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Ceria nanoparticles immobilized with self-assembling peptide for biocatalytic applications†‡

Moumita Halder, Vatan Chawla and Yashveer Singh *

Peptide-based artificial enzymes exhibit structure and catalytic mechanisms comparable to natural enzymes but they suffer from limited reusability due to their existence in homogenous solutions. Immobilization of self-assembling peptides on the surface of nanoparticles can be used to overcome limitations associated with artificial enzymes. A high, local density of peptides can be obtained on nanoparticles to exert cooperative or synergistic effects, resulting in an accelerated rate of reaction, distinct catalytic properties, and excellent biocompatibility. In this work, we have immobilized a branched, self-assembled, and nanofibrous catalytic peptide, (C₁₂-SHD)₂KK(Alloc)-NH₂, onto thiolated ceria nanoparticles to generate a heterogeneous catalyst with an enhanced number of catalytic sites. This artificial enzyme mimics the activities of esterase, phosphatase, and haloperoxidase enzymes and the catalytic efficiency remains nearly unaltered when reused. The enzyme-mimicking property is investigated for pesticide detection, bone regeneration, and antibiofouling applications. Overall, this work presents a facile approach to develop a multifunctional heterogeneous biocatalyst that addresses the challenges associated with unstable peptide-based homogeneous catalysts and, thus, shows a strong potential for industrial applications.

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Introduction

Biocatalysis refers to the use of natural catalysts, such as enzymes or whole cells, to perform chemical transformations.¹ These catalysts accelerate the rate of chemical reactions by providing an alternative pathway with lower activation energy, allowing reactions to occur under milder conditions. Enzymes are proteins that act as highly efficient biocatalysts in living organisms.² They possess specific active sites that can bind to substrates and facilitate their conversion into products. Enzymes are incredibly selective, often catalyzing specific reactions and producing specific products, which makes them valuable tools in industries.³ Biocatalysis offers several advantages over traditional chemical catalysis. Enzymes exhibit high selectivity, which minimizes the formation of by-products⁴ and they work under mild conditions, including lower temperatures and pressures, which can reduce energy consumption

and production costs. Enzymes are also more environment friendly as they avoid the use of toxic or hazardous chemicals, and they find application in a wide range of industries, including pharmaceuticals, agriculture, food and beverage, biofuels, and manufacturing.^{5,6} Examples of biocatalytic reactions include the production of antibiotics, synthesis of fine chemicals, degradation of environmental pollutants, and conversion of biomass into biofuels. Overall, biocatalysis harnesses the power of biological catalysts to enable more sustainable and efficient chemical processes, with significant potential for advancing industries and reducing environmental impact.

While biocatalysis offers numerous benefits, there are also several challenges associated with the use of biocatalysts. The lack of availability of these enzymes in large quantity is the major concern for the industrial use.⁷ Some enzymes may be derived from rare or difficult-to-cultivate organisms, making them less accessible, which significantly increases their costs. Many biocatalysts, especially enzymes, can be sensitive to environmental conditions, such as temperature and pH.⁸ They may undergo denaturation or loss of activity over time, which reduces their stability and shelf-life. There are issues related to the use of enzymes in the development of heterogeneous catalysts.^{9,10} The enzymes are attached to solid surfaces either through adsorption or covalent bonding. Adsorption-based immobilizations invariably result in a loss of adsorbed proteins during the course of subsequent cycles, whereas the covalent linkages generated through chemical processes may

Department of Chemistry, Indian Institute of Technology Ropar, Rupnagar-140 001, Punjab, India. E-mail: 2018cyz0011@iitrpr.ac.in, vatan.19cyz0016@iitrpr.ac.in, yash@iitrpr.ac.in

† This article is dedicated to Prof. Santanu Bhattacharya on his 65th birthday. We wish him many years of active research ahead and hope that he continues to inspire and motivate research groups around the world. YS carried out postdoctoral research under his mentorship.

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cause the connected enzyme to become denatured.¹¹ As a result, it is increasingly difficult to reuse catalysts and the cost-effectiveness of process is also reduced. Artificial enzymes or compounds mimicking enzymes can be used to surmount these challenges.^{1,12,13} An enzyme's catalytic activity comes from supramolecular interactions that form a three-dimensional structure within the protein molecule. The correct spatial arrangement of amino acids inside these structures produces catalytic pockets needed to bind to a particular substrate and carry out chemical reactions.¹⁴ The minimalistic approach to develop artificial biocatalysts with same catalytic efficiency, stereoselectivity, and specificity as enzymes is to imitate their catalytic sites by incorporating amino acids found in catalytic pockets.

Self-assembling peptides can act as biocatalysts by organizing into well-defined structures or scaffolds that provide an environment for catalytic reactions to occur.^{15,16} These peptides have the unique ability to spontaneously form ordered structures through non-covalent interactions. The self-assembling peptides can be designed to incorporate catalytic residues or functional groups within their sequence, allowing them to perform specific enzymatic reactions. The catalytic activity of self-assembling peptides can be influenced by factors such as the peptide sequence, secondary structure, and the microenvironment created by the self-assembled structure.¹⁷ Proline-containing peptides, for instance, are known to catalyze Mannich, Michael, aldol, and acyl transfer processes.^{18,19} In a similar way, ester hydrolysis reactions are known to be catalyzed by histidine (His).^{20,21} For the regeneration of hydrolase and esterase activities, a variety of His-containing hydrogels and nanofibrils have been developed. Similarly, hydrogel-trapped hemin chloride has been found to have peroxidase-like activity.²² In this regard, hydrolytic enzymes, such as lipase and α -chymotrypsin, have drawn a lot of attention due to their prevalence in living things and growing economic significance.²³ Therefore, much effort has been made to develop artificial esterase and phosphatase.²⁴

Several design techniques have been used to generate peptide-based artificial enzymes. For instance, Korendovych *et al.* developed a set of heptapeptides that formed β -sheets, and by self-assembling these peptides, they obtained a Zn(II)-coordinated structure,²⁵ which mimicked the catalytic center of natural carbonic anhydrase and demonstrated catalytic activity towards carbonic anhydrase substrates. Numata *et al.* developed an enzyme-like catalyst by incorporating the conventional serine-protease catalytic triad to peptides that can self-assemble into fibrils and resemble amyloids.²⁶ The formation of β -sheet backbone was essential to imitate the protease binding site. Gulseren *et al.* reported esterase-mimicking catalytic nano-system with essential residues, Ser, His, and Asp in the peptide sequence.²⁷ Mikolajczak *et al.* developed peptide-gold nanoparticle conjugates as sequential cascade catalyst for the hydrolysis followed by the hydrogenation reaction.²⁸ Dowari *et al.* reported the immobilization of peptide amphiphile containing Asp(Ser)His triad on silica surface for hydrolase-mimicking properties.²⁹ Reja *et al.* developed nanotubes

from self-assembled lipidated short peptide (C₁₀-FFVK) for aldolase-mimicking properties.³⁰ Overall, self-assembling peptide provide a versatile platform for creating functional materials with catalytic capabilities and expanding the scope of biocatalysis beyond traditional enzymes.

There has been a lot of reports available in literature, where self-assembled peptides have been extensively explored to develop artificial enzymes. The major limitation associated with such system is that they are in homogenous solution with reaction mixture and cannot be removed once the reaction is over. Unbranched, linear peptides reported so far can only expose their single binding pocket to the substrate, which limits the rate of reaction. Additionally, there are very few reports on the use of artificial enzymes exhibiting multiple catalytic activities simultaneously. Developing multifunctional peptide catalysts that can perform diverse reactions or exhibit additive/synergistic effects has the potential to open new possibilities in biocatalysis.

The main aim of this work was to develop multifunctional, heterogenous catalyst with esterase, phosphatase, and haloperoxidase-mimicking activities and use them for pesticide detection (organophosphorus and carbamate), bone regeneration, and anti-biofouling applications. In this work, we have immobilized self-assembled, nanofibrous catalytic peptide on ceria nanoparticles and evaluated their potential to act as a biocatalyst. Ceria nanoparticles (CeNP) were fabricated using hydrothermal method and thiolated (TC) using (3-mercaptopropyl)trimethoxysilane, which was further conjugated to peptide amphiphile (PA) by "thiol-ene" reaction, thus, forming thioether bonds. The branched peptide amphiphile contains catalytic triad 'Ser-His-Asp,' which is also present in the active site of serine protease.³¹ These residues work together to perform various hydrolytic reactions by facilitating binding to the substrate and catalysis. This peptide was immobilized on thiolated ceria nanoparticle surface to generate a heterogenous catalyst (TCP) with a greater number of catalytic sites. CeNPs were selected due to their haloperoxidase-mimicking properties, which involves catalyzing the oxidation of halide ions by hydrogen peroxide.³² Peptide conjugated ceria nanoparticles were thoroughly characterized and they were found to mimic esterase, phosphatase, and haloperoxidase enzymes. This enzyme-mimicking activities were used to elucidate their role in pesticide detection, bone regeneration, and anti-biofouling material preparation.

Experimental

Materials and methods

All solvents and chemicals were of high analytical quality and utilized without further purification unless otherwise stated. Rink amide AM resins (0.80 mmol g⁻¹ loading) was procured from Novabiochem. Fmoc-protected amino acids, such as Fmoc-Lys(Alloc)-OH, Fmoc-Lys(Fmoc)-OH, Fmoc-His(trt)-OH, Fmoc-Ser(trt)-OH, Fmoc-Asp(OtBu)-OH were purchased from BLD Pharm. The 1-[bis(dimethylamino)methylene]-1*H*-1,2,3-

triazolo[4,5-*b*] pyridinium 3-oxide hexafluorophosphate (HATU), trifluoroacetic acid (TFA), *N,N*-diisopropylethylamine (DIEA), triisopropyl silane (TIS), lauric acid, 5,5'-dithiobis-2-nitrobenzoic acid (DTNB), and 3-mercaptopropyl trimethoxysilane (MPTMS) were bought from TCI Chemicals. Dimethyl sulfoxide (DMSO) and HPLC grade solvents, such as acetonitrile, methanol, and isopropyl alcohol were purchased from Merck and used for reverse phase high-pressure liquid chromatography (RP-HPLC). Piperidine, *N,N* dimethyl formamide (DMF), and dichloromethane (DCM) were bought from Spectrochem and Rankem Laboratories. *p*-Nitrophenyl phosphate (*p*NPP) and *p*-nitrophenyl acetate (*p*NPA) were procured from Sisco Research Laboratories Pvt. Ltd (SRL). Cerium nitrate hexahydrate 99.9% AR and Irgacure 2959 were purchased from Loba Chemie and Sigma Aldrich. All pesticides (azamethiphos, profenofos, and carbendazim) were procured from Sigma-Aldrich. PolyPrep chromatography columns from Bio-Rad were used for solid-phase peptide synthesis (SPPS). Deionized water (DI, 18.2 MΩ cm) was obtained from a Milli-Q system and used in all experiments. MC3T3-E1 cells (CRL-2593, Subclone-4) were purchased from ATCC. Minimum Essential Medium alpha (MEM-α), 0.25% trypsin/EDTA, pen-strep, SYBR™ Green Master Mix, and trizol were purchased from Thermo Fisher Scientific. The iScript cDNA synthesis kit was bought from Bio-Rad. The 3-(4,5-dimethylthiazol)-2,5-diphenyltetrazolium bromide (MTT reagent), ascorbic acid, and fetal bovine serum (FBS) were procured from HiMedia. *E. coli* (MTCC 1687) was purchased from CSIR- IMTECH, Chandigarh. For antibacterial studies, Luria broth (HiMedia) was used as culture media.

Synthesis of branched peptide amphiphile (PA) and characterization

The self-assembled, branched peptide amphiphile was synthesized using standard Fmoc-based SPPS method. Rink amide AM resin was employed as a solid support. The amino acids were coupled using a mixture of HATU (2.85 equiv.), and DIEA (5.7 equiv.) and 20% piperidine in DMF (3 mL) were used to deprotect Fmoc group. The synthesized peptide was cleaved from resin by using a cocktail comprising of 3 mL of 95% TFA, 2.5% TIS, and 2.5% water (v/v). Subsequently, the amide terminated peptide/PA (C₁₂-SHD)₂KK(Alloc)-NH₂ was cleaved, and precipitated from 30 mL of cold diethyl ether, which was followed by centrifugation and drying under vacuum. The purity of these peptides was determined using a RP-HPLC equipped with Xbridge BEH C₁₈ column (250 × 4.6 mm, 5 μm) and a gradient flow of acetonitrile/water for 35 min with 0.1% TFA as a mobile phase at a flow rate of 1 mL min⁻¹. The peptides were further characterized by mass (XEVO G2-XS QTOF) and ¹H NMR (JEOL JNM-ECS, 400 MHz) analysis.

Secondary structure of peptide

The secondary structure of peptide was determined by circular dichroism (CD) and Thioflavin T (ThT) assay. CD analysis was performed on a CD spectrometer (JASCO J-1500) using a quartz cuvette with a 0.1 cm path length. In order to induce

the formation of secondary structures, peptide solutions were prepared in DI water and incubated at room temperature for five days prior to investigation. Spectra of peptides were recorded at a concentration of 0.5 mM in the region of 195–350 nm at a continuous scanning rate of 200 nm min⁻¹. Fluorescence spectra was used to analyze the binding of peptides with ThT according to a previously described method.³³ The fluorescence was recorded using a Tecan multimode microplate reader at an excitation wavelength of 440 nm. ThT solution without peptides was used as a blank.

Preparation of cerium oxide nanoparticles (CeNPs)

A conventional hydrothermal technique was used for the preparation of CeNPs.³⁴ A 60 mL of 4 M NaOH was used to make a suspension of 0.4 g of cerium nitrate hexahydrate (1 mmol) followed by stirring for 30 min. The mixture was transferred to a 100 mL Teflon-lined stainless-steel autoclave and incubated for 24 h at 100 °C. After centrifuging, the CeNPs were washed with water and ethanol, and dried at 70 °C to get a distinctive dark yellow powder.

Thiol functionalization of CeNPs (TC)

The surface of CeNP was activated by dehydrating the sample at 80 °C overnight. Around 75 mg of CeNPs were dispersed in 15 mL of toluene in a round bottom flask and ultrasonicated for 30 min.³⁵ To this suspension, MPTMS (0.6 mL) was added, and the suspension was refluxed at 110 °C overnight followed by centrifugation. The resulting thiol functionalized ceria nanoparticles (TC) were washed with toluene and ethanol and air-dried at 50 °C overnight.

Conjugation of peptide to TC (TCP)

The branched peptide amphiphile was conjugated to thiol functionalized ceria *via* thiol-ene reaction.^{36–38} The 20 mg of TC was dispersed in 1 mL of methanol and sonicated for 15 min, followed by the addition of 2 mM, 5 mL solution of PA. Radical initiator (Irgacure 2959, 0.2 equiv., 2.81 mg) was added to the mixture and suspension was exposed to UV light (365 nm) with stirring at room temperature for 1 h. The peptide-functionalized ceria nanoparticles (TCPs) were centrifuged at 6000 rpm, washed thoroughly with methanol, and finally dried overnight at 50 °C.

Characterization

All materials (CeNPs, TCs, and TCPs) were characterized as follows:

Chemical properties. Crystallinity of materials were determined by PXRD using Rigaku Miniflex diffractometer in the range of 10–80 with Cu Kα (λ = 0.154 nm) radiation. EDX (Bruker Splash 6130) and XPS (Thermo Fisher Scientific, K-alpha) analysis were performed to confirm the elemental distribution and electronic state of elements. Thermogravimetric analysis (TA Instruments, USA, SDT-650) was done under N₂ flow (100 mL min⁻¹) from 40–800 °C at a ramp of 10 °C min⁻¹ to validate the thiol modification on the surface of CeNPs and

the conjugation of peptide amphiphile on thiolated ceria surface.

Physical properties. Particle size distribution was measured on a DLS Microtrac/Nanotrak Flex instrument. Surface charge was determined by measuring the zeta potential on Particle Metrix zetazizer. Field emission scanning electron microscope (FESEM) was used to study the morphological characteristics of nanomaterials by employing the drop-casting method on silicon wafer followed by drying and platinum sputtering. Using N₂ adsorption analysis (Quantachrome), surface area and pore size were determined. Samples were degassed at 120 °C prior to the examination. Using a Quantachrome device, the surface area was calculated using the Brunauer–Emmett–Teller (BET) method.

Thiol group quantification

Ellman's assay was used to quantify the functionalization of CeNPs with thiol group.³⁹ Working solution of DTNB (Ellman's reagent) was prepared by dissolving 4 mg of DTNB and 20.5 g of sodium acetate in 5 mL of DI water. About 50 μL of this Ellman's reagent was mixed with 2.5 mL of tris buffer (1 M, pH: 8) to make blank solution. Samples were suspended in tris buffer and 250 μL of each sample (1 mg mL⁻¹) was added to blank solution and incubated for 15 min at room temperature. The absorbance was measured at 412 nm, and the quantity of surface thiol group was determined by dividing absorbance by the extinction co-efficient of the reagent (13 600 M⁻¹ cm⁻¹).

Hydrolase-mimicking activity

The esterase-mimicking activity was measured using *p*-nitro phenyl acetate (*p*NPA) as a substrate.²⁹ All nanomaterials (CeNP, TC, TCP, and PA) were dispersed in 20 mM phosphate buffer of pH 7 at a concentration of 1 mg mL⁻¹. The suspension was thoroughly vortexed and sonicated for 10 min to get a homogeneous suspension. The catalytic suspension (100 μL) was added into the well of a 96-well plate, followed by the addition of 6 μL of *p*NPA solution (12.5 mM in ACN). The absorbance was measured at 405 nm, every 15 min for 150 min. Catalyst without *p*NPA was considered as the reference and its absorbance was recorded simultaneously. The absorbance value of reference was subtracted from that of sample to eliminate the contribution of catalyst. Absorbance of substrate without catalyst was observed to check their self-hydrolysis potential and these values were also subtracted to obtain the values exclusively from catalysis. The calibration curve of *p*-nitrophenol (*p*NP) was used to determine the amount of *p*NP formed and the rate of hydrolysis was calculated using the extinction coefficient of *p*NP, which is 13 014 M⁻¹ cm⁻¹. The phosphatase-mimicking activity was measured using *p*-nitro phenyl phosphate (*p*NPP) as a substrate and its working solution was prepared (12.5 mM) in 25 mM tris buffer of pH 8. The experiment was performed in a similar way as mentioned above. In order to calculate kinetic parameters, *p*NPA or *p*NPP concentrations between 0.5 and 10 mM were used and the data was fitted to Michaelis–Menten equation.

Catalytic rate constant of biocatalyst (k_{cat}) was obtained from linear Lineweaver–Burk plots.

Reusability of catalyst

We used a different method to test the reusability of catalyst because the 96-well plate was not suitable for it. In this instance, catalysts were suspended (1 mg mL⁻¹) in 940 μL of 20 mM phosphate buffer, pH 7 using the same procedure as previously described. Substrate (60 μL of 12.5 mM solution of *p*NPA and *p*NPP in ACN or Tris buffer) was added to this suspension, and the sample was agitated on a mechanical shaker for 45 min. The sample was immediately centrifuged, supernatant was transferred to cuvette, and solution's absorbance was measured at 405 nm. The absorbances of catalyst (same concentrations as used for samples and similar treatment but without adding the substrate) or the substrate (same concentrations as used for samples) were recorded at 405 nm and subtracted from data obtained for samples in order to remove any absorbance that might come from catalyst or substrate. The entire process was carried out five times. PNP's calibration curve was used to quantify PNP formed in each case. The catalysts were cleaned five times with water and ACN, and vacuum-dried at 50 °C for 12 h after each cycle. The dried catalyst was used in additional cycles using the same procedure. In order to confirm the morphology and structure of recycled TCP, FT-IR and FESEM studies were performed after fifth cycle.

Haloperoxidase-mimicking activity

The haloperoxidase-mimicking potential of catalyst was investigated by a protocol reported earlier with little modifications.⁴⁰ Phenol red (0.6 mg, 50 μM) and ammonium bromide (0.1 mg, 25 μM) were dissolved in water, which was followed by the addition of H₂O₂ (1 μL, 300 μM). The pH of solution was maintained between 5 and 6 and each sample (1 mg mL⁻¹) was incubated with reactant solution for 5 h at room temperature and absorbance was recorded in the range of 400–600 nm using a UV-Vis spectrophotometer.

Acetylcholine esterase (AChE)-mimicking activity

Stock solutions of acetylthiocholine and DTNB (40 mg, 100 mM) were prepared in 1 mL DMSO. About 1 mg of each material (CeNP, TC, TCP, and PA) were suspended in 1 mL of PBS, and 10 μL of DTNB and acetylthiocholine were added to the suspension of catalysts. The absorbance data was collected at 405 nm, every 15 min for 150 min. The rate of reaction was obtained by using the molar extinction coefficient of 13 600 M⁻¹ cm⁻¹.

Detection of pesticides

Pesticides, such as organophosphorus (OP) and carbamates, can be detected by nanocatalysts using colorimetric assay. Stock solution of acetylthiocholine and DTNB were prepared as mentioned above and 10 μL of both solutions were added to the suspension of CeNP, TC, TCP, and PA (1 mg mL⁻¹). To this control solution, 10 mM (10 μL) of pesticide (azamethiphos, profenofos, or carbendazim) was added and incubated at

37 °C for 15 min followed by the measurement of absorbance at 405 nm.

***In vitro* biomineralization**

The capability of catalyst to imitate alkaline phosphatase (ALP) enzyme was assessed by Ca^{2+} mineralization study.^{24,41} Wells of a 24-well plate were coated with 1 mg mL⁻¹ suspension of materials (CeNP, TC, TCP, and PA) and dried over night at 37 °C. An osteogenic solution was prepared containing 15 mM of β -glycerophosphate as a source of Pi and 20 mM of CaCl_2 as a source of Ca^{2+} in Tris-HCl buffer (25 mM, pH 7.4). About 800 μL of this solution was added into each well and incubated at 37 °C for 7 days. Every day, osteogenic solution was replenished with fresh osteogenic solution. At 7th day, wells were washed after removing the solution with distilled water and allowed to air dry. The mineralized calcium developed due to the ALP-mimicking activity was further measured by using 0.1% alizarin red S staining. After staining, the excess dye was removed and wells were washed with DI water and 10% cetylpyridinium chloride was added to dissolve calcium-bound dye. Finally, absorbance was measured at 562 nm to quantify the mineralized calcium using a Tecan multimode microplate reader. Wells containing only osteogenic solution without any sample were treated as a control.

Cell culture study

The MC3T3-E1, passage 5–6 were cultured in MEM- α , supplemented with 10% FBS and 1% Pen-Strep antibiotic solution at 37 °C with 5% CO_2 in a humidified atmosphere. For experiments involving long term cell culture studies, medium was changed every 72 h and cells were harvested using a solution of trypsin-EDTA at around 70–80% confluency. The cells were further redispersed in complete media and seeded in dishes or cell culture plates to conduct various cell culture studies. Osteogenic media (OM) composed of complete MEM- α containing 100 nM dexamethasone, 50 $\mu\text{g mL}^{-1}$ ascorbic acid, and 10 mM β -glycerophosphate disodium salt hydrate was considered as a positive control, while complete MEM- α without sample was treated as a negative control.

Cytocompatibility study. The cytocompatible nature of nanomaterials (CeNP, TC, TCP, and PA) was evaluated *via* MTT assay as per protocols reported earlier.⁴² Pre-osteoblast cells were seeded with a cell density of 1×10^5 cells into a 96 well plate (100 μL per well) and after an incubation for 24 h, media was replaced with the suspension of nanomaterials in varying amounts (50, 100, 300, and 500 μg). After 2, 7, and 14 days of incubation, treated cells were incubated with 20 μL of MTT solution (5 mg mL⁻¹ in PBS) for 3.5 h to evaluate their cyto-compatibility. The absorbance of solubilized formazan in DMSO was measured at 595 nm, which is directly correlated to the cellular viability. The experiment was performed in triplicate and data were presented as mean \pm SD.

Alkaline phosphatase (ALP) activity. *In vitro* ALP activity was estimated following the protocol described earlier in literature with little modifications.^{35,43} Cells with a density of 1×10^5 cells per mL were cultured in a 48-well plate for 24 h. After

aspirating the media, cells were treated with a suspension of nanomaterials (50, 100, 300, and 500 μg) in complete MEM- α and incubated for 7 days. The cell media was replenished with nanoparticle suspension every 72 h. Media was removed at day 7 and cells were washed with DPBS, which was followed by lysis of cells with 200 μL of 0.2% Triton X for 30 min in order to rupture the cell membrane and release ALP molecules. Subsequently, 200 μL of *p*-nitrophenyl phosphate (*p*NPP) solution was added to each well containing cell lysates and incubated at 37 °C for another 90 min in dark. The final absorbance was measured at 405 nm using a plate reader.

Calcium deposition assay. The ALP activity was further confirmed by calcium deposition study on MC-3T3 cells by alizarin red S staining. The media was replaced with the suspension of nanomaterials and osteogenic media every 72 h. On day 7 of cell culturing, cells were washed with DPBS and fixed with 4% paraformaldehyde solution for 10 min at 4 °C. Followed by washing with DI water, fixed cells were stained with 500 μL of alizarin red S solution (40 mM, pH = 4.1–4.3) for 1 h in dark. After staining, cells were washed with DI water to remove the excess dye, and cells were visualized under an inverted microscope to observe red crystals. To further quantify the total amount of calcium deposited in mineralized cells, 200 μL of 10% cetylpyridyl chloride solution was added to each well and incubated for 30 min in dark condition, and 100 μL of dissolved crystals was transferred to 96-well plate and absorbance was measured at 562 nm.

Study of osteogenic markers. To assess the impact of nanomaterials on osteogenic differentiation, MC3T3-E1 cells were used to measure the mRNA expression of markers related to osteogenesis, such as alkaline phosphatase (ALP), runt-related transcription factor 2 (RUNX2), osteocalcin (OCN), and osteopontin (OPN). Cells were seeded at a density of 1×10^4 cells per mL in a 6-well plate and cultivated for a duration of 24 h. After removing media, 1 mL of fresh media containing 300 μg of nanomaterial was added. Every 72 h, medium was changed, and on days 7 and 14, total cellular RNA was extracted by lysis in TRIzol and then subjected to cDNA synthesis. Quantitative real-time polymerase chain reaction was used to measure the expression of particular genes (qRT-PCR). The β -actin gene was utilized as a housekeeping gene to standardize expression levels.

Anti-biofouling activity

The anti-biofouling property of nanomaterials was evaluated using bacteria *E. coli*. Suspension (300 μL) of nanomaterials (1 mg mL⁻¹) were casted on 24-well plates and air dried for 24 h.^{44,45} Meanwhile, *E. coli* was grown till mid-log phase in Luria broth (LB) media overnight and optical density was adjusted to 0.1 at 600 nm. Samples were incubated with 500 μL of bacterial suspension, 4.4 mM NH_4Br , and 0.42 mM H_2O_2 for 24 h at 37 °C inside an incubator. Wells containing bacterial suspension, NH_4Br , and H_2O_2 without samples were considered as controls. After 24 h of incubation, suspension was removed and wells were gently washed with distilled water to remove the weakly attached bacteria. Next, the biofilm was

air-dried and stained with crystal violet (0.1%) for 15 min and washed again with DI water to remove excess dye. Finally, 33% acetic acid solution was added to dissolve crystal violet bound to biofilm and absorbance was measured at 590 nm to quantify biofilm formation.

Statistical analysis

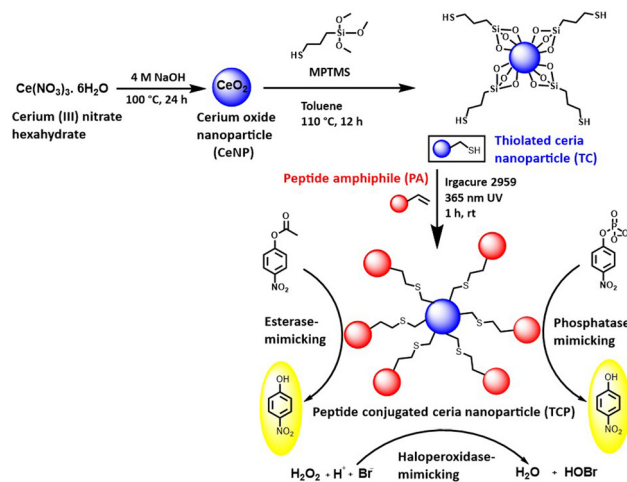
The studies were performed in triplicates and presented as average values and standard deviations from the mean value. The data were analyzed using Student's *t*-test. **p* value ≤ 0.05 was considered significant, whereas ns represents non-significant difference.

Results and discussion

The aim of this work was to develop multifunctional, artificial enzyme to mimic esterase, phosphatase, and haloperoxidase activity for pesticide detection, bone regeneration, and antibiofouling material application. In order to achieve this goal, we have synthesized self-assembled, branched peptide amphiphile (PA) containing the catalytic triad (Ser: S; His: H; and Asp: D) present in serine protease enzyme. Histidine acts as a base to abstract proton from the side chain of serine and nucleophilic oxygen of serine attacks carbonyl carbon of a substrate to form a covalent bond during catalysis. The negatively charged Asp stabilizes the positive charge that forms on His residue through hydrogen bonding and helps in the orientation of substrate within the active site. These three amino acid residues together create a microenvironment within enzyme's active site that promotes specific chemical reactions, thus, making peptide-based artificial enzymes capable of catalyzing various biochemical transformations.²⁷ Immobilization of peptide on the surface of thiol-modified ceria nanoparticles (TC) helped us in developing a recyclable, heterogenous biocatalyst. The self-assembled peptide rendered the hydrolase-mimicking activity, whereas CeNPs imparted the haloperoxidase-mimicking activity. Ceria nanoparticles undergo reversible oxidation and reduction reactions, and shuttle between Ce^{4+} and Ce^{3+} oxidation states. This redox cycling property allows ceria nanoparticles to act as catalysts for the oxidation of halide ions by hydrogen peroxide. The multifunctional, enzyme mimic was explored for pesticide detection, bone regeneration, and antibiofouling applications (Scheme 1).

Design, synthesis, and characterization of PA

We have rationally designed the self-assembled, peptide amphiphile (PA) by incorporating a catalytic triad (S, H, and D) to mimic the catalytic site of serine protease, which belongs to the class of hydrolase enzyme (Fig. 1A). The branched peptide was synthesized in order to provide two catalytic sites in a single molecule. Based on literature review, we hypothesized that participating amino acids from nearby peptides will work together to increase the activity of peptide in its self-assembled state and local substrate concentration will be higher at the surface of aggregates. Therefore, we incorporated hydrophobic



Scheme 1 Fabrication of thiol-functionalized ceria nanoparticles and their conjugation to branched peptide amphiphile via thiol–ene conjugation for esterase, phosphatase, and haloperoxidase-mimicking activities.

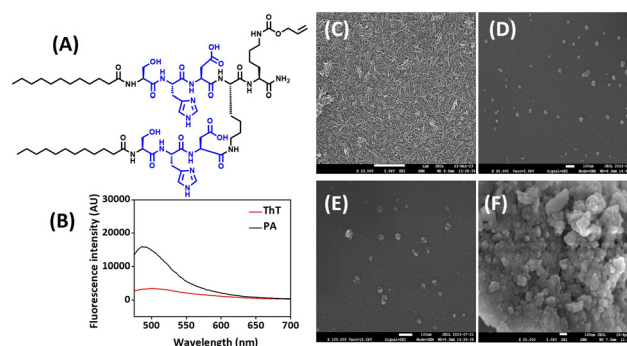


Fig. 1 (A) Structure of self-assembled, branched peptide amphiphile (PA). (B) ThT assay. (C–F) FESEM images of: (C) PA, (D) CeNP, (E) TC, and (F) TCP. Scale bar: (C) 1 μm , and (D–F) 100 nm.

chain (C_{12}) at the N-terminus of peptide to make it amphiphilic. As our aim was to conjugate the peptide to nanoparticle by thiol–ene reaction, we did not deprotect the Alloc-protecting group in peptide sequence.

Fmoc-based SPPS technique was used to synthesize peptide and rink amide resin was used as a solid support to generate carboxamide group at C-terminus. The peptide was characterized by RP-HPLC, ^1H NMR, and mass spectrometry. The RP-HPLC showed that peptide is 90–95% pure and retention time was 4.6 min (Fig. S1 \ddagger). The MW derived from the mass spectrum matched to its theoretical value (Fig. S2 \ddagger). The ^1H NMR spectrum of peptide along with assigned peaks is included in the ESI (Fig. S3 \ddagger). The secondary structure of self-assembled peptide was investigated by CD spectroscopy and ThT assay. Negative peaks around 200 and 215 nm indicated the formation of α -helical morphology of peptide (Fig. S4 \ddagger). ThT assay demonstrated the enhancement of fluorescence intensity in the range of 490–500 nm, thereby implying the

binding of ThT to the hydrophobic region of peptide (Fig. 1B). Unbound ThT without any sample was considered as a control. FT-IR spectra further indicated the presence of α -helical structure as it showed characteristic peak at 1673 cm^{-1} (Fig. S5 \ddagger). The nanoscale architecture of self-assembled peptide was evaluated by FESEM investigations. The microscopic image suggested the formation of long nanofibers by peptide in its self-assembled form (Fig. 1C). The width of nanofibers was calculated by ImageJ software and was found to be below 20 nm, which can be attributed to long hydrophobic carbon chain attached at the N-terminus of peptide.

Preparation and functionalization of CeNPs

CeNPs were fabricated using hydrothermal method by hydrolyzing a ceria precursor, cerium nitrate hexahydrate, in a basic environment. The surface was modified post synthesis using MPTMS to obtain thiol modified ceria nanoparticles (TCs). The thiol grafting on surface was quantified by Ellman's assay, which suggested a $1306.88\text{ }\mu\text{mol g}^{-1}$ of thiol density. The nanoparticles were characterized by XRD, TGA, BET, elemental mapping, and XPS. The morphology and size of nanoparticles were confirmed by FESEM and DLS, as discussed below.

Conjugation of peptide amphiphile to ceria

Our objective was to develop a multifunctional peptide-based catalyst in which the hydrolase-mimicking PA was immobilized on a ceria surface to produce a heterogeneous catalyst. Immobilization on solid surface is expected to maintain its catalytic efficiency. The assembly of catalytic peptides on a solid surface increases the peptide catalyst's robustness and reusability features in an integrated manner. With this aim, the PA was conjugated on the surface of thiolated ceria (TC) *via* thiol-ene click reaction. The successful grafting was confirmed by Ellman's assay, which showed that free thiol density on ceria surface has reduced after conjugation to the peptide.

Characterization

All nanomaterials (CeNP, TC, and TCP) were characterized by PXRD, TGA, elemental mapping, FESEM, and DLS. FESEM images revealed that CeNPs have a spherical morphology and grafting of thiol groups and peptide on the surface of CeNPs did not change its morphology (Fig. 1D–F). It is interesting to note that the conjugation of branched PA on the surface of CeNPs reduced their aggregation, thus, increasing their stability. The size of nanoparticles increased a little after functionalization. The particle size distribution is evident from DLS study. The size of CeNPs, TC, and TCP were found to be 41.6, 49.7, and 54.3 nm (Fig. S6 \ddagger). Zeta potential of nanoparticles was also investigated to measure their surface charge and found to be 0.5, -1.5 , and 5.9 mV for CeNP, TC, and TCP. It was observed that zeta potential becomes more negative after thiolation of CeNPs and it increased after conjugation to peptide. XRD patterns of materials were investigated to determine their crystal structure (Fig. 2A). CeNPs showed characteristic sharp peaks at 28.38 , 32.82 , 47.16 , and 56.1° corresponding to 111, 200, 220, and 311 planes.⁴⁰ This data

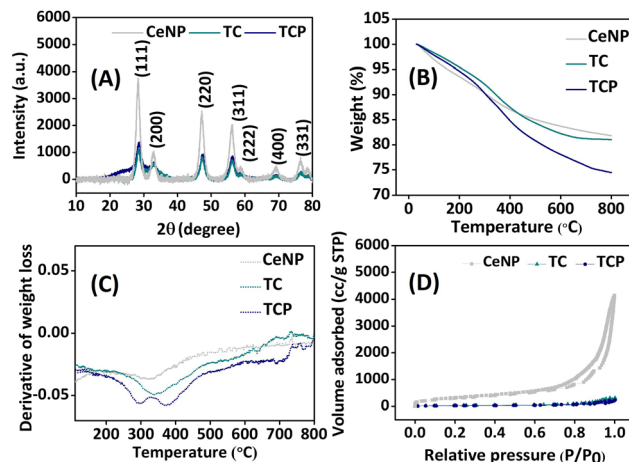


Fig. 2 Characterization of nanoparticles: (A) PXRD patterns, (B) TGA study, (C) derivative of thermogravimetry (DTG) curve, and (D) N₂ adsorption/desorption isotherm.

suggested that CeNPs have a fluorite-type crystalline nature as per JCPDS file no. 34-0394.⁴⁶ Neither surface modification with thiol groups nor the integration of peptide on it changed the PXRD pattern or induced any new peak. However, the intensity of peaks was significantly reduced. This data suggests that crystallinity decreased after modification of ceria surface.

Successful grafting of thiol group and PA on ceria surface was further confirmed by TGA. The percentage mass loss was observed from TGA curve (Fig. 2B), whereas derivative thermogravimetry curve (DTG) demonstrated the rate of material weight change upon heating (Fig. 2C). The weight loss at $100\text{ }^\circ\text{C}$ signified the loss of adsorbed water molecules from nanomaterials. The weight loss from 250 – $400\text{ }^\circ\text{C}$ in DTG curve was due to the loss of organic moiety from ceria surface.²⁹ The NPs obtained at different stages were also investigated for pore size measurements and surface area using Brunauer–Emmett–Teller (BET) method. The pore size parameters and specific surface area are tabulated in Table S1 (ESI) \ddagger . The BET surface area analysis revealed type IV isotherm, which indicated mesoporous structures for all NPs (Fig. 2D). The BET surface area of CeNP was found to be $1120\text{ m}^2\text{ g}^{-1}$, which sharply decreased to $105\text{ m}^2\text{ g}^{-1}$ and $76\text{ m}^2\text{ g}^{-1}$ after thiol-functionalization (TC) and subsequent peptide conjugation (TCP). This considerable reduction in surface area implies that surface functions are blocking pores. All data confirm the functionalization of CeNP with thiol ($-\text{SH}$) and PA. EDX spectra and elemental mapping helped in understanding the elemental distribution of all NPs (Fig. 3A and S7 \ddagger). It was observed from the elemental mapping that CeNP surfaces were functionalized with thiol and PA.

The oxidation state of components of nanocomposite was revealed by XPS spectra (Fig. 3B–G). High-resolution Ce 3D spectrum of TCP showed spin-orbit coupling peaks corresponding to Ce $3d_{3/2}$ and Ce $3d_{5/2}$. The peaks designated at II (884.9 eV), IV (898.0 eV), and VI (903.0 eV) corresponds to Ce^{3+} oxidation state, whereas peaks at I (882.1 eV), III (888.5 eV), V (900.6 eV), VII (907.0 eV), and VIII (916.3 eV) corresponds to

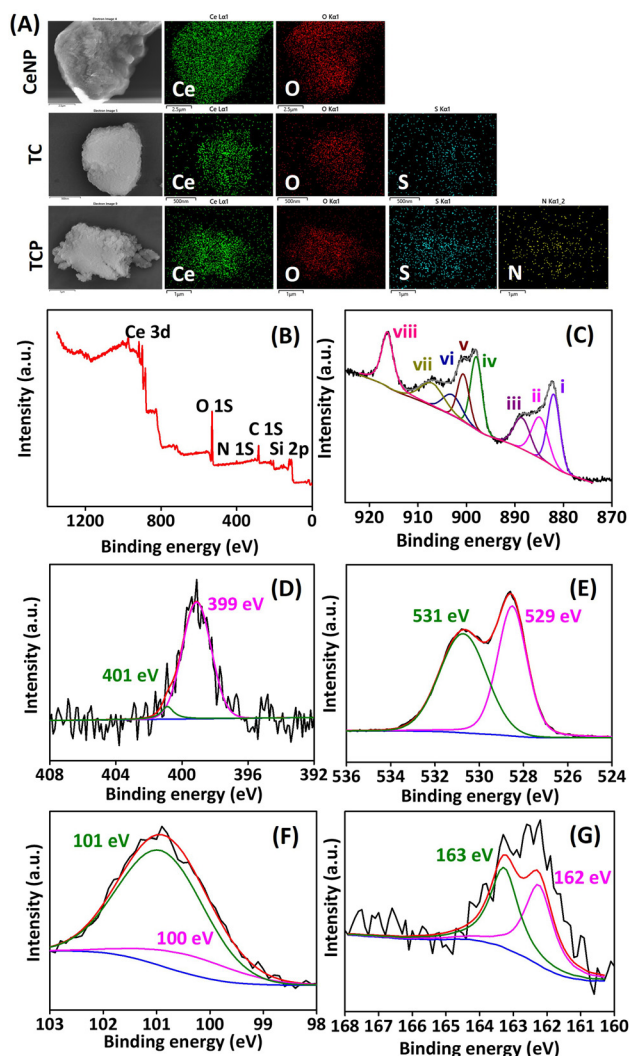


Fig. 3 (A) Elemental mapping of CeNP, TC, and TCP. (B–G) XPS spectra of TCP: (B) surface survey spectrum, (C) Ce 3d, (D) N 1s, (E) O 1s, (F) Si 2p, and (G) S 2p.

Ce⁴⁺ oxidation state.⁴⁰ The ratio of Ce³⁺ to Ce⁴⁺ was calculated as 0.57 (Table S2, ESI†). The oxygen vacancy in material was assessed using the expression $\text{Ce}^{3+}/(\text{Ce}^{3+} + \text{Ce}^{4+}) \times 100$, and found to be 37%. TCPs with 37% oxygen vacancy can demonstrate efficient catalytic activity, as this activity is enhanced by oxygen vacancy.

Catalytic activity

The individual and cooperative role of amino acids in catalytic pockets of enzymes are best understood using peptide nano catalysts, which are simple to synthesize and modify. His residues can be used to create PA nano catalysts with catalytic domains. Here, we developed branched, amphiphilic peptide with two catalytic domains, composed of Ser–His–Asp. The artificial enzyme can undergo self-assembly to mimic the folding mechanism of natural enzyme. Accordingly, we integrated a catalytic peptide on the surface of CeNPs to evaluate

their multiple enzyme-mimicking properties. The esterase-mimicking activity was assessed using *p*NPA as a substrate, which upon hydrolysis generates yellow colored product, *p*NP ($\lambda_{\text{max}} = 405 \text{ nm}$). Autohydrolysis of *p*NPA was considered as a blank and its absorbance value was subtracted from that of final absorbance (Fig. 4A). The PA exhibited maximum absorbance, which signifies highest amount of *p*-NP production due to the hydrolysis of substrate by esterase-mimicking enzyme. The activity was reduced when PA was conjugated to ceria NP surface but it was better than CeNPs and TCs. The overall velocity was also determined and it was found to be 0.57, 0.64, 0.82, and 1.42 $\mu\text{M min}^{-1}$ for CeNP, TC, TCP, and PA (Fig. 4B). The esterase-mimicking activity was comparable to catalyst reported by Gulseren *et al.*²⁷

The phosphatase-mimicking activity was determined using *p*NPP as a substrate. It was fascinating to note that CeNPs exhibited the highest rate of hydrolysis of phosphoester bond (Fig. 4E). This dephosphorylation by CeNPs was attributed to the Lewis acidity of metal, which facilitates the interaction with negatively charged phosphate moiety. It results in a series of phenomenon like, Lewis acid activation and nucleophilic attack, followed by the removal of phosphate group. In particular, Ce⁴⁺ tend to exhibit higher catalytic reaction than Ce³⁺ due to its oxygen vacancy and better efficacy in attracting electron.^{47–49} After the functionalization of CeNPs with thiol and PA, the phosphoesterase-mimicking activity was reduced. The overall velocity of TCP was better than PA (Fig. 4F). Herein, the catalytic activity of peptide conjugated ceria nanoparticles (TCP) toward *p*NPA and *p*NPP was assessed. The initial hydrolysis rates V_0 -*p*NPA for the substrate *p*NPA are shown as a function of *p*NPA concentration (0.5–10 mM) in Fig. 4C and V_0 -*p*NPP was measured at different concentrations (0.5–10 mM) of *p*NPP (Fig. 4G). The reactions followed conventional Michaelis–Menten model, confirming that TCP exhibited catalytic behavior. An enzyme's catalytic rate constant (k_{cat}) represents the highest quantity of substrate molecules transformed into product for each active site in a unit of time. This value indicates the rate at which an enzyme–substrate complex proceeds with its forward process. The k_{cat} was measured based on linear Lineweaver–Burk plot and it was calculated as 0.041 min^{-1} and 0.01 min^{-1} for TCP to catalyse *p*NPA and *p*NPP (Fig. 4D and H). A comprehensive parameter to assess an enzyme's capacity to catalyse a particular substrate is its catalytic efficiency ($k_{\text{cat}}/K_{\text{m}}$). The catalytic activity of TCP toward *p*NPA and *p*NPP was examined in this study and calculated as 0.0160 $\mu\text{M}^{-1} \text{min}^{-1}$ and 0.0082 $\mu\text{M}^{-1} \text{min}^{-1}$. This data validates the superior esterase-like activity of TCP compared to its phosphatase-mimicking property.

A catalyst's recyclability is one of main issues for the industrial use. We carried out a single batch of catalysis followed by the removal of catalyst by filtration, washing, and drying in order to determine whether the heterogenous catalyst (TCP) can be recycled after each usage. The catalyst was reused for subsequent batch after drying. The same procedure was carried out five times (Fig. S8†), and no significant loss of activity was observed even during the fifth cycle. This demon-



Fig. 4 (A–D) Catalytic activity for *p*NPA hydrolysis. (A) Absorbance of *p*NP vs. time, (B) overall velocity of reaction catalyzed by CeNP, TC, TCP, and PA, and (C and D) catalytic reaction rate. (E–H) Similar curves for *p*NPP hydrolysis. The data in (C), (D), (G), and (H) are fitted with Michaelis–Menten equation and Lineweaver–Burk plot.

strates that catalyst can be recycled using a simple and affordable process. The catalyst retained its morphology and structure after fifth cycle, as evident from FESEM and FT-IR analysis (Fig. S9, ESI†).

A class of peroxidase enzyme called haloperoxidase catalyzes the oxidation of halides.⁴⁵ The material's haloperoxidase-like activity was assessed utilizing the phenol red bromination assay. In the presence of halide ions and peroxide, phenol red can be brominated to generate bromophenol blue using a catalyst that resembles haloperoxidase. A successful reaction is indicated by a drop in absorbance intensity of phenol red ($\lambda_{\max} = 432$ nm) and a corresponding increase in absorbance intensity of bromophenol blue ($\lambda_{\max} = 590$ nm). It can be observed from Fig. 5A that CeNPs exhibited highest haloperoxidase-

mimicking activity with a shift of wavelength from 430 to 590 nm and absorbance was highest at 590 nm among other nanomaterials. TCP also exhibited significant potential in bromination of phenol red while showing haloperoxidase-mimicking activity.

Applications

The heterogenous biocatalyst, developed in this work was explored for several applications.

Acetylcholinesterase-mimicking activity.

Acetylcholinesterase is an enzyme found on postsynaptic membrane and at neuromuscular junction.⁵⁰ Its primary function is to hydrolyze acetylcholine into choline and acetate. The breakdown of acetylcholine by acetylcholinesterase allows for precise control over nerve impulses, preventing overstimulation or prolonged activation of postsynaptic receptors. By clearing acetylcholine from synapse, acetylcholinesterase prepares postsynaptic neuron for subsequent signals, ensuring efficient synaptic communication. The two main types of pesticides that have been frequently used in agricultural are organophosphates (OPs) and carbamates, which irreversibly inhibit AChE and pose a serious threat to human nervous, respiratory, and cardiovascular systems. Therefore, it is highly desirable to identify OPs and carbamates selectively.⁵¹ In order to estimate the AChE-mimicking property of catalyst, we carried out experiment on acetylthiocholine, which is an analogue of acetylcholine and can produce thiocholine upon hydrolysis by AChE. The generated thiocholine can be estimated by colorimetric method using Ellman's reagent (Fig. 5B). PA displayed highest efficiency in hydrolysis of acetylthiocholine followed by TCP. CeNP and TC exhibited very less activity, which suggests the importance of catalytic triad in the artificial enzyme. It is evident from Fig. 5C that carbendazim, an acetylcholine esterase inhibitor, inhibited TCP the most, which reduced absorbance of thiocholine. After confirming the acetylcholine esterase (AChE)-mimicking potential of catalyst, it was investigated

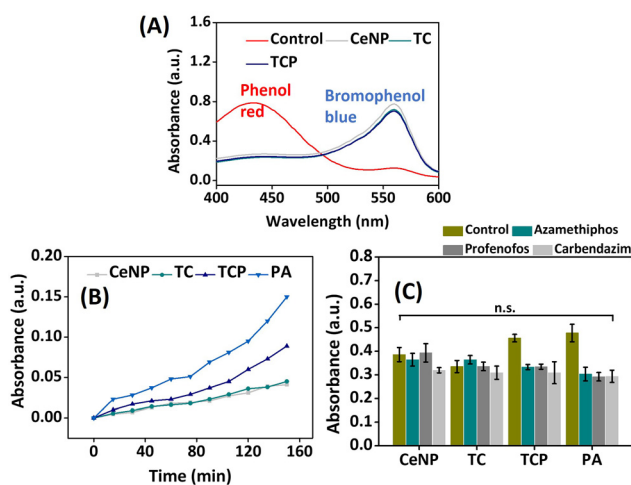


Fig. 5 (A) UV-Vis spectra of haloperoxidase-like activity of CeNP, TC, and TCP, (B) comparison of catalytic activity of nanomaterials using acetylthiocholine as the substrate, and (C) response of nanomaterials towards the detection of pesticides. Data are presented as mean \pm SD, $n = 3$ and ns indicate non-significant difference.

for pesticide (organophosphorus and carbamate) detection. The mechanism of pesticide detection has been elucidated in Fig. S10 (ESI).[†]

Alkaline-phosphatase mimicking activity. Alkaline phosphatase (ALP) is an enzyme, which hydrolyses phosphoester bond found in various tissues throughout the body, including the liver, bone, intestines, and placenta.^{24,41,52} It plays an important role in biological processes, such as energy production, signal transduction, bone formation, and gene maintenance. It is particularly important in bone formation and mineralization. ALP is produced by osteoblasts, which are responsible for building and maintaining bone. Alkaline phosphatase catalyzes the removal of phosphate groups from various molecules, thus, allowing the deposition of calcium and phosphate ions to form hydroxyapatite, the mineralized matrix of bone. Therefore, we assessed the potential of our catalyst to imitate ALP to further use it for bone tissue regeneration applications.

We evaluated the cell viability of mesenchymal stem cells (MC3T3-E1) at different concentrations of nanomaterials (50, 100, 300 and 500 $\mu\text{g mL}^{-1}$). It was evident from the data shown in Fig. 6A that CeNPs exhibited toxicity at higher concentrations, whereas PA helped in the proliferation of cells at increased concentration. Surface modification of CeNPs with thiol and PA decreased its toxicity and enhanced the cell viability. The same trend was observed in long-term cell viability analysis, where toxicity increased with increase in the concentration of nanomaterials but it was masked after conjugating

the peptide (Fig. S11[†]). As far as ALP assay was concerned, CeNPs exhibited highest ALP activity on MSCs at 300 $\mu\text{g mL}^{-1}$ concentration, which can be attributed to their inherent phosphatase-mimicking properties owing to their reversible $\text{Ce}^{3+}/\text{Ce}^{4+}$ redox pair and oxygen vacancy in the outer shell⁴⁷ (Fig. 6B). Modification of their surface with thiol and peptide interferes with their oxidation state and deteriorate their ALP-mimicking potential.

Considering the phosphatase like catalytic behavior of nanomaterials, we anticipated that they will have a great potential in stimulating osteogenic activity of natural ALP. In order to verify this, 24-well plates were coated with nanomaterials and incubated with osteogenic solution containing β -glycerophosphate and CaCl_2 for 7 days. Alizarin red S staining was carried out to quantify the amount of calcium deposition on the surface of well. As shown in Fig. 6C, the deposited calcium on the surface of well coated with nanomaterial was higher compared to the control (osteogenic media without any sample), thus, confirming the ALP-like catalytic property of nanomaterials, which accelerated the hydrolysis of β -glycerophosphate. The phosphatase-mimicking activity was again evaluated on MC3T3-E1 cells and similar trend was observed where TCP allowed a significant calcium deposition (Fig. 6D). Controls used were cells treated with complete media and those treated with osteogenic media (OM). Inverted microscopy revealed red-stained calcium nodules representing the extracellular calcium accumulation



Fig. 6 (A) Cytocompatibility study on MC3T3-E1 cells in the presence of nanomaterials (CeNPs, TC, TCP, and PA) at different concentrations (50, 100, 300, and 1 $\mu\text{g mL}^{-1}$) using MTT assay at 48 h, (B) alkaline phosphatase (ALP) activity of nanomaterials on MC3T3-E1 cells on 7th day, and (C–E) evaluation of calcium deposition using alizarin red S staining: (C) on the surface of well plate, (D) on the surface of MC3T3-E1 cells, and (E) microscopic images of MC3T3-E1 cells after 7 days of treatment with nanomaterials. Red precipitates illustrate calcium nodules. Data are presented as mean \pm SD, $n = 3$, and $*p < 0.05$ indicates statistically significant data and ns non-significant data. Scale bar: 100 μm .

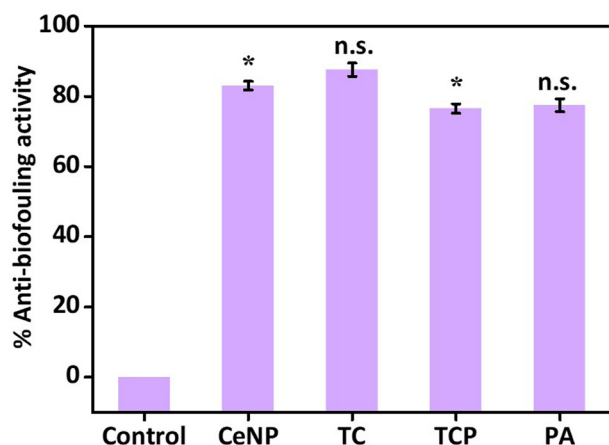


Fig. 7 Antibiofouling activity of CeNPs, TC, TCP, and PA against *E. coli*. Data are presented as mean \pm SD, $n = 3$, and * $p < 0.05$ indicates statistically significant data and ns non-significant data.

(Fig. 6E). The results are comparable to earlier reports by Wang and coworkers and Gulseren *et al.*⁴¹ Understanding the expression of genes that regulate bone tissue regeneration is essential for developing therapies and interventions to enhance bone healing in various clinical contexts, including fractures, bone defects, and osteoporosis. Therefore, we studied the expression of genes (ALP, OPN, OCN, and RUNX2) in the presence of CeNP, TC, TCP, and PA on MC3TC-E1 cell lines for 7 and 14 days by RTPCR. We observed that TCP was able to upregulate expressions of ALP, OPN, OCN, and RUNX2 by 5, 7, 8, and 16-fold (Fig. S12†). The data was also compared to positive control (OM). The results suggest that this material holds significant potential for advancing bone regeneration therapies.

Finally, the haloperoxidase-like activity of nanomaterials was exploited for the development of anti-biofouling materials.⁵³ The non-specific surface adherence of microorganisms is known as biofouling, which can harm materials' performance and decrease its effectiveness by causing corrosion on metallic surfaces and bacterial infections on surgically implanted medical equipment.^{32,54} Haloperoxidase can catalyze the oxidation of halides with hydrogen peroxide to generate hypohalous acids, such as HOBr, which is known to disrupt the quorum sensing of microorganisms, thus inhibiting biofilm formation, the initial step for biofouling of a surface. We determined the inhibition of biofilm formation by crystal violet staining method (Fig. 7) and heterogenous catalyst (TCP) demonstrated around 80% anti-biofouling activity. Therefore, the material can be a good candidate for integration on the surface of implant or as a coating material to prevent the adhesion of bacteria.

Conclusions

In summary, we have developed self-assembling, branched peptide amphiphile containing catalytic triad, 'Ser-His-Asp',

to mimic the active site of serine protease, which can effectively bind with esters and hydrolyze the bond. This hydrolase mimetic catalyst was immobilized on the surface of ceria nanoparticles using thiol-ene click reaction, which exhibits haloperoxidase-mimicking properties due to their reversible change between two oxidation states, Ce^{3+}/Ce^{4+} . The nanofibrous morphology of self-assembling peptide was not retained upon conjugation with ceria nanoparticles and the heterogenous catalyst adopted a spherical shape, with a diameter of 54.3 nm. Integration of peptide on nanoparticle surface minimized the aggregation of nanoparticles and made the catalyst recyclable without the loss of activity. The peptide-based nanomaterial showed multifunctional catalytic activity by mimicking the esterase, phosphatase, and haloperoxidase enzymes. The enzyme mimicking potential of biocatalyst was further explored for various applications. The acetylcholinesterase (AChE)-mimicking property was used for the detection of pesticides, like organophosphates and carbamates. Alkaline phosphatase (ALP)-like activity was exploited to develop material for bone regeneration and anti-biofouling applications. Thus, the catalyst created in this work can be quite valuable for industrial applications. Overall, this work showcases the benefits of developing multifunctional heterogenous biocatalyst to reduce the problem of instability associated with peptide-based homogenous catalyst. We also exploited the potential enzyme-mimicking role of catalyst in different fields of human welfare.

Author contributions

The manuscript was written through contributions of all authors, and they have given their approval to the final version of the manuscript.

Data availability

The data supporting this article have been included as part of the ESI.†

Conflicts of interest

Authors declare no competing financial interest.

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References

- W. Hamley, *Biomacromolecules*, 2021, **22**, 1835–1855.
- D. Yi, T. Bayer, C. P. S. Badenhorst, S. Wu, M. Doerr, M. Höhne and U. T. Bornscheuer, *Chem. Soc. Rev.*, 2021, **50**, 8003–8049.
- E. L. Bell, W. Finnigan, S. P. France, A. P. Green, M. A. Hayes, L. J. Hepworth, S. L. Lovelock, H. Niikura, S. Osuna, E. Romero, K. S. Ryan, N. J. Turner and S. L. Flitsch, *Nat. Rev. Methods Primers*, 2021, **1**, 46.
- A. Schmid, J. S. Dordick, B. Hauer, A. Kiener, M. Wubboldt and B. Witholt, *Nature*, 2001, **409**, 258–268.
- Y. R. Maghraby, R. M. El-Shabasy, A. H. Ibrahim and H. M. E.-S. Azzazy, *ACS Omega*, 2023, **8**, 5184–5196.
- R. Singh, S. Langyan, B. Rohtagi, S. Darjee, A. Khandelwal, M. Shrivastava, R. Kothari, H. Mohan, S. Raina, J. Kaur and A. Singh, *Mater. Sci. Energy Technol.*, 2022, **5**, 294–310.
- F. Rudroff, M. D. Mihovilovic, H. Gröger, R. Snajdrova, H. Iding and U. T. Bornscheuer, *Nat. Catal.*, 2018, **1**, 12–22.
- G. Alvarez, T. Shahzad, L. Andanson, M. Bahn, M. D. Wallenstein and S. Fontaine, *Global Change Biol.*, 2018, **24**, 4238–4250.
- N. F. Mokhtar, R. N. A. Rahman, N. D. Md. Noor, F. Md. Shariff and M. S. Md. Ali, *Catalysts*, 2020, **10**, 744.
- A. Basso and S. Serban, *Mol. Catal.*, 2019, **479**, 110607.
- S. Sabater, J. A. Mata and E. Peris, *ACS Catal.*, 2014, **4**, 2038–2047.
- H. J. Federsel, T. S. Moody and S. J. C. Taylor, *Molecules*, 2021, **26**, 2822.
- A. M. Garcia, M. Kurbasic, S. Kralj, M. Melchionna and S. Marchesan, *Chem. Commun.*, 2017, **53**, 8110–8113.
- C. M. Rufo, Y. S. Moroz, O. V. Moroz, J. Stöhr, T. A. Smith, X. Hu, W. F. DeGrado and I. V. Korendovych, *Nat. Chem.*, 2014, **6**, 303–309.
- M. O. Guler and S. I. Stupp, *J. Am. Chem. Soc.*, 2007, **129**, 12082–12083.
- Y. Lou, B. Zhang, X. Ye and Z.-G. Wang, *Mater. Today Nano*, 2023, **21**, 100302.
- J. N. B. D. Pelin, B. B. Gerbelli, C. J. C. Edwards-Gayle, A. M. Aguilar, V. Castelletto, I. W. Hamley and W. A. Alves, *Langmuir*, 2020, **36**, 2767–2774.
- B. List, R. A. Lerner and C. F. Barbas, *J. Am. Chem. Soc.*, 2000, **122**, 2395–2396.
- S. J. Miller, *Acc. Chem. Res.*, 2004, **37**, 601–610.
- C. Zhang, X. Xue, Q. Luo, Y. Li, K. Yang, X. Zhuang, Y. Jiang, J. Zhang, J. Liu, G. Zou and X.-J. Liang, *ACS Nano*, 2014, **8**, 11715–11723.
- Z. Huang, S. Guan, Y. Wang, G. Shi, L. Cao, Y. Gao, Z. Dong, J. Xu, Q. Luo and J. Liu, *J. Mater. Chem. B*, 2013, **1**, 2297.
- D. W. Watkins, J. M. X. Jenkins, K. J. Grayson, N. Wood, J. W. Steventon, K. K. Le Vay, M. I. Goodwin, A. S. Mullen, H. J. Bailey, M. P. Crump, F. MacMillan, A. J. Mulholland, G. Cameron, R. B. Sessions, S. Mann and J. L. R. Anderson, *Nat. Commun.*, 2017, **8**, 358.
- S. C. Patel, L. H. Bradley, S. P. Jinadasa and M. H. Hecht, *Protein Sci.*, 2009, **18**, 1388–1400.
- Y. Wang, L. Yang, M. Wang, J. Zhang, W. Qi, R. Su and Z. He, *ACS Catal.*, 2021, **11**, 5839–5849.
- O. Zozulia and I. V. Korendovych, *Angew. Chem., Int. Ed.*, 2020, **59**, 8108–8112.
- Y.-M. Wong, H. Masunaga, J.-A. Chuah, K. Sudesh and K. Numata, *Biomacromolecules*, 2016, **17**, 3375–3385.
- G. Gulseren, M. A. Khalily, A. B. Tekinay and M. O. Guler, *J. Mater. Chem. B*, 2016, **4**, 4605–4611.
- D. J. Mikolajczak and B. Koksich, *ChemCatChem*, 2018, **10**, 4324–4328.
- P. Dowari, M. K. Baroi, T. Das, B. K. Das, S. Das, S. Chowdhuri, A. Garg, A. Debnath and D. Das, *J. Colloid Interface Sci.*, 2022, **618**, 98–110.
- A. Reja, S. P. Afrose and D. Das, *Angew. Chem., Int. Ed.*, 2020, **59**, 4329–4334.
- L. Hedstrom, *Chem. Rev.*, 2002, **102**, 4501–4524.
- K. Herget, P. Hubach, S. Pusch, P. Deglmann, H. Götz, T. E. Gorelik, I. A. Gural'skiy, F. Pfitzner, T. Link, S. Schenk, M. Panthöfer, V. Ksenofontov, U. Kolb, T. Opatz, R. André and W. Tremel, *Adv. Mater.*, 2017, **29**, 1603823.
- M. Halder, Y. Bhatia and Y. Singh, *Biomater. Sci.*, 2022, **10**, 2248–2262.
- I. Trenque, G. C. Magnano, M. A. Bolzinger, L. Roiban, F. Chaput, I. Pitault, S. Briançon, T. Devers, K. Masenelli-Varlot, M. Bugnet and D. Amans, *Phys. Chem. Chem. Phys.*, 2019, **21**, 5455–5465.
- N. Rasool, D. Negi and Y. Singh, *ACS Biomater. Sci. Eng.*, 2023, **9**, 3535–3545.
- A. A. Aimetti, R. K. Shoemaker, C.-C. Lin and K. S. Anseth, *Chem. Commun.*, 2010, **46**, 4061.
- Y. Liu, W. Hou, H. Sun, C. Cui, L. Zhang, Y. Jiang, Y. Wu, Y. Wang, J. Li, B. S. Sumerlin, Q. Liu and W. Tan, *Chem. Sci.*, 2017, **8**, 6182–6187.
- Y. Li, M. Yang, Y. Huang, X. Song, L. Liu and P. R. Chen, *Chem. Sci.*, 2012, **3**, 2766.
- M. Kalantari, T. Ghosh, Y. Liu, J. Zhang, J. Zou, C. Lei and C. Yu, *ACS Appl. Mater. Interfaces*, 2019, **11**, 13264–13272.
- N. Wang, W. Li, Y. Ren, J. Duan, X. Zhai, F. Guan, L. Wang and B. Hou, *Colloids Surf., A*, 2021, **608**, 125592.
- G. Gulseren, I. C. Yasa, O. Ustahuseyin, E. D. Tekin, A. B. Tekinay and M. O. Guler, *Biomacromolecules*, 2015, **16**, 2198–2208.
- M. Halder, M. Narula and Y. Singh, *Bioconjugate Chem.*, 2023, **34**, 645–663.
- X. Cui, L. Xu, Y. Shan, J. Li, J. Ji, E. Wang, B. Zhang, X. Wen, Y. Bai, D. Luo, C. Chen and Z. Li, *Sci. Bull.*, 2024, **69**, 1895–1908.
- H. Meng, S. Cheng, L. Wang, Y. L. Balachandran, W. Zhang and X. Jiang, *Mater. Adv.*, 2022, **3**, 8137–8140.

- 45 M. Hu, K. Korschelt, M. Viel, N. Wiesmann, M. Kappl, J. Brieger, K. Landfester, H. Thérien-Aubin and W. Tremel, *ACS Appl. Mater. Interfaces*, 2018, **10**, 44722–44730.
- 46 T. S. Sreeremya, K. M. Thulasi, A. Krishnan and S. Ghosh, *Ind. Eng. Chem. Res.*, 2012, **51**, 318–326.
- 47 A. Dhall, A. Burns, J. Dowding, S. Das, S. Seal and W. Self, *Environ. Sci.: Nano*, 2017, **4**, 1742–1749.
- 48 F. Tan, Y. Zhang, J. Wang, J. Wei, Y. Cai and X. Qian, *J. Mass Spectrom.*, 2008, **43**, 628–632.
- 49 Y. Bai, Y. Li, Y. Li and L. Tian, *ACS Omega*, 2024, **9**, 8601–8614.
- 50 J. S. Sidhu, K. Rajendran, A. B. Mathew, T. Iqbal, D. K. Saini and D. Das, *Anal. Chem.*, 2023, **95**, 7594–7602.
- 51 S. Qian and H. Lin, *Anal. Chem.*, 2015, **87**, 5395–5400.
- 52 C. Uslu, S. Narin, Z. Demirsoy, H. B. Öksüz and G. Gülseren, *Int. J. Biol. Macromol.*, 2023, **233**, 123604.
- 53 E. Pütz, I. Tutzschky, H. Frerichs and W. Tremel, *Nanoscale*, 2023, **15**, 5209–5218.
- 54 H. Xiong, X. He, T. Lou and X. Bai, *Biochem. Eng. J.*, 2023, **195**, 108931.