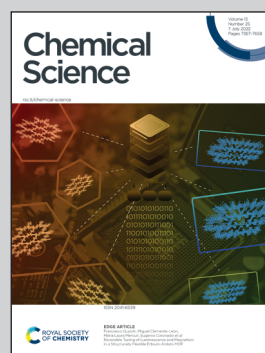


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Biocatalytic cascades and intercommunicated biocatalytic cascades in microcapsule systems

Biomolecule-loaded nucleic acid-functionalized hydrogel microcapsule are introduced as cell-like containments emulating native cells. Intracapsular biocatalytic cascades and intercapsular biocatalytic cascades and switchable biocatalytic cascades are demonstrated by the systems. Substantially enhanced biocatalytic transformations proceed in the cell-like containment (protocells), as compared to a homogeneous phase.

As featured in:



See Itamar Willner *et al.*,
Chem. Sci., 2022, **13**, 7437.

Cite this: *Chem. Sci.*, 2022, 13, 7437

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Biocatalytic cascades and intercommunicated biocatalytic cascades in microcapsule systems†

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Biomolecule-loaded nucleic acid-functionalized carboxymethyl cellulose hydrogel-stabilized microcapsules (diameter ca. 2 μm) are introduced as cell-like containments. The microcapsules are loaded with two DNA tetrahedra, T_1 and T_2 , functionalized with guanosine-rich G-quadruplex subunits, and/or with native enzymes (glucose oxidase, GOx, and/or β -galactosidase, β -gal). In the presence of K^+ -ions and hemin, the T_1/T_2 tetrahedra constituents, loaded in the microcapsules, assemble into a hemin/G-quadruplex bridged tetrahedra dimer DNAzyme catalyzing the oxidation of Amplex Red to Resorufin by generating H_2O_2 . In the presence of co-loaded GOx or GOx/ β -gal, the GOx// T_1/T_2 hemin/G-quadruplex cascade catalyzing the glucose-mediated oxidation of Amplex Red to Resorufin, and the three-biocatalysts cascade consisting of β -gal//GOx//hemin/G-quadruplex bridged T_1/T_2 catalyzing the lactose-driven oxidation of Amplex Red to Resorufin proceed in the microcapsules. Enhanced biocatalytic transformations in the microcapsules, as compared to the performance of the reactions in a homogeneous phase, are observed, due to the proximity of the biocatalysts in a confined volume. As the synthetic methodology to prepare the microcapsules yields boundaries functionalized with complementary nucleic acid tethers, the dynamic association of different microcapsules, loaded selectively with biomolecular catalysts, proceeds. The dynamic dimerization of GOx-loaded microcapsules and hemin/G-quadruplex bridged T_1/T_2 DNAzyme-loaded microcapsules yields effective intercommunicated microcapsules driving the GOx//hemin/G-quadruplex bridged T_1/T_2 DNAzyme cascade. In addition, the dynamic dimerization of GOx-loaded microcapsules with β -gal//hemin/G-quadruplex bridged T_1/T_2 -loaded microcapsules enables the bi-directional intercommunicated operation of the lactose-stimulated three catalysts β -gal//GOx//hemin/G-quadruplex bridged T_1/T_2 DNAzyme cascade. The guided separation and formation of dynamic supramolecular dimer microcapsular containments, and the dictated switchable operation of intercommunicated biocatalytic cascades are demonstrated.

Received 17th March 2022
Accepted 29th April 2022DOI: 10.1039/d2sc01542k
rsc.li/chemical-science

Introduction

The construction of artificial cell-like assemblies emulating the complexity of native cells, cell-like containments, is a scientific “holy grail” attracting continuous research efforts in the area of Systems Chemistry and Systems Biology.¹ Development of cellular containment systems addresses variety of challenging sub-goals that need to be integrated into the artificial cell-like

machinery. These includes: (i) The development of cell-like containments that allow the immobilization and operation of cell-mimicking reactions and functionalities. (ii) The integration of chemically engineered ensembles into the cell-like containments capable of inducing dictated catalytic transformations (metabolism), ability to sense and response to environmental triggers, and dynamically adapt the structure, composition and chemical performance in response to external stimuli. (iii) The influx of chemical agents and the efflux of metabolites from the cell-like containments, and the guided operation of chemical transformations in the artificial containments using chemical fuels, auxiliary energy sources, such as light or redox potential, should be feasible. (iv) Appropriate chemical networks and machineries allowing motility, differentiation, exchange constituents and information transfer between the artificial cells should be integrated into the artificial containments, allowing the operation of complex catalytic cascades. Needless to state, the availability of cell-like

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† Electronic supplementary information (ESI) available. See <https://doi.org/10.1039/d2sc01542k>

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containments combining all of these features (or more) characterizing native cells are non-existent, yet functional assemblies mimicking some of those functions were realized, and several excellent review articles summarized the advances in the field.²

The design of organized biocatalytic cascades on supramolecular scaffolds or biocatalytic cascades operating in confined micro- or nano-environments attracts substantial research efforts aiming to emulate biocatalytic networks in nature.³ The spatial proximity between the enzymes in these organized media leads to effective channeling of the product of one enzyme, acting as substrate for the neighboring enzyme, thereby, overcoming diffusional limitations and producing high local concentrations for effective intercommunication between the biocatalysts.⁴ For example, the assembly of spatially organized enzymes on DNA frameworks, such as strips,⁵ bundles⁶ or origami tiles⁷ or the organization of enzymes, on peptide ligands⁸ or protein scaffolds⁹ led to structurally ordered assemblies guiding efficient biocatalytic cascades. Similarly, the integration of multi-enzyme systems in confined environments such as microdroplets,¹⁰ polymersomes,¹¹ or metal-organic-framework nanoparticles¹² generated engineered containments of spatially proximate assemblies of biocatalysts for efficient operation of biocatalytic cascades. Such organized assemblies of biocatalytically active cascades were suggested as functional systems for effective sensing,¹³ bioreactors for driving efficient catalytic transformations and the synthesis of drugs.¹⁴

Different cell-like containments were developed including liposomes,¹⁵ polymersomes,¹⁶ proteinosomes,¹⁷ colloidosomes,¹⁸ dendrosomes,¹⁹ coacervate microdroplets,²⁰ and were suggested as models for cells compartments. Enzymes were integrated in cell-like containments and the advantages of driving biocatalytic cascades in confined nano/micro-environments were demonstrated.²¹ Cell-like containments demonstrating motility,²² chemical signaling²³ proliferation,²⁴ and passive and active transport²⁵ were reported. Genetic circuits²⁶ and RNA catalytic networks were embedded in cell-like containments²⁷ and DNA-guided communication between cell-like containments²⁸ and gene expression²⁹ were realized. Compartmentalization of cell-like containments and control over cascaded reactions in the compartmentalized assemblies were achieved.³⁰ Also, adaptive temporal pH-responsive polymersome nanoreactor structures³¹ or transient chemical fuel-triggered microgel systems³² were reported, and compartmentalization of native cells was emulated by cell-like containment model systems and these revealed increased functional complexity,³³ and control over cascaded reactions.³⁴

The information encoded in the base sequence of nucleic acids provides structural and functional feature that could be utilized to design active constituents in artificial cells. For example, the base pairing guided formation of duplex DNAs provides means to construct programmed duplex structures that can be dynamically interchanged by strand displacement processes controlled by the relative stabilities of the duplexes.³⁵ In addition, sequence dictated dynamic reconfiguration of nucleic acid strands provides versatile means to develop

switchable DNA structures, *e.g.* G-quadruplexes,³⁶ triplexes³⁷ or i-motif assemblies,³⁸ and pre-designed nucleic acid structures, such as hairpin DNAs, or circular DNAs, introduced functional motives to operate dynamic DNA machineries, such as the hybridization chain reaction³⁹ or the rolling circle amplification.⁴⁰ Moreover, the sequence specific-nucleic acid constructs reveal selective recognition properties (aptamers)⁴¹ or catalytic properties (DNAzymes or nucleozymes),⁴² such as the hemin/G-quadruplex, horseradish peroxidase mimicking DNAzyme⁴³ or the metal-ion-cofactor dependent DNAzymes.⁴⁴ The nucleic acid biopolymers could substitute proteins or native enzymes, or alternatively be coupled to native enzymes, as functional constituents in artificial cells. Furthermore, the guided hybridization properties of nucleic acids led to the supramolecular assembly of 2D and 3D DNA structures, such as Y-shaped,⁴⁵ crossover-junctions⁴⁶ or origami frameworks.⁴⁷ Within this context, DNA tetrahedra 3D nanostructures revealing tunable sizes and high stabilities attract substantial recent research efforts.⁴⁸ The DNA tetrahedra nanostructures are formed upon the assembly of four appropriately sequence-engineered strands,⁴⁹ and reveal size tunability⁵⁰ and structural stability.⁵¹ By the conjugation of nucleic acid tethers to the corners of the tetrahedra or by encoding sequence specific domains into the edges of the tetrahedra structures, the functionalization of the nanostructures with nanoparticles,⁵² proteins⁵³ and functional nucleic acids (DNAzymes, aptamers)⁵⁴ were demonstrated. Various applications of DNA tetrahedra for sensing,⁵⁵ imaging,⁵⁶ drug carrying,⁵⁷ and scaffolds for the organization of chiroplasmonic structures⁵⁸ were reported. The dimensions of the DNA tetrahedra, 2.4–12.6 nm, are controlled by the length of the strands comprising the structures. As the dimensions of DNA tetrahedra are comparable to the sizes of low-molecular-weight proteins, they were suggested as structural modules that emulate proteins.⁵⁹

In the present study, we introduce DNA-based hydrogel microcapsules as model systems for cell-like containments. The synthesis of DNA-based microcapsules and particularly the preparation of stimuli-responsive drug-loaded microcapsules has been a subject of extensive research in our laboratory.⁶⁰ We have demonstrated the design of pH,⁶¹ aptamer,⁶² light⁶³ and microRNA responsive⁶⁴ microcapsules and applied the systems for programmed release of anti-cancer drugs or insulin.⁶⁵ The study will make use of the following properties of the microcapsules to introduce the cell mimicking functions: (i) The method to prepare the microcapsules allows the programmed loading of the containment with pre-designed constituents. (ii) The methodology to synthesize the microcapsule introduces surface functionalities that allow the dynamic intercommunication between microcapsules. (iii) The hydrogel coating membrane reveals cell-like permeation and efflux properties. The resulting DNA crosslinked hydrogel coating reveals in/out permeation and efflux of low molecular weight agents, *ca.* < 5 kDa,⁶⁵ yet higher molecular weight substance is non-permeable acrossing the membrane boundaries and reveal confined stabilities for at least three days. These properties enable the control over the catalytic transformations within the microcapsules. (iv) The introduction of catalytic DNA tetrahedra



nanostructures and native enzyme into the microcapsule containments will be presented. The intracapsular DNA tetrahedra/enzyme cascades and the dynamic intercommunication of microcapsules loaded with catalytic tetrahedra/enzyme constituents and the guided operation of catalytic cascades will be introduced. (v) The advantages of dynamic intercommunication of cell-like assemblies driving biocatalytic cascades over analog biocatalytic cascades in aqueous environments are presented.

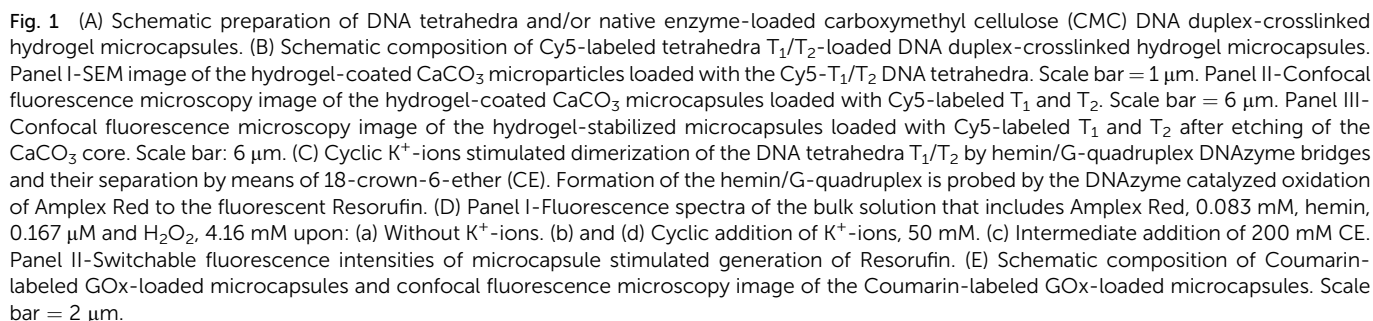
Results and discussion

The preparation of the loaded hydrogel microcapsules followed a synthetic path developed in our laboratory and is exemplified with the loading of a DNA tetrahedra T_1 and/or a native enzyme, E, Fig. 1 (A). CaCO_3 microparticles were impregnated with the DNA tetrahedra T_1 and/or the enzyme and acted as template for the construction of the microcapsules. The T_1 /E-impregnated microparticles were coated with polyallylamine hydrochloride (PAH) and the polymer coating was modified with the promoter nucleic acid strand (1). The (1)-functionalized particles were interacted with the carboxymethyl cellulose (CMC) polymer chains P_1 and P_2 modified with the hairpins H_1 and the conjugate (2)/ H_2 . The hairpin H_1 and H_2 were pre-engineered to allow the (1)-triggered opening of H_1 followed by the opening of hairpin H_2 by the single strand stem domain of opened H_1 , and subsequently, opened stem domain of H_2 hybridizes with hairpin H_1 . That is, the interaction of the (1)-modified particles with the two polymers induces the cross-opening of the hairpins H_1/H_2 and crosslinking of the polymer chains by the duplex H_1/H_2 bridges while forming a hydrogel coating on the particles. Formation of the hydrogel coating is terminated when the deposition of the polymer chains on the core particles was completed. Note, however, that the cross-opening of the hairpin constituents associated with the coating leads to surface-bound tether functionalities p and q that correspond to unused open hairpin domains of H_1 and H_2 , respectively. (Note that hairpin H_2 was hybridized to the strand (2) associated with P_2 in order to follow directionality constrains for cross-opening of the two hairpins). The resulting hydrogel coated microparticles were etched with EDTA to dissolve the CaCO_3 cores, resulting in the T_1 /E-loaded microcapsules, acting as the cell-model containment units in the present study. The resulting loaded hydrogel microcapsules reveal the following features that will be used in the present study: (i) Within the preparation procedure, the microcapsules can be loaded with mixtures of different tetrahedra, mixtures of different enzymes or mixtures of DNA tetrahedra and enzymes. (ii) As will be demonstrated, the DNA tetrahedra structures retain their structural features after the etching procedure. Also, the enzyme encapsulated in the microcapsules retain over >80% of their native activities, after the etching process. (iii) The tethers p and q associated with the hydrogel coating exhibit inherent complementary and inter-tether recognition features. As a result, the equilibrated dimerization (or oligomerization of the microcapsules is possible, or alternatively the equilibrated dimerization of microcapsules containing different loads is feasible). (iv) By

appreciate labeling of the loads with fluorophores, the stability and confinement of the loads within the microcapsules can be evaluated. (v) Specifically, we apply two different tetrahedra nanostructures, T_1 and T_2 , modified with G-quadruplex subunits (x) and (y) that self-assemble, in the presence of K^+ -ions, into G-quadruplex-bridged dimer constituents. In the presence of hemin, the association of the hemin ligand to the G-quadruplex yields the hemin/G-quadruplex DNAzyme-crosslinked tetrahedra T_1/T_2 dimer.

For the characterization of the loaded microcapsules, we performed a series of background experiments: (i) We loaded the microcapsules with the Cy5-labeled tetrahedra T_1 and non-labeled T_2 , Fig. 1(B), and probed the confinement of the tetrahedra to microcapsule containment. Fig. 1(B), panel I, depicts the SEM image of the Cy5-labeled tetrahedra T_1 -loaded microparticles, and the confocal fluorescence microscopy images of the resulting hydrogel-coated Cy5-modified T_1 loaded microparticles before etching, panel II, and of the Cy5-modified T_1 -loaded microcapsules after etching, panel III. Microcapsules revealing an average diameter corresponding to 2–3 μm are formed. (ii) To validate the intact structure of the tetrahedra loads in microcapsular containment after the synthesis and etching of the core CaCO_3 microcapsules, we performed a control experiment where the CaCO_3 particle were impregnated with the T_1 -tetrahedra and subsequently etched with EDTA, at similar condition used for the preparation of the microcapsules. The resulting solution containing the tetrahedra was analyzed by electrophoresis and compared to a standard tetrahedra solution. The result and accompanying discussion are presented in Fig. S1.† Pure, intact tetrahedra structure were observed suggesting that the subsequent etching of the core CaCO_3 have no perturbing effect on the tetrahedra nanostructures. (For further characterization of the tetrahedra structures by electrophoretic separation, see ESI, Fig. S2,† and the characterization of the monomer tetrahedra or dimer tetrahedra nanostructures by atomic force microscopy (AFM) imaging, see ESI, Fig. S3†). (iii) The resulting tetrahedra-loaded microcapsules are stable for at least three days, Fig. S4.† No leakage of the tetrahedra from the microcapsule containments could be detected within this time-interval. (iv) The loading of the tetrahedra T_1 and T_2 in the microcapsules was evaluated to be 0.1 μM and 0.09 μM , respectively, by separate labeling T_1 with Cy5 and T_2 with Cy3 in the loaded mixture of T_1/T_2 and by using appropriate calibration curves of Cy5-labeled T_1 and Cy3-labeled T_2 (For details, see ESI Fig. S5†). (v) The two tetrahedra structures, T_1/T_2 , were loaded in the microcapsular containments, treated with K^+ -ions, in presence of hemin as cofactor, to probe the formation of the hemin/G-quadruplex crosslinked tetrahedra dimer structure and its catalytic activity to stimulate the catalyzed oxidation of Amplex Red to the fluorescent Resorufin product, Fig. 1(C). The results in Fig. 1(D), panel I demonstrate the permeation of K^+ -ions and hemin into the T_1/T_2 loaded microcapsules, and the formation of the hemin/G-quadruplex DNAzyme in the presence of Amplex Red and H_2O_2 , results in the activation of the hemin/G-quadruplex DNAzyme that catalyzes the oxidation of Amplex Red to Resorufin. Furthermore, Fig. 1(D), panel II, demonstrates





appropriate fluorophores, the fluorophore-functionalized enzyme-loaded microcapsules allow the fluorescence imaging of the resulting biocatalytic capsules. For example, Fig. 1(E) depicts the resulting coumarin-modified glucose oxidase (GOx)-loaded microparticles and the confocal fluorescence microscopy image of the microcapsules. Using an appropriate calibration curve relating the fluorescence intensities to the concentration of the modified enzyme and knowing the number of microcapsules, the loading of the labeled enzyme in the microcapsules is evaluated. (For the specific system, a loading corresponding to 0.026 nM coumarin-labeled GOx per microcapsule was evaluated). Furthermore, by the integration of a non-modified enzyme in the microcapsules, and assuming

a similar loading degree to that of the labeled enzymes, and the activity of the enzyme entrapped in the microcapsules was followed, thus allowing the evaluation of the respective enzyme activity after the preparation and etching of the microcapsules. Along the study, mixtures of the catalytic DNA tetrahedra and the respective enzymes will be integrated in the microcapsular containments to stimulate catalytic cascade and inter-communicated catalytic cascades. The methods described above will be used to characterize different assemblies.

The first system to be addressed is displayed in Fig. 2. The two tetrahedra T_1/T_2 and GOx were loaded in the microcapsules. The loading of T_1/T_2 and GOx in the microcapsules were evaluated by the alternate loading of the capsules with mixtures that include the fluorophore Cy5-labeled T_1 , Cy5-labeled T_2 , and coumarine-labeled GOx to be 0.063 μM , 0.07 μM and 0.12 μM , respectively. (The calibration curve of coumarin-labeled GOx is shown in Fig. S9†). Treatment of the T_1/T_2 and GOx-loaded microcapsules with K^+ -ions and hemin resulted in the formation of the hemin/G-quadruplex bridged T_1/T_2 DNAzyme in the capsules. The subsequent treatment of the capsules with glucose and Amplex Red resulted in the intracapsular operation of the GOx/hemin/G-quadruplex cascade where GOx catalyzed the aerobic oxidation of glucose to gluconic acid and H_2O_2 , and

the resulting H_2O_2 mediated the hemin/G-quadruplex catalyzed oxidation of Amplex Red to the fluorescent product Resorufin. The biocatalytic cascade was probed by following the time-dependent formation of Resorufin in the microcapsules, Fig. 2(B), curve (i). Fig. 2(B), curve (ii), shows the time-dependent fluorescence intensities of the same system upon excluding K^+ -ions. No fluorescence changes are observed indicating that the K^+ -ions stabilized formation of the hemin/G-quadruplex DNAzyme is essential for operating the DNAzyme cascade in the microcapsules. Fig. S10† shows the fluorescence spectra of Resorufin at different time-intervals upon treatment of the cell-like microcapsules with different concentrations of glucose. As the concentration of glucose increases, the biocatalytic cascade is intensified. In addition, the progress of the biocatalytic cascade in the microcapsular heterogeneous system was compared to the operation of the biocatalysts hemin/G-quadruplex bridged T_1/T_2 and GOx in a homogeneous buffer solution that included the catalysts at bulk concentrations identical to the concentrations of the catalysts in the microcapsules. (All other constituents, e.g. K^+ -ions, hemin, Amplex Red and glucose were identical to those present in the microcapsular system). Fig. 2(C), curve (ii), presents the time-dependent formation of Resorufin in the homogeneous buffer solution, in comparison to the formation of Resorufin in the microcapsular system, curve (i). The biocatalytic cascade in the microcapsular system is 6-fold enhanced as compared to the homogeneous system. This is attributed to generation of the biocatalyst cascade in the confined volume of the microcapsules. That is, the GOx-biocatalyzed formation of H_2O_2 in spatial proximity to the hemin/G-quadruplex DNAzyme that leads to the effective channeling of H_2O_2 to the hemin/G-quadruplex DNAzyme. Similar rate enhancement of biocatalysts cascades in confined microenvironments were previously demonstrated,³ for example, by the spatial proximate organization of two enzymes on DNA templates,^{5,6} DNA origami structures,⁷ microdroplets¹⁰ or other cell-like containments.¹¹ The confinement effect of the microcapsules on enhancing biocatalytic cascades was further demonstrated in a three biocatalyst-loaded microcapsule containment assembly. In this system, the hemin/G-quadruplex bridged T_1/T_2 horseradish peroxidase mimicking DNAzyme was coupled to GOx and β -galactosidase, β -gal, using lactose as the parent substrate. In this system, Fig. S11† (A), β -gal hydrolyses lactose to glucose and galactose, the resulting glucose is aerobically oxidized gluconic acid and H_2O_2 , and the resulting H_2O_2 acts as intracapsular oxidant for the hemin/G-quadruplex bridged T_1/T_2 DNAzyme-catalyzed oxidation of Amplex Red to Resorufin. The characterization of this three biocatalysts-loaded microcapsule system and the results corresponding to the operation of the three catalysts are presented and discussed in detail in Fig. S12–S15.† As for the two-catalysts-loaded microcapsule system, we find that the three biocatalysts-loaded microcapsules reveal a 5-fold rate enhancement, as compared to the three catalysts cascade in a homogeneous buffer phase, due to the operation of the cascaded catalysis in the confined microenvironment of the microcapsule.

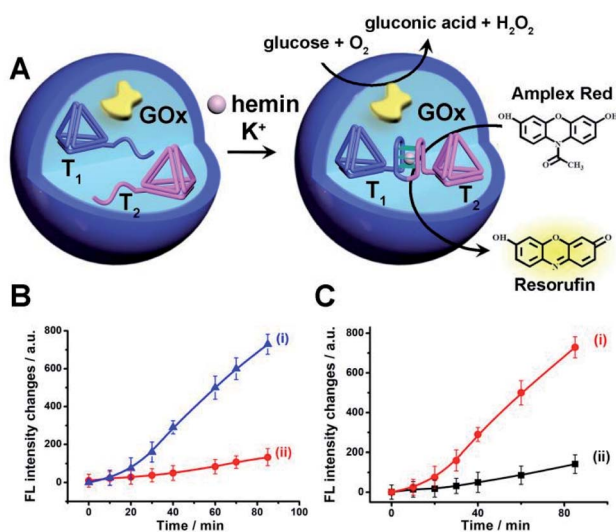


Fig. 2 (A) Schematic operation of a catalytic cascade consisting of GOx and hemin/G-quadruplex bridged tetrahedra dimer T_1/T_2 loaded in a microcapsule. The GOx catalyzed aerobic oxidation of glucose yields H_2O_2 that drives the oxidation of Amplex Red to the fluorescent Resorufin product. (B) Time-dependent fluorescence changes of Resorufin generated by: (i) The microcapsules loaded with GOx and the separated tetrahedra, T_1 and T_2 , in the presence of glucose, 10 mM, Amplex Red, 0.083 mM, hemin, 0.167 μM , in the presence of K^+ -ions, 50 mM. (ii) The reaction mixture described in (i) in the absence of K^+ -ions. (C) Time-dependent fluorescence intensities of Resorufin generated by: (i) The GOx/hemin/G-quadruplex bridged tetrahedra T_1/T_2 confined to the microcapsules. (ii) The GOx/hemin/G-quadruplex constituents in a homogeneous phase at the same concentrations of the catalysts present in the microcapsules. GOx, 0.063 μM , hemin/G-quadruplex bridged tetrahedra T_1/T_2 dimer 0.07 μM , glucose, 10 mM, Amplex Red, 0.083 mM, hemin, 0.167 μM and K^+ -ions 50 mM. Error bars derived from $N = 3$ experiments.



The introduction of DNA-based hydrogel microcapsules as cell-like containments, allows, however, the use of the microcapsules as functional units to establish intercommunication between the microcapsules, while the discussion till now addressed the loading and catalytic cascade in the microcapsule containments, the method fabricating the microcapsules provides structural features that eventually allow the design of intercommunicating microcapsules. The cross-opening hairpins H_1 and H_2 associated with the polymer chains P_1 and P_2 leads to a duplex-bridged hydrogel boundary that stabilizes the microcapsules. The synthesized hydrogel matrices yield, however, microcapsules that contain non-hybridized free, self-complementary tethers of H_1 and H_2 at the microcapsule membrane boundaries. Albeit, the surface coverage of these tethers is low, the tethers could provide important means to intercommunicate between microcapsules that include different loads. This is schematically outlined in Fig. 3, where two hydrogel microcapsules M_1 and M_2 potentially, loaded with different loads are interacted. The tethers p , q , exhibiting half complementarity represent segment of the H_1 and H_2 opened hairpin tethers, and thus dynamic supramolecular dimerization (or eventually enhanced oligomerization) may proceed to yield M_1 and M_2 monomers and M_1 and M_2 microcapsular dimers. The spatial proximity between the microcapsules M_1/M_2 , and the permeability of the microcapsular boundaries to low-molecular-weight agents might allow effective transport across the boundaries from microcapsules M_1 and M_2 or from M_2 to M_1 without diffusion to the bulk medium. Such mechanism could, then, provide the basis for the intercommunication biocatalytic processes occurring in neighboring microcapsules.

Accordingly, a mixture of two microcapsules M_1 , loaded with GOx (loading content of 0.12 μM), and microcapsules M_2 that were loaded with the K^+ -ions-stabilized hemin/G-quadruplex-bridged tetrahedra T_1/T_2 (loading content of 0.1 μM) was subjected to glucose, with the attempt to drive the intercommunicated biocatalytic cascade in the presence of Amplex Red, Fig. 4(A). The inter-microcapsule biocatalytic cascade proceeds effectively, Fig. 4(B), curve (i), where the aerobic oxidation of glucose in microcapsule M_1 yields H_2O_2 that diffuses through the microcapsules into the spatially proximate

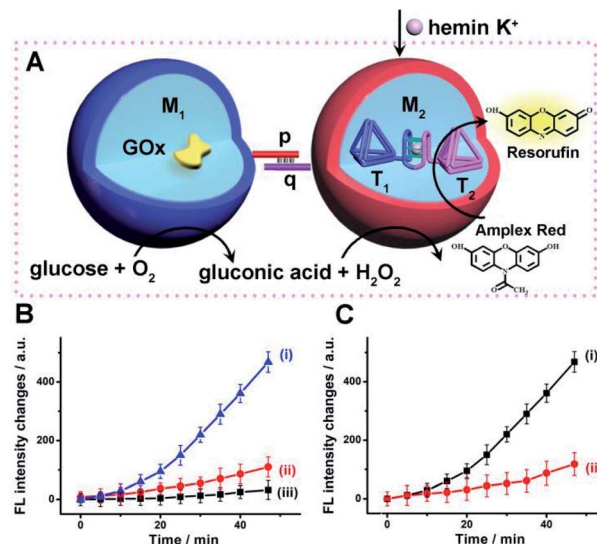


Fig. 4 (A) Schematic intercommunication between two types of loaded microcapsules: M_1 -microcapsule loaded with GOx; M_2 -microcapsule loaded with the hemin/G-quadruplex bridged T_1/T_2 dimer. Intercommunication leads to the activation of the biocatalytic cascade. (B) Time-dependent fluorescence changes of Resorufin generated by: (i) The intercommunicated M_1/M_2 microcapsules consisting of GOx and hemin/G-quadruplex bridged T_1/T_2 loads. Microcapsules, in the presence of K^+ -ions, 50 mM. (ii) The intercommunicated M_1/M_2 microcapsules in the absence of K^+ -ions. (iii) Upon addition of CE to the K^+ -ions containing M_1/M_2 microcapsule mixture. In all experiments, hemin, 0.167 μM , glucose, 30 mM, and Amplex Red, 0.083 mM, are added to the system. (C) Time-dependent fluorescence changes of Resorufin generated by: (i) The intercommunicated M_1/M_2 microcapsules loaded with GOx, 0.12 μM , and the hemin/G-quadruplex bridged T_1/T_2 dimer, 0.1 μM , in the presence of K^+ -ions, 50 mM. (ii) The homogeneous mixture of GOx and hemin/G-quadruplex-bridged tetrahedra T_1/T_2 at concentration identical to those in the M_1/M_2 microcapsules. All experiments included K^+ -ions, 50 mM, hemin, 0.167 μM , glucose, 30 mM, and Amplex Red, 0.083 mM. Error bars derived from $N = 3$ experiments.

microcapsule M_2 , where the hemin/G-quadruplex catalyzed oxidation of Amplex Red to the fluorescent Resorufin proceeds. The effectiveness of the inter-microcapsule cascade is controlled by presence of K^+ -ions. Fig. 4(B), curve (ii), shows the

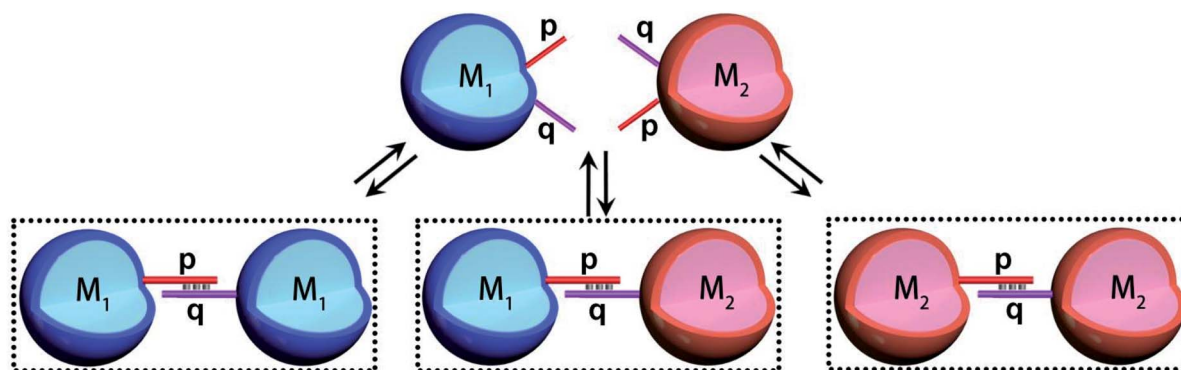


Fig. 3 Schematic dynamic dimerization paths of the different types of microcapsules, M_1 and M_2 loaded with Coumarin-labeled GOx (M_1) and Cy5-labeled T_1 and unlabeled T_2 (M_2) using surface-linked nucleic acid tethers (p and q) residues of the HCR hairpins H_1/H_2 used to generate the microcapsules.

cascaded catalysis of the intercommunicated M_1/M_2 microcapsules in the absence of K^+ -ions. Upon addition of CE to the K^+ -ions containing M_1/M_2 microcapsule mixture, the cascaded catalysis was inhibited due to the separation of the hemin/G-quadruplex-bridged tetrahedra T_1/T_2 , Fig. 4(B), curve (iii). The effectiveness of the inter-microcapsule cascade is, also, controlled by the concentration of glucose, Fig. S16† shows the fluorescence spectra of Resorufin at different time-intervals upon treatment the microcapsules with different concentrations of glucose. Control experiments revealed that no leakage of the microcapsules constituents into the bulk solution occurred, as evidenced by precipitation of the microcapsules mixture (after three days) and subjecting the solution to K^+ -ions, glucose, hemin and Amplex Red (Fig. S17,† curve (i)). No Resorufin could be detected in the bulk solution, implying that no leakage of the constituents from the microcapsules occurred (Fig. S17,† curve (ii)). Therefore, the loaded microcapsules M_1 , M_2 mixture retained their intact compositions and activities after a time-interval of three days. For further experiments demonstrating that the biocatalytic cascade originates from intimate contact between the microcapsules M_1/M_2 , *vide infra*. Fig. 4(C) curve (ii) depicts the rate of oxidation of Amplex Red to Resorufin by a homogeneous mixture of GOx and the hemin/G-quadruplex tetrahedra, at similar concentrations present in the microcapsular microcapsules, upon subjecting to the biocatalysts glucose 30 mM, as compared to the biocatalytic cascade in the microcapsular system, Fig. 4(C), curve (i). Evidently, the biocatalytic cascade driven by the intercommunicating microcapsules is 5-fold enhanced as compared to homogeneous mixture of biocatalytic constituents.

To further support the spatial intercommunication of the microcapsules through the dynamic interlinking of the microcapsules, a set of complementary experiments was performed. A mixture of coumarin-labeled GOx loaded microcapsules, 20 μ L (ca. 4500 microcapsules per μ L) and microcapsules loaded with Cy5-labeled T_1 and T_2 , 20 μ L (ca. 4500 microcapsules per μ L) were allowed to equilibrate. The mixture was subjected to confocal fluorescence imaging to probe the presence of monomer microcapsules loaded with coumarin-labeled GOx (blue fluorescence), monomer Cy5- T_1/T_2 -loaded microcapsules (red fluorescence), and dimer microcapsules consisting of dimer coumarin-labeled GOx microcapsules (blue-blue), dimer Cy5- T_1/T_2 -loaded microcapsules (red-red) and mixed coumarin-loaded microcapsules and Cy5- T_1/T_2 -loaded microcapsules (blue-red). Fig. 5(A) exemplifies images of the respective microcapsules. By analyzing ten different fluorescent domains of three different mixture samples (total 30 field frames), the respective monomer and dimer constituent were counted, and the average contents of the respective constituents were evaluated and presented in Fig. 5(B). The mixture includes ca. 51% monomer microcapsules and 37% dimer microcapsules and 12% of non-defined structures. The importance of this experiment is the demonstration of monomer and dimer microcapsules and the quantitative evaluation of the different constituents in the microcapsules mixture. Analog experiments were performed on the respective coumarin-labeled GOx-loaded microcapsules only, and on the Cy5- T_1/T_2 -loaded microcapsules

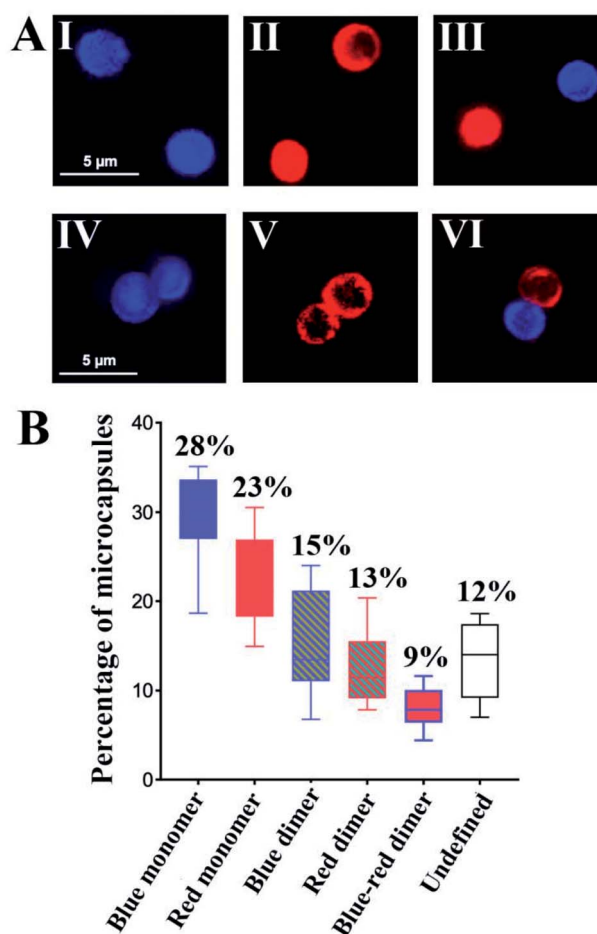


Fig. 5 (A) Examples of confocal microscopy images corresponding to: upper row-monomer microcapsules. Panel I-Coumarin-labeled GOx-loaded microcapsules (blue fluorescence). Panel II-Cy5- T_1/T_2 -loaded microcapsules (red fluorescence). Panel III-Separated coumarin-labeled GOx-loaded microcapsules (blue fluorescence) and Cy5- T_1/T_2 -loaded microcapsules (red fluorescence). Bottom row-dimer microcapsules. Panel IV-Coumarin-labeled GOx-loaded microcapsules (blue fluorescence). Panel V-Cy5- T_1/T_2 -loaded microcapsules (red fluorescence). Panel VI-Interlinked Coumarin-labeled GOx-loaded microcapsules and Cy5- T_1/T_2 -loaded microcapsules (blue-red fluorescence). (B) Statistical analysis of confocal fluorescence microscopy fields of the population of different monomer/dimer microcapsules assemblies, in a graphic form consisting of boxes and whiskers as error bar types, (total 30 fields). (For blue fluorescence: $E_x = 405$ nm, $E_m = 430-470$ nm. For red fluorescence: $E_x = 640$ nm, $E_m = 650-700$ nm). Error bars derived from $N = 3$ experiments.

only, and these are presented in the ESI† (For quantitative evaluation of the monomer/dimer mixture, see Fig. S18 and S19† and accompanying discussion).

To further support the equilibrated monomers/dimers composition in the mixture, fluorescence-activated cell sorting (FACS) experiments were performed, Fig. S20,† and accompanying discussion. These experiments indicated ca. 35% of Cy5- T_1/T_2 -loaded microcapsules including monomers and dimers, 35% of coumarin-labeled GOx loaded microcapsules containing monomer and dimer, and ca. 7% population of the dimer structure of the coumarin-labeled GOx loaded microcapsules



and Cy5- T_1/T_2 -loaded microcapsules in microcapsular monomer/dimer mixture, which is in good agreement with the confocal fluorescence microscopy analysis.

The confocal microscopy fluorescence images of the microcapsules demonstrated the feasibility of the nucleic acid-functionalized microcapsules to form dimer structures enabling the spatial intimate contacting of the microcapsules for intercommunication of the microcapsules and operation of the biocatalytic cascade, Fig. 6(A), state I. The duplex nucleic acid bridging units generating the functional dimer microcapsule structure include, however, encoded information that allows the guided inhibition of the intercommunicated microcapsules biocatalytic cascade, Fig. 6(A). The p/q duplex generating the intercommunicated dimer dictating the biocatalytic cascade, can be separated by the addition of two fuel strands p' and q', state II. Thus, in the presence of the fuel strands p' and q', the active microcapsule configuration activating the intercommunicated biocatalytic cascade is blocked due to the separation of dimer microcapsules. That is, the blocker-induced inhibition of the biocatalytic cascade not only introduces a means to inhibit the intercell communication, but provides a proof that the intercommunication of the microcapsule originates from the intermediary formation of the dimer microcapsules. Furthermore, by the addition of counter-fuel strands p'' and q'' displacing the blocker units p' and q', the re-assembly of the active dimer structures is stimulated, resulting in the switched "ON" GOx/hemin/G-quadruplex tetrahedra biocatalytic cascade, state I. Fig. 6(B), curve (i), shows the time-dependent fluorescence changes of Resorufin formed by the intercommunicated dimer-microcapsule biocatalytic cascade driven by GOx and the hemin/G-quadruplex bridged DNA tetrahedra in state I. Fig. 6(B), curve (ii), shows the time-dependent fluorescence changes of Resorufin by the mixture

consisting of the GOx-loaded microcapsules and the hemin/G-quadruplex bridged tetrahedra dimer-loaded microcapsule, treated with the fuel strands p' and q' and triggered by glucose. Inefficient Resorufin formation is observed, consistent with the inhibited formation of the dimer microcapsules, where the intimate spatial structure for communicating substrate transfer is perturbed. Treatment of the inhibited microcapsule mixture with the counter-fuels p'' and q'' reactivates the intercommunication between the microcapsule, Fig. 6(B), curve (iii). That is, the displacement of the p'/p'' and q'/q'' duplexes uncages the microcapsules functionalized with the p and q tethers, allowing regeneration of the intercommunicating dimer structures of the microcapsules. It should be noted that the residual very low cascaded biocatalytic activity of the p', q' separated GOx-loaded microcapsules and hemin/G-quadruplex-bridged T_1/T_2 -loaded microcapsules is attributed to incomplete separation of the dimer microcapsule structure, rather than the possible exchange of loads within the dimer microcapsule structure. This is supported by an experiment, Fig. S21,[†] allowing the dimer-microcapsules mixture to interact for a time-interval of ten days and subsequent treatment of the mixture after this time-interval with p', q' to induce separation. The resulting separated microcapsule mixture demonstrated the same residual biocatalytic performance depicted in Fig. 6(B), curve (ii), indicating that no content exchange occurred even within a time-interval of ten days.

It should be noted that the rate of the biocatalytic cascade in the mixture of M_1/M_2 intercommunicated microcapsules is *ca.* 5-6-fold enhanced as compared to the mixture of the homogenous biocatalysts, even though only *ca.* 9% of the microcapsules are in the "correct" configuration to allow the efficient intercommunicated cascade. This means that the "effective" rate enhancement of cascaded catalysis demonstrated by the

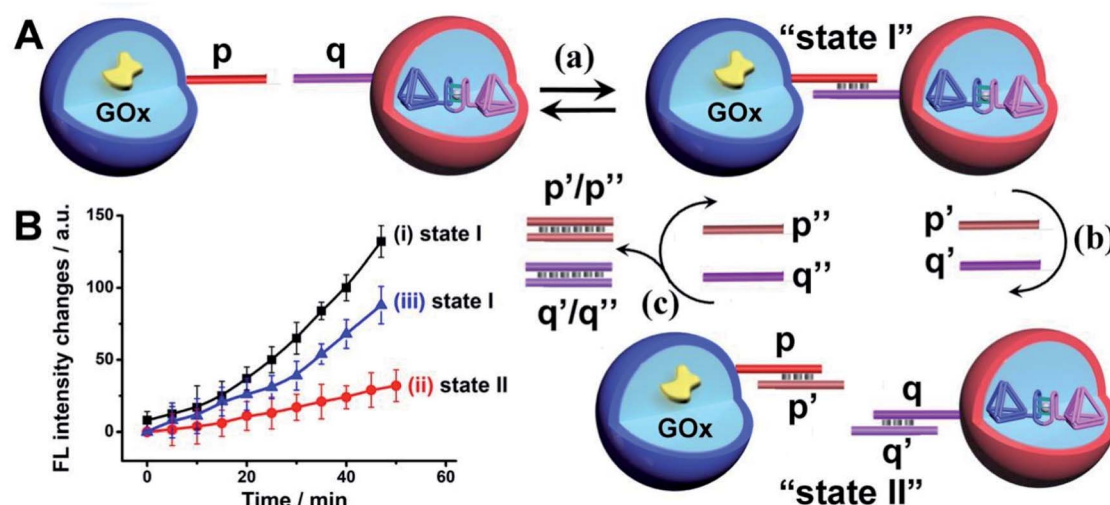


Fig. 6 (A) Switchable inhibition of the intercommunicated microcapsule GOx/hemin/G-quadruplex biocatalyst cascade using a "fuel" inhibiting strands p', q' and "counter-fuel strands" p'', q''. (B) Time-dependent fluorescence changes corresponding to Resorufin generated by the intercommunicated GOx/hemin/G-quadruplex bridged T_1/T_2 cascade as a result of (a)-equilibrated p/q interlinked microcapsule dimers, state I, curve (i). (b) Upon separation of the dimers using p' and q' fuel strands, state II, curve (ii), and (c) upon treatment of the separated capsules, state II with the counter fuel strands p'' and q'' to regenerate state I, curve (iii). Error bars derived from $N = 3$ experiments.



intercommunicated pairs of microcapsules, as compared to the homogenous biocatalysts is significantly higher. Also, note that the rate enhancement by the intercommunicated microcapsules as compared to the homogenous mixture of biocatalysts is comparable to the rate enhancement demonstrated by the GOx//hemin/G-quadruplex in a single microcapsule, despite that only 9% of the microcapsules exist in the appropriate dimer configuration. This is explained by the substantially lower loading of the GOx and hemin/G-quadruplex catalysts in single microcapsules that can be accomplished in the course of the microcapsules preparation (the number of microcapsules in the different systems is approximately similar).

The successful operation of the biocatalyst cascade through the dynamic formation of dimer microcapsules assemblies allowed us to establish a three-catalysts cascade through the bi-directional intercommunication of the microcapsules, Fig. 7(A). The system consisted of the M_1 microcapsules loaded with GOx, 0.12 μM , and the second type of microcapsules, M_3 -loaded with β -gal, 0.058 μM , and the hemin/G-quadruplex bridged tetrahedra T_1/T_2 dimer constituent, 0.07 μM . Since the microcapsular containments are functionalized with the tethers p and q, the dynamic assembly of M_1/M_3 dimer allows the bi-directional

intercommunication of the microcapsules. In the presence of lactose, the β -gal catalyzed hydrolysis of lactose in microcapsules M_3 yields glucose that is being transported through the dimer boundary into microcapsules M_1 . The aerobic oxidation of glucose in microcapsules M_1 yields H_2O_2 that is back channelled through the dimer boundary into microcapsules M_3 , where the hemin/G-quadruplex bridged DNA tetrahedra T_1/T_2 catalyzes the oxidation of Amplex Red to Resorufin by H_2O_2 . Fig. 7(B), curve (i), depicts the time-dependent fluorescence changes of Resorufin upon subjecting to two-capsule system in the presence of lactose, K^+ -ions, hemin and Amplex Red.

The three-catalysts cascade is activated through the bi-directional intercommunication of the two microcapsules. Further experiments demonstrated that exclusion of K^+ -ions from the system that resulted in the inhibition of the hemin/G-quadruplex bridged T_1/T_2 catalyst, prohibited the formation of Resorufin, Fig. 7(B), curve (ii). Also, addition of CE to the intercommunicating microcapsules separated the hemin/G-quadruplex bridged T_1/T_2 and eliminated the formation of Resorufin, Fig. 7(B), curve (iii). Finally, elimination of lactose from the system, blocked the entire intercommunicated biocatalytic cascade, Fig. 7(B), curve (iv). Fig. 7(C), compare the time-dependent fluorescence changes of Resorufin by the intercommunicated microcapsular system, curve (i), to the time-dependent fluorescence changes of Resorufin generated by the three catalysts in a homogeneous solution, where the catalysts are present at identical concentrations present in microcapsules, curve (ii). The biocatalytic cascade in the homogeneous mixture is very inefficient, and the three-catalyst cascade operating in the intercommunicating microcapsule assembly is at least 10-fold enhanced as compared to the homogeneous mixture. These results highlight the significance of bi-directional communication between the catalysts-loaded microcapsules that originates from the dynamic formation of dimer microcapsules bridged by the duplex nucleic acid p/q tethers.

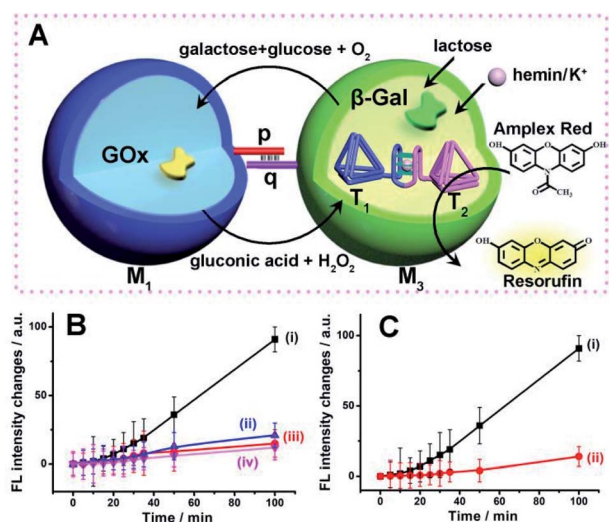


Fig. 7 (A) Schematic intercommunication of two microcapsular microcapsules activating the three-biocatalysts cascade consisting of the lactose-driven β -gal//GOx//hemin/G-quadruplex bridged T_1/T_2 dimer catalytic cascade yielding the fluorescent Resorufin. (B) Time-dependent fluorescence intensities of Resorufin generated by: (i) The intercommunicated M_1/M_3 capsules in the presence of lactose, 40 mM, hemin, 0.167 μM , K^+ -ions, 50 mM, and Amplex Red, 0.083 mM. (ii) Lactose, 40 mM, hemin, 0.167 μM and Amplex Red, 0.083 mM, yet in the absence of K^+ -ions. (iii) As in (i), yet in the presence of CE, 200 mM. (iv) As in (i), yet in the absence of lactose. (C) Time-dependent fluorescence changes of Resorufin generated by: (i) The M_1/M_3 capsules, loaded with GOx, 0.12 μM , β -gal, 0.07 μM , and the hemin/G-quadruplex bridged T_1/T_2 dimer, 0.058 μM , in the presence of lactose, 40 mM, hemin, 0.167 μM , K^+ -ions, 50 mM, and Amplex Red, 0.083 mM. (ii) A homogeneous mixture consisting of GOx, 0.12 μM , β -gal, 0.07 μM , and the hemin/G-quadruplex bridged T_1/T_2 dimer, 0.058 μM , in the presence of lactose, 40 mM, hemin, 0.167 μM , K^+ -ions, 50 mM, and Amplex Red, 0.083 mM. Error bars derived from $N = 3$ experiments.

Conclusions

We introduced nucleic acid-crosslinked carboxymethyl cellulose hydrogel-stabilized microcapsules as a versatile containment for the construction of microcapsules. The microcapsule containments reveal advantages over the present art of microcapsule assemblies: (i) The hydrogel-stabilized microcapsules provide stable non-leakable containments for constituents with molecular weights, ≥ 5 kDa while low-molecular-weight agents freely cross the microcapsule boundaries.⁶⁵ This allows to trigger chemical transformations in the microcapsule assemblies and to monitor the chemical transformations proceeding in the microcapsules by analyzing the chemical composition of the bulk solution. (ii) The method applied to construct the nucleic acid-modified microcapsules yields microcapsules functionalized with nucleic acid tethers revealing self-recognition properties. This unique feature provides means to yield dimer structures of two different microcapsules of programmed pre-engineered constitutional compositions. The spatial proximity of the interlinked microcapsules provide means to intercommunicate between the microcapsules and



operate inter-cell catalytic cascades. (iii) The nucleic acid-guided formation of the dimer microcapsules can be switched off and re-activated by applying “fuel” and “counters-fuel” strands, thereby allowing the programmed activation and deactivation of intercommunicating microcapsule cascades. Specifically, the present study demonstrated the integration of enzymes and/or hemin/G-quadruplex-crosslinked DNA tetrahedra as biocatalysts or synthetic models for protein–protein nanostructure exhibiting catalytic (DNAzyme) activities. Besides the basic structural characterization of the catalysts-loaded microcapsules, important functionalities of the microcapsules were achieved including: (i) The integration of the hemin/G-quadruplex-crosslinked DNA tetrahedra T_1/T_2 dimer structure into the microcapsules and the application of the nanostructure as catalytic protein–protein model system and switchable catalytic systems. (ii) The integration of glucose oxidase and hemin/G-quadruplex crosslinked DNA tetrahedra T_1/T_2 DNAzyme in the microcapsules or the integration of β -galactosidase, glucose oxidase, and the hemin/G-quadruplex crosslinked DNA tetrahedra T_1/T_2 unit as catalytic assemblies driving biocatalyst and three-catalysts cascades. Superior catalytic cascades in the confined environments of microcapsules, as compared to the operation of the catalytic cascades in homogeneous aqueous mixtures were demonstrated. (iii) The intercommunication of microcapsules and the guided directional and bi-directional operation of biocatalytic and three-catalytic cascades were highlighted. A unique property of the nucleic acid-functionalized hydrogel microcapsules was highlighted by demonstrating the dynamic formation of dimer microcapsule assemblies that leads to spatial proximity of the microcapsules and to a mechanistic pathway to intercommunicate microcapsules.

We believe, however, that these results and the nucleic acid-based microcapsule assemblies introduce a substantially broader impact on the development of functional microcapsules; the introduction of other catalytic constituents into the microcapsules, such as DNA machineries⁶⁶ or transcription machineries,⁶⁷ the integration of dynamic catalytic networks⁶⁸ into the hydrogel microcapsules, and the further functionalization of the nucleic acid-modified microcapsular hydrogel boundaries with stimuli-responsive nucleic acids that switch the chemical transformations in the microcapsules are interesting paths to follow. Furthermore, the study has emphasized the superior functions of enzyme or multi-enzyme assemblies loaded in microcapsules to drive biocatalytic transformations. Other applications of such enzyme-loaded microcapsules for therapeutic or sensing applications may be envisaged. For example, the intracapsule integration of a biocatalytic agent⁶⁹ might yield effective carriers for drug release. Also, the amplification phenomenon of biocatalytic cascades, in the confined microcapsular environment, and the possibility to engineer nucleic acid tethers on the boundaries of the microcapsules provide means to anchor the microcapsules onto sensing recognition events as amplifying transducers. For example, by anchoring microcapsules loaded with biocatalytic cascades generating redox active compounds, to recognition events occurring on electrode surfaces, amplified electrochemical sensors can be realized.⁷⁰

Author contributions

P. Z., A. F. and I. W. formulated the concepts and methodology of the study. The manuscript was written through the contributions of all authors. All authors participated in the experimental work and have given approval regarding the final version of the manuscript.

Conflicts of interest

There are no conflicts to declare.

Acknowledgements

This research is supported by the Israel Science Foundation (ISF)/National Natural Science Foundation of China-NSFC Joint Scientific Research Program (Grant No. 3542/21).

Notes and references

- (a) R. J. R. W. Peters, I. Louzao and J. C. M. van Hest, *Chem. Sci.*, 2012, **3**, 335–342; (b) Y. Tu, F. Peng, A. Adawy, Y. Men, L. K. Abdelmohsen and D. A. Wilson, *Chem. Rev.*, 2016, **116**, 2023–2078.
- (a) B. C. Buddingh and J. C. M. van Hest, *Acc. Chem. Res.*, 2017, **50**, 769–777; (b) A. J. Dzieciol and S. Mann, *Chem. Soc. Rev.*, 2012, **41**, 79–85; (c) T. Trantidou, M. Friddin, Y. Elani, N. J. Brooks, R. V. Law, J. M. Seddon and O. Ces, *ACS Nano*, 2017, **11**, 6549–6565; (d) P. Stano and P. L. Luisi, *Curr. Opin. Biotechnol.*, 2013, **24**, 633–638.
- (a) M. Vázquez-González, C. Wang and I. Willner, *Nat. Catal.*, 2020, **3**, 256–273; (b) J. Fu, M. Liu, Y. Liu and H. Yan, *Acc. Chem. Res.*, 2012, **45**, 1215–1226; (c) K. S. Rabe, J. Müller, M. Skoupi and C. M. Niemeyer, *Angew. Chem., Int. Ed.*, 2017, **56**, 13574–13589; (d) J. Shi, Y. Wu, S. Zhang, Y. Tian, D. Yang and Z. Jiang, *Chem. Soc. Rev.*, 2018, **47**, 4295–4313.
- G. A. Ellis, W. P. Klein, G. Lasarte-Aragones, M. Thakur, S. A. Walper and I. L. Medintz, *ACS Catal.*, 2019, **9**, 10812–10869.
- O. I. Wilner, Y. Weizmann, R. Gill, O. Lioubashevski, R. Freeman and I. Willner, *Nat. Nanotechnol.*, 2009, **4**, 249–254.
- J. Fu, Y. R. Yang, A. Johnson-Buck, M. Liu, Y. Liu, N. G. Walter, N. W. Woodbury and H. Yan, *Nat. Nanotechnol.*, 2014, **9**, 531–536.
- T. A. Ngo, E. Nakata, M. Saimura and T. Morii, *J. Am. Chem. Soc.*, 2016, **138**, 3012–3021.
- J. Qu, S. Cao, Q. Wei, H. Zhang, R. Wang, W. Kang, T. Ma, L. Zhang, T. Liu, S. Wing-Ngor Au and F. Sun, *ACS Nano*, 2019, **13**, 9895–9906.
- C. You, S. Myung and Y. H. P. Zhang, *Angew. Chem., Int. Ed.*, 2012, **51**, 8787–8790.
- Y. Elani, R. V. Law and O. Ces, *Nat. Commun.*, 2014, **5**, 5305.
- D. M. Vriezema, P. M. Garcia, N. Sancho Oltra, N. S. Hatzakis, S. M. Kuiper, R. J. Nolte, A. E. Rowan and J. C. Van Hest, *Angew. Chem., Int. Ed.*, 2007, **46**, 7378–7382.



- 12 (a) J. Liang, F. Mazur, C. Tang, X. Ning, R. Chandrawati and K. Liang, *Chem. Sci.*, 2019, **10**, 7852–7858; (b) W. H. Chen, M. Vázquez-González, A. Zoabi, R. Abu-Reziq and I. Willner, *Nat. Catal.*, 2018, **1**, 689–695.
- 13 (a) C. Y. Jeong, Y. D. Han, J. H. Yoon and H. C. Yoon, *J. Biotechnol.*, 2014, **175**, 7–14; (b) Y. Guo, H. Wang, F. Zhang, J. Xia and Z. Wang, *Anal. Chim. Acta*, 2020, **1140**, 10–17.
- 14 (a) S. Z. Wang, Y. H. Zhang, H. Ren, Y. L. Wang, W. Jiang and B. S. Fang, *Crit. Rev. Biotechnol.*, 2017, **37**, 1024; (b) I. Wheeldon, S. D. Minter, S. Banta, S. C. Barton, P. Atanassov and M. Sigman, *Nat. Chem.*, 2016, **8**, 299; (c) P. N. Devine, R. M. Howard, R. Kumar, M. P. Thompson, M. D. Truppo and N. J. Turner, *Nat. Rev. Chem.*, 2018, **2**, 409–421.
- 15 (a) L. Martini and S. S. Mansy, *Chem. Commun.*, 2011, **47**, 10734–10736; (b) L. Sun, Y. Gao, Y. Wang, Q. Wei, J. Shi, N. Chen, D. Li and C. Fan, *Chem. Sci.*, 2018, **9**, 5967–5975; (c) Z. Nourian and C. Danelon, *ACS Synth. Biol.*, 2013, **2**, 186–193; (d) K. Kurihara, Y. Okura, M. Matsuo, T. Toyota, K. Suzuki and T. Sugawara, *Nat. Commun.*, 2015, **6**, 8352; (e) R. S. Rikken, H. Engelkamp, R. J. Nolte, J. C. Maan, J. C. van Hest, D. A. Wilson and P. C. Christianen, *Nat. Commun.*, 2016, **7**, 12606; (f) Y. Elani, R. V. Law and O. Ces, *Nat. Commun.*, 2014, **5**, 5305.
- 16 (a) Z. Wang, M. C. van Oers, F. P. Rutjes and J. C. van Hest, *Angew. Chem., Int. Ed. Engl.*, 2012, **51**, 10746–10750; (b) D. M. Vriezema, P. M. L. Garcia, N. Sancho Oltra, N. S. Hatzakis, S. M. Kuiper, R. J. M. Nolte, A. E. Rowan and J. C. M. van Hest, *Angew. Chem.*, 2007, **119**, 7522–7526.
- 17 (a) X. Huang, A. J. Patil, M. Li and S. Mann, *J. Am. Chem. Soc.*, 2014, **136**, 9225–9234; (b) X. Huang, M. Li, D. C. Green, D. S. Williams, A. J. Patil and S. Mann, *Nat. Commun.*, 2013, **4**, 2239; (c) L. Wang, P. Wen, X. Liu, Y. Zhou, M. Li, Y. Huang, L. Geng, S. Mann and X. Huang, *Chem. Commun.*, 2017, **53**, 8537–8540.
- 18 (a) M. Li, X. Huang and S. Mann, *Small*, 2014, **10**, 3291–3298; (b) M. Li, R. L. Harbron, J. Weaver, B. P. Binks and S. Mann, *Nat. Chem.*, 2013, **5**, 529–536; (c) M. Li, R. L. Harbron, J. V. Weaver, B. P. Binks and S. Mann, *Nat. Chem.*, 2013, **5**, 529–536.
- 19 C. M. Paleos, D. Tsiourvas, Z. Sideratou and A. Pantos, *J. Controlled Release*, 2013, **170**, 141–152.
- 20 (a) D. S. Williams, A. J. Patil and S. Mann, *Small*, 2014, **10**, 1830–1840; (b) Y. Qiao, M. Li, R. Booth and S. Mann, *Nat. Chem.*, 2017, **9**, 110–119; (c) Y. Yin, L. Niu, X. Zhu, M. Zhao, Z. Zhang, S. Mann and D. Liang, *Nat. Commun.*, 2016, **7**, 10658; (d) J. Fothergill, M. Li, S. A. Davis, J. A. Cunningham and S. Mann, *Langmuir*, 2014, **30**, 14591–14596; (e) T. Y. Dora Tang, C. Rohaida Che Hak, A. J. Thompson, M. K. Kuimova, D. S. Williams, A. W. Perriman and S. Mann, *Nat. Chem.*, 2014, **6**, 527–533.
- 21 (a) S. Liu, Y. Zhang, M. Li, L. Xiong, Z. Zhang, X. Yang, X. He, K. Wang, J. Liu and S. Mann, *Nat. Chem.*, 2020, **12**, 1165–1173; (b) A. Kuchler, M. Yoshimoto, S. Luginbuhl, F. Mavelli and P. Walde, *Nat. Nanotechnol.*, 2016, **11**, 409–420; (c) M. Vázquez-González, C. Wang and I. Willner, *Nat. Catal.*, 2020, **3**, 256–273.
- 22 (a) B. Kumar, A. J. Patil and S. Mann, *Nat. Chem.*, 2018, **10**, 1154–1163; (b) W. S. Jang, H. J. Kim, C. Gao, D. Lee and D. A. Hammer, *Small*, 2018, **14**, e1801715; (c) D. A. Wilson, R. J. Nolte and J. C. van Hest, *Nat. Chem.*, 2012, **4**, 268–274.
- 23 (a) L. Tian, M. Li, J. Liu, A. J. Patil, B. W. Drinkwater and S. Mann, *ACS Cent. Sci.*, 2018, **4**, 1551–1558; (b) S. Sun, M. Li, F. Dong, S. Wang, L. Tian and S. Mann, *Small*, 2016, **12**, 1920–1927.
- 24 K. Kurihara, M. Tamura, K. Shohda, T. Toyota, K. Suzuki and T. Sugawara, *Nat. Chem.*, 2011, **3**, 775–781.
- 25 Z. Xu, T. Hueckel, W. T. M. Irvine and S. Sacanna, *Nature*, 2021, **597**, 220–224.
- 26 K. P. Adamala, D. A. Martin-Alarcon, K. R. Guthrie-Honea and E. S. Boyden, *Nat. Chem.*, 2017, **9**, 431–439.
- 27 R. R. Poudyal, R. M. Guth-Metzler, A. J. Veenis, E. A. Frankel, C. D. Keating and P. C. Bevilacqua, *Nat. Commun.*, 2019, **10**, 490.
- 28 A. Joesaar, S. Yang, B. Bogels, A. van der Linden, P. Pieters, B. Kumar, N. Dalchau, A. Phillips, S. Mann and T. F. A. de Greef, *Nat. Nanotechnol.*, 2019, **14**, 369–378.
- 29 M. Schwarz-Schilling, L. Aufinger, A. Muckl and F. C. Simmel, *Integr. Biol.*, 2016, **8**, 564–570.
- 30 M. S. Long, C. D. Jones, M. R. Helfrich, L. K. Mangeney-Slavin and C. D. Keating, *Proc. Natl. Acad. Sci. U. S. A.*, 2005, **102**, 5920–5925.
- 31 H. Che, S. Cao and J. C. M. van Hest, *J. Am. Chem. Soc.*, 2018, **140**, 5356–5359.
- 32 H. Che, B. C. Buddingh and J. C. M. van Hest, *Angew. Chem., Int. Ed. Engl.*, 2017, **56**, 12581–12585.
- 33 M. Marguet, C. Bonduelle and S. Lecommandoux, *Chem. Soc. Rev.*, 2013, **42**, 512–529.
- 34 L. Rodriguez-Arco, M. Li and S. Mann, *Nat. Mater.*, 2017, **16**, 857–863.
- 35 F. C. Simmel, B. Yurke and H. R. Singh, *Chem. Rev.*, 2019, **119**, 6326–6369.
- 36 C. H. Lu, X. J. Qi, R. Orbach, H. H. Yang, I. Mironi-Harpaz, D. Seliktar and I. Willner, *Nano Lett.*, 2013, **13**, 1298–1302.
- 37 Y. W. Hu, A. Ceconello, A. Idili, F. Ricci and I. Willner, *Angew. Chem., Int. Ed.*, 2017, **56**, 15210–15233.
- 38 K. Gehring, J.-L. Leroy and M. Gueron, *Nature*, 1993, **363**, 561–565.
- 39 R. M. Dirks and N. A. Pierce, *Proc. Natl. Acad. Sci. U. S. A.*, 2004, **101**, 15275–15278.
- 40 F. Wang, C.-H. Lu, X. Liu, L. Freage and I. Willner, *Anal. Chem.*, 2014, **86**, 1614–1621.
- 41 I. Willner and M. Zayats, *Angew. Chem., Int. Ed. Engl.*, 2007, **46**, 6408–6418.
- 42 G. F. Joyce, *Angew. Chem., Int. Ed. Engl.*, 2007, **46**, 6420–6436.
- 43 P. Travascio, A. J. Bennet, D. Y. Wang and D. Sen, *Chem. Biol.*, 1999, **6**, 779–787.
- 44 R. R. Breaker and G. F. Joyce, *Chem. Biol.*, 1994, **1**, 223–229.
- 45 C. Mao, W. Sun and N. C. Seeman, *J. Am. Chem. Soc.*, 1999, **121**, 5437–5443.
- 46 X. Wang, A. R. Chandrasekaran, Z. Shen, Y. P. Ohayon, T. Wang, M. E. Kizer, R. Sha, C. Mao, H. Yan, X. Zhang, S. Liao, B. Ding, B. Chakraborty, N. Jonoska, D. Niu,



- H. Gu, J. Chao, X. Gao, Y. Li, T. Ciengshin and N. C. Seeman, *Chem. Rev.*, 2019, **119**, 6273–6289.
- 47 P. W. Rothmund, *Nature*, 2006, **440**, 297–302.
- 48 R. P. Goodman, R. M. Berry and A. J. Turberfield, *Chem. Commun.*, 2004, **12**, 1372–1373.
- 49 H. Pei, X. Zuo, D. Zhu, Q. Huang and C. Fan, *Acc. Chem. Res.*, 2014, **47**, 550–559.
- 50 M. Lin, J. Wang, G. Zhou, J. Wang, N. Wu, J. Lu, J. Gao, X. Chen, J. Shi, X. Zuo and C. Fan, *Angew. Chem., Int. Ed. Engl.*, 2015, **54**, 2151–2155.
- 51 H. Pei, L. Liang, G. Yao, J. Li, Q. Huang and C. Fan, *Angew. Chem., Int. Ed.*, 2012, **51**, 9020–9024.
- 52 V. E. Ferry, J. M. Smith and A. P. Alivisatos, *ACS Photonics*, 2014, **1**, 1189–1196.
- 53 N. Y. Wong, C. Zhang, L. H. Tan and Y. Lu, *Small*, 2011, **7**, 1427–1430.
- 54 Z. Zhou, Y. S. Sohn, R. Nechushtai and I. Willner, *ACS Nano*, 2020, **14**, 9021–9031.
- 55 H. Pei, N. Lu, Y. Wen, S. Song, Y. Liu, H. Yan and C. Fan, *Adv. Mater.*, 2010, **22**, 4754–4758.
- 56 Z. Qing, J. Hu, J. Xu, Z. Zou, Y. Lei, T. Qing and R. Yang, *Chem. Sci.*, 2019, **11**, 1985–1990.
- 57 A. S. Walsh, H. Yin, C. M. Erben, M. J. Wood and A. J. Turberfield, *ACS Nano*, 2011, **5**, 5427–5432.
- 58 A. J. Mastroianni, S. A. Claridge and A. P. Alivisatos, *J. Am. Chem. Soc.*, 2009, **131**, 8455–8459.
- 59 Z. Zhou, J. Wang and I. Willner, *J. Am. Chem. Soc.*, 2021, **143**, 241–251.
- 60 W.-C. Liao and I. Willner, *Adv. Funct. Mater.*, 2017, **27**, 1702732.
- 61 W.-C. Liao, M. Riutin, W. J. Parak and I. Willner, *ACS Nano*, 2016, **10**, 8683–8689.
- 62 W.-C. Liao, Y. S. Sohn, M. Riutin, A. Cecconello, W. J. Parak, R. Nechushtai and I. Willner, *Adv. Funct. Mater.*, 2016, **26**, 4262–4273.
- 63 F. Huang, W.-C. Liao, Y. S. Sohn, R. Nechushtai, C.-H. Lu and I. Willner, *J. Am. Chem. Soc.*, 2016, **138**, 8936–8945.
- 64 W.-H. Chang, Y.-F. Lee, Y.-W. Liu, I. Willner and W.-C. Liao, *Nanoscale*, 2021, **13**, 16799–16808.
- 65 A. Fischer, S. Lilienthal, M. Vazquez-Gonzalez, M. Fadeev, Y. S. Sohn, R. Nechushtai and I. Willner, *J. Am. Chem. Soc.*, 2020, **142**, 4223–4234.
- 66 Z. Cheglakov, Y. Weizmann, A. B. Braunschweig, O. I. Wilner and I. Willner, *Angew. Chem., Int. Ed. Engl.*, 2008, **47**, 126–130.
- 67 Y. W. Yin and T. A. Steitz, *Science*, 2002, **298**, 1387–1395.
- 68 L. Yue, S. Wang and I. Willner, *J. Am. Chem. Soc.*, 2020, **142**, 14437–14442.
- 69 C. Chen, M. Vázquez-González, M. P. O'Hagan, Y. Ouyang, Z. Wang and I. Willner, *Small*, 2022, **18**, 2104420.
- 70 L. Alfonta, A. Bardea, O. Khersonsky, E. Katz and I. Willner, *Biosens. Bioelectron.*, 2001, **16**, 675–687.

