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## A switchable magnetic resonance imaging nanoplatform for *in situ* microRNA imaging†

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Aberrant microRNA (miRNA) expression is associated with various types of carcinogenesis, making miRNA a promising candidate for diagnostic and therapeutic biomarkers. However, *in situ* miRNA diagnostics remains a significant challenge owing to the various biological barriers. Herein, we report a novel miRNA imaging probe consisting of PEG-polylysine-PNIPAM polymer matrix-modified small Fe<sub>3</sub>O<sub>4</sub> (PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP) nanoparticles with an improved circulatory half-life, efficient tissue permeability, and enhanced tumor accumulation, for *in situ* miRNA magnetic resonance imaging (MRI). In this strategy, we employed large size PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP to improve circulatory time and utilized PEG-polylysine-PNIPAM as a GSH-responsive moiety to dissociate PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP and release small size PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA for enhanced tumor permeability. Specifically, the target miRNA acts as a cross-linker for PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA, forming larger assemblies that not only amplify the MRI signal for detection but also enhance retention for prolonged observation. Both the *in vitro* and *in vivo* results validate that the imaging probe exhibits an enhanced MRI signal with 3.69-fold amplification for tumor interior miRNA detection, allowing the dynamic changes in miRNA to be monitored by the probe. Given its long circulation, efficient penetration, and enhanced tumor accumulation, the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP probe holds great promise for *in situ* miRNA imaging and spatial genomics analysis *in situ*.

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## Introduction

*In situ* biomarker imaging that utilizes advanced imaging techniques such as microscopy, magnetic resonance imaging (MRI), or positron emission tomography (PET) to identify and monitor these biomarkers within their natural environment, holds great promise for personalized medicine in cancer diagnosis and advancing our understanding of the complex pathological process. However, several challenges and limitations persist in efficiently delivering imaging agents to tumor tissue to achieve optimal efficacy. As reported, nanoparticles (NPs) with a diameter of 100 nm can be stable in the circulation, but not efficiently pass through tumor vessels.<sup>1</sup> In contrast, small-size NPs with an approximate size of 4–20 nm easily penetrate through deep tumor tissues, yet they are more inclined to

clearance and lack retention at tumor sites.<sup>2</sup> Therefore, rational designs of NPs with size-tunable performance under various biological conditions to achieve effective tumor accumulation and enhanced tumor permeability have been continuously developed.<sup>3–5</sup>

As a significant small non-coding RNA molecular, microRNA (miRNA) is the post-transcriptional regulator in biological and pathological effects, especially serving as a crucial biomarker for tumor classification.<sup>6–8</sup> In our previous studies, selective and sensitive detection of miRNA in living cells based on nanoparticle systems has been developed.<sup>9–11</sup> *In situ* miRNA detection is vital for biological function and mechanism research and monitoring of cancerous pathological conditions; however, it is still a huge challenge because of their low-abundant expression level and complex biological environments. Photoacoustic imaging (PAI) of miRNA at the tumor site was realized *via* miRNA-triggered self-assembly of DNA-coated nanostructures to form particle aggregation with enhanced PAI signals.<sup>11</sup> A cancer cell membrane camouflaged nanoprobe with homotypic-targeting capability and remarkable amplification was also designed for PAI of miRNA in living mice.<sup>12</sup> Smartly designed hierarchical nanoparticles with size-tunable properties in response to physiological conditions to satisfy long circulation, efficacy permeability and tumor site-specific miRNA activatable signal enhancement have never been reported.

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Magnetic resonance imaging (MRI) has been proven as a mighty appliance for clinical diagnosis because of its resultful tissue contrast, high spatial resolution (about 50  $\mu\text{m}$ ), low ionizing radiation, and unstinted signal penetration depth.<sup>13–16</sup> Thereinto, ultrasmall superparamagnetic iron oxide nanoparticles (IONPs) with appreciable biodistribution and metabolism have been widely used as  $T_1$ -weighted MRI contrast agents.<sup>17,18</sup> Interestingly, previous studies reported that the assemblies of IONP contrast agents dramatically affected the magnetic properties due to the aggregation-induced enhancement of the magnetic field resulting in  $T_2$  contrast imaging promotion.<sup>19–22</sup> Such properties provided great opportunities to establish conditional activatable contrast agents that could further amplify the imaging contrast by abating the interference generated from nonspecific adsorption or endogenous iron-derived imaging contrast.<sup>23</sup> A series of transformations of IONPs exhibited an excellent capacity for penetration and aggregation in the physiological environment and have been developed to evaluate deep tumor imaging in clinical applications.<sup>24</sup> For example, “turn-on” MRI nanoprobes were fabricated that showed dark  $T_2$  contrast imaging in the vessels and transformed into bright  $T_1$  contrast imaging owing to GSH or pH-triggered cleavage in the tumor microenvironment (TME).<sup>25–28</sup> Smart switchable MRI probes based on the self-assembly of IONPs were fabricated as a transformation from  $T_2$ -weighted into  $T_1$ -weighted contrast imaging at the tumor site under low pH-induced disassembly.<sup>29</sup> However, multistep switchable MRI probes for *in situ* miRNA imaging have scarcely been explored.

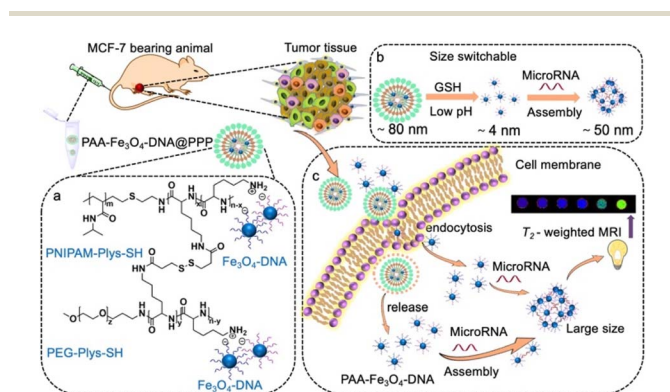
Therefore, for the first time, we designed a “large-small-large” size-switchable PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP nanoplatform, consisting of PEG-polylysine-PNIPAM polymer matrix-modified small  $\text{Fe}_3\text{O}_4$ , for *in situ* miRNA imaging. As shown in Scheme 1, PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP maintains a large size for prolonged circulation time; it decomposes to a small size (4–5 nm) PAA- $\text{Fe}_3\text{O}_4$ -DNA exposing to TME GSH conditions for deep

penetration. After the internalization of PAA- $\text{Fe}_3\text{O}_4$ -DNA, the small-size nanoparticles would be linked by the intracellular miRNA to self-assemble to form stable larger-size nanoparticles (more than 50 nm) that consequently enhance the  $T_2$ -weighted MRI in the tumor for miRNA *in situ* detection. Meanwhile, fluorescent dye Cy5 was conjugated on the DNA strand to track the nanosystem accurately. The size switchable imaging system is anticipated to be applied to plenty of *in situ* biomolecular detections by rational detection probe design.

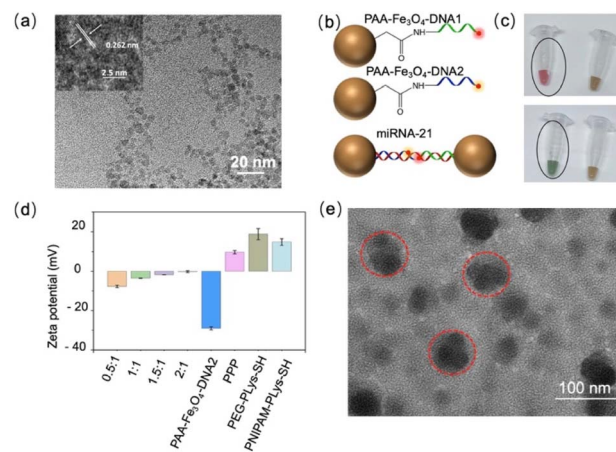
## Results and discussion

### “Small-to-large” PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP assembly

The construction procedure of PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP is presented in Scheme 1. First, polyacrylic acid-modified  $\text{Fe}_3\text{O}_4$  (PAA- $\text{Fe}_3\text{O}_4$ ) nanoparticles were synthesized according to the previous method with the hydrothermal decomposition in the existence of polyacrylic acid and ferric chloride raw materials.<sup>30</sup> The average diameter of PAA- $\text{Fe}_3\text{O}_4$ , as measured by transmission electron microscopy (TEM) is about  $4.43 \pm 0.6407$  nm (Fig. 1a and S1†). The high resolution TEM measurement of PAA- $\text{Fe}_3\text{O}_4$  shows that the lattice fringe spacing value is 0.262 nm, which is assigned to the (311) planes of  $\text{Fe}_3\text{O}_4$  (inset of Fig. 1a). Fourier transform infrared spectrometry (FTIR) was chosen to confirm the carboxyl group of PAA conjugated on  $\text{Fe}_3\text{O}_4$ . The peak at  $1639\text{ cm}^{-1}$  is due to the C=O bond in the PAA.<sup>31</sup> On the other hand, the vibration absorption peaks ( $3300\text{ cm}^{-1}$ ) for the O–H bond demonstrated that PAA- $\text{Fe}_3\text{O}_4$  was successfully synthesized (Fig. S2†). Additionally, the peak at around  $580\text{ cm}^{-1}$  represented the vibration of FeO groups.<sup>32</sup> Two fluorescent dye-labeled single-strand DNA sequences complementary to target miRNA were conjugated on PAA- $\text{Fe}_3\text{O}_4$  *via* the



**Scheme 1** Illustration of size-flexible PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP for MRI activated by miRNA. (a) Large-size PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP is formed by coating the polymer PPP in the core of DNA-modified  $\text{Fe}_3\text{O}_4$  nanoparticles. (b) PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP degrades into small size PAA- $\text{Fe}_3\text{O}_4$ -DNA exposed to GSH and low pH in the TME. (c) The small size PAA- $\text{Fe}_3\text{O}_4$ -DNA is taken up *via* cellular endocytosis and then assembled into a larger complex linked by the miRNA with an enhanced  $T_2$  MR signal.



**Fig. 1** Preparation and characterization of PAA- $\text{Fe}_3\text{O}_4$ -DNA and the miRNA assemble aggregate. (a) TEM image of the PAA- $\text{Fe}_3\text{O}_4$  nanoparticles; the inserted panel shows the distribution profile of the average size of PAA- $\text{Fe}_3\text{O}_4$  nanoparticles. (b) Scheme illustration for miRNA-21 triggering the PAA- $\text{Fe}_3\text{O}_4$ -DNA assembly. (c) Photos of Cy5 and Cy3 labeled DNA conjugated PAA- $\text{Fe}_3\text{O}_4$ . (d) The  $\zeta$  potential of PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP based samples. 0.5:1, 1:1, 1.5:1 and 2:1 groups represent the N/P ratios of PPP and PAA- $\text{Fe}_3\text{O}_4$ -DNA. (e) TEM image of PAA- $\text{Fe}_3\text{O}_4$ -DNA@PPP with a 1.5 N/P ratio.



amino-carboxyl group crosslinking reaction to form PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1 and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2 (Fig. 1b to c). After the conjugation, the extra fluorescent DNA was removed with the help of a magnet (Fig. S3†). The successful modification of DNA molecules on PAA-Fe<sub>3</sub>O<sub>4</sub> was determined *via* UV spectra at 260 nm (Fig. S4†). The amount of DNA1 and DNA2 on PAA-Fe<sub>3</sub>O<sub>4</sub> was estimated to be 13.74 μmol g<sup>-1</sup> Fe and 11.88 μmol g<sup>-1</sup> Fe, respectively, according to the standard curve of fluorescence and concentration (Fig. S5†). These results suggested that the DNA was successfully conjugated on PAA-Fe<sub>3</sub>O<sub>4</sub>. To protect the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA nanoparticles, multiblock cationic polymer PEG-polylysine-PNIPAM (PPP) was synthesized subsequently. Polyethyleneglycol (PEG) modified polycationic polylysine (Plys) (PEG-PLys-SH) (Fig. S6†) and poly(*N*-isopropyl acrylamide) (PNIPAM) grafted Plys (PNIPAM-Plys-SH) (Fig. S7†) were prepared according to the previously reported method.<sup>33</sup> The resulting polymers were characterized by <sup>1</sup>H NMR, and the number of -SH was calculated to be 1.25 and 7 for PEG-PLys-SH (Fig. S8†) and PNIPAM-Plys-SH (Fig. S9†).

The assembly of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was optimized with different N/P ratios derived from PPP and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA. The zeta potential analysis confirmed that the PPP polymer was positive, and the negative charge of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA was neutralized as the N/P ratios increased (Fig. 1d). Agarose gel electrophoresis showed that the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP samples were gradually trapped in the beginning line of the gel as the N/P ratios increased, and no free PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA was visualized at a ratio of 1.5 : 1 (Fig. S10†). These results indicated that the negatively charged DNA has effectively condensed into the PPP polymer. As shown in Fig. 1e, the average diameter of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP is 82 ± 20 nm while the N/P ratio is 1.5. As reported, the PPP polymer would have a relatively smaller size when at a physiological temperature of 37 °C because of thermally responsive PNIPAM.<sup>34</sup> When the temperature increases, PNIPAM could transform into the hydrophobic phase that provides an additional drive force to stimulate further condensation of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP and protect PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA from enzymatic degradation in the physical environment. The high-resolution TEM image and the element mapping analysis indicated that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA was encapsulated into the PPP polymer structure (Fig. S11 & S12†).

### “Large-to-small” and “small-to-large” transformation

PPP was obtained by disulfide cross-linking formation of PEG-PLys-SH and PNIPAM-Plys-SH, and the resulting PPP could pack the small DNA-functionalized nanoparticles through electrostatic assemblage between cationic Plys-SH and anionic nucleic acid of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA to form PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP. The disulfide cross-linking was readily cleaved under acidic and reductive conditions with glutathione (GSH). The existence of an ester bond and amide bond in PPP would accelerate the hydrolysis of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP nanocapsule in the acidic TME. The degradation of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was evaluated by exposing it to GSH with different concentrations of 0 mM, 10 mM, 20 mM, 40 mM, and 80 mM (Fig. 2a). The gel electrophoresis results confirmed that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP gradually

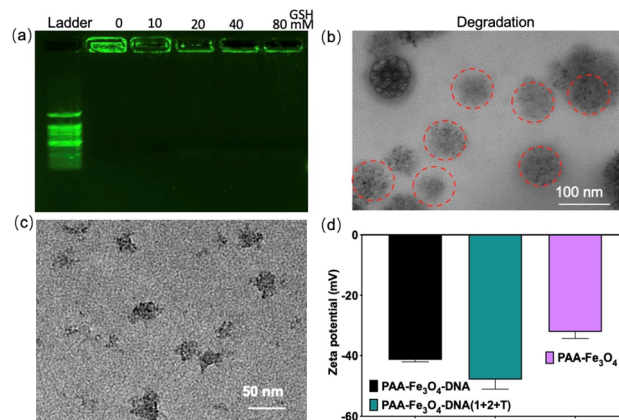


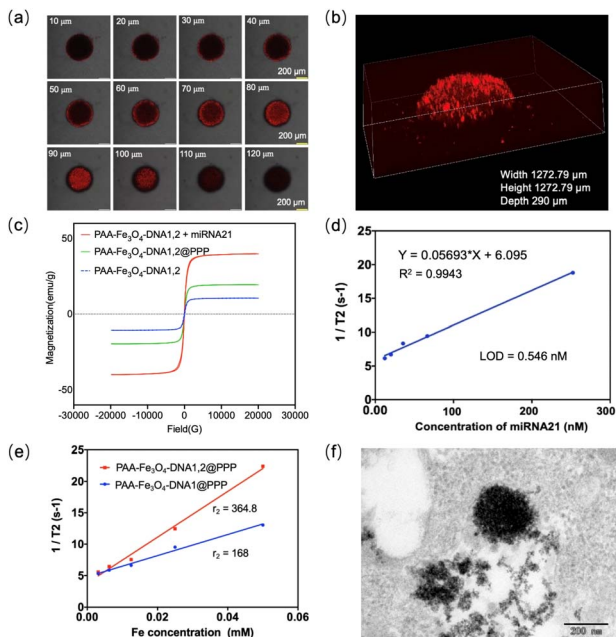
Fig. 2 Evaluation of the size change from large to small. (a) Degradation process of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP. Gel electrophoresis of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP after incubating with different concentrations of GSH (0 mM, 10 mM, 20 mM, 40 mM, and 80 mM). (b) TEM image of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP after degradation in the presence of 20 mM GSH for 24 hours. (c) TEM image of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA after self-assembly *via* miRNA-21. (d) Zeta potential of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA and miRNA (target) triggered aggregation of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA (1 + 2 + T).

degraded with the increase in the GSH concentration. In addition, the TEM image showed the disassembly morphology of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP which was marked with red circles (Fig. 2b). These results confirmed the efficient degradation of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP to small PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA, which was useful for enhanced tumor depth penetration. Using miRNA-21, a tumor-related up-regulated miRNA, as a model target miRNA, we investigated the feasibility of target miRNA-mediated PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA assembly, leading to a conversion process from small to large size. As anticipated, the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA (1 + 2 + T) aggregates exhibited an average size of 50 nm when the concentration ratio of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1, PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2, and miRNA-21 was maintained at 1 : 1 : 1 (Fig. 2c). The resulting aggregates displayed more negative zeta potential (Fig. 2d). As the ratio increases from 1 : 1 : 1 to 1 : 1 : 10, the size increases from about 50 nm to 80 nm (Fig. S13†). These results verified that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1 and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2 could be assembled through the bridge miRNA-21, implementing small-size to larger-size transformation.

### Penetration evaluation and *in vitro* magnetic resonance imaging

The penetration of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was investigated by confocal imaging. MCF-7 multicellular tumor spheroids were formed in the agarose-based microwells and reached a diameter of more than 400 μm after culturing for several days according to previous methods.<sup>35,36</sup> PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP labeled with Cy5 in the DNA has efficiently penetrated the MCF-7 multicellular tumor spheroids characterized by confocal imaging through a layer scan model within 10 μm intervals (Fig. 3a). According to the confocal imaging, cross-sectional fluorescence can be observed when the scanning depth of the Z-axis is 100 μm. The three-dimensional fluorescence reconfiguration of multicellular





**Fig. 3** Cy5-labeled PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP to penetrate the MCF-7 multicellular tumor spheroids measured using a confocal imaging microscope. (a) Confocal imaging of the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP penetration evaluation from 10 μm to 120 μm. The scale bar is 200 μm. (b) The 3D image was stacked using the software. (c) Field-dependent magnetization hysteresis loops of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1 and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2 incubated with miRNA-21 or not. The ratio of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1, PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2, and miRNA-21 is 1 : 1 : 1. (d) The  $r_2$  relaxivity of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA treated with various concentrations of miRNA-21. (e) The  $r_2$  relaxivity of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP incubated with MCF-7 cells and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1@PPP as a negative control. (f) The bioTEM image of the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP treated MCF-7 cells. The scale bar is 200 nm.

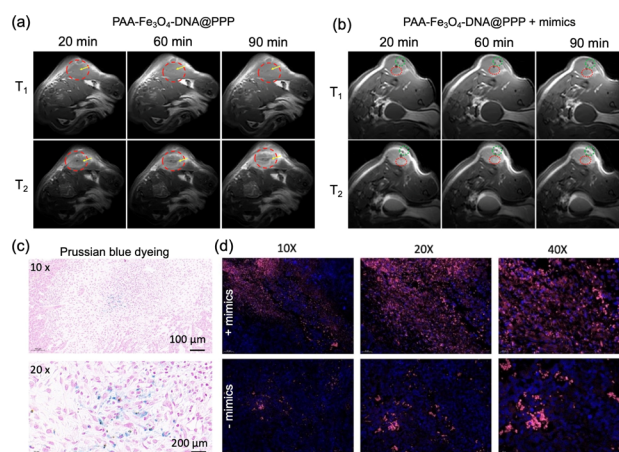
tumor spheroids exposed to PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP confirmed the deep penetration (Fig. 3b).

This reported that ultras-small Fe<sub>3</sub>O<sub>4</sub> nanoparticles with a diameter of less than 5 nm have good T<sub>1</sub>-MRI contrast function. In contrast, the self-assembled ultras-small Fe<sub>3</sub>O<sub>4</sub> would enhance the magnetic size inducing a higher saturation magnetization, leading to an improved dark T<sub>2</sub> MRI contrast.<sup>37–39</sup> As seen in , the value of saturated magnetic intensity for PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1 increased from 10.5 to 39.8 meu g<sup>-1</sup> when assembled with miRNA-21. Interestingly, the magnetic saturation value of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was 19.5 meu g<sup>-1</sup>, indicating that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1 and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA2 encapsulated into PPP. Furthermore, we measured the T<sub>1</sub> and T<sub>2</sub> relaxation times of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2 with or without miRNA-21. The  $r_1$  and  $r_2$  relaxivity of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2 in the presence of miRNA-21 was calculated to be 0.28 mM<sup>-1</sup> s<sup>-1</sup>, 250.1 mM<sup>-1</sup> s<sup>-1</sup>, while  $r_1$  and  $r_2$  relaxivity of the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2 mixture without miRNA-21 was 0.83 mM<sup>-1</sup> s<sup>-1</sup> and 101.0 mM<sup>-1</sup> s<sup>-1</sup> (Fig. S14 & S15<sup>†</sup>). These results suggested that the assembly PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2 triggered by miRNA-21 showed enhanced T<sub>2</sub> suitable for T<sub>2</sub> MRI contrast imaging. As the miRNA-21 concentration increased from 0 to 250 nM, the  $r_2$  increased from 6 to 20.4 s<sup>-1</sup> (Fig. 3d), and the limit of detection

was calculated to be 0.546 nM using T<sub>2</sub>-MRI. We further investigated its performance in living cells. PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP was incubated with the MCF-7 cells for 8 hours, and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1@PPP was a negative control. Then the cells were all harvested and washed with PBS. After that, the cells were fixed using agarose and were detected under a 3.0 T MR (Fig. S16<sup>†</sup>). Fig. 3e shows a higher  $r_2$  relaxivity in the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP treated cells compared with the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1@PPP incubated cell group. Meanwhile, the  $r_1$  relaxivity was also measured (Fig. S17<sup>†</sup>), and the  $r_2/r_1$  value was respectively calculated to be 46.0 and 112.7 for the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1@PPP and PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP treated groups. As shown in Fig. 3f, the large aggregation structures were visualized in the bioTEM image of the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA1,2@PPP treated MCF-7 cells, suggesting that target miRNA-21 induced aggregation. The results illustrated that miRNA-21 in the cells would contribute to the aggregation of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA.

### *In vivo* magnetic resonance imaging

To investigate the *in vivo* MRI of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP. PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was tail intravenously injected into the MCF-7-bearing mice. The signal was observed after 20 min of injection (Fig. 4a). As indicated in the T<sub>2</sub> weighted MRI, the assembled PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP derived T<sub>2</sub> signal reached the maximum at 90 minutes post-injection. Notably, the ratio of the T<sub>2</sub>/T<sub>1</sub> signal was less than 1.0 at 20 min after injection (Fig. S18<sup>†</sup>), and the signal increased as time increased, verifying that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was a good T<sub>2</sub>-weighted contrast reagent. Furthermore, to investigate the monitoring ability of miRNA change by PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP, miRNA-21 mimics were injected intratumorally for 4 h before the imaging



**Fig. 4** (a) T<sub>1</sub>-MRI and T<sub>2</sub>-MRI images of mice bearing MCF-7 tumor cells according to the intravenous injections of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP. (b) The miRNA-21 mimics were transferred before the intravenous injections of PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP, and then the T<sub>1</sub>-MRI and T<sub>2</sub>-MRI were measured. (c) The Prussian blue dyeing of the tumor tissues after MRI observation. The microscope objective is 10× and 40× for the upper and lower figures. (d) The tumor slices were observed under a fluorescence microscope. The Cy5 FL indicates Cy5 labeled PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP. 10×, 20× and 40× represent objectives with different magnification times.



experiment. The mimic DNA sequence has the same function as miRNA-21. The  $T_1$  and  $T_2$  MRI signals were evaluated after being injected for 20 min, 40 min, 60 min, and 90 min. As shown in the  $T_2$  MRI images displayed in Fig. 4b, the hole in the green circle was the injection site. Many dark dots derived from the miRNA-21-induced PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA assembly were observed (in red circles).

The tumors were collected for sectioning and staining to verify whether PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP entered into the tissues. Prussian blue dyeing was employed to observe iron. As exhibited in Fig. 4c, the blue granules were in the tumor cells, indicating that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP could reach the tumor site. Because PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP was labeled with Cy5, the slices were also observed under a fluorescence microscope (Fig. 4d). The mimic miRNA-21-treated tissues displayed strong Cy5 fluorescence compared with that without treatment, which proved that PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA was trapped in the tumor tissues with the assistance of the miRNA-induced assembly.

## Conclusions

In this work, we report the construction of a switchable miRNA MRI imaging probe, PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP, for *in situ* miRNA imaging. PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP undergoes a “large-small-large” switchable process to cross various biological barriers, accumulate in tumor tissue, and achieve effective *in situ* miRNA MRI imaging. It maintains a large size for prolonged circulation time and then decomposes into small-sized (4–5 nm) PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA in response to the TME GSH conditions, allowing for deep penetration. After cellular internalization, the small-sized nanoparticles are linked by intracellular miRNA, self-assembling into stable larger-sized nanoparticles (over 50 nm). This assembly enhances  $T_2$ -weighted MRI signals in the tumor, enabling *in situ* miRNA detection and allowing *in situ* monitoring of the dynamic changes in miRNA. The long circulation time, efficient penetration, and enhanced tumor accumulation make the PAA-Fe<sub>3</sub>O<sub>4</sub>-DNA@PPP probe highly promising for *in situ* miRNA imaging and spatial genomics analysis.

## Data availability

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

## Author contributions

Conceptualization: YT, HFD, and XJZ; experiments and formal analysis: YT, JRW, and QW; methodology: JLY, JKH, and ZJZ; project administration: HFD and XJZ; writing the original draft: YT; writing – review & editing: YT, JRW and HFD. All authors have read and agreed to the final version of this manuscript.

## Conflicts of interest

There are no conflicts to declare.

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