

Cite this: *J. Mater. Chem. B*,  
2024, 12, 4289

## Self-assembly of peptides in living cells for disease theranostics

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The past few decades have witnessed substantial progress in biomedical materials for addressing health concerns and improving disease therapeutic and diagnostic efficacy. Conventional biomedical materials are typically created through an *ex vivo* approach and are usually utilized under physiological environments *via* transfer from preparative media. This transfer potentially gives rise to challenges for the efficient preservation of the bioactivity and implementation of theranostic goals on site. To overcome these issues, the *in situ* synthesis of biomedical materials on site has attracted great attention in the past few years. Peptides, which exhibit remarkable biocompatibility and reliable noncovalent interactions, can be tailored *via* tunable assembly to precisely create biomedical materials. In this review, we summarize the progress in the self-assembly of peptides in living cells for disease diagnosis and therapy. After a brief introduction to the basic design principles of peptide assembly systems in living cells, the applications of peptide assemblies for bioimaging and disease treatment are highlighted. The challenges in the field of peptide self-assembly in living cells and the prospects for novel peptide assembly systems towards next-generation biomaterials are also discussed, which will hopefully help elucidate the great potential of peptide assembly in living cells for future healthcare applications.

Received 21st February 2024,  
Accepted 25th March 2024

DOI: 10.1039/d4tb00365a

rsc.li/materials-b

### 1. Introduction

Biomedical materials serve as the basic platform for most healthcare, ranging from diagnosis to rehabilitation and treatment. Lying at the intersection of biology, chemistry, and materials science, biomedical materials can implement different healthcare processes, primarily including drug delivery, biosensors, and tissue engineering.<sup>1,2</sup> Over the past decades, biomaterials have been classified based on their components

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into different types, such as metals, ceramics, polymers, and hybrids.<sup>3,4</sup> In particular, fabricating biomaterials using bottom-up approaches opens a new avenue to improve their performance in healthcare and revolutionize the approaches for improving human health.

Conventionally, biomaterials utilized in healthcare are created in an *ex vivo* manner. In typical processes, guided by healthcare goals, the materials are formulated from individual components with corresponding functions. The bioactivity of the resulting materials is evaluated through *in vitro* and *in vivo* studies. Despite the enormous progress achieved in this field, conventional biomaterials prepared under *ex vivo* conditions still face challenges arising from their preparation under simplified solution environments, whereby their utilization *via* transfer to the physiological environment can lead to biosafety issues associated with off-targeting or a loss of bioactivity. Therefore, currently available biomaterials consequently exhibit poor treatment outcomes and limited selection between healthy and pathological regions.<sup>5,6</sup> To control

bioactivity, stimulus-responsive biomedical materials have been developed as second-generation counterparts.<sup>7–9</sup> Employing either external or internal stimuli, the bioactivity of stimulus-responsive biomaterials can be precisely regulated at pathological sites. However, preformed biomaterials still suffer from problems in blood circulation and site accumulation. As an extension of stimulus-responsive categories, third-generation biomaterials can be created on site, termed as *in situ*-formed biomaterials.<sup>10,11</sup> Generally, stimulus-responsive materials undergo various responses upon exposure to stimuli, broadly ranging from structural changes to functional implementation. *In situ*-formed materials are usually created through stimulus-responsive processes. Therefore, *in situ*-formed materials are, in principle, one category of stimulus-responsive materials. However, the definition of *in situ*-formed materials emphasizes the formulation on targeting sites and allows for a precise clarification of stimulus-responsive processes. In addition, compared to traditional strategies, biomedical materials prepared in living environments obviously allows for satisfying the demand for healthcare with real-time characterization and detection under physiological conditions.<sup>12,13</sup> Accuracy and the precision activation of bioactivity at specific sites could thereby alleviate the possible biosafety issues of such materials.<sup>14–16</sup>

*In situ* biomaterials are strongly dependent on stimulus-responsive reactions to regulate their bioactivities. In principle, *in situ* biomedical materials are typically involve stimulus-formation and stimulus-responsive activation aspects. Stimulus-responsive formation refers to the creation of a biomaterial's individual components, thereby generating the bioactivity.<sup>17–21</sup> Stimulus-responsive activation involves the removal of caged functional groups and self-immolative moieties. Considering the advantages of self-assembly for generating nanostructures, bottom-up fabrication has great potential for the creation of *in situ* biomaterials in living systems.



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Peptides consisting of amino acids exhibit great biocompatibility and reliable noncovalent interactions.<sup>22,23</sup> The self-assembly of peptides into well-defined nanostructures therefore has been considered as one of the versatile strategies for creating biomedical materials.<sup>24–29</sup> Due to their great biocompatibility and reliable noncovalent interactions, peptides have been widely utilized as components for the development of biomedical materials. The resulting peptide-based materials have been broadly utilized in biomedical engineering, ranging from bioimaging, drug delivery, cell culture, and tissue regeneration, to antibacterial therapy.<sup>30–34</sup> In addition, expanding knowledge about the peptide assembly mechanism has promoted the development of a considerable number of strategies to manipulate the self-assembly of peptides under various conditions.<sup>35–42</sup> While either external or internal stimuli could be employed as the sources to control peptide assembly, stimulus-responsive reactions upon amino acid sequences or individual residues enrich the approaches to efficiently modulate the assembly driving forces for peptides.<sup>43,44</sup> Hence, the formation of assembled peptide materials with distinct structural features and bioactivities in living conditions becomes possible. The versatility of peptide-based biomedical materials leads to a vast space for the development of next-generation biomaterials and precise medication.

*In situ* peptide biomaterials refer to biomaterials formed at targeted sites caused by appropriate stimuli. Only considering the biomaterials utilized for body healthcare, the targeting sites could be defined under different aspects. At the macroscale level, the targeting sites could be on the skin or inside the body; while at the microscale level, the targeting sites could be classified into the surface or interior of tissues; and at the cellular level, the *in situ* biomaterials could be extracellularly or intracellularly formed. However, peptide-based materials in living cells usually originate from either the internalization of *ex vivo*-assembled biomaterials or *in situ* assembly. On the basis of these considerations, *in situ* peptide biomaterials and peptide-based materials assembled in living cells are different, while exhibiting some overlap between each other. Analogous to *ex vivo* formed materials, combining an *in situ* assembling propensity with antimicrobial activity can lead to the *in situ* formulation of peptide biomaterials with great potential in various areas, such as antibacterial agents.

This review presents an overview of the progress in the self-assembly of peptides in living cells toward biomedical agents that have been achieved within the past few years. Starting with a brief introduction of biomedical materials, we introduce the design principles for peptide assembly in living cells with respect to the stimulus sources, bioactivity generation, and assembly regulation. Afterwards, the corresponding applications of peptide-based biomaterials formed in living environments, predominately including bioimaging, disease diagnosis, and treatment, will be highlighted. Eventually, the challenges faced in the self-assembly of peptides in living cells and the prospects of novel peptide assembly systems toward next-generation biomaterials will be also discussed, thereby helping elucidate their great potential for healthcare in the future.

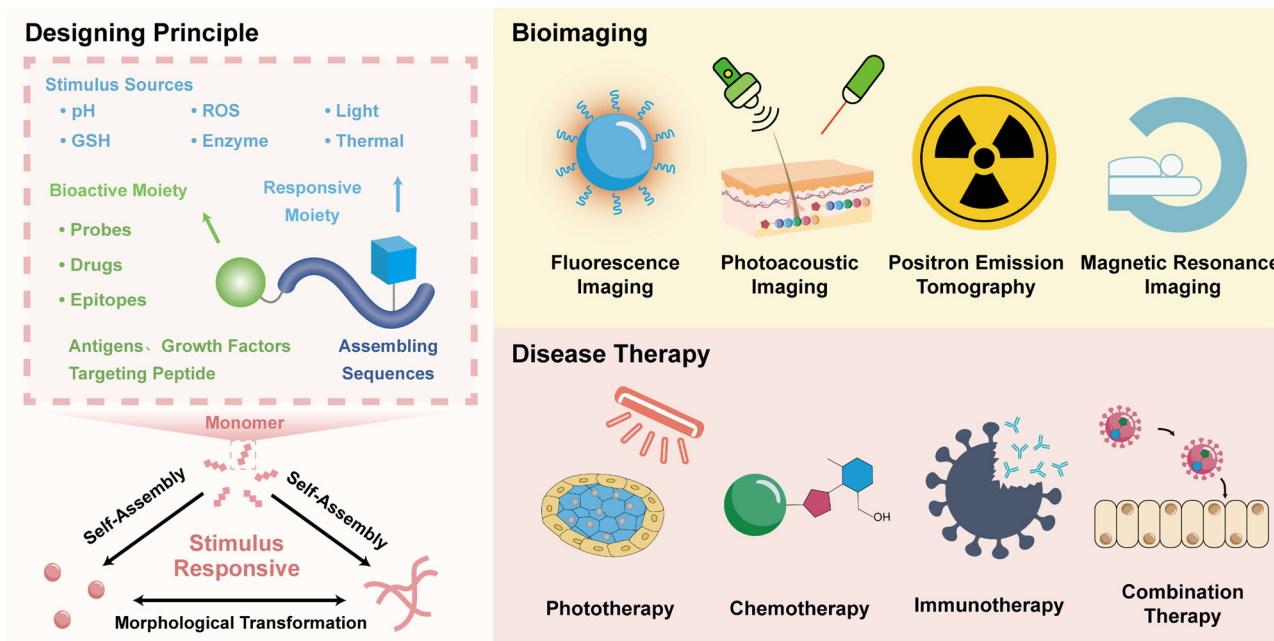
## 2. Self-assembly of peptides in living cells

The self-assembly of peptides could be manipulated by external and internal stimuli,<sup>45</sup> precisely forming biomedical materials.<sup>46</sup> Thus far, various stimulus-responsive strategies, including protonation, cleavage, cross-linking, redox, and isomerization promoted by enzymes, photons, acids, ROS, and glutathione (GSH), have been developed to establish peptide controllable assembly systems.<sup>47–49</sup> These responsive reactions provide various options for the formulation of biomedical materials under living conditions.<sup>50–52</sup> The *in situ* formation of peptide-based biomaterials is primarily based on stimulus-assembly (including the formation of less-ordered aggregates) and stimulus-responsive morphological transformation (Scheme 1).

The responsive assembling systems undergo stimulus-responsive assembly prompted by precursor monomers, in which the precursors are converted to assembling monomers and subsequently assemble into bioactive assemblies for disease theranostics. On the contrary, morphological transformation systems are based on biomaterials undergoing a stimulus-responsive morphological transition. Peptide assemblies exhibit morphological diversity and they may be present in diverse forms, including nanoparticles, nanofibers, nanoribbons, and nanotubes. Upon exposure to stimuli, these assemblies can undergo morphological transition, leading to structural changes and thereby affecting the assembly driving forces. Therefore, these assembly systems allow for the development of assembled peptide biomaterials *via* stimulus-responsive morphological transformation, which is distinct from the case of peptide materials formed through assembly from monomers. Thus far, there are several reference studies that have reported such kinds of peptide materials exhibiting responsive morphological transformation.<sup>53–55</sup>

The design of peptide assembly systems for biomedical materials is predominately based on the incorporation of stimulus-responsive and functional moieties into peptides. The stimulus-responsive units allow for the formulation of materials at targeting sites, while the functional moieties account for the bioactivity, which many be either simultaneously generated during the self-assembly or activated through specific processes.<sup>29,53</sup> The stimulus-induced self-assembly or morphological transformation of peptides is usually promoted by a sequence cleavage or by reactions at the side chains.<sup>55</sup> For instance, the selective hydrolysis of amino acid sequences by enzymes allows for regulating the assembly *via* sequence cleavage, while stimulus-responsive reactions on the side chains of amino acids provide the versatility for the assembly of peptides in living cells. Accompanied with natural amino acids, several noncanonical amino acids have been synthesized and incorporated in peptide sequences.

The incorporation of functional moieties in biomedical materials depends on the healthcare objectives. In most cases, imaging probes, drugs, or epitopes as targeting moieties and growth factors have been covalently attached with peptide



**Scheme 1** Schematic illustration of the self-assembly of peptides in living cells toward biomedical materials with an emphasis on the design principles and applications in bioimaging and disease therapy.

sequences and eventually integrated into materials utilized in tissue engineering, bioimaging, and disease therapy. Besides offering possibilities for attaching intrinsically bioactive moieties, stimulus-responsive functional moieties also offer the possibility for the *in situ* activation of biomaterials, particularly for bioimaging agents. In some special cases, the bioactivity of materials could be enhanced due to formation of ordered nanostructures. In the following sections, we outline some examples of peptide assembly in living cells according to biomedical applications in bioimaging and disease therapy with an emphasis on the design principles for materials formulation and activation of the biomedical functions.

### 3. Peptide assembly in living cells for bioimaging

Self-assembled peptide probes, which can be classified as either “always-on” or “responsive-activated” types, are considered as new bioimaging agents due to their advantages for addressing the need for accumulation and retention at the target, as well as enhancing signal intensity.<sup>56</sup> “Always-on” peptide probes generate fluorescent signals through accumulation and retention at the targeting sites.<sup>57</sup> For example, Pu’s group developed semiconducting polymer nanoprobe for protein sulfonic acids without the influence of fluorescence and PA properties.<sup>58</sup> In contrast, “responsive-activated” probes can interact with the detecting targets to activate fluorescence emission. Regardless of the type of self-assembled peptide probes, stimulus-responsive assembly processes can contribute to optimizing the pharmacokinetic parameters and avoid the rapid degradation of peptide probes. Self-assembly into

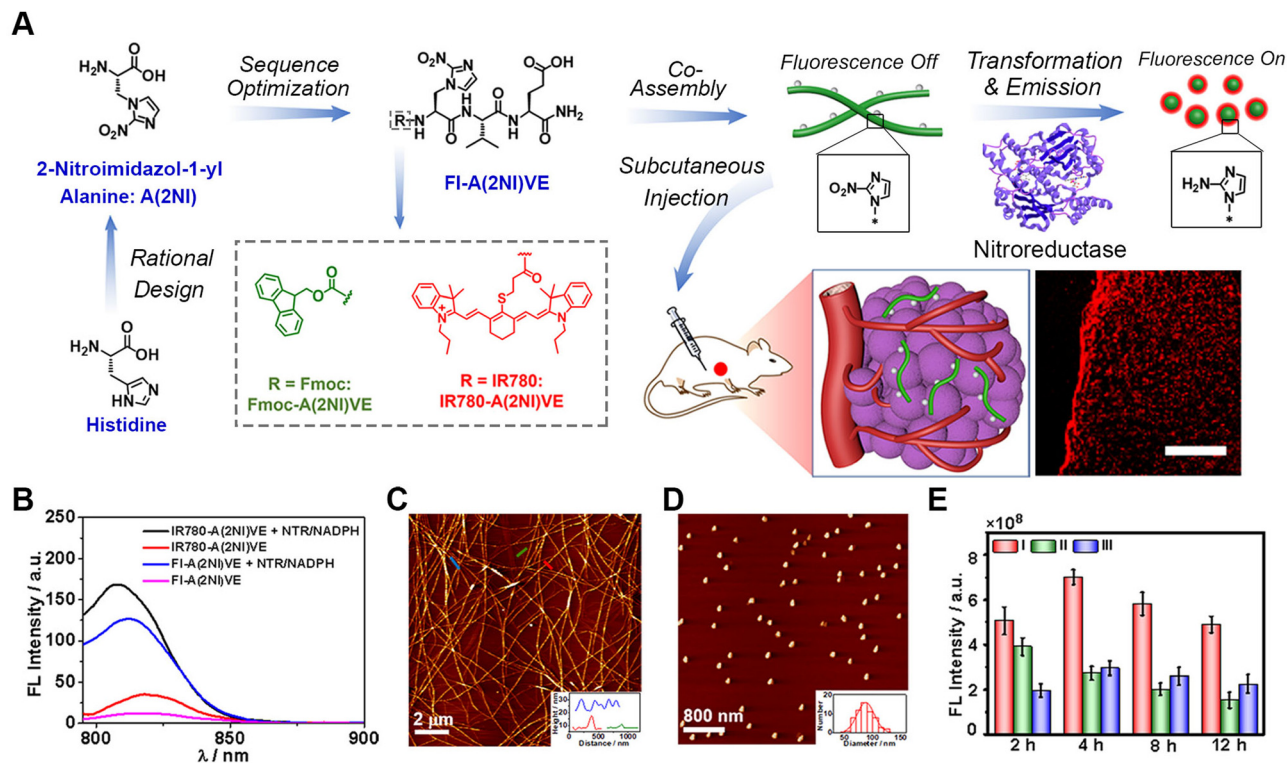
ordered supramolecular structures can also potentially amplify the signal intensity, thereby improving signal-to-noise ratios.<sup>59–64</sup> In addition, the stimulated-activation of peptide probes allows for the improvement of imaging sensitivity and for the real-time monitoring of physiological information in response to intrinsic biomarkers at pathological sites.<sup>65,66</sup> This section discusses peptide self-assembly in living cells with bioimaging functions based on various imaging models, ranging from fluorescence, to photoacoustic, and magnetic resonance, as well as multi-model systems.

#### 3.1. Fluorescence imaging

Fluorescence imaging is one of the most conventional methods for bioimaging due to its affordability, simplicity, and capability for non-invasive and minimally invasive surgeries.<sup>67–71</sup> The fluorescence imaging of pathological tissues depends on the precise generation of fluorescence signals at specific sites, for example fluorescence generation in living cells.

Recently, our group designed and synthesized an enzyme-activated peptide fluorescence probe (FI-A(2NI)VE) that can undergo morphological transformation in the tumour hypoxic region induced by nitroreductase (NTR) (Fig. 1),<sup>72</sup> which is a typical reductase overexpressed in solid tumour hypoxic regions.<sup>73</sup> Both the morphological transformation and fluorescence activation were promoted in the living environment by the NTR-reduction of a noncanonical amino acid 2-nitroimidazol-1-yl alanine [A(2NI)]. Conversion of the hydrophobic nitro group into a hydrophilic amino group within the amino acid simultaneously weakened the assembling driving forces and prevented fluorescence quenching. By rationally incorporating A(2NI) into the peptide sequence, the FI-A(2NI)VE probe





**Fig. 1** Nitroreductase-induced enzyme-activated transformable supramolecular probes for fluorescence imaging in the tumour hypoxic region. (A) Schematic illustration of the design of NTR-responsive amino acid A(2NI) and the mechanism for efficient hypoxia fluorescent imaging by the morphology-transformable fluorescent supramolecular probe FI-A(2NI)VE. Scale bar: 200 μm. (B) Fluorescence spectra of peptide IR780-A(2NI)VE and probe FI-A(2NI)VE in the presence or absence of NTR (excitation: 780 nm). AFM images of FI-A(2NI)VE (C) and FI-A(2NH<sub>2</sub>)VE (D). (E) Quantitative *in vivo* fluorescence intensity of tumour-bearing mice treated with FI-A(2NI)VE (I), FI-A(2NH<sub>2</sub>)VE (II), or FI-A(2NI)VE + dicoumarin (III). Reproduced with permission from ref. 72. Copyright 2021, American Chemical Society.

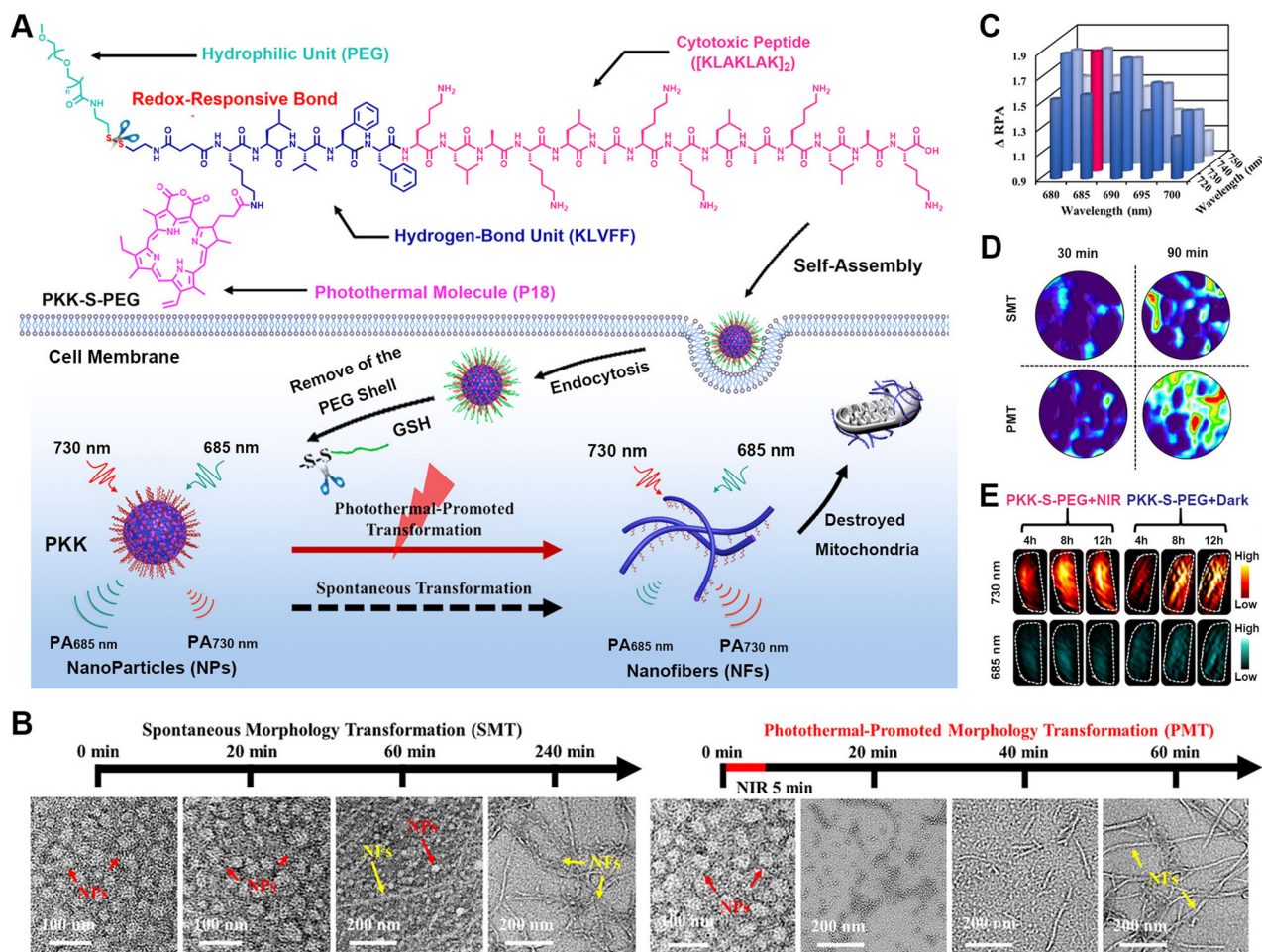
was created *via* co-assembling the peptide A(2NI)VE with the dye-labelled derivative IR780-A(2NI)VE. *In vivo* results demonstrated that the FI-A(2NI)VE probe could efficiently accumulate at the periphery of the solid periphery compared to spherical probes and could also penetrate into the deep region of solid tumours compared to the morphology-persistent fibrous probes. This hypoxic-activated morphological transformable peptide-based fluorescence probe provides a new approach for the efficient imaging of solid tumours, and may potentially be particularly useful for application with tissues with rapid metabolizing characteristics.

### 3.2. Photoacoustic imaging

To overcome the limited tissue penetration and signal interference drawbacks of fluorescence imaging, photoacoustic imaging (PAI) has been developed based on a photoacoustic effect, in which input light energy is converted into heat, subsequently leading to transient thermoelastic expansion and wideband ultrasonic emission.<sup>74–76</sup> The resulting ultrasonic waves are then detected by transducers and analysed to generate PA images. Compared to other imaging modes, PAI combines the high contrast of optical imaging with the good tissue penetration of ultrasonic waves, thus allowing it to fill the scale gap between microscopic and macroscopic imaging techniques. Therefore, in the past few years, many PAI contrast

agents have been developed for *in vivo* PAI imaging.<sup>77–81</sup> Among these probes, responsive-formed peptide-based PAI probes exhibit real-time, non-invasive, and continuous monitoring features and are therefore considered a promising strategy for designing intelligent *in vivo* probes.<sup>82–84</sup>

In addition to responsive formation, peptide-based PA probes could also undergo reactive morphological transformation to modulate the bioactivity and signal intensity.<sup>85–87</sup> Wang and colleagues designed a photothermal-promoted morphological transformation (PMT) strategy to accelerate the responsive formation of polymer-peptide conjugates into ratiometric PA probes (Fig. 2).<sup>87</sup> The polymer-peptide conjugate consisted of a mitochondrial-targeting therapeutic peptide [KLAKLAK]<sub>2</sub>, an assembly segment KLVFF, a polymer-peptide conjugate PKK-S-PEG, a GSH-responsive unit tailored with hydrophilic PEG, and a photothermal/optoacoustic motif purpurin-18 (P18). While the conjugate formed nanoparticles coated with hydrophilic PEG tails, GSH-induced cleavage of disulphide bonds led to the formation of nanofibers driven by the strong assembling propensity of KLVFF. Upon exposure to light irradiation, the morphological transformation from nanoparticles to nanofibers was accelerated by a photothermal effect associated with P18. The morphological transformation also altered the molecular arrangement of P18 along the nanofibers, thus leading to a red-shift of the UV-vis absorption for P18 and remarkably



**Fig. 2** Photothermal-promoted morphological transformation (PMT) of ratiometric probes for PA imaging. (A) Chemical structure of PKK-S-PEG and the process for the photothermal-accelerated responsive transformation of nanofibers monitored by PA imaging. (B) Time-course TEM images of PKK-S-PEG treated with GSH in the dark (left) and upon NIR irradiation (right). (C) RPA difference between NFs and NPs formed by PKK-S-PEG and GSH-reduced PKK-S-PEG at different wavelengths. (D) RPA<sub>730/685</sub> images of HeLa cells treated by PKK-S-PEG in either PMT or spontaneous morphology transformation (SMT) process. (E) PA images at 730 or 685 nm in transverse sections of the tumour tissues from the mice treated with PKK-S-PEG for 4 to 12 h. Reproduced with permission from ref. 87. Copyright 2018, American Chemical Society.

different ratiometric photoacoustic (RPA) values at 730/685 nm between the nanoparticles and nanofibers. Compared to spontaneous morphology transformation, the responsive PMT-accelerated morphology transformation in living cells or at target sites was also confirmed by PA images of HeLa cells and tumour-bearing mice treated by the conjugates *via* recording the RPA signals at 730/685 nm. This work demonstrated the potential of photothermal-promoted morphology transformation as an efficient strategy for the responsive formulation of peptide-based nanostructure for biomedical engineering in the future.

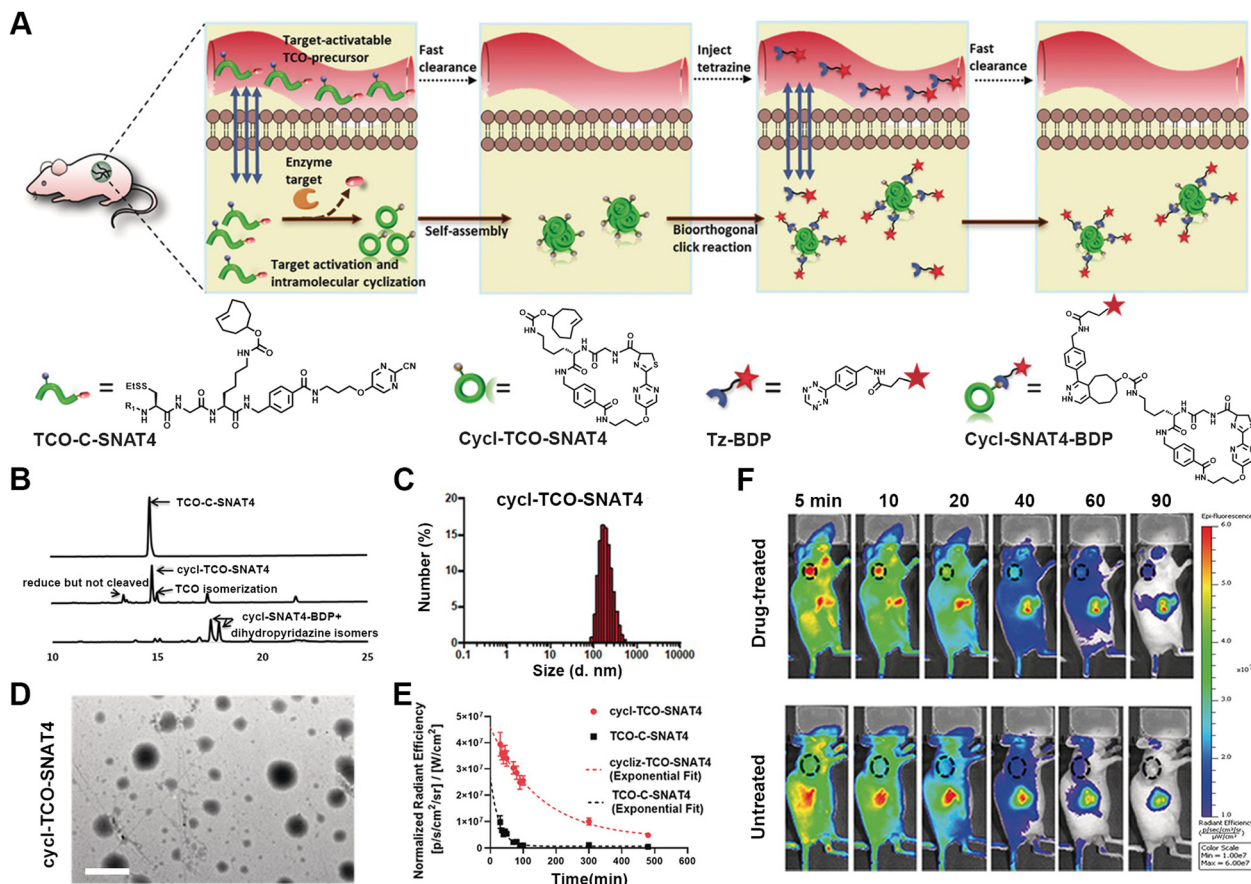
### 3.3. Positron emission tomography

Positron emission tomography (PET) is a powerful non-invasive imaging technique for disease scanning *via* utilizing radioactive probes and then collecting the annihilating radiation information arising from the probes.<sup>88–90</sup> Efficient PET imaging strongly depends on the target accumulation of the probes at

the disease sites. However, the limitation of conventional small molecular probes and nanoprobe for target accumulation significantly prevents further contrast improvement. For instance, while small-molecule probes with deep tissue penetration are rapidly metabolized, nanoprobe with high signal-to-background ratios tend to exhibit poor tissue penetration.<sup>91,92</sup> Therefore, responsive-formed PET probes have been developed, as exemplified by the hallmark work reported by Rao's group, as well as other research groups.<sup>93–95</sup>

In their early work, Rao *et al.* reported a strategy of target-enabled *in situ* ligand aggregation (TESLA) to develop responsive-formed PET tracers through a biorthogonal cyclization between 2-cyano-benzothiazole (CBT) and cysteine (Cys).<sup>96,97</sup> The PET tracer was composed of one <sup>18</sup>F-labelled tag and one DVED-caged cysteine residue and underwent caspase-3 induced cyclization and subsequent aggregation in doxorubicin-treated tumour cells, which ultimately enhanced retention of the <sup>18</sup>F-labelled tracer in cells for imaging caspase-3 activity.





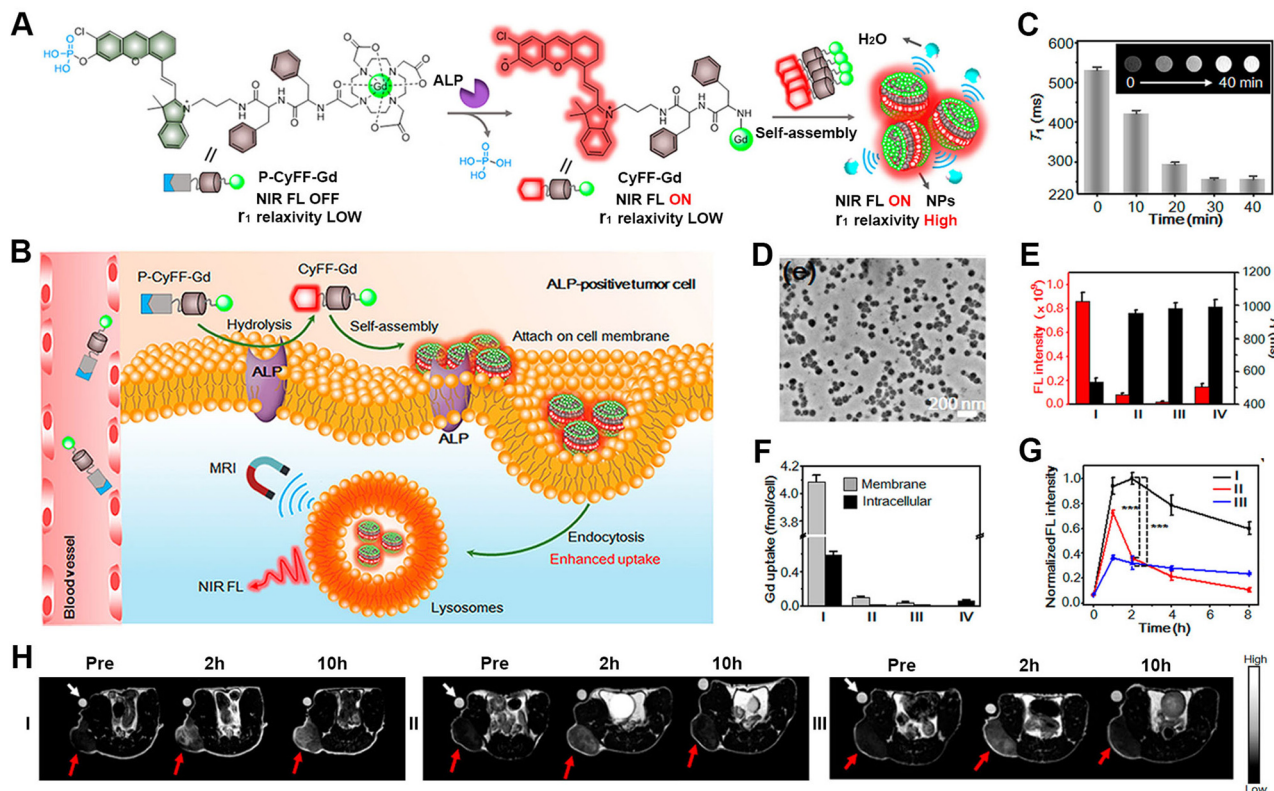
**Fig. 3** Pre-targeted PET imaging of caspase-3/7 activity by the controllable assembly of nanoparticles. (A) Illustration of the pre-target strategy for imaging enzyme activity based on reactive aggregation and the chemical structures of the corresponding ligands. (B) HPLC traces of TCO-C-SNAT4 along with the incubation with caspase-3 or addition of Tz-BDP to cycl-TCO-SNAT4. (C) DLS analysis of the nanoparticle sizes formed by cycl-TCO-SNAT4. (D) TEM image of nanoparticles of cycl-TCO-SNAT4. Scale bar: 500 nm. (E) Decay of the fluorescence signals at the injection sites of cycl-TCO-SNAT4 and TCO-C-SNAT4 over 8 h. (F) Time-course fluorescence imaging of cisplatin-pretreated H460 tumour-bearing mice injected with TCO-C-SNAT4. Reproduced with permission from ref. 99. Copyright 2020, Wiley-VCH GmbH.

A targeted-enabled responsive formulation of PET tracers was also developed by utilizing the intramolecular cyclization between CBT and disulfide-caged cysteine for enabling the imaging activity of different enzymes.<sup>98</sup> Recently, Rao and co-authors developed a pre-targeted imaging strategy for the responsive formation of PET tracers based on two biorthogonal reactions, *i.e.* the condensation between pyrimidinecarbonitrile and cysteine and the inverse-electron demand Diels–Alder (IEDDA) reaction between tetrazine (Tz) and *trans*-cyclooctene (TCO) (Fig. 3).<sup>99</sup> The precursor molecule TCO-C-SNAT4 consisted of a TCO moiety as the handle to rapidly capture imaging probes and could be converted into Cycl-TCO-SNAT4, which underwent responsive aggregation into TCO-labelled nanoparticles. Subsequently, either the fluorescence tag Tz-BDP or PET imaging tag <sup>64</sup>Cu-chelated Tz-DOTA was administered to label the nanoparticles through an IEDDA reaction between Tz and TCO. The caspase-3/7-promoted cleavage of the caged segment and responsive formation of nanoparticles were thoroughly confirmed by a combination of HPLC, dynamic light scattering, and TEM studies. *In vivo* experiments demonstrated the pre-targeted imaging strategy for both the fluorescence and

PET imaging of caspase-3/7 activity, due to the decoupling of enzyme activation and imaging tag immobilization.

### 3.4. Magnetic resonance imaging

Magnetic resonance imaging (MRI) is based on the principles of magnetic resonance to generate *in vivo* signals with high resolution and deep tissue penetration.<sup>100–103</sup> Due to its mature and robust characteristics, MRI has become a powerful diagnostic tool in clinical medicine. The responsive formulation of MRI contrast agents allows for broadening the difference between pathological and healthy tissues and thus improving the sensitivity of MRI.<sup>104–110</sup> Liang's group developed several unique approaches for the responsive formation of gadolinium (Gd)-based MRI contrast agents instructed by enzyme-catalysed reactions.<sup>111</sup> For instance, the group designed a  $\gamma$ -glutamyl-transpeptidase (GGT)-instructed CBT-Cys cyclization for the formulation of Gd-based contrast agents with enhanced  $T_2$ -weighted MRI under a high magnetic field. Upon GGT cleavage and GSH reduction, the probe, Glu-Cys(StBu)-Lys(DOTA-Gd)-CBT, underwent intermolecular CBT-Cys condensation and formed nanoparticles in GGT-overexpressed tumour cells. Both



**Fig. 4** ALP-activatable NIR fluorescence/MRI bimodal probes for *in vivo* imaging. (A) Chemical structure of P-CyFF-Gd and its ALP-induced assembly into nanoparticles exhibiting enhanced fluorescence and  $r_1$  relaxivity. (B) Mechanism of P-CyFF-Gd for the bimodality imaging of ALP-positive tumour cells *in vivo*. (C)  $T_1$ -weighted MR images and  $T_1$  values for P-CyFF-Gd upon the addition of ALP. (D) TEM image of nanoparticles collected from HeLa cell membranes incubated with P-CyFF-Gd for 1 h. (E) Average FL intensity (red) and  $T_1$  value (black) of cell pellets incubated with different treatments. (F) ICP-MS analysis of the Gd contents on the cell membrane or intracellularly after incubation with the indicated treatments. I–IV: HeLa cells incubated with P-CyFF-Gd (I), PCy-Gd (II), P-CyFF-Gd (III), IV: P-Cy-Gd (IV). (G) Normalized FL intensity of HeLa tumour-bearing mice with different treatments. (H)  $T_1$ -weighted MR images of HeLa tumour-bearing mice with different treatments. In (G) and (H), I–III: HeLa cells incubated with P-CyFF-Gd (I), PCy-Gd (II), or P-CyFF-Gd (III). Reproduced with permission from ref. 113. Copyright 2019, American Chemical Society.

*in vitro* and *in vivo* experiments confirmed the enhanced  $T_2$ -weighted MRI signal in HeLa cells and tumour-bearing mice, due to the improved cellular uptake and formation of the nanoparticles.

In addition, combining MRI with other imaging methods enables further improving the sensitivity for pathological tissue detection.<sup>112</sup> In this context, Ye's group established strategies for the responsive formulation of probes with multimodality imaging functions, including NIR fluorescence (FL)/MRI and NIR FL/MRI/PET imaging (Fig. 4).<sup>113</sup> The primary challenge for the formulation of such probes lies in integrating the stimulus-responsive self-assembling process with the signal-generating units. Therefore, the authors integrated activatable NIR and MR probes with a dipeptide motif, leading to the bimodal probe P-CyFF-Gd containing a phosphorated merocyanine dye. The probe assembled into nanoparticles at the tumour sites following an ALP-cleavage reaction within the dye, thereby amplifying and accumulating the imaging signals. In particular, the formation of nanoparticles prevented molecular rotation and generated bright  $T_1$ -weighted MR *via* increasing the  $r_1$  relaxation rate and shortening the water protons'  $T_1$  relaxation. The HeLa cell experiments and tumour-bearing mice studies

showed a robust NIR FL signal, a shortened  $T_1$  relaxation, and the enhanced fluorescence and MR contrast, thus evidencing the efficient activation of P-CyFF-Gd through ALP dephosphorylation and stimulus-responsive self-assembly.

## 4. Peptide assembly in living cells for disease therapy

The efficient delivery of therapeutic agents to pathological sites is the basis for achieving a perfect curative effect in disease therapy. Conventional delivering vehicles have been demonstrated with limited capability to overcome the physiological barriers present in the delivery pathway, including the blood flow, tumour cell interstitial compartment, and cell membranes.<sup>114–117</sup> The application of stimulus-activated peptide assemblies towards drug delivery allows for overcoming the physiological barriers in delivery *via* the morphological transforming or assembling process.<sup>118–121</sup> For example, the assembly process bestows peptide-based nanomedicines with stabilization during blood circulation, whereas the transformation from nanofibers into small nanoparticles facilitates tumour



penetration.<sup>122</sup> Therefore, due to the precise control of stimulus-responsive self-assembly or morphological transformations, self-assembled peptide nanostructures in living cells have great potential for application in various disease treatments.<sup>123,124</sup> This section discusses several recent examples utilizing stimulus-responsive self-assembled peptides in different therapeutic methods, including phototherapy, chemotherapy, immunotherapy, and combination therapy.

#### 4.1. Phototherapy

Phototherapy predominately includes photodynamic therapy (PDT) and photothermal therapy (PTT).<sup>125–129</sup> With respect to PDT, the exposure of photosensitizers to light leads to the production of ROS, such as hydrogen peroxide ( $H_2O_2$ ) and hydroxyl radicals ( $HO^\bullet$ ) that usually elicit cell death *via* the oxidation of biomolecules.<sup>130</sup> In contrast, the activation of photosensitizers utilized in PTT by light irradiation releases

vibrational energy, rather than transferring into ROS species, and subsequently kills cells by local heating. However, high temperature can also cause thermal damage to adjacent healthy tissues. Therefore, mild-temperature PTT strategies that minimize the associated adverse effects have been developed, but suffer from limited therapeutic efficacy or the thermoresistance of cancer cells.<sup>131</sup> This can be overcome by combining PTT with other therapeutic modalities, such as immunotherapy or chemotherapy, which is discussed later in the combination therapy section.<sup>132–136</sup> It is worth noting that within phototherapy, the integration of photosensitizers with peptide assemblies not only facilitates the target delivery of photosensitizers, it most probably also enhances the production of ROS or heat energy through organization within peptide nanostructures.<sup>137–141</sup>

As an interesting example, our group developed reactive transformable drug delivering vehicles for PDT tumour therapy, termed as tumour microenvironment-adaptable self-assembly

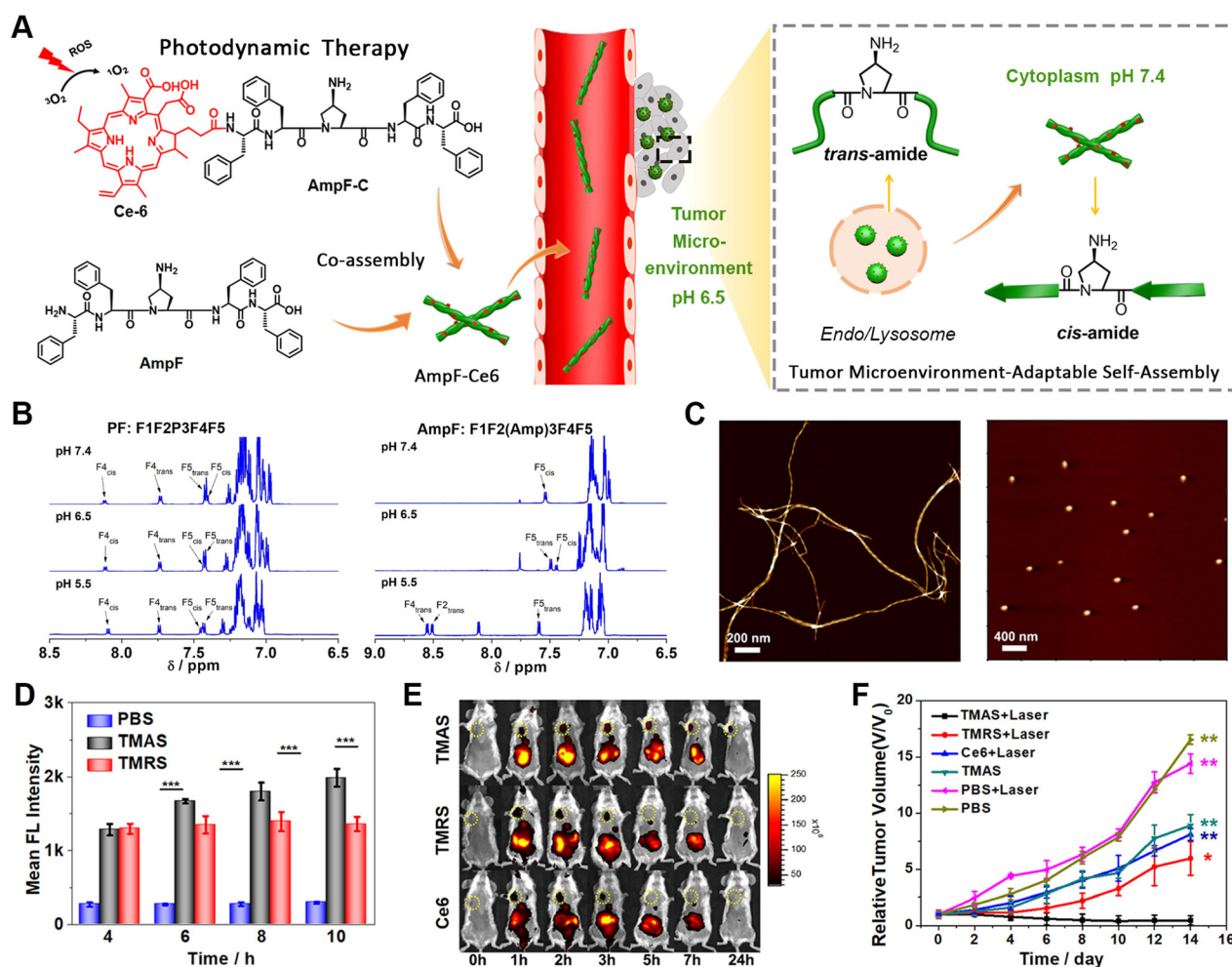


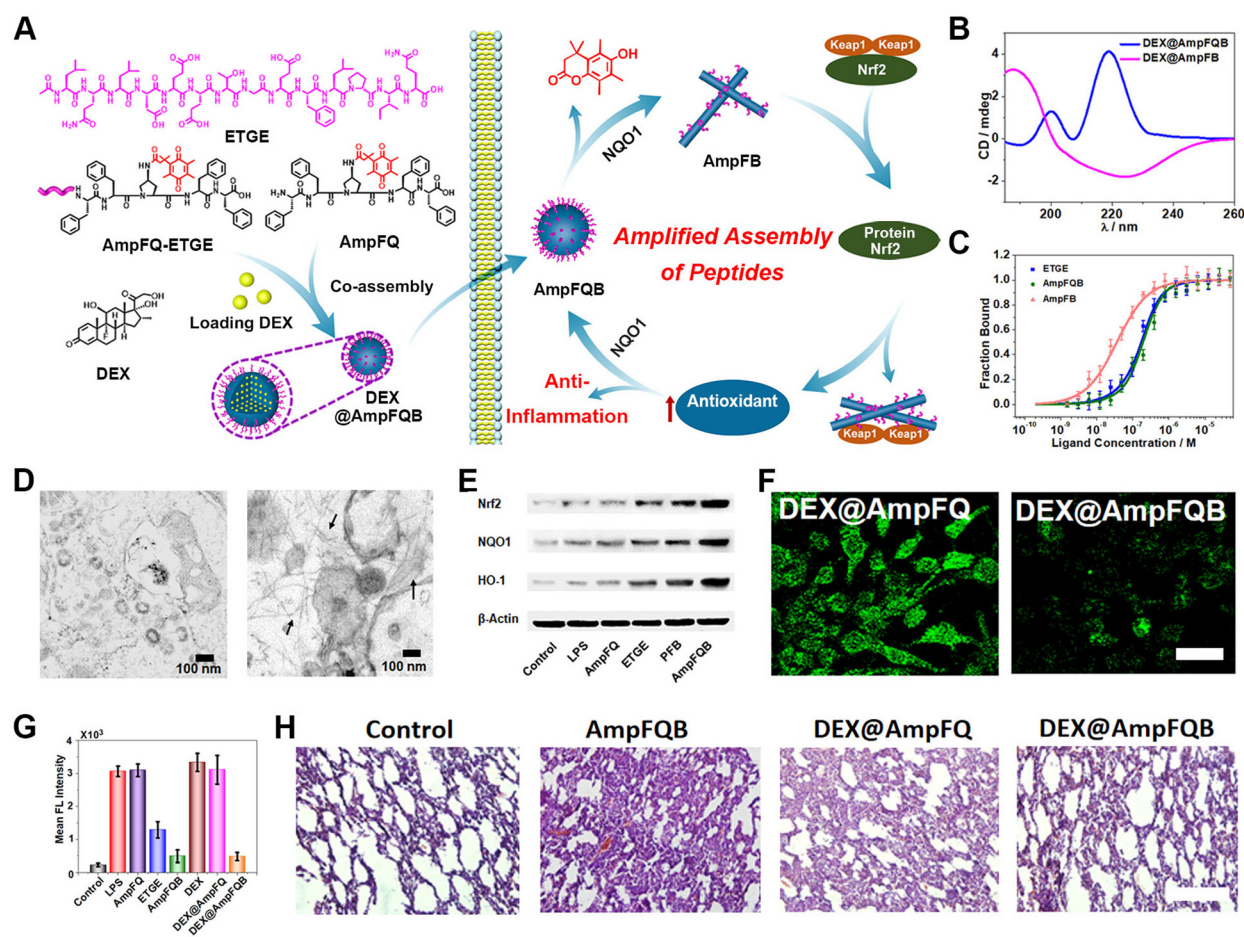
Fig. 5 (A) Schematic illustration of the tumour microenvironment-adaptable self-assembly (TMAS) of pentapeptide AmpF and its co-assembly with Ce6-modified AmpF (AmpF-C) into the nanomedicine TMAS, in which AmpF underwent a reversible morphological transition between nanofibers and nanoparticles due to the acid-sensitive *cis/trans*-isomerization of Amp amide bonds. (B)  $^1H$  NMR spectra of the peptides PF (left) and AmpF (right) under various pH conditions. (C) AFM images of the assemblies formed by AmpF at pH 7.4 (left) and 6.5 (right). (D) Quantification of intracellular ROS generation upon the laser irradiation of TMAS and TMRS by 4T1 cells with time. (E) *In vivo* fluorescence imaging of 4T1 breast tumour-bearing mice administrated with TMAS, TMRS (formed by PF and its PF-C derivative), and Ce6. (F) Relative tumour volume change in tumour-bearing mice upon different treatments. Reproduced with permission from ref. 142. Copyright 2019, American Chemical Society.

(TMAS), based on the reversible acid-responsive self-assembly of the pentapeptide AmpF (Fig. 5).<sup>142</sup> While AmpF contains a central 4-aminoproline (Amp) residue and two diphenylalanine fragments on each termini, TMAS is composed of AmpF and its derivative (AmpF-C) modified with the photosensitizer Ce6. The Xaa-Amp amide bond in AmpF adopts a *cis*- or *trans*-conformation under neutral and acidic conditions, respectively, thus resulting in the pH-dependent self-assembly of AmpF into superhelices and nanoparticles. Analogous to AmpF, TMAS exhibited a similar reversible morphological transformation between superhelices and nanoparticles triggered by alternating exposure to neutral and acidic environments. This unique pH-responsive property renders the morphology of TMAS adaptable to the pH gradient during delivery pathway and can eventually improve the accumulation of photosensitizers in cancer cells. Both cellular and animal experiments confirmed the enhanced cellular internalization, tumour accumulation, penetration, and retention of photosensitizers due to the

adaptable morphology of TMAS, accompanied by improved ROS generation. These results demonstrate the great potential of TMAS for efficient drug delivery to tumour sites for cancer therapy.

#### 4.2. Chemotherapy

Despite being one of most conventional disease-treatment methods, chemotherapy suffers from drug resistance and adverse effect arising from off-target activity.<sup>143–146</sup> Self-assembling peptide-based biomedical materials in living cells represents one promising approach to address these challenges.<sup>147–151</sup> Due to the capability of peptide nanostructures, including nanoparticles, nanofibers, and nanofibrous networks, to interfere in physiological processes, stimulus-activatable peptide assemblies have been directly utilized as therapeutic agents.<sup>152–157</sup> Xu's group and other colleagues performed pioneering work on rationally controlling the pericellular or intracellular self-assembly of peptides.<sup>158–160</sup>



**Fig. 6** NQO1-activated self-amplifying assembly of peptides in macrophages. (A) Self-amplifying assembly of peptide AmpFQB due to the amplified relationship between peptide assembly and enzyme expression, and its application as a delivery vehicle for dexamethasone (DEX) for improving inflammatory therapy. (B) CD spectra of DEX@AmpFQ before after QPA reduction. (C) MST studies on the free peptide ETGE, AmpFQB, and AmpFB with the protein Keap1. (D) Bio-TEM images of stimulated macrophages incubated with AmpFQ (left) and AmpFQB (right). (E) WB assays of the expression of Nrf2, NQO1, and HO-1 in macrophages treated with AmpFQB and other groups. (F) CLSM images of stimulated macrophages incubated with various treatment stained with DCFH-DA. Scale bar: 10  $\mu$ m. (G) Quantitative fluorescence intensity associated with DCFH-DA in the treated macrophages. (H) H&E-stained images of lung tissues dissected from injured mice with various treatments. Scale bar: 40  $\mu$ m. Reproduced with permission from ref. 161. Copyright 2022, American Chemical Society.



Dexamethasone (Dex) is a chemical agent commonly used for cancer therapy. Here, we used intracellular peptide materials in combination with Dex for inflammation treatment, broadening the chemotherapy scope for disease treatment. Recently, our group reported the development of an stimulus-activated self-amplifying assembling system, AmpFQB, for inflammation therapy based on the robust relationship between the self-assembly of AmpFQB and the expression of the enzyme NAD(P)H quinone dehydrogenase 1 (NQO1) (Fig. 6).<sup>161</sup> AmpFQB consists of quinone propionate (QPA)-modified pentapeptide AmpF (AmpFQ) and its derivative AmpFQ-ETGE, which contains an ETGE domain. NQO1 is a critical cytosolic reductase mediating redox homeostasis in cells, which is regulated by protein nuclear factor erythroid-related factor 2 (Nrf2).<sup>162,163</sup> Containing the ETGE sequence, which was derived from Nrf2, AmpFQB coassembles compete with Nrf2-Keap1 complexes and release Nrf2, thereby promoting NQO1 expression.<sup>164</sup> While the modification of AmpFQ with QPA supports its NQO1 responsive conversion to AmpF, the ETGE sequence derived from the Nrf2 domain

enables binding with the Keap1 receptor and the release of Nrf2 from Keap1-Nrf2 complexes. The liberated Nrf2 increases the level of NQO1 and thus conversely contributes to the transition from AmpFQB nanoparticles to AmpFB nanofibers. Microscale thermophoresis (MST) revealed a four-fold affinity of AmpFB nanofibers to Keap1 compared to AmpFQB nanoparticles or ETGE. Combining the results from the WB assay of the intracellular protein levels and bio-TEM images of macrophages treated with AmpFQB, the amplifying relationship between the reactive assembly of AmpFQB into nanofibers and the expression of enzyme NQO1. AmpFQB could be utilized to deliver the drug dexamethasone (DEX) to treat acute injured lungs, leading to the nanomedicine DEX@AmpFQB. Both *in vitro* and *in vivo* studies confirmed the capability of DEX@AmpFQB to enhance the anti-inflammatory efficacy of DEX *via* alleviating ROS-associated side effects and downregulating the proinflammatory cytokines. The self-amplifying assembly of peptides in living cells with a regular enzyme level provides a unique strategy for the responsive creation of theranostic agents.

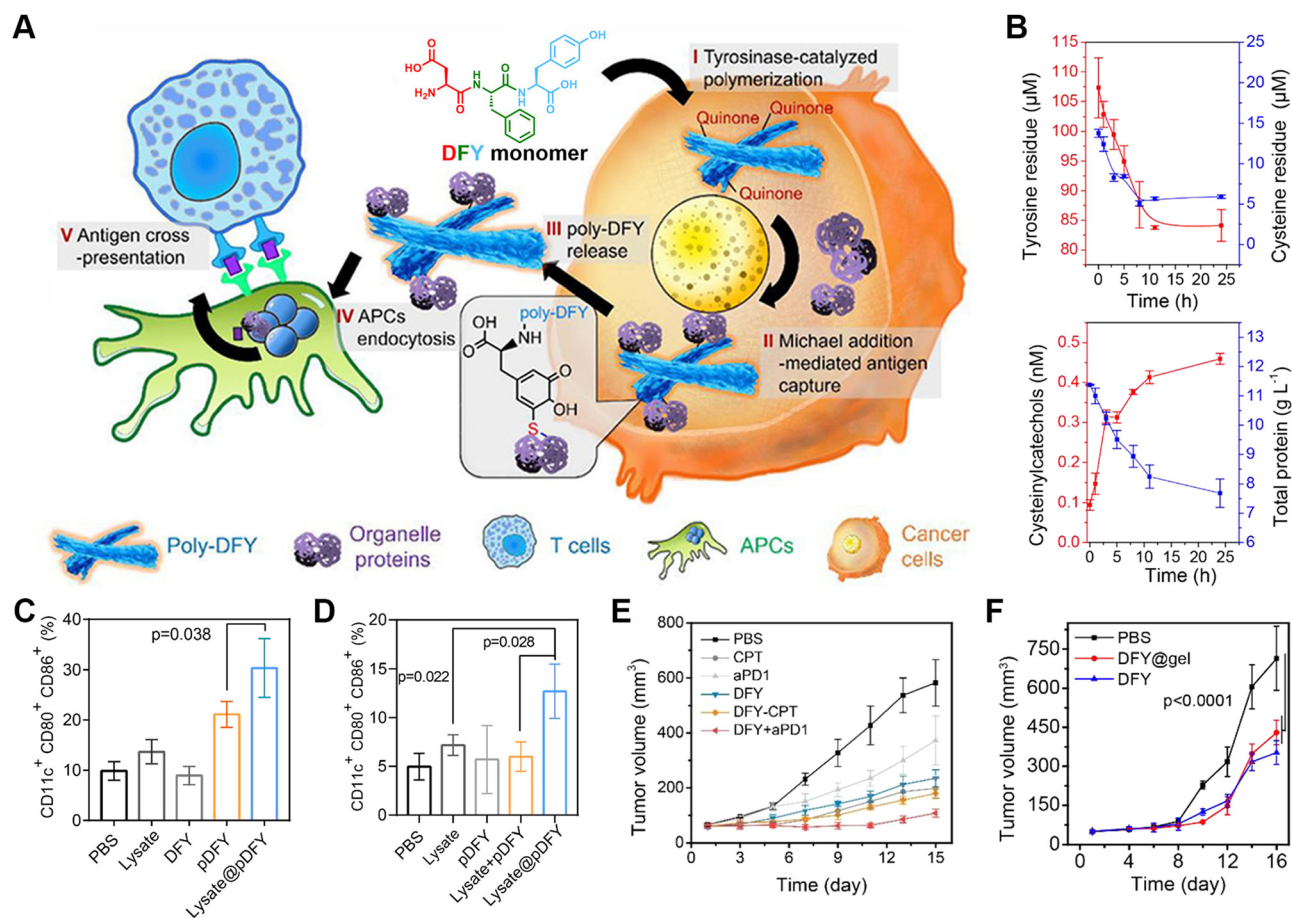


Fig. 7 Enzyme-catalyzed polymerization of tripeptide DFY for cancer immunotherapy. (A) Schematic illustration of the underlying mechanism for the tyrosinase-induced reactive polymerization of DFY and integration with antigens to form nanofibers. (B) Concentration profiles for tyrosine, cysteine, cysteinylcatechol, and proteins in the polymerizing solution of DFY and tumour lysate. (C) *In vitro* efficacy of antigen-loaded pDFY in promoting DC maturation. (D) *In vivo* effect of antigen-loaded pDFY in immunological activation. (E) Tumour volume of B16 tumour-bearing mice from different treated groups. (F) Comparison of therapeutic efficacy between injected DFY and DFY-loaded transdermal hydrogels. Reproduced with permission from ref. 173. Copyright 2021, American Chemical Society.



### 4.3. Immunotherapy

Immunotherapy is considered one of the most promising tumour-treatment strategies due to its advanced curative efficacy for both local and remote tumour tissues *via* eliciting host systemic immune responses.<sup>165–167</sup> Combining the advantages of responsive assembled peptides for targeting delivery with immunotherapy, peptide-based cancer immunotherapy represents one of main topics in disease therapy.<sup>168–172</sup> Zhang's group designed a tumour protein-engineering system based on the tyrosinase-catalysed polymerization of tripeptide DFY into quinone-rich microfibrils and a reactive collection of immunogenic tumour proteins *via* a Michael addition between quinone and cysteine (Fig. 7).<sup>173</sup> The resulting tumour-protein-captured microfibrils delivered proteins to antigen-presenting cells, which eventually boosted the immune response. Decreases in the levels of tyrosine residues and cysteine residues, as well as an increase in the content of cysteinylcatechol groups, indicated the oxidative polymerization of tyrosine residues and the addition between quinone with protein sulfhydryl groups. In comparison to the lysate from untreated tumour cells, lysate protein-cross-linked pDFY (lysate@pDFY) treatment resulted in a 2.3-fold increase in DC maturation,

while tumour-bearing mice subjected to the lysate@pDFY treatment demonstrated a 2.8-fold increase in the maturation of DCs. The co-administration of B16 tumour-bearing mice with DFY and anti-PD1 antibody suppressed tumour growth by 91% compared to monotherapy alone in other treatment groups. Furthermore, a transdermal delivery system incorporating DFY exhibited comparable therapeutic efficacy to that achieved by peritumoural DFY injection. This reactive polymerization of peptides and selective capture of tumour-specific proteins could overcome the systemic inhibition of antigen presentation, providing a promising avenue for modulating the immune system during cancer therapy.

### 4.4. Combination therapy

Given the inherent limited therapeutic efficacy and the side effects of individual treating modalities, treating diseases *via* a combination of different pharmaceutical agents may achieve synergistic therapeutic efficacy while minimizing undesirable adverse effects.<sup>174–178</sup> Some examples of such combination therapies include unitary combinations from individual therapeutic modality, binary combinations of phototherapy, chemotherapy, radiotherapy, and immunotherapy, as well as ternary or even more combinations.<sup>179–182</sup>

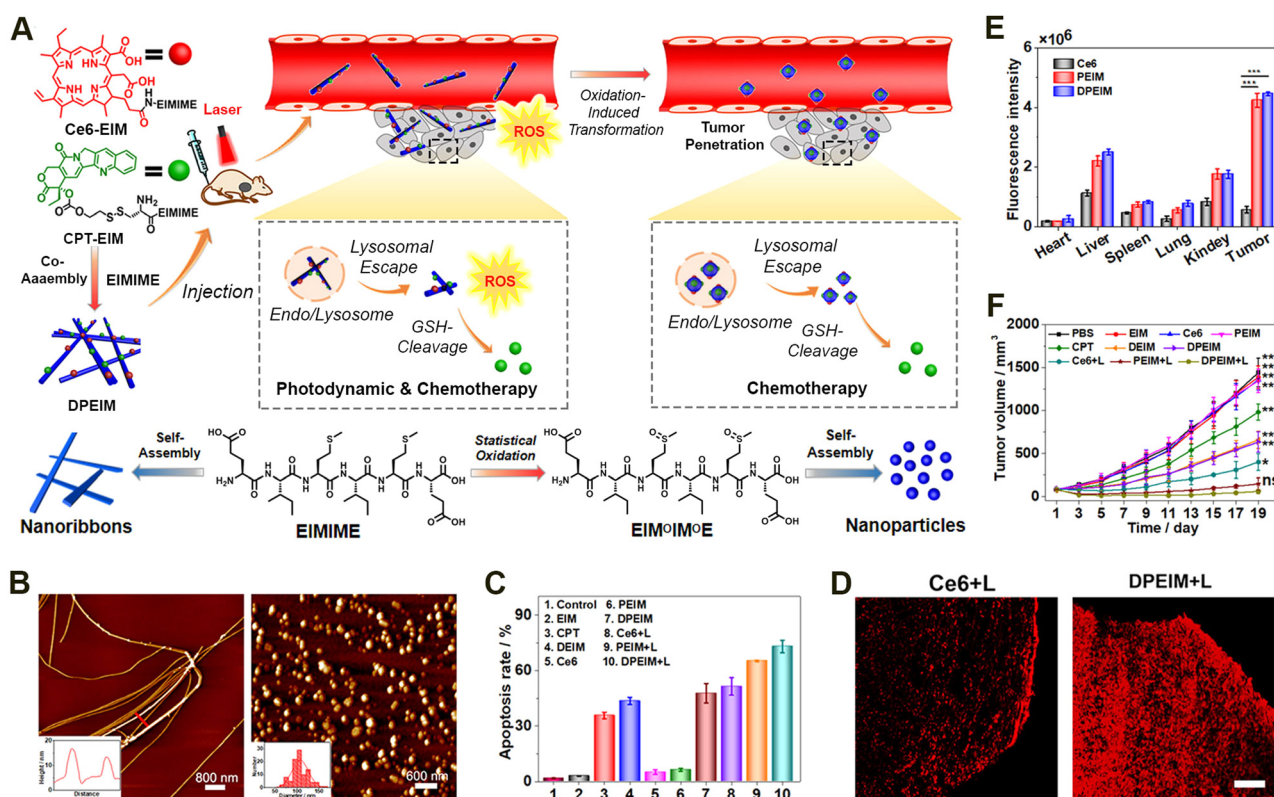


Fig. 8 Oxidation-regulated transformation of peptide nanomedicines. (A) Graphic illustration of the mechanism for the creation of the oxidation-regulated reactive transformable peptide nanomedicine DPEIM and its facilitated tumour penetration and cascade combinatorial cancer therapy. (B) AFM images of DPEIM before (left) and after (right) methionine oxidation. (C) Cellular apoptosis studies of 4T1 cells induced with different treatments estimated by flow cytometric analysis. (D) CLSM images of the transverse sections of the tumour tissues from mice administrated with free Ce6 or DPEIM in the presence of laser irradiation. Scale bar: 250 μm. (E) Quantitative fluorescence intensity of ex vivo signals at tumour sites or major organs from mice administrated with free Ce6, PEIM, and DPEIM. (F) Growth profiles for the tumour tissues of mice administrated with different treatments with or without laser irradiation. Reproduced with permission from ref. 183. Copyright 2021, Elsevier.

In terms of binary combination therapy, our group reported an reactive morphological transformation of the peptide scaffold DPEIM for implementing photodynamic and chemodrug combination cancer therapy (Fig. 8).<sup>183</sup> The oxidation-responsive peptide scaffold DPEIM was created *via* the co-assembly of hexapeptide EIMIME with its derivatives Ce6-EIM and CPT-EIM, which were functionalized with the photosensitizer Ce6 and drug CPT, respectively. EIMIME contained two methionine residues, in which the hydrophobic thioether was quantitatively converted to hydrophilic sulfoxide groups upon the addition of H<sub>2</sub>O<sub>2</sub>, and thereby promoting nanofibrillar-to-nanoparticle morphological transition for the peptide assemblies. While the GSH-cleavage of disulfide bonds led to the release of the carried drug CPT, the generation of ROS by Ce6 under laser irradiation resulted in an oxidation of methionine residues and also implemented photodynamic cancer therapy. Treating cancer cells by DPEIM led to a higher apoptosis rate compared to individual treatments. *In vivo* experiments indicated that the laser irradiation-induced morphological transformation of DPEIM improved its tumour accumulation and penetration into the tumour tissue, thereby significantly inhibiting the growth of tumour tissues. Both *in vitro* and *in vivo* studies demonstrated the effectiveness of cascade combinatorial photodynamic therapy and chemotherapy arising from CPT and Ce6 moieties within DPEIM.

## 5. Conclusions and outlook

In this review, we summarized the progress in the stimulus-responsive self-assembly of peptides into biomedical agents in living cells that have been achieved during the past decades. After a brief introduction to the research background, we discussed the design principles of stimulus-responsive peptide materials. We highlighted the achievements of stimulus-responsive peptide biomaterials for implementing disease diagnosis and therapy, providing several examples reported recently to demonstrate their great potential towards precise and real-time healthcare.

Due to the remarkable biocompatibility and robust controllable assembling propensity of peptides, the stimulus-responsive self-assembly of peptides in living cells presents a versatile strategy for the controllable formulation of biomaterials at the target site. A simultaneous incorporation of bioactive moieties into peptides usually leads to responsive-formed biomaterials for implementing healthcare objectives.<sup>46,184</sup> The bioactivity of the resulting stimulus-reactive biomaterials not only depends on the incorporated functional moieties, but is also associated with the organization and dynamic natures of the peptide assemblies.<sup>185–187</sup> In addition, the accumulation and retention of the bioactive moieties at targeting sites could be optimized by the stimulus-responsive assembly processes of peptides.

In terms of peptide assembly-based biomedical materials, the gap between the few number of stimulus-responsive peptide assembling strategies and the complex biological conditions significantly limits their biomedical applications. Hence, the development of stimulus-responsive self-assembling

systems is critical for its applications in different fields. In this context, our group intensively engaged in developing noncanonical amino acids containing stimulus-responsive moieties through organic synthesis.<sup>188</sup> The rational incorporation of these amino acids into peptides allows for establishing stimulus-responsive assembling systems for the formulation of peptide-based materials in living cells or the body. In addition, exploiting new enzymes as internal stimulus sources for regulating peptide assembly is another direction to expand researchers and designers toolkits.<sup>44</sup> This strongly depends on understanding the pathological mechanism, which reveals the biological pathways and involved enzymes corresponding to abnormal physiological behaviour. The finding of new stimulus sources usually exponentially increases the stimulus-responsive units for regulating peptide assembly. Simultaneously, the development of *in situ* characterizing approaches also facilitates a direct understanding of the assembly process in living systems, which thus far suffers from a lack of appropriate approaches to confirm the assembling process compared to solution conditions.

Despite the achieved progress in stimulus-responsive biomaterials, including peptide assembly-based systems, several critical concerns remain to be addressed with respect to the development of reactive synthetic approaches, the understanding of the underlying mechanism as well as the need for preclinical studies to pave the way towards their clinical usage. Thus far, the approaches available for obtaining stimulus-responsive materials are confined to a few number of versatile methods that are broadly utilized under solution conditions. The utilization of solution methods in responsive biomedical materials studies has been significantly hindered by the limited accessibility of the internal biological structures and signals or the low biocompatibility of the conditions. For instance, mass spectroscopy and scattering techniques could potentially facilitate elucidating the biological structures from the molecular and microscopic levels, respectively.<sup>189–194</sup> The stimulus-responsive formulation of biomaterials in living environments might benefit from the development of synthetic approaches involving novel physical phenomena. For example, liquid droplet formation in living cells based on liquid–liquid phase transition provides a new approach for the responsive formulation of biomaterials at targeting sites.<sup>195,196</sup> In addition, due to the versatility of machine learning in scanning functional monomers, designing reactive biomaterials based on machine learning is a potentially powerful method to enrich experimental findings.<sup>197,198</sup>

## Conflicts of interest

There are no conflicts to declare.

## Acknowledgements

This work was financially supported by the Haihe Laboratory of Synthetic Biology (22HHSWSS00010), the Beijing-Tianjin-Hebei Basic Research Cooperation Project (22JCZXC00010), and the

National Natural Science Foundation of China (52073148 & 52273130).

## Notes and references

- 1 Y. Fang, L. Meng, A. Prominski, E. N. Schaumann, M. Seebald and B. Tian, *Chem. Soc. Rev.*, 2020, **49**, 7978–8035.
- 2 O. Valenzuele, M. Cannataro, I. Rusur, J. Wang, Z. Zhao and I. Rojas, *MBC Bioinformatics*, 2023, **24**, 361.
- 3 N. Huebsch and D. J. Mooney, *Nature*, 2009, **462**, 426–432.
- 4 M. J. Webber, E. A. Appel, E. W. Meijer and R. Langer, *Nat. Mater.*, 2016, **15**, 13–26.
- 5 F. Ajallouecian, G. Lemon, J. Hilborn, I. S. Chronakis and M. Fossum, *Nat. Rev. Urol.*, 2018, **15**, 155–174.
- 6 D. J. Lipomi, M. Vosgueritchian, B. C. K. Tee, S. L. Hellstrom, J. A. Lee, C. H. Fox and Z. N. Bao, *Nat. Nanotechnol.*, 2011, **6**, 788–792.
- 7 S. Mura, J. Nicolas and P. Couvreur, *Nat. Mater.*, 2013, **12**, 991–1003.
- 8 M. Karimi, A. Ghasemi, P. Sahandi Zangabad, R. Rahighi, S. M. Moosavi Basri, H. Mirshekari, M. Amiri, Z. Shafaei Pishabad, A. Aslani, M. Bozorgomid, D. Ghosh, A. Beyzavi, A. Vaseghi, A. R. Aref, L. Haghani, S. Bahrami and M. R. Hamblin, *Chem. Soc. Rev.*, 2016, **45**, 1457–1501.
- 9 P. Yang, F. Zhu, Z. Zhang, Y. Cheng, Z. Wang and Y. Li, *Chem. Soc. Rev.*, 2021, **50**, 8319–8343.
- 10 H. J. He, W. Y. Tan, J. Q. Guo, M. H. Yi, A. N. Shy and B. Xu, *Chem. Rev.*, 2020, **120**, 9994–10078.
- 11 J. Kim, S. Lee, Y. Kim, M. Choi, I. Lee, E. Kim, C. G. Yoon, K. Pu, H. Kang and J. S. Kim, *Nat. Rev. Mater.*, 2023, **8**, 710–725.
- 12 X. Wu, R. Wang, N. Kwon, H. Ma and J. Yoon, *Chem. Soc. Rev.*, 2022, **51**, 450–463.
- 13 J. R. Sempionatto, J. A. Lasalde-Ramírez, K. Mahato, J. Wang and W. Gao, *Nat. Rev. Chem.*, 2022, **6**, 899–915.
- 14 A. K. Gaharwar, I. Singh and A. Khademhosseini, *Nat. Rev. Mater.*, 2020, **5**, 686–705.
- 15 D. Wu, J. Lei, Z. Zhang, F. Huang, M. Buljan and G. Yu, *Chem. Soc. Rev.*, 2023, **52**, 2911–2945.
- 16 H. Qin, T. Zhang, N. Li, H.-P. Cong and S.-H. Yu, *Nat. Commun.*, 2019, **10**, 2202.
- 17 Z. Ge and S. Liu, *Chem. Soc. Rev.*, 2013, **42**, 7289–7325.
- 18 Y. Lu, A. A. Aimetti, R. Langer and Z. Gu, *Nat. Rev. Mater.*, 2016, **2**, 16075.
- 19 L. Cui, S. Vivona, B. R. Smith, S.-R. Kothapalli, J. Liu, X. Ma, Z. Chen, M. Taylor, P. H. Kierstead, J. M. J. Fréchet, S. S. Gambhir and J. Rao, *J. Am. Chem. Soc.*, 2020, **142**, 15575–15584.
- 20 E. Fuentes, M. Gerth, J. A. Berrocal, C. Matera, P. Gorostiza, I. K. Voets, S. Pujals and L. Albertazzi, *J. Am. Chem. Soc.*, 2020, **142**, 10069–10078.
- 21 H. Yan, Y. Wang, F. Huo and C. Yin, *J. Am. Chem. Soc.*, 2023, **145**, 3229–3237.
- 22 C. J. Bowerman and B. L. Nilsson, *Biopolymers*, 2012, **98**, 169–184.
- 23 S. Zhang, *Interface Focus*, 2017, **7**, 20170028.
- 24 J. Zhou, X. Du, C. Berciu, H. He, J. Shi, D. Nicastro and B. Xu, *Chem*, 2016, **1**, 246–263.
- 25 S. Eskandari, T. Guerin, I. Toth and R. J. Stephenson, *Adv. Drug Delivery Rev.*, 2017, **110–111**, 169–187.
- 26 G. Wei, Z. Su, N. P. Reynolds, P. Arosio, I. W. Hamley, E. Gazit and R. Mezzenga, *Chem. Soc. Rev.*, 2017, **46**, 4661–4708.
- 27 F. Sheehan, D. Sementa, A. Jain, M. Kumar, M. Tayarani-Najjaran, D. Kroiss and R. V. Ulijn, *Chem. Rev.*, 2021, **121**, 13869–13914.
- 28 J. A. Raskatov, J. P. Schneider and B. L. Nilsson, *Acc. Chem. Res.*, 2021, **54**, 2488–2501.
- 29 S. Panja and D. J. Adams, *Chem. Soc. Rev.*, 2021, **50**, 5165–5200.
- 30 R. V. Ulijn and A. M. Smith, *Chem. Soc. Rev.*, 2008, **37**, 664–675.
- 31 C. Yan and D. J. Pochan, *Chem. Soc. Rev.*, 2010, **39**, 3528–3540.
- 32 N. Svensen, J. G. A. Walton and M. Bradley, *Trends Pharmacol. Sci.*, 2012, **33**, 186–192.
- 33 T. Lehto, K. Ezzat, M. J. A. Wood and S. E. Andaloussi, *Adv. Drug Delivery Rev.*, 2016, **106**, 172–182.
- 34 R. Chang, L. Zhao, R. Xing, J. Li and X. Yan, *Chem. Soc. Rev.*, 2023, **52**, 2688–2712.
- 35 J. Wang, K. Liu, R. Xing and X. Yan, *Chem. Soc. Rev.*, 2016, **45**, 5589–5604.
- 36 M. P. Hendricks, K. Sato, L. C. Palmer and S. I. Stupp, *Acc. Chem. Res.*, 2017, **50**, 2440–2448.
- 37 E. G. Baker, G. J. Bartlett, K. L. Porter Goff and D. N. Woolfson, *Acc. Chem. Res.*, 2017, **50**, 2085–2092.
- 38 C. Yuan, W. Ji, R. Xing, J. Li, E. Gazit and X. Yan, *Nat. Rev. Chem.*, 2019, **3**, 567–588.
- 39 M. Hebel, A. Riegger, M. M. Zegota, G. Kizilsavas, J. Gačanin, M. Pieszka, T. Lückerrath, J. A. S. Coelho, M. Wagner, P. M. P. Gois, D. Y. W. Ng and T. Weil, *J. Am. Chem. Soc.*, 2019, **141**, 14026–14031.
- 40 Q. Song, Z. Cheng, M. Kariuki, S. C. L. Hall, S. K. Hill, J. Y. Rho and S. Perrier, *Chem. Rev.*, 2021, **121**, 13936–13995.
- 41 J. Zhu, N. Avakyan, A. Kakkis, A. M. Hoffnagle, K. Han, Y. Li, Z. Zhang, T. S. Choi, Y. Na, C.-J. Yu and F. A. Tezcan, *Chem. Rev.*, 2021, **121**, 13701–13796.
- 42 C. Wu, H. Zhang, N. Kong, B. Wu, X. Lin and H. Wang, *Angew. Chem., Int. Ed.*, 2023, **62**, e202303455.
- 43 Y. Cong, Z. Qiao and H. Wang, *Adv. Thermoelectr.*, 2018, **1**, 1800067.
- 44 J. Gao, J. Zhan and Z. Yang, *Adv. Mater.*, 2020, **32**, e1805798.
- 45 Y. Marciano, V. del Solar, N. Nayeem, D. Dave, J. Son, M. Contel and R. V. Ulijn, *J. Am. Chem. Soc.*, 2023, **145**, 234–246.
- 46 D. W. P. M. Löwik, E. H. P. Leunissen, M. V. D. Heuvel, M. B. Hansen and J. C. M. V. Hest, *Chem. Soc. Rev.*, 2010, **39**, 3394–3412.
- 47 Y. Shen, X. Fu, W. Fu and Z. Li, *Chem. Soc. Rev.*, 2015, **44**, 612–622.



- 48 R. Zou, Q. Wang, J. Wu, J. Wu, C. Schmuck and H. Tian, *Chem. Soc. Rev.*, 2015, **44**, 5200–5219.
- 49 N. Habibi, N. Kamaly, A. Memic and H. Shafiee, *Nano Today*, 2016, **11**, 41–60.
- 50 S. Lee, J. Xie and X. Chen, *Chem. Rev.*, 2010, **110**, 3087–3111.
- 51 M. Zelzer and R. V. Uljin, *Chem. Soc. Rev.*, 2010, **39**, 3351–3357.
- 52 K. Tao, A. Levin, L. Adler-Abramovich and E. Gazit, *Chem. Soc. Rev.*, 2016, **45**, 3935–3953.
- 53 I. W. Hamley, *Chem. Rev.*, 2017, **117**, 14015–14041.
- 54 L. L. Li, Z. Y. Qiao, L. Wang and H. Wang, *Adv. Mater.*, 2019, **31**, e1804971.
- 55 S. Chagri, D. Y. W. Ng and T. Weil, *Nat. Rev. Chem.*, 2022, **6**, 320–338.
- 56 Z. Liu, G. Liang and W. Zhan, *Chem. Res. Chin. Univ.*, 2021, **37**, 889–899.
- 57 Y. Lyu and K. Pu, *Adv. Sci.*, 2017, **4**, 1600481.
- 58 Y. Lyu, X. Zhen, Y. Miao and K. Pu, *ACS Nano*, 2017, **11**, 358–367.
- 59 A. S. Mahadevi and G. N. Sastry, *Chem. Rev.*, 2016, **116**, 2775–2825.
- 60 P. Zhang, Y. Cui, C. F. Anderson, C. Zhang, Y. Li, R. Wang and H. Cui, *Chem. Soc. Rev.*, 2018, **47**, 3490–3529.
- 61 X. X. Zhao, L. L. Li, Y. Zhao, H. W. An, Q. Cai, J. Y. Lang, X. X. Han, B. Peng, Y. Fei, H. Liu, H. Qin, G. Nie and H. Wang, *Angew. Chem., Int. Ed.*, 2019, **58**, 15287–15294.
- 62 B. Sun, K. Tao, Y. Jia, X. Yan, Q. Zou, E. Gazit and J. Li, *Chem. Soc. Rev.*, 2019, **48**, 4387–4400.
- 63 G. Kwek, T. C. Do, X. Lu, J. Lin and B. Xing, *ACS Appl. Bio Mater.*, 2021, **4**, 2192–2216.
- 64 Y. Zhong, J. Zhan, G. Xu, Y. Chen, Q. Qin, X. Liao, S. Ma, Z. Yang and Y. Cai, *Angew. Chem., Int. Ed.*, 2021, **60**, 8121–8129.
- 65 D. J. Ye, A. J. Shuhendler, L. N. Cui, L. Tong, S. S. Tee, G. Tikhomirov, D. W. Felsher and J. H. Rao, *Nat. Chem.*, 2014, **6**, 519–526.
- 66 Q. C. Jiang, X. Y. Liu, G. L. Liang and X. B. Sun, *Nanoscale*, 2021, **13**, 15142–15150.
- 67 S. He, J. Song, J. Qu and Z. Cheng, *Chem. Soc. Rev.*, 2018, **47**, 4258–4278.
- 68 S. A. Hilderbrand and R. Weissleder, *Curr. Opin. Chem. Biol.*, 2010, **14**, 71–79.
- 69 H. Kobayashi and P. L. Choyke, *Acc. Chem. Res.*, 2011, **44**, 83–90.
- 70 P. Sarder, D. Maji and S. Achilefu, *Bioconjugate Chem.*, 2015, **26**, 963–974.
- 71 L. Yuan, W. Y. Lin, K. B. Zheng, L. W. He and W. M. Huang, *Chem. Soc. Rev.*, 2013, **42**, 622–661.
- 72 B. Hu, N. Song, Y. Cao, M. Li, X. Liu, Z. Zhou, L. Shi and Z. Yu, *J. Am. Chem. Soc.*, 2021, **143**, 13854–13864.
- 73 M. Ikeda, T. Tanida, T. Yoshii, K. Kurotani, S. Onogi, K. Urayama and I. Hamachi, *Nat. Chem.*, 2014, **6**, 511–518.
- 74 R. E. Borg and J. Rochford, *Photochem. Photobiol.*, 2018, **94**, 1175–1209.
- 75 J. G. Huang and K. Y. Pu, *Angew. Chem., Int. Ed.*, 2020, **59**, 11717–11731.
- 76 Y. Liu, L. Teng, B. Yin, H. Meng, X. Yin, S. Huan, G. Song and X. B. Zhang, *Chem. Rev.*, 2022, **122**, 6850–6918.
- 77 X. L. Dean-Ben, S. Gottschalk, B. Mc Larney, S. Shoham and D. Razansky, *Chem. Soc. Rev.*, 2017, **46**, 2158–2198.
- 78 P. K. Upputuri and M. Pramanik, *Wiley Interdiscip. Rev.: Nanomed. Nanobiotechnol.*, 2020, **12**, e1618.
- 79 W. T. Yang, W. S. Guo, W. J. Le, G. X. Lv, F. H. Zhang, L. Shi, X. L. Wang, J. Wang, S. Wang, J. Chang and B. B. Zhang, *ACS Nano*, 2016, **10**, 10245–10257.
- 80 Q. Chen, C. Liang, X. Q. Sun, J. W. Chen, Z. J. Yang, H. Zhao, L. Z. Feng and Z. Liu, *Proc. Natl. Acad. Sci. U. S. A.*, 2017, **114**, 5343–5348.
- 81 X. J. Cheng, R. Sun, L. Yin, Z. F. Chai, H. B. Shi and M. Y. Gao, *Adv. Mater.*, 2017, **29**, 1604894.
- 82 Y. Wang, X. Hu, J. Weng, J. Li, Q. Fan, Y. Zhang and D. Ye, *Angew. Chem., Int. Ed.*, 2019, **58**, 4886–4890.
- 83 C. Wang, W. Du, C. Wu, S. Dan, M. Sun, T. Zhang, B. Wang, Y. Yuan and G. Liang, *Angew. Chem., Int. Ed.*, 2022, **61**, e202114766.
- 84 X. Wen, R. Zhang, Y. Hu, L. Wu, H. Bai, D. Song, Y. Wang, R. An, J. Weng, S. Zhang, R. Wang, L. Qiu, J. Lin, G. Gao, H. Liu, Z. Guo and D. Ye, *Nat. Commun.*, 2023, **14**, 800.
- 85 C. Wu, R. Zhang, W. Du, L. Cheng and G. Liang, *Nano Lett.*, 2018, **18**, 7749–7754.
- 86 Y. Sun, Y. Liang, W. Dai, B. He, H. Zhang, X. Wang, J. Wang, S. Huang and Q. Zhang, *Nano Lett.*, 2019, **19**, 3229–3237.
- 87 X. H. Zhang, D. B. Cheng, L. Ji, H. W. An, D. Wang, Z. X. Yang, H. Chen, Z. Y. Qiao and H. Wang, *Nano Lett.*, 2020, **20**, 1286–1295.
- 88 B. Foster, U. Bagci, A. Mansoor, Z. Y. Xu and D. J. Mollura, *Comput. Biol. Med.*, 2014, **50**, 76–96.
- 89 K. Lameka, M. D. Farwell and M. Ichise, *Handb. Clin. Neurol.*, 2016, **135**, 209–227.
- 90 S. Preshlock, M. Tredwell and V. Gouverneur, *Chem. Rev.*, 2016, **116**, 719–766.
- 91 Y. Geng, P. Dalhaimer, S. Cai, R. Tsai, M. Tewari, T. Minko and D. E. Discher, *Nat. Nanotechnol.*, 2007, **2**, 249–255.
- 92 L. L. Lock, C. D. Reyes, P. Zhang and H. Cui, *J. Am. Chem. Soc.*, 2016, **138**, 3533–3540.
- 93 B. Shen, J. Jeon, M. Palner, D. Ye, A. Shuhendler, F. T. Chin and J. Rao, *Angew. Chem., Int. Ed.*, 2013, **52**, 10511–10514.
- 94 L. Qiu, W. Wang, K. Li, Y. Peng, G. C. Lv, Q. Z. Liu, F. Gao, Y. Seimille, M. H. Xie and J. G. Lin, *Theranostics*, 2019, **9**, 6962–6975.
- 95 Y. Wang, H. Bai, Y. Miao, J. Weng, Z. Huang, J. Fu, Y. Zhang, J. Lin and D. Ye, *Angew. Chem., Int. Ed.*, 2022, **61**, e202200369.
- 96 T. H. Witney, A. Hoehne, R. E. Reeves, O. Ilovich, M. Namavari, B. Shen, F. T. Chin, J. Rao and S. S. Gambhir, *Clin. Cancer Res.*, 2015, **21**, 3896–3905.
- 97 M. Palner, B. Shen, J. H. Jeon, J. G. Lin, F. T. Chin and J. H. Rao, *J. Nucl. Med.*, 2015, **56**, 1415–1421.
- 98 J. Lin, D. Gao, S. Wang, G. Lv, X. Wang, C. Lu, Y. Peng and L. Qiu, *J. Am. Chem. Soc.*, 2022, **144**, 7667–7675.

- 99 Z. Chen, M. Chen, K. Zhou and J. Rao, *Angew. Chem., Int. Ed.*, 2020, **59**, 7864–7870.
- 100 E. T. Ahrens and J. W. M. Bulte, *Nat. Rev. Immunol.*, 2013, **13**, 755–763.
- 101 J. Estelrich, M. J. Sanchez-Martin and M. A. Busquets, *Int. J. Nanomed.*, 2015, **10**, 1727–1741.
- 102 J. J. Futterer, A. Briganti, P. De Visschere, M. Emberton, G. Giannarini, A. Kirkham, S. S. Taneja, H. Thoeny, G. Villeirs and A. Villers, *Eur. Urol.*, 2015, **68**, 1045–1053.
- 103 D. Ni, W. Bu, E. B. Ehlerding, W. Cai and J. Shi, *Chem. Soc. Rev.*, 2017, **46**, 7438–7468.
- 104 G. Liang, J. Ronald, Y. Chen, D. Ye, P. Pandit, M. L. Ma, B. Rutt and J. Rao, *Angew. Chem., Int. Ed.*, 2011, **50**, 6283–6286.
- 105 D. Ye, A. J. Shuhendler, P. Pandit, K. D. Brewer, S. S. Tee, L. Cui, G. Tikhomirov, B. Rutt and J. Rao, *Chem. Sci.*, 2014, **5**, 3845–3852.
- 106 C. Diaferia, E. Gianolio, P. Palladino, F. Arena, C. Boffa, G. Morelli and A. Accardo, *Adv. Funct. Mater.*, 2015, **25**, 7003–7016.
- 107 H. Nejadnik, D. J. Ye, O. D. Lenkov, J. S. Donig, J. E. Martin, R. Castillo, N. Derugin, B. Sennino, J. H. Rao and H. Daldrup-Link, *ACS Nano*, 2015, **9**, 1150–1160.
- 108 Y. Yuan, J. Zhang, X. Qi, S. Li, G. Liu, S. Siddhanta, I. Barman, X. Song, M. T. McMahon and J. W. M. Bulte, *Nat. Mater.*, 2019, **18**, 1376–1383.
- 109 G. Gambino, T. Gambino, L. Connah, F. La Cava, H. Evrard and G. Angelovski, *J. Med. Chem.*, 2021, **64**, 7565–7574.
- 110 L. J. Luo, X. M. Liu, X. Zhang, J. Liu, Y. Gao, T. Y. Sun and L. L. Li, *Nano Lett.*, 2022, **22**, 1694–1702.
- 111 Z. Hai, Y. Ni, D. Saimi, H. Yang, H. Tong, K. Zhong and G. Liang, *Nano Lett.*, 2019, **19**, 2428–2433.
- 112 Y. Hu, J. Zhang, Y. Miao, X. Wen, J. Wang, Y. Sun, Y. Chen, J. Lin, L. Qiu, K. Guo, H. Y. Chen and D. Ye, *Angew. Chem., Int. Ed.*, 2021, **60**, 18082–18093.
- 113 R. Yan, Y. Hu, F. Liu, S. Wei, D. Fang, A. J. Shuhendler, H. Liu, H. Y. Chen and D. Ye, *J. Am. Chem. Soc.*, 2019, **141**, 10331–10341.
- 114 Q. Sun, Z. Zhou, N. Qiu and Y. Shen, *Adv. Mater.*, 2017, **29**, 1606628.
- 115 M. T. Manzari, Y. Shamay, H. Kiguchi, N. Rosen, M. Scaltriti and D. A. Heller, *Nat. Rev. Mater.*, 2021, **6**, 351–370.
- 116 M. J. Mitchell, M. M. Billingsley, R. M. Haley, M. E. Wechsler, N. A. Peppas and R. Langer, *Nat. Rev. Drug Discovery*, 2021, **20**, 101–124.
- 117 Z. X. Zhou, M. Vazquez-Gonzalez and I. Willner, *Chem. Soc. Rev.*, 2021, **50**, 4541–4563.
- 118 K. Li, C. J. Liu and X. Z. Zhang, *Adv. Drug Delivery Rev.*, 2020, **160**, 36–51.
- 119 G. Li, T. Sasaki, S. Asahina, M. C. Roy, T. Mochizuki, K. Koizumi and Y. Zhang, *Chem*, 2017, **2**, 283–298.
- 120 M. T. Jeena, L. Palanikumar, E. M. Go, I. Kim, M. G. Kang, S. Lee, S. Park, H. Choi, C. Kim, S.-M. Jin, S. C. Bae, H. W. Rhee, E. Lee, S. K. Kwak and J.-H. Ryu, *Nat. Commun.*, 2017, **8**, 26.
- 121 G. Li, X. Hu, X. Wu and Y. Zhang, *Nano Lett.*, 2021, **21**, 3052–3059.
- 122 M. Li, Z. Wang, X. Liu, N. Song, Y. Song, X. Shi, J. Liu, J. Liu and Z. Yu, *J. Controlled Release*, 2021, **340**, 35–47.
- 123 C. Zhang, W. Wu, R. Q. Li, W. X. Qiu, Z. N. Zhuang, S. X. Cheng and X. Z. Zhang, *Adv. Funct. Mater.*, 2018, **28**, 1804492.
- 124 G.-B. Qi, Y.-J. Gao, L. Wang and H. Wang, *Adv. Mater.*, 2018, **30**, e1703444.
- 125 Y. J. Liu, P. Bhattarai, Z. F. Dai and X. Y. Chen, *Chem. Soc. Rev.*, 2019, **48**, 2053–2108.
- 126 J. C. Li and K. Y. Pu, *Chem. Soc. Rev.*, 2019, **48**, 38–71.
- 127 T. C. Pham, V. N. Nguyen, Y. Choi, S. Lee and J. Yoon, *Chem. Rev.*, 2021, **121**, 13454–13619.
- 128 Q. Y. Zheng, X. M. Liu, Y. F. Zheng, K. W. K. Yeung, Z. D. Cui, Y. Q. Liang, Z. Y. Li, S. L. Zhu, X. B. Wang and S. L. Wu, *Chem. Soc. Rev.*, 2021, **50**, 5086–5125.
- 129 B. Sun, J. Y. Teo, J. Wu and Y. Zhang, *Acc. Chem. Res.*, 2023, **56**, 1143–1155.
- 130 Z. J. Zhou, J. B. Song, L. M. Nie and X. Y. Chen, *Chem. Soc. Rev.*, 2016, **45**, 6597–6626.
- 131 G. Gao, X. Sun and G. Liang, *Adv. Funct. Mater.*, 2021, **31**, 2100738.
- 132 Y. W. Chen, Y. L. Su, S. H. Hu and S. Y. Chen, *Adv. Drug Delivery Rev.*, 2016, **105**, 190–204.
- 133 D. de Melo-Diogo, C. Pais-Silva, D. R. Dias, A. F. Moreira and I. J. Correia, *Adv. Healthcare Mater.*, 2017, **6**, 1700073.
- 134 X. Y. Huang, Y. Lu, M. X. Guo, S. Y. Du and N. Han, *Theranostics*, 2021, **11**, 7546–7569.
- 135 M. Y. Chang, Z. Y. Hou, M. Wang, C. X. Li and J. Lin, *Adv. Mater.*, 2021, **33**, 2004788.
- 136 J. Qin, X. Wang, G. Fan, Y. Lv and J. Ma, *Adv. Thermoelectr.*, 2023, **6**, 2200218.
- 137 M. Abbas, Q. Zou, S. Li and X. Yan, *Adv. Mater.*, 2017, **29**, 1605021.
- 138 W. Ma, S. N. Sha, P. L. Chen, M. Yu, J. J. Chen, C. B. Huang, B. Yu, Y. Liu, L. H. Liu and Z. Q. Yu, *Adv. Healthcare Mater.*, 2020, **9**, e1901100.
- 139 L. Zhang, Y. Wu, X. Yin, Z. Zhu, T. Rojalin, W. Xiao, D. Zhang, Y. Huang, L. Li, C. M. Baehr, X. Yu, Y. Ajena, Y. Li, L. Wang and K. S. Lam, *ACS Nano*, 2021, **15**, 468–479.
- 140 T. Ma, R. Chen, N. Lv, Y. Li, Z. R. Yang, H. Qin, Z. Li, H. Jiang and J. Zhu, *Small*, 2022, **18**, e2204759.
- 141 X. Liu, W. Zhan, G. Gao, Q. Jiang, X. Zhang, H. Zhang, X. Sun, W. Han, F. G. Wu and G. Liang, *J. Am. Chem. Soc.*, 2023, **145**, 7918–7930.
- 142 M. Li, Y. Ning, J. Chen, X. Duan, N. Song, D. Ding, X. Su and Z. Yu, *Nano Lett.*, 2019, **19**, 7965–7976.
- 143 A. Overbeek, M. H. van den Berg, F. E. van Leeuwen, G. J. L. Kaspers, C. B. Lambalk and E. van Dulmen-den Broeder, *Cancer Treat. Rev.*, 2017, **53**, 10–24.
- 144 R. Oun, Y. E. Moussa and N. J. Wheate, *Dalton Trans.*, 2018, **47**, 6645–6653.
- 145 X. Wang, H. Y. Zhang and X. Z. Chen, *Cancer Drug Resist.*, 2019, **2**, 141–160.

- 146 T. T. Hu, H. L. Gong, J. Y. Xu, Y. Huang, F. B. Wu and Z. Y. He, *Pharmaceutics*, 2022, **14**, 1606.
- 147 E. De Santis and M. G. Ryadnov, *Chem. Soc. Rev.*, 2015, **44**, 8288–8300.
- 148 M. Pieszka, S. Han, C. Volkmann, R. Graf, I. Lieberwirth, K. Landfester, D. Y. W. Ng and T. Weil, *J. Am. Chem. Soc.*, 2020, **142**, 15780–15789.
- 149 N. J. Sinha, M. G. Langenstein, D. J. Pochan, C. J. Kloxin and J. G. Saven, *Chem. Rev.*, 2021, **121**, 13915–13935.
- 150 J. Wang, Y. Li and G. Nie, *Nat. Rev. Mater.*, 2021, **6**, 766–783.
- 151 X.-J. Wang, J. Cheng, L.-Y. Zhang and J.-G. Zhang, *Drug Delivery*, 2022, **29**, 1184–1200.
- 152 J. Zhan, Y. Cai, S. He, L. Wang and Z. Yang, *Angew. Chem., Int. Ed.*, 2018, **57**, 1813–1816.
- 153 L. Zhang, D. Jing, N. Jiang, T. Rojalín, C. M. Baehr, D. Zhang, W. Xiao, Y. Wu, Z. Cong, J. J. Li, Y. Li, L. Wang and K. S. Lam, *Nat. Nanotechnol.*, 2020, **15**, 145–153.
- 154 F. Wang, D. Xu, H. Su, W. Zhang, X. Sun, M. K. Monroe, R. W. Chakroun, Z. Wang, W. Dai, R. Oh, H. Wang, Q. Fan, F. Wan and H. Cui, *Sci. Adv.*, 2020, **6**, eaaz8985.
- 155 C. Keum, J. Hong, D. Kim, S.-Y. Lee and H. Kim, *ACS Appl. Mater. Interfaces*, 2021, **13**, 14866–14874.
- 156 L. Xie, Y. Ding, Y. Wang, X. Li, Z. Yang and Z.-W. Hu, *Adv. Funct. Mater.*, 2022, **32**, 2206969.
- 157 L. Hu, Y. Li, X. Lin, Y. Huo, H. Zhang and H. Wang, *Angew. Chem., Int. Ed.*, 2021, **60**, 21807–21816.
- 158 R. A. Pires, Y. M. Abul-Haija, D. S. Costa, R. Novoa-Carballal, R. L. Reis, R. V. Ulijn and I. Pashkuleva, *J. Am. Chem. Soc.*, 2015, **137**, 576–579.
- 159 S. Liu, Q. Zhang, H. He, M. Yi, W. Tan, J. Guo and B. Xu, *Angew. Chem., Int. Ed.*, 2022, **61**, e202210568.
- 160 W. Tan, Q. Zhang, M. C. Quiñones-Frias, A. Y. Hsu, Y. Zhang, A. Rodal, P. Hong, H. R. Luo and B. Xu, *J. Am. Chem. Soc.*, 2022, **144**, 6709–6713.
- 161 Y. Song, M. Li, N. Song, X. Liu, G. Wu, H. Zhou, J. Long, L. Shi and Z. Yu, *J. Am. Chem. Soc.*, 2022, **144**, 6907–6917.
- 162 D. Ross and D. Siegel, *Redox Biol.*, 2021, **41**, 101950.
- 163 J. Chang, W. Cai, C. Liang, Q. Tang, X. Chen, Y. Jiang, L. Mao and M. Wang, *J. Am. Chem. Soc.*, 2019, **141**, 18136–18141.
- 164 Z.-Y. Jiang, M.-C. Lu and Q.-D. You, *J. Med. Chem.*, 2016, **59**, 10837–10858.
- 165 D. J. Irvine, M. C. Hanson, K. Rakhra and T. Tokatlian, *Chem. Rev.*, 2015, **115**, 11109–11146.
- 166 A. Ribas and J. D. Wolchok, *Science*, 2018, **359**, 1350–1355.
- 167 A. D. Waldman, J. M. Fritz and M. J. Lenardo, *Nat. Rev. Immunol.*, 2020, **20**, 651–668.
- 168 M. Li, X. Zhao, J. Dai and Z. Yu, *Sci. China Mater.*, 2019, **62**, 1759–1781.
- 169 R. Chang and X. Yan, *Small Struct.*, 2020, **1**, 2000068.
- 170 M. D. Wang, G. T. Lv, H. W. An, N. Y. Zhang and H. Wang, *Angew. Chem., Int. Ed.*, 2022, **61**, e202113649.
- 171 Y. Ding, D. Zheng, L. Xie, X. Zhang, Z. Zhang, L. Wang, Z. W. Hu and Z. Yang, *J. Am. Chem. Soc.*, 2023, **145**, 4366–4371.
- 172 L. Zhang, Z. Jiang, X. Yang, Y. Qian, M. Wang, S. Wu, L. Li, F. Jia, Z. Wang, Z. Hu, M. Zhao, X. Tang, G. Li, H. Shang, X. Chen and W. Wang, *Adv. Mater.*, 2023, **35**, 2207330.
- 173 Q. L. Zhang, D. Zheng, X. Dong, P. Pan, S. M. Zeng, F. Gao, S. X. Cheng and X. Z. Zhang, *J. Am. Chem. Soc.*, 2021, **143**, 5127–5140.
- 174 W. P. Fan, B. Yung, P. Huang and X. Y. Chen, *Chem. Rev.*, 2017, **117**, 13566–13638.
- 175 S. A. Patel and A. J. Minn, *Immunity*, 2018, **48**, 417–433.
- 176 X. W. Wang, X. Y. Zhong, Z. Liu and L. Cheng, *Nano Today*, 2020, **35**, 100946.
- 177 J. C. Dawson, A. Serrels, D. G. Stupack, D. D. Schlaepfer and M. C. Frame, *Nat. Rev. Cancer*, 2021, **21**, 313–324.
- 178 S. G. Lim, T. F. Baumert, C. Boni, E. Gane, M. Levrero, A. S. Lok, M. K. Maini, N. A. Terrault and F. Zoulim, *Nat. Rev. Gastroenterol. Hepatol.*, 2023, **20**, 238–253.
- 179 G. Gao, Y. W. Jiang, W. Zhan, X. Liu, R. Tang, X. Sun, Y. Deng, L. Xu and G. Liang, *J. Am. Chem. Soc.*, 2022, **144**, 11897–11910.
- 180 Z. Zhang, K. Wang, M. Liu, P. Hu, Y. Xu, D. Yin, Y. Yang, X. Dong, C. Qu, L. Zhang, J. Ni and X. Yin, *Drug Delivery*, 2022, **29**, 1608–1619.
- 181 W. Jiang, C. Cheng, X. Qiu, L. Chen, X. Guo, Y. Luo, J. Wang, J. Wang, Z. Xie, P. Li, Z. Wang, H. Ran, Z. Zhou and J. Ren, *Adv. Sci.*, 2023, **10**, e2204989.
- 182 Y. Sun, B. Lyu, C. Yang, B. He, H. Zhang, X. Wang, Q. Zhang and W. Dai, *Bioact. Mater.*, 2023, **22**, 47–59.
- 183 N. Song, Z. Zhou, Y. Song, M. Li, X. Yu, B. Hu and Z. Yu, *Nano Today*, 2021, **38**, 101198.
- 184 Y. Li, G. Yang, L. Gerstweiler, S. H. Thang and C. X. Zhao, *Adv. Funct. Mater.*, 2022, **33**, 2210387.
- 185 S. Lou, X. Wang, Z. Yu and L. Shi, *Adv. Sci.*, 2019, **6**, 1802043.
- 186 P. Makam and E. Gazit, *Chem. Soc. Rev.*, 2018, **47**, 3406–3420.
- 187 D. M. Raymond and B. L. Nilsson, *Chem. Soc. Rev.*, 2018, **47**, 3659–3720.
- 188 Y. Song, Z. Zhang, Y. Cao and Z. Yu, *ChemBioChem*, 2023, **24**, e202200497.
- 189 R. Kubota, W. Tanaka and I. Hamachi, *Chem. Rev.*, 2021, **121**, 14281–14347.
- 190 J. Hofmann, H. S. Hahm, P. H. Seeberger and K. Pagel, *Nature*, 2015, **526**, 241–244.
- 191 M. A. Walti, F. Ravotti, H. Arai, C. G. Glabe, J. S. Wall, A. Bockmann, P. Guntert, B. H. Meier and R. Riek, *Proc. Natl. Acad. Sci. U. S. A.*, 2016, **113**, E4976–E4984.
- 192 L. Wei, Z. X. Chen, L. X. Shi, R. Long, A. V. Anzalone, L. Y. Zhang, F. H. Hu, R. Yuste, V. W. Cornish and W. Min, *Nature*, 2017, **544**, 465–470.
- 193 R. Hervas, M. J. Rau, Y. Park, W. Zhang, A. G. Murzin, J. A. J. Fitzpatrick, S. H. W. Scheres and K. Si, *Science*, 2020, **367**, 1230–1234.
- 194 Y. Yang, D. Arseni, W. Zhang, M. Huang, S. Lövestam, M. Schweighauser, A. Kotecha, A. G. Murzin, S. Y. Peak-Chew, J. Macdonald, I. Lavenir, H. J. Garringer, E. Gelpi, K. L. Newell, G. G. Kovacs, R. Vidal, B. Ghetti, B. Ryskeldi-



## Review

- Falcon, S. H. W. Scheres and M. Goedert, *Science*, 2022, **375**, 167–172.
- 195 Y. Sun, S. Y. Lau, Z. W. Lim, S. C. Chang, F. Ghadessy, A. Partridge and A. Miserez, *Nat. Chem.*, 2022, **14**, 274–283.
- 196 J. Liu, E. Spruijt, A. Miserez and R. Langer, *Nat. Rev. Mater.*, 2023, **8**, 139–141.
- 197 J. Huang, Y. Xu, Y. Xue, Y. Huang, X. Li, X. Chen, Y. Xu, D. Zhang, P. Zhang, J. Zhao and J. Ji, *Nat. Biomed. Eng.*, 2023, **7**, 797–810.
- 198 M. Zhu, J. Zhuang, Z. Li, Q. Liu, R. Zhao, Z. Gao, A. C. Midgley, T. Qi, J. Tian, Z. Zhang, D. Kong, J. Tian, X. Yan and X. Huang, *Nat. Nanotechnol.*, 2023, **18**, 657–666.