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Supramolecular Regulation of Bioorthogonal Catalysis in Cells Using Nanoparticle-Embedded Transition Metal Catalysts

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Abstract

Bioorthogonal catalysis broadens the functional possibilities of intracellular chemistry. Effective delivery and regulation of synthetic catalytic systems in cells is challenging due to the complex intracellular environment and catalyst instability. Here, we report the fabrication of protein-sized bioorthogonal nanozymes through the encapsulation of hydrophobic transition metal catalysts into the monolayer of water-soluble gold nanoparticles. The activity of these catalysts can be reversibly controlled by binding a supramolecular cucurbit[7]uril 'gate-keeper' onto the monolayer surface, providing a biomimetic control mechanism that mimics the allosteric regulation of enzymes. The potential of this gated nanozyme for use in imaging and therapeutic applications was demonstrated through triggered cleavage of allylcarbamates for pro-fluorophore activation and propargyl groups for prodrug activation inside living cells.

TOC image

Regulation of bioorthogonal catalysis in living systems is challenging because of the complex intracellular environment. Now, the activity of protein-sized bioorthogonal nanozymes has been regulated by binding a supramolecular cucurbit[7]uril 'gate-keeper' onto the monolayer surface, providing controlled activation of pro-fluorophores and prodrugs inside living cells for imaging and therapeutic applications.

Author contributions

Additional information

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G.Y.T., Y.J., and V.M.R. conceived and designed the experiments. G.Y.T., Y.J., B.D., T.M., R.M., R.D., S.T.K., Y,-C.Y., B.Y., and S.H. performed the experiments. All authors analyzed and discussed the data. G.Y.T., Y.J., B.D., and V.M.R. co-wrote the paper. V.M.R. revised the paper.

G.Y.T. and Y.J. contributed equally to this work.

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Bioorthogonal chemistry^{1–3} is a promising strategy for the intracellular generation of molecules for therapeutic⁴ and imaging applications^{5,6} unattainable through naturally occurring bioprocesses^{7,8}. Transition metal catalysts (TMCs) are excellent candidates for use in bioorthogonal processes $9-11$, rapidly catalyzing transformations that cannot be performed via enzymatic processes^{12–15}. However, the application of TMC-mediated reactions in living cells is challenging due to issues of biocompatibility, water solubility, catalyst stability, and rapid efflux of catalysts from living cells 12,13 .

Loading of TMCs into nanomaterial scaffolds can be used to provide water solubility and a protective environment for TMCs. Bradley *et al.*¹³ and Unciti-Broceta *et al.*¹⁶ used palladium-catalyst loaded polystyrene beads to catalyze reactions such as Suzuki-Miyaura coupling and alkylcarbamate/N-propargyl cleavage inside and outside cells, respectively. The particles used in these studies, however, were far larger than normal proteins, creating potential interference in cellular processes. Additionally, these particles did not provide the capability of mimicking allosteric regulation of enzymes, a key component in cellular homeostasis¹⁷. The integration of biomimetic size and controlled response into a bioorthogonal catalysis platform would provide new avenues for both therapeutics and integrated biological/abiotic cellular systems.

We have developed a family of gold nanoparticles (AuNPs) based on \sim 2 nm core size^{18–20} that feature biomimetic size, possess diverse functional properties^{21,22}, and are efficiently transported into cells^{23,24}. We report here the use of this AuNP structural motif to encapsulate25 hydrophobic TMCs, providing **NP_Ru** (Fig. 1) or **NP_Pd**. The resulting nanozymes^{26–30} feature surface moieties that can be reversibly functionalized using hostguest chemistry31,32 to provide **NP_Ru_CB[7]** or **NP_Pd_CB[7]**. Complexation of the monolayer terminal functionalities by cucurbit[7]uril $(CB[7])^{33-35}$ in this system blocks access to the catalytic site, resulting in essentially complete inhibition of catalytic activity. The gatekeeper molecules $36,37$ can then be released from the AuNPs using competitive guests38, restoring catalytic activity (Fig. 1). The efficacy of this system was demonstrated in solution and in cells through two applications: 1) the gated generation of a fluorophore through deallylation of a non-fluorescent precursor and 2) the gated activation of a prodrug by cleaving the propargyl functionality that has been introduced to block the active side of original drug. Such gated control of catalysis has not been demonstrated in cells to date and is important because it allows the potential for multiple useful capabilities, such as switching on therapeutics at target tissues and regulation of activity to maintain homeostasis for longterm therapeutics. To date, supramolecular machines based on gating strategy were designed

to entrap the guest molecules in the pore reservoir of silica nanoparticles and their release was studied mostly in test tubes³⁹ or intracellularly³⁷ in a few studies however, no gated control of substrate activation was demonstrated inside cells.

Results and discussion

Design and synthesis of nanozymes

We used AuNPs with core diameters of \sim 2 nm as the scaffold for our catalysts, with the goal of creating protein-sized systems with functional monolayers. The monolayer coverage of the **NP** nanoreactor scaffold features three crucial components: 1) a hydrophobic alkane segment for catalyst encapsulation, 2) a tetra(ethylene glycol) unit to provide biocompatibility⁴⁰, and 3) a dimethylbenzylammonium group to impart water solubility and bind the nontoxic CB[7] gatekeeper^{41,42} (Fig. 1f).

We chose ruthenium-catalyzed deallylation^{43,44} as a model bioorthogonal process to regenerate the fluorescence from an allylcarbamate caged fluorophore in solution and inside cells. To this end, we immobilized $[Cp*Ru(cod)Cl]$ ($Cp* = pentamethylcyclopentadienyl$, cod = 1,5-cyclooctadiene) in the hydrophobic portion of the AuNP monolayer to provide **NP_Ru**. Transmission electron microscopy (TEM) images of the AuNPs before and after encapsulation of the catalyst indicated that no aggregation or decomposition of AuNP structure occurred after encapsulation (Supplementary Fig. S9), a result that was also confirmed by dynamic light scattering (Supplementary Fig. S10). The amount of ruthenium catalyst relative to AuNP was quantified using inductively coupled plasma mass spectrometry (ICP-MS, Supplementary Fig. S11)⁴⁵ of ¹⁰¹Ru relative to ¹⁹⁷Au, with 42 \pm 3 catalyst molecules encapsulated per AuNP.

Catalytic efficacy of NP_Ru nanozymes in solution

The catalytic efficacy of the nanozymes in solution was assessed using the allylcarbamate cleavage of $bis-N$, N' -allyloxycarbonyl rhodamine 110 (Fig. 1g). Fluorescence started to increase immediately for **NP_Ru** after the addition of substrate, while no significant change in fluorescence was observed for **NP_Ru_CB[7]** (Supplementary Fig. S12). After 24 h bright fluorescence was observed using the **NP_Ru** nanozymes (Fig. 2d). However, only minimal fluorescence was observed for the **NP_Ru_CB[7]**, which originated from the background of caged fluorophore as observed at 0 h (Supplementary Fig. S13). As expected no reaction occurred with the control particle **NP** that lacked embedded catalysts (Supplementary Fig. S14), as the [Cp*Ru(cod)Cl] was insoluble in water, preventing study of the catalyst alone. Additionally, fluorogenesis of **NP_Ru** and **NP_Ru_CB[7]** taken after 5 days confirmed the long term stability of the gated catalysis system in solution (Supplementary Fig. S15).

Supramolecular control of catalysis through CB[7] complexation

With catalytic efficacy of the nanozymes established, we next explored CB[7] complexation of the ligand headgroups to act as the gatekeepers in **NP_Ru_CB[7]**. Isothermal titration calorimetry (ITC)⁴⁶ indicated that 60 ± 5 CB[7] molecules bound per NP, with K_D = 11.3 \pm 2.4 μM (Supplementary Fig. S16). Complexation with CB[7] effectively shut down

catalysis in **NP_Ru_CB[7]**, as minimal fluorogenesis was observed with nanozymes after addition of CB[7] (Fig. 2d). Kinetic studies (Fig. 2a, c) verified this observation, indicating essentially complete inhibition of catalysis by CB[7] complexation. This inhibition was reversible: after the addition of 1-adamantylamine (ADA), a competitive guest molecule for CB[7], catalytic activity was completely restored (Fig. 2b, c). Control studies of the free catalyst in an acetone/water solution showed that catalyst efficiency was unaffected by either CB[7] or ADA (Supplementary Fig. S17), demonstrating that particle-CB[7] gating controls the catalytic process.

Kinetic analysis of the NP_Ru nanozymes using Lineweaver–Burk analysis

The analogy between controlled regulation of the nanozymes and their enzyme counterparts was explored through kinetic analysis of the nanozymes using Lineweaver–Burk analysis $(LBA)^{47,48}$. These studies indicate that CB[7] complexation results in competitive inhibition of reactor activity (Fig. 3 and Supplementary Fig. S18), presumably by blocking access to the [Cp*Ru(cod)Cl] "active site". Little change in activity was observed for CB[7]:NPs ratios greater than ~80:1, consistent with the ITC results (Supplementary Fig. S19 and S20). Likewise, the value of K_i from the LBA (11.4 \pm 4.3 μ M) is essentially identical to the affinity obtained by ITC (11.3 \pm 2.4 μ M). Taken together, these studies demonstrate the direct correlation between CB[7]-AuNP equilibrium binding processes and kinetic behavior of the nanozyme.

Catalytic efficacy of NP_Pd nanozymes in solution

To demonstrate the versatility of this catalysis platform, we chose a second catalyst: hydrophobic palladium catalyst: (1,1′- Bis(diphenylphosphino) ferrocene)palladium(II)dichloride. This palladium catalyst was encapsulated in the hydrophobic portion of the AuNP monolayer to provide **NP_Pd**. The amount of palladium catalyst relative to AuNP was quantified using ICP-MS of ^{106}Pd relative to ^{197}Au . It was calculated that 32 ± 1 catalyst molecules encapsulated per AuNP (Supplementary Fig. S21).

Similar to **NP_Ru** nanozyme, **NP_Pd** can also cleave the allylcarbamate of bis-N,N′ allyloxycarbonyl rhodamine 110^{13} . After 6 h, fluorescence was generated for the **NP_Pd** (Supplementary Fig. S22), however; **NP_Pd_CB[7]** showed only a slight background fluorescence of caged fluorophore (Supplementary Fig. S22). Kinetic studies for **NP_Pd** and **NP_Pd_CB[7]** indicated the inhibition of catalysis by CB[7] complexation. Followed by the addition of ADA, the catalytic activity of nanozyme was restored (Supplementary Fig. S23). These findings supported that gated catalysis is efficiently working for different catalyst systems.

Supramolecular regulation of catalysis inside living cells

Having characterized the activity and gating of nanozyme catalysis in solution, we next studied the intracellular behavior of these nanozymes using HeLa cells. The cellular uptake of the nanozymes was quantified by tracking [197Au] using ICP-MS, with **NP_Ru_CB[7]** particles demonstrating a slightly more efficient (1.7-fold greater) uptake than uncomplexed **NP-Ru** (Supplementary Fig. S24). Significantly, no toxicity was observed at the concentrations used for our studies with either **NP** or **NP_Ru** (Supplementary Fig. S25).

We then probed the catalytic activity of the nanozymes inside the living cells. HeLa cells were incubated with the nanozyme for 24 h in serum-containing media and then washed multiple times to remove adsorbed particles on the cell surface. Fresh media containing the substrate was added, followed by 24 h incubation and washing. As shown in Fig. 4a, flow cytometry indicated that there was a significant increase in fluorescence with **NP_Ru** relative to the control **NP**. Confocal microscopy (Fig. 4d) showed that the cells treated with **NP_Ru** had bright punctate fluorescence. This fluorescence co-localized with LysoTracker® (Supplementary Fig. S26), indicating that the deallylation reaction occurred in the endosomes and cleaved fluorophore stayed in the endosomes. This outcome is expected given the endosomal uptake pathway^{49,50} previously observed for similar nanoparticles, coupled with the limited membrane permeability of the cleaved dye⁵¹.

We next investigated the intracellular gating of the nanozyme by CB[7] complexation. Flow cytometry showed that the **NP_Ru_CB[7]** particles were completely inhibited, indicating intracellular stability of the complexes (Fig. 4a, e). Treatment with ADA (400 μM) restored nanozyme activity, with even higher activity observed in the cells after treatment with ADA than was observed with **NP_Ru** (Fig. 4b). This increased catalytic efficiency is echoed in the micrograph (Fig. 4f) and potentially arises from enhanced protection of the catalyst by the CB[7] coverage in serum and inside the cell.

We further tested the control of gated catalysis through the treatment of cells with free CB[7]. First, HeLa cells were incubated with the **NP_Ru** for 24 h and then after multiple washings, free CB[7] in serum-containing media was treated to cells for 24 h before the addition of the substrate. A significant decrease in the fluorescence was observed indicating the complexation of CB[7] with **NP_Ru** and blocking access to the catalytic site (Supplementary Fig. S27).

Palladium catalyst embedded nanozymes were likewise used for intracellular controlled activation of the caged fluorophore. **NP_Pd** effectively performed the intracellular catalysis, while as expected the gated **NP_Pd_CB[7]** did not. ADA addition into cells that were previously incubated with **NP_Pd_CB[7]** resulted in the recovery of intracellular catalysis (Supplementary Fig. S28).

Intracellular prodrug activation using NP_Pd nanozymes

Bioorthogonal activation of prodrugs using TMC-loaded nanomaterials was recently demonstrated by Unciti-Broceta and co-workers¹⁶ Although prodrug activation was successfully achieved, no gated control over this activation was shown. We have bioorthogonally demonstrated gated-activation of a prodrug inside cells using **NP_Pd/ NP_Pd_CB[7].** 5-fluorouracil (5FU) is a chemotherapeutic drug used in cancer treatment including breast, stomach, pancreatic, and skin cancers. Although it has an established history in cancer treatment, it shows toxic side effects due to its limited safety profile⁵². These side effects can be eliminated or at least minimized if a prodrug strategy coupled with gated-intracellular catalysis were to be available. By this gated catalysis, activation of prodrugs will be achieved on demand upon reaching to site of action thus eliminating the off target effects.

5FU can be converted into a prodrug via functionalization on its N1 position, as modification at this position will block the activity of 5FU and render it non-toxic¹⁶. We introduced a propargyl moiety to turn 5FU into an inactive prodrug 'pro-5FU' (Fig. 5a). First, through palladium mediated chemocatalysis the propargyl masking unit is intracellularly cleaved to yield 5FU followed by the enzymatic reactions to convert 5FU into the cytotoxic nucleotidic fluorouridine monophosphate via functionalization at the N1 position53. These active metabolites work through misincorporation into RNA and DNA molecules and irreversibly inhibit the nucleotide synthetic enzyme thymidylate synthase to disrupt cell functions and induce cytotoxicity. Overall, this process is an example of the integration of chemocatalysis and enzymatic biochemistry in a one-pot chemical sequence^{54,55}.

Cleavage of propargyl group on pro-5FU (1 mM) in the presence of **NP_Pd** (100 nM) was monitored using matrix-assisted laser desorption/ionization (MALDI)-MS. Most of the pro-5FU was converted into 5FU after 48 h (Supplementary Fig. S29). After confirming the cleavage, the toxicity profile of 5FU and pro-5FU were investigated by performing a cell viability assay. Cells were treated with various concentrations from 10 nM to 1 mM. While 5-FU showed toxicity as concentration increased, pro-5FU retained high cellular viability at all concentrations studied (Fig. 5b).

Cell viability studies for prodrug system in the presence of nanozymes were carried out using HeLa cells in a 96 well plate. Cells were first incubated with **NP_Pd** or **NP_Pd_CB[7]** at a concentration of 100 nM in serum-containing media for 24 h. Then, after multiple washings, cells were incubated with different concentrations of pro-5FU (0, 0.05, 0.1, 0.25, 0.5 and 1 mM) while some of the cells that were incubated with **NP_Pd_CB[7]**, were treated with pro-5FU and ADA at the same time. Cells that were incubated with **NP_Pd** and **NP_Pd_CB[7]** + ADA showed elevated toxicity at higher concentration of pro-5FU while **NP_Pd_CB[7]** retained ~100 % cell viability even at higher concentration of pro-5FU (Fig. 5c). As expected, pro-5FU was not toxic at any concentrations used. Likewise, **NP_Pd**, **NP_Pd_CB[7]**, and **NP_Pd_CB[7]** + ADA at zero pro-drug concentration were not toxic indicating toxicity was coming from the intracellular conversion of pro-5FU into 5FU using gated-catalysis but not the nanozyme itself.

In conclusion, we have described the fabrication of a AuNP based, bioorthogonal nanozyme that uses transition metal catalysis to effect transformations of imaging and therapeutic relevance without biological counterparts. These catalysts were built upon a platform featuring biomimetic size and surface functionality, making them attractive components for both *in vitro* and *in vivo* applications. The ability to control the activity of these nanozymes through host-guest interactions of CB[7] molecules with the benzyl headgroup of the AuNP ligands likewise provides an efficient and reversible means of regulating catalysis. This bioorthogonal catalysis can be employed not only in therapeutic applications such as the activation of prodrugs at the site of action but also in treating non-cancerous chronic diseases where the goal is not to kill the cell but to kill an infective agent or restore a malfunctioning pathway. This gated platform enables the introduction of an inactive reservoir of a bioorthogonal catalyst that would remain dormant between successive administrations of the substrate, thus reducing potential interference of the catalyst with cell

constituents and protecting the metal from poisoning, release, systemic distribution and/or clearance. Furthermore, our protein-sized system demonstrates biomimetic behavior and yet performs totally abiotic chemistry that can be controlled intracellularly through a very simple host-guest feature. This system integrates biomimetic and bioorthogonal design elements to provide a new platform for imaging and therapeutic applications as well as pharmacological treatments integrating biological activity with man-made synthetic tools.

Methods

Synthesis of the ligands and the AuNPs (Fig. 1f) for the nanozyme are described in the Supplementary Information (Supplementary Fig. S1–S8). Details of the particle characterization methods, catalyst encapsulation process, ICP-MS sample preparation, kinetic studies and additional cell culture experiments can be found in the Supplementary Information.

Catalyst encapsulation into the monolayer of AuNPs

The catalyst ($[CP^*Ru(cod)Cl]$ or 1,1[']-Bis(diphenylphosphino) ferrocene)palladium(II)dichloride) and the AuNP were dissolved in an acetone/water mixture and then the acetone was slowly removed by evaporation. During the evaporation the catalyst was encapsulated in the particle monolayer, with excess catalyst precipitating and removed by filtration. The AuNPs were purified by multiple filtrations and dialyzed against water to remove free catalysts. The amount of encapsulated catalyst was measured by ICP-MS by tracking ¹⁰¹Ru relative to ¹⁹⁷Au for NP_Ru and ¹⁰⁶Pd relative to ¹⁹⁷Au for NP_Pd.

Kinetic studies of nanozymes using Lineweaver–Burk analysis

In the titration of the reaction rate of the nanozyme using the CB[7], 400 nM of the AuNPs, 80 μM of the substrate, and set molar ratios of CB[7]s were used in sodium phosphate buffer (5 mM, pH 7.4). For the Lineweaver-Burk analysis, 400 nM of the AuNPs, 2, 4, 20, 40, and 80 μM of the substrate, and 0, 4, 16, and 80 μM of CB[7] were used. The concentration of the products was calculated based on the fluorescence standard curve of rhodamine 110 (Supplementary Fig. S30).

Catalytic activity of the nanozymes inside living cells

HeLa cells were grown in a cell culture flask using low-glucose Dulbecco's modified Eagle medium supplemented with 10% fetal bovine serum at 37°C in a humidified atmosphere of 5% CO2. For flow cytometry and confocal analysis, HeLa cells were seeded at 20,000 and 80,000 cells in 0.5 mL per well in 24-well plates and 1 ml in confocal dish 24 h prior to the experiment. During the experiment old media were replaced by 200 nM of the nanozymes and the nanozyme-bound CB[7] in serum-containing media and the cells were incubated for 24 h and washed with PBS buffer three times. 100 μM of the substrates were then added to the cells and incubated for 24 h. The cells were then washed with PBS three times. In the case of ADA treatment, cells were treated with 0.4 mM ADA solution (0.5 mL) for 24 h. For the cytometry, a Becton Dickinson LSR cytometer was used. A total of 10,000 events per sample were analyzed. A530/30 bandpass filter (FITC) was used for rhodamine 110.

Confocal microscopy images were obtained on a Zeiss LSM 510 Meta microscope by using a 63× objective. The settings of the confocal microscope were as follows unless otherwise specified: green channel: λ_{ex} =488 nm and λ_{em} =BP 505–530 nm; red channel: λ_{ex} =543 nm and λ_{em} =LP 650 nm. Emission filters: BP=band pass, LP=high pass. LysoTracker® Red DND-99 was obtained from Invitrogen. 100 nM of LysoTracker® was incubated with cells for 30 min prior to the microscopy experiments.

Cytotoxicity measurements of 5FU versus pro-5FU

HeLa cells were seeded at 10,000 in 0.2 mL per well in 96-well plates 24 h prior to the experiment. Cells were washed with PBS buffer and then treated with 11 different concentrations ranging from 1 nM to 1 mM of 5FU or pro-5FU in triplicate. After 72 h of incubation, the cells were then completely washed off with PBS buffer three times and 10% Alamar Blue in serum containing media was added to each well (220 μL) and further incubated at 37 °C for 4 h. The cell viability was then determined by measuring the fluorescence intensity at 570 nm using a SpectraMax M5 microplate spectrophotometer.

Intracellular prodrug activation

HeLa cells were seeded at 10,000 in 0.2 mL per well in 96-well plates 24 h prior to the experiment. Cells were washed off and incubated with **NP_Pd** (100 nM) or **NP_Pd_CB[7]** (100 nM) in 10% serum containing media. After 24 h, cell were washed with PBS buffer three times and treated with pro-5FU at a concentration of 0, 0.05, 0.1, 0.25, 0.5 and 1 mM for 48 h. Some cells treated with **NP_Pd_CB[7]** were incubated with both pro-5FU and ADA (0.05 mM). After 48 h of incubation, the cells were then completely washed off and 10% Alamar Blue in serum containing media was added to each well (220 μL) and further incubated at 37 °C for 4 h. The cell viability was then determined by measuring the fluorescence intensity at 570 nm using a SpectraMax M5 microplate spectrophotometer.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Figure 1. Bioorthogonal nanozyme design and supramolecular regulation of intracellular catalysis

a, AuNP, catalyst-embedded AuNP, and CB[7]-capped catalyst-embedded AuNPs used in study. **b**, Endosomal uptake of nanozymes. **c**, Intracellular catalysis with **NP_Ru** converting substrate into product. **d**, CB[7] complexation with the ligand headgroup to provide **NP_Ru_CB[7]** inhibits catalyst activity. **e**, Nanozyme activity is restored through addition of the competitive guest 1-adamantylamine (ADA). **f**, Structures of the **NP** platform with the surface ligand bearing a dimethylbenzylammonium headgroup, CB[7] gatekeeper, and ADA, a competitive guest molecule for CB[7] binding. **f**, Structures of the pro-fluorescent substrate (rhodamine 110 derivative), fluorescent product (rhodamine 110) obtained after catalysis, and embedded catalyst for allylcarbamate cleavage.

Figure 2. Catalytic activity of nanozymes in solution

a, Fluorescence was generated by **NP_Ru** after the cleavage of profluorophore bis-Allocrhodamine 110, while **NP_Ru_CB[7]** showed no significant change. **b**, After adding ADA, catalytic activity of **NP_Ru_CB[7]** was restored and no significant effect was observed for the activity of **NP_Ru**. **c**, The reaction rates of **NP_Ru** and **NP_Ru_CB[7]** before and after adding ADA showing catalytic activity for **NP_Ru_CB[7]** was fully recovered after the addition of ADA. The reaction rate experiments were performed in triplicate. Error bars represent standard deviations of these measurements. **d**, Photo of the reaction mixtures in water with **NP_Ru** and **NP_Ru_CB[7**] under UV light.

Figure 3. Lineweaver-Burk plot showing competitive binding of CB[7] to nanozyme Kinetic studies of **NP_Ru** and **NP_Ru_CB[7]** in sodium phosphate buffer (5 mM, pH 7.4) indicate that CB[7] inhibits catalyst activity through a competitive inhibition mechanism, with CB[7] affinity and stoichiometry consistent with ITC binding studies. Numbers calculated on a per particle basis. Kinetic experiments with each nanozyme were repeated in triplicate. Error bars represent standard deviations of these measurements.

Figure 5. Prodrug activation in living cells using supramolecularly controlled nanozymes a, Structures of pro-5FU, 5FU and the palladium catalyst used for prodrug activation. **b**, Viability of cells treated with 5FU and pro-5FU at various concentrations, showing a nice therapeutic window was obtained between 5FU and pro-5FU. **c, NP_Pd** and ADA treated **NP_Pd_CB[7]** showed increasing intracellular toxicity as a result of more conversion of prodrug into 5FU drug at higher pro-5FU concentrations, while **NP_Pd_CB[7]** did not show any toxicity at any prodrug concentration used due to the blocking catalysis. Also, only prodrug, **NP_Pd**, **NP_Pd_CB[7]**, and **NP_Pd_CB[7]** + ADA did not cause any toxicity into the system at zero prodrug concentration. Cell viability experiments were performed in triplicate. Error bars represent standard deviations of these measurements.