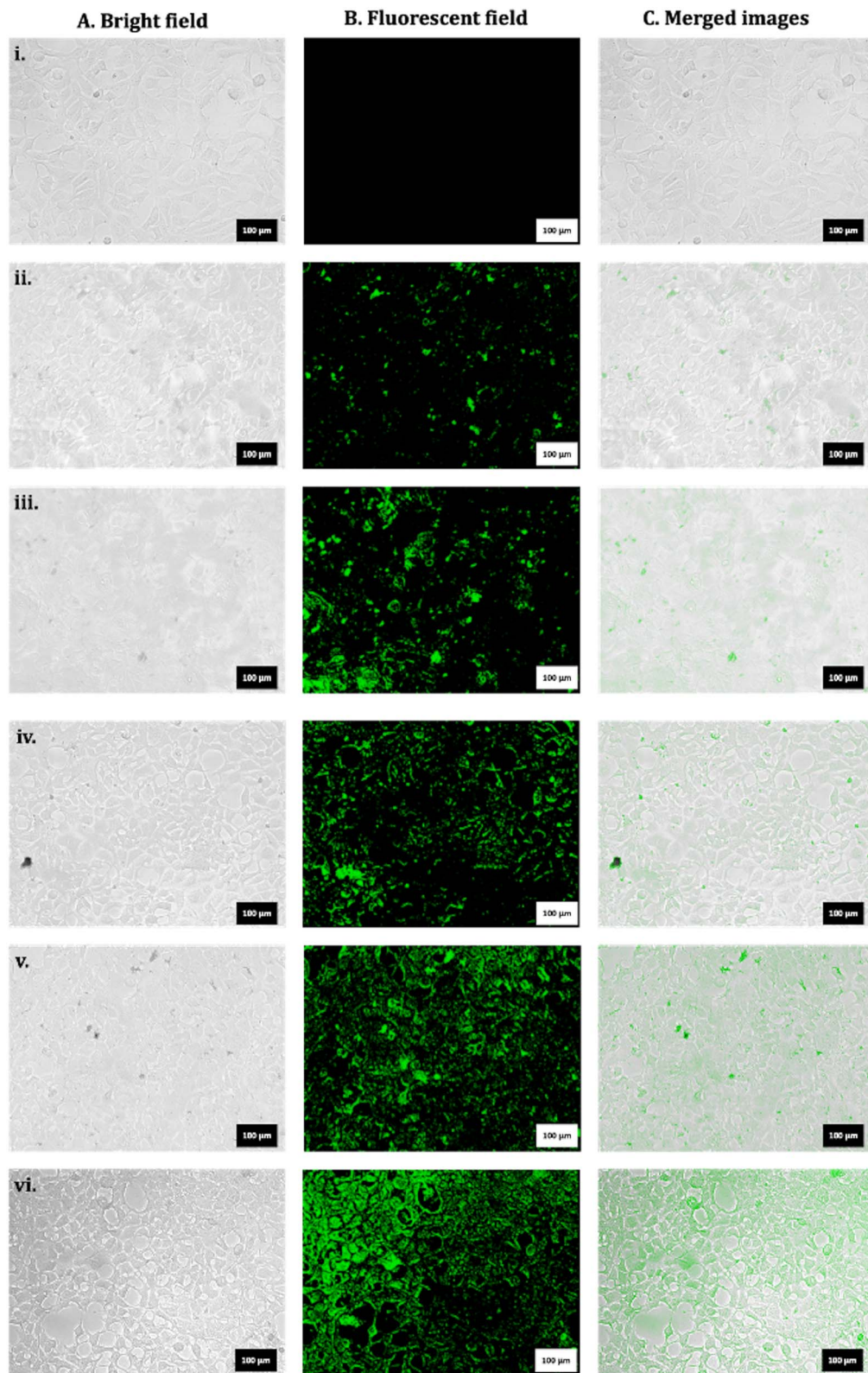


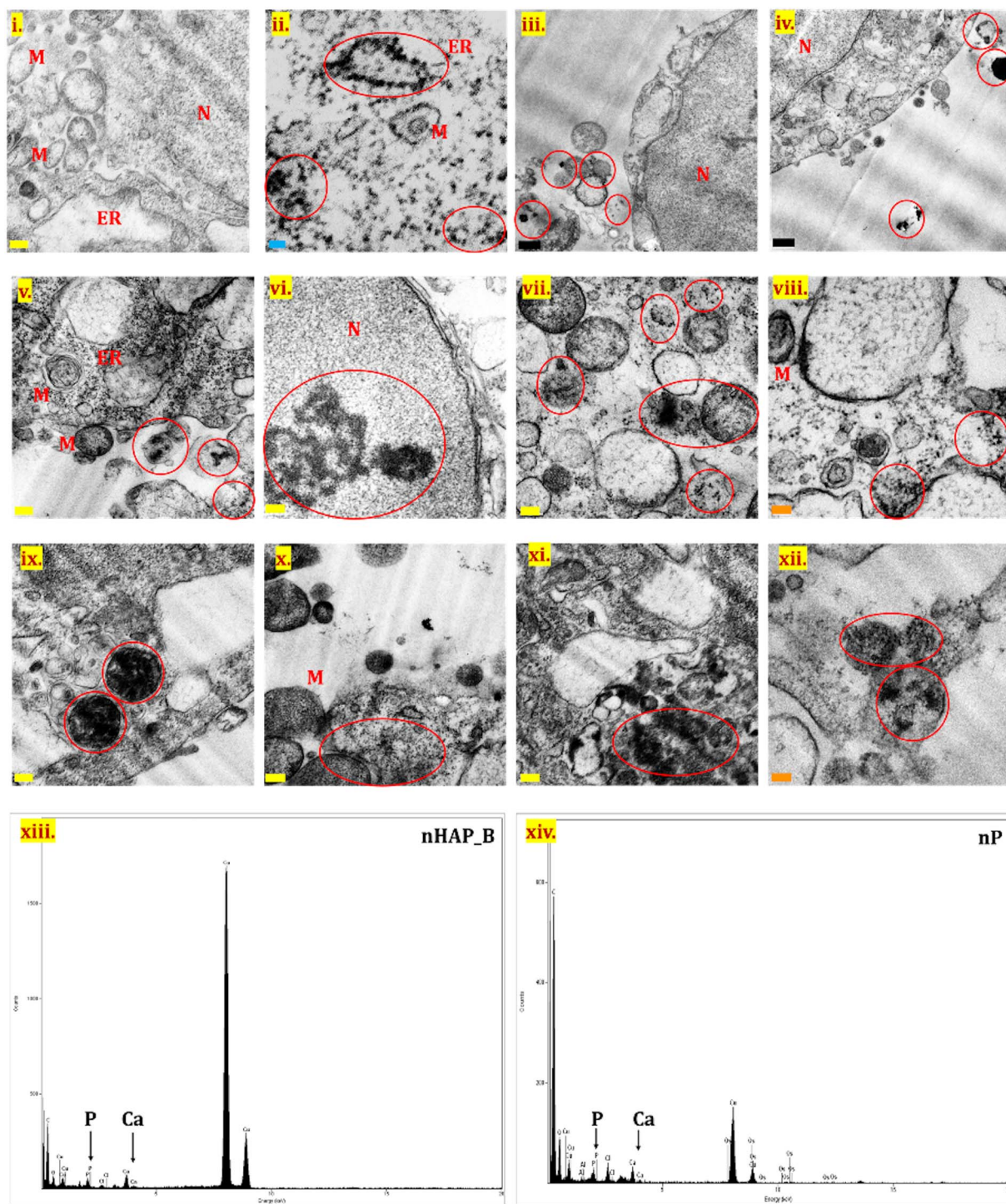
Fig. 7 Representative microscopic images for HEK293 for (i) untreated control. Exposure to FITC\_nHAP\_B at (ii) 2 h, (iii) 4 h, (iv) 6 h, (v) 8 h, and (vi) 24 h (scale: 100  $\mu\text{m}$ ).

respectively, and thus have higher sizes than nHAP\_Sigma (spherical) or nHAP\_B (platelet-shaped). From their TEM images, it was apparent that the media formed a film

surrounding the surface of NMs. It can be suggested that the serum components form a capping over these NMs. The negative zeta potential values for all the NMs were decreased upon







**Fig. 8** Representative TEM micrographs for HEK293 for (i) untreated control (scale: 0.2  $\mu\text{m}$ ), (ii) internalized nHAP\_B inside the cells (scale: 50 nm), (iii and iv) images for samples processed just after the addition of nHAP\_B and nP, respectively (scale: 0.5  $\mu\text{m}$ ), (v–viii) images for cells treated with nHAP\_B (scale: 0.2  $\mu\text{m}$  for (v) to (vii), 0.1  $\mu\text{m}$  for (viii)), (ix–xii) images for cells treated with nP (scale: 0.2  $\mu\text{m}$  for (ix) to (xi), 0.1  $\mu\text{m}$  for (xii)). (xiii) EDX for cells treated with nHAP\_B and (xiv) EDX for cells treated with nP. Red circles show the prominent presence of NMs.

interaction with FBS. This could be due to the absorption of positively charged amine groups on the surface of NMs. A similar observation was made by Casals *et al.* (2010) for Au NMs.<sup>53</sup> Similar observations were also made for nP, where the hydrodynamic diameter increased after dispersal in the serum-

supplemented media. It was also discussed recently by Zhao *et al.* (2021) that serum protein can be deposited more on particles with smaller diameters.<sup>54</sup> This agrees with the TEM profile obtained for nP where the pristine diameter was  $\sim 10$  nm (ref. 55) and upon interaction with serum-supplemented media,





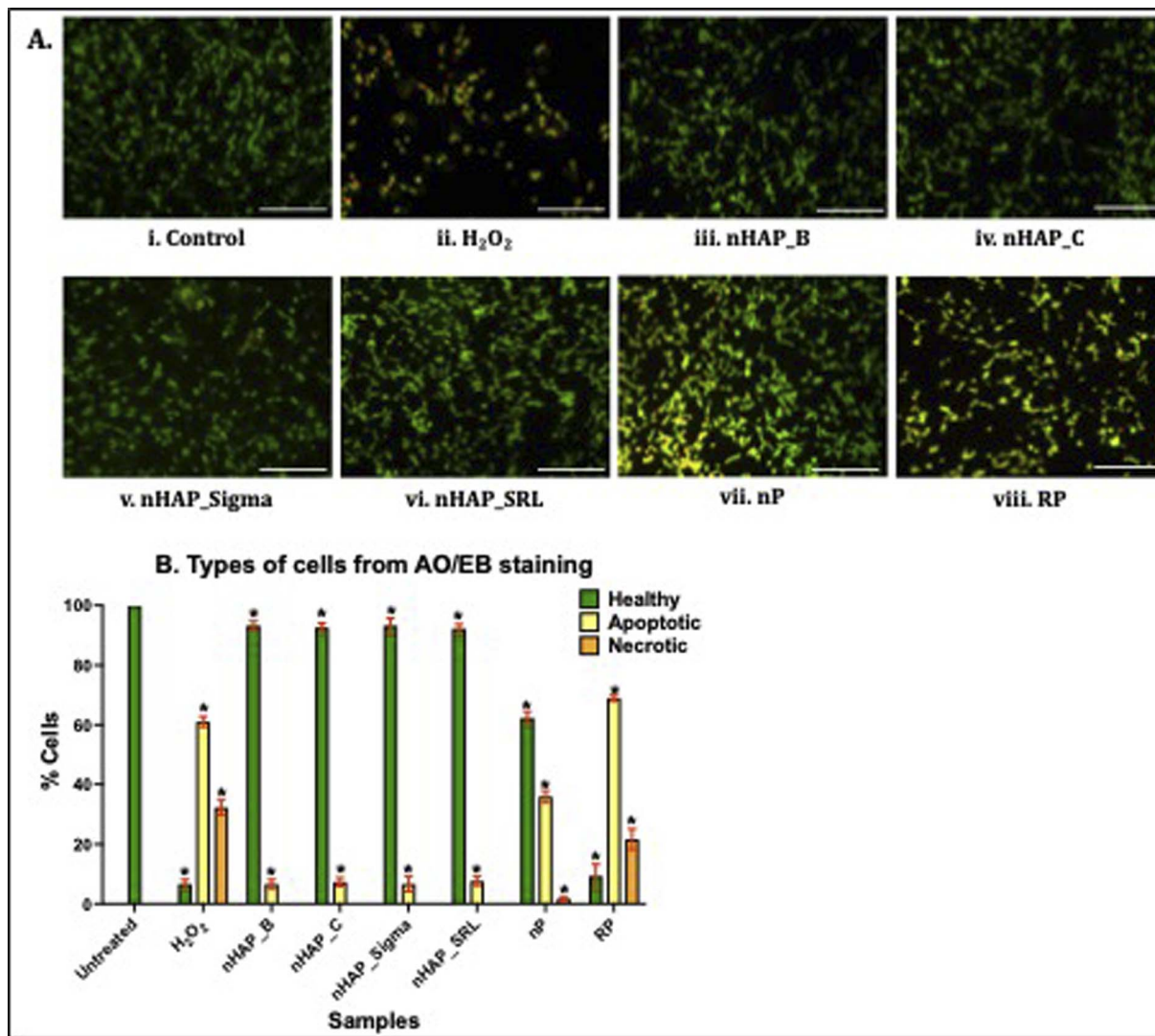


Fig. 9 (A) Microscopic images (scale: 200 μm) for AO/EtBr staining in case of (i) untreated control, (ii) 300 μM of H<sub>2</sub>O<sub>2</sub>, 100 μg mL<sup>-1</sup> of (iii) nHAP\_B, (iv) nHAP\_C, (v) nHAP\_Sigma, (vi) nHAP\_SRL, (vii) nP, and (viii) RP. (B) Quantitative data for AO/EB indicating % of healthy, apoptotic, and necrotic cells (\* denote significance for  $p < 0.01$  w.r.t. to the untreated control cells).

a large visible matrix was observed with small-embedded particles of nP (~18 nm).

In this study, the investigations on protein corona revealed that the protein adsorption from the CCM onto the surfaces of P-based NMs was dependent on their inherent shapes and physicochemical properties. The proteins on hard corona were profiled in the FPLC analysis. These were then identified by referring to the peaks reported previously for a similar analysis.<sup>56</sup> The main proteins identified in the hard corona were lipoprotein, β-lactoglobulin, albumin (BSA), and immunoglobulins (IgG and IgA). The presence of the protein corona on the NMs could affect the uptake and intracellular trafficking of the NMs.<sup>57,58</sup> It has been shown previously that phagocytosis can be influenced by immunoglobulins in alveolar macrophages,<sup>59,60</sup> and lipoproteins<sup>61</sup> have been shown to affect the intracellular transport and localisation of the NMs in the rat epithelial cells<sup>62</sup> and bone marrow-derived macrophages.<sup>63</sup>

Several interesting findings were observed when HEK293 cells were exposed to P-based NMs in a dose-dependent manner. It was also observed that the P-release from the biologically synthesised NMs was slow and minimal over the maximum exposure time of 24 h. Thus, the effects on the cells were dominantly due to interaction with NMs only. Exposure of HEK293 to nP derived from RP significantly decreased the mitochondrial and lysosomal activity, enhanced membrane disintegrity, and increased ROS and cellular apoptosis. However, these effects were further increased when the cells were exposed to the bulk RP at the same doses. The elemental composition of RP might be the reason for its severe cytotoxicity. Fagiya and colleagues<sup>64</sup> have previously shown that RP is a heterogeneous mix of different elements: 20–28% P<sub>2</sub>O<sub>5</sub>, 43.9% O, 34.1% Ca, 13.6% P, 3.57% F, 1.43% Fe, 1.26% Si, and 2.14% other heavy elements such as Pb, Zn, and Al.<sup>64</sup> It is however interesting to observe from this study that the nano-





also been previously confirmed using TEM. In one of the studies with Ag NMs, uptake in murine melanoma B16 cells was shown where the particles localise to intracellular vesicles and late endosomes, and clathrin-mediated internalisation was suggested.<sup>81</sup> In another study on Ag NMs and human hepatocyte carcinoma HepG2, the authors reported localisation of particles in endosomes and lysosomes after being taken up by micropinocytosis and clathrin-mediated endocytosis.<sup>82</sup> A study with Au NMs in RAW267.1 cells has shown the localisation of NMs in the perinuclear region and vesicles and has reported the internalisation *via* clathrin-mediated and dynamin-dependent endocytosis.<sup>83</sup> In another study demonstrating the uptake of Fe NMs in human ovarian cancer (SKOV-3) and RAW 264.7 cells, it was shown that the NMs localise in endosomes and lysosomes suggesting the possible uptake by endocytosis.<sup>84</sup> Another study found the time-dependent uptake of ZnO NMs in human mesenchymal stem cells after repetitive treatments and the localisation into the nucleus and mitochondria.<sup>85</sup> Considering these previous studies, and the results from this study where P-based NMs were located around the lysosomes and the other membrane-bound organelles, it can be hypothesized that their internalisation would have been either *via* adsorptive or receptor-mediated endocytosis. However, it will be interesting to further investigate if the internalisation of P-based NMs in human cells is through multiple pathways or a single energy-dependent pathway.

The impacts of exposure to P-based NMs on acute cellular behaviour are significant. In a nutshell, for P-based NMs to be used potentially as nanofertilisers, it is important to know the specific safe doses for manufacturers and end users to handle these NMs. These doses are dependent on the physicochemical properties of nHAPs and nP. Although, some sub-lethal effects including mitochondrial and lysosomal activities were observed at the higher NM doses and were more evident after exposure to nP and RP compared to nHAPs. For environmentally relevant concentration of  $100 \mu\text{g mL}^{-1}$ , the safer NMs follow the order as nHAP\_B  $\approx$  nHAP\_Sigma > nHAP\_SRL > nHAP\_C > nP. However, there were no LC<sub>50</sub> values obtained in either of the assays for any of the P-based NMs up to  $1000 \mu\text{g mL}^{-1}$ . This highest tested concentration is hypothesized to be significantly higher than the environmentally relevant concentration for NM bioactivity, providing further support for these P-based NMs being further advanced for use as safe nanofertilisers for agriculture. Findings such as the transformation of NMs from the pristine state, NM colloidal interactions, and the effects on cellular metabolic activities, as discussed above, affirm that the effects of NMs on mammalian cells are due to multiple mechanisms and physicochemical factors. This further supports the aim of our study where the emphasis is to understand the nano-bio interactions with varying physicochemical properties of P-based NMs.

## 5 Conclusions

In this study, *in vitro* nano-bio interactions of five physicochemically different P-based NMs were evaluated by changes in size, P content, colloidal interactions, protein corona

estimation, and effects on HEK293 cells. The colloidal interactions and protein hard corona composition were shape-dependent. Biogenic nHAP and nP were shown to be taken up by the cells and localise intracellularly in time-dependent manner. None of the variants of P-based NMs had LC<sub>50</sub> up to  $1000 \mu\text{g mL}^{-1}$  when cell viability, mitochondrial and lysosomal activity; and membrane disintegrity were evaluated. This highest tested concentration is hypothesized to be significantly higher than the environmentally relevant concentration for NM bioactivity, suggesting that the tested P-based NMs can be considered safe as nano-agromaterials. At the environmentally relevant dose of  $100 \mu\text{g mL}^{-1}$  platelet (biogenic) and spherical-shaped nHAPs are safer as compared to the rod (chemically synthesised) and needle-shaped NMs. This study aims to provide experimental data on safer doses for the new generation of biogenic P-based NMs to be used as nanofertilisers to supplement P in agricultural soils.

## List of abbreviations

nHAP_B	Nanohydroxyapatite biologically synthesized
nHAP_C	Nanohydroxyapatite chemically synthesized
nHAP_Sigma	Nanohydroxyapatite procured from Sigma-Aldrich
nHAP_SRL	Nanohydroxyapatite procured from SRL Pvt Ltd
nP	Nanophosphorus
RP	Rock phosphate
CCM	Cell culture media
DPBS	Dulbecco's phosphate buffer saline
FBS	Fetal bovine serum
TEM	Transmission electron microscopy
DLS	Dynamic light scattering
FITC	Fluorescein isothiocyanate

## Ethics for approval

All experiments were approved by Deakin University Human Research Ethics Committee (DUHREC reference no. 2021-095), Australia.

## Consent for publication

All the authors have read and approved the manuscript for publication.

## Data availability

The datasets used in the present study are available from the corresponding author upon reasonable request.

## Author contributions

Conceiving and designing the experiments: AP, AKD, PPS. Wet-lab work and data compilation: AP. Data analyses and manuscript writing: AP and PPS. Critical inputs and finalization of the manuscript: AP, LOA, AGS, AKD, and PPS.





## Conflicts of interest

The authors declare the following conflict of interest: nano-phosphorus (nP) is the nanomaterial developed at TERI-Deakin Nanobiotechnology Centre (TDNBC), Gurgaon, India, and the synthesis and characterization of nP as the intellectual property of TDNBC will be filed with the Indian patent office.

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