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Superparamagnetic Iron Oxide Nanoparticle–Aptamer Bioconjugates for Combined Prostate Cancer Imaging and Therapy

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Over the past two decades, molecular targeted diagnostic and therapeutic agents have dramatically improved cancer diagnosis and treatment.^[1-8] Targeting allows the preferential delivery of therapeutic, diagnostic, or imaging agents to the intended site. Advances in nanotechnology have enabled the development of a variety of targeted nanoparticle platforms for diagnostic and therapeutic applications.^[9-11] Preclinical data have shown that targeted nanoparticle systems accumulate preferentially in the target tissue, demonstrating the vast potential of targeted nanoparticles.[12–14] In addition, the development of multifunctional nanoparticle platforms, with both diagnostic and therapeutic capabilities, may allow in vivo monitoring of both biodistribution of the nanocarriers and tumor response to therapy.^[11, 15-17] Therefore, research efforts have been focused on the further development of multifunctional molecular agents for the diagnosis and treatment of cancer.

One of the most promising diagnostic agents is superparamagnetic iron oxide nanoparticles (SPION).^[18] SPION have several important advantages over traditional gadolinium-based magnetic resonance (MR) contrast agents: lower toxicity, stronger enhancement of proton relaxation, and lower detection limit.^[19, 20] Ferumoxtran-10 (Combidex), a dextran-coated SPION with a mean diameter of \sim 30 nm, is currently in phase III clinical trials for prostate cancer (PCa) imaging.[21] Combidex has a 90.5% sensitivity and 97.8% specificity for detecting PCa lymph node disease by passively accumulating in metastatic nodes.[22] The major shortcoming of Combidex is its inability to detect PCa disease outside of the lymph nodes.

Herein, we report the development of a novel, multifunctional, thermally cross-linked SPION (TCL-SPION) that can both detect PCa cells, and deliver targeted chemotherapeutic agents directly to the PCa cells. We previously reported the use of the A10 RNA aptamer (Apt), which binds the extracellular domain of the prostate-specific membrane antigen (PSMA), to engineer targeted nanoparticles for PCa therapy and imaging.^[12, 13, 23] PSMA is a well-established marker for PCa cells, with relatively low levels of expression in normal prostate, kidney, brain, and small intestine tissue.^[24] The percentage of PCa cells that express PSMA approaches 100% with highest expression in androgen-independent PCa cells.^[25,26] Additionally, we have shown that the A10 aptamer can be used to deliver doxorubicin (Dox), a chemotherapeutic agent, by intercalation of Dox into the CG sequence in the aptamer.^[23, 27, 28] By combining the above concepts, we have formulated SPION–Apt bioconjugates for combined PCa imaging and therapy. The components of the nanoparticle include: a) N-terminated A10 aptamer, a 57 bp nuclease-stabilized 2'-fluoropyrimidine RNA molecule modified with C_{18} -amine at the 3' end, for targeting PSMA-expressing PCa cells, and acting as a carrier for Dox; b) TCL-SPION coated with a carboxylic acid-PEG-derived, anti-biofouling polymer,^[29] which acts as both a MR contrast agent and as a carrier for Dox; and c) Dox, a chemotherapeutic agent that is intercalated in the aptamer and complexed with the TCL-SPION through charge interactions. The hydroxy and carbonyl groups on the surface of the TCL-SPION make them apt for the formulation of targeted nanoparticle platforms. The PEGylated surface prevents protein and cell adsorption, while the carboxyl groups allow conjugation of targeting moieties, like the A10 aptamer. TCL-SPIONs are also well suited for therapeutic delivery because of their low toxicity profiles.^[30-32]

Conjugation of the TCL-SPION with an A10 aptamer, using standard coupling chemistry, gave the TCL-SPION–Apt bioconjugate formulation (Figure 1 a); conjugation led to an increase in both size $(60.8 \pm 1.9$ to 66.4 ± 1.5 nm), and ζ -potential

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Figure 1. a) Schematic illustration of the TCL-SPION–Apt bioconjugate system; b) confirmation of TCL-SPION–Apt bioconjugate formation by gel electrophoresis (1. 100-bp ladder; 2. A10 aptamer; 3. TCL-SPION–Apt bioconjugate; 4. TCL-SPION).

 $(-36.0 \pm 1.8$ to -42.7 ± 3.8 mV) of the nanoparticles. The conjugation of the A10 aptamer to TCL-SPION was confirmed using agarose gel electrophoresis (Figure 1 b); the free A10 aptamers matched the 60-bp band in the 100-bp ladder, and the TCL-SPION–Apt bioconjugate lane showed a band at a much higher molecular weight, confirming the conjugation of Apt to TCL-SPION.

Differential uptake of the TCL-SPION–Apt bioconjugate by PSMA-expressing PCa cells (LNCaP), compared with non-PSMAexpressing PCa cells (PC3), was then confirmed in whole-cell assays by comparison with the uptake of TCL-SPION . Monitoring the uptake at regular time intervals (1, 3, 6, 12, 18 and 24 h) and using the Prussian blue reaction to visualize uptake, intracellular TCL-SPION–Apt bioconjugate uptake in LNCaP cells was detected as early as 3 h after dosing and progressively increases in a time-dependent manner. In contrast, TCL-SPION–Apt bioconjugates incubated with PC3 cells, and TCL-SPION incubated with LNCaP and PC3 cells, did not show intracellular uptake of the nanoparticles (Figure 2). These results confirm that TCL-SPION–Apt bioconjugates can differentially target PSMA-expressing PCa cells.

Figure 2. Prussian blue stained LNCaP and PC3 cells after incubation with TCL-SPION–Apt bioconjugate and non-targeted TCL-SPION.

The potential of the TCL-SPION–Apt bioconjugate as a targeted MR contrast agent was investigated by NMR studies. The longitudinal relaxation time (T_1) , and the transverse relaxation time (T_2) of the TCL-SPION–Apt bioconjugate and non-targeted TCL-SPION was measured using a single-sided NMR probe, after incubation with LNCaP and PC3 cells (6 h), and resuspension in Matrigel to simulate prostate tumors. Only a small change in T_1 and T_2 was observed for the non-targeted TCL-SPION in LNCaP cells (T_1 : 1939 \pm 116 to 1521 \pm 201 ms; T_2 : 104.2 ± 1.4 to 89.8 ± 1.1 ms), however, the TCL-SPION–Apt bioconjugate led to a dramatic decrease in T_1 and T_2 (T_1 : 1939 \pm 116 to 263 ± 23 ms; T_2 : 104.2 \pm 1.4 to 26.6 \pm 0.4 ms). As expected, TCL-SPION–Apt bioconjugates did not lead to significant reduction of T_1 and T_2 relaxation times in PC3 cells (Figure 3). These data suggest that TCL-SPION–Apt bioconjugates can detect PSMA-expressing PCa cells with high sensitivity.

After demonstrating the TCL-SPION–Apt bioconjugate's potential as a targeted MRI contrast agent, its potential as a therapeutic carrier was investigated. Firstly, the amount of Dox that can bind to the TCL-SPION–Apt bioconjugate through intercalation into the aptamer, and adsorption into the negatively charged polymer surface of the nanoparticle was determined. We had previously shown that the conjugation of Dox results in the quenching of its fluorescence.^[27,33] Using a spectrofluorophotometer, we titrated increasing concentrations of TCL-SPION and TCL-SPION–Apt bioconjugate against a fixed amount of Dox. As seen in Figure 4, the amount of TCL-SPION and TCL-SPION-Apt bioconjugates needed to quench 12 µg of Dox were 0.52 and 0.44 mg, respectively, giving loading efficiencies of 23.1 μ g Dox mg⁻¹ and 27.3 μ g Dox mg⁻¹ respectively. From the titration data, approximately 15% of Dox was intercalated in the aptamer and approximately 85% was bound to the polymer by electrostatic interactions.

The Dox-loaded TCL-SPION–Apt bioconjugates were evaluated for antiproliferation activity against both the LNCaP and PC3 cell lines (Figure 5). While free Dox was equipotent against both LNCaP and PC3 cell lines, the Dox-loaded TCL-SPION–Apt bioconjugate was significantly more potent against the PMSA-

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Figure 3. a) T_1 longitudinal relaxation times of LNCaP (\blacksquare) and PC3 (\blacksquare) cells incubated with TCL-SPION and TCL-SPION–Apt bioconjugates; b) T_2 transverse relaxation times of LNCaP (\blacksquare) and PC3 (\blacksquare) cells incubated with TCL-SPION and TCL-SPION–Apt bioconjugates.

expressing LNCaP cells relative to the non-targeted PC3 cells (cell viability: LNCaP 47.3 \pm 1.4% vs. PC3 69.3 \pm 1.7%). The data also showed that the cytotoxicity of Dox-loaded TCL-SPION– Apt bioconjugates was nearly as potent as free Dox. The observed cytotoxicity of Dox-loaded TCL-SPION–Apt bioconjugates to PC3 cells (69.3 \pm 1.7% compared with 95.7 \pm 1.9% of control) was likely due to uptake of Dox released after the dissociation of Dox from TCL-SPION–Apt bioconjugates.

In summary, a novel multifunctional TCL-SPION–Apt bioconjugate was synthesized, and shown to detect and treat PCa cells in vitro. However, the potential of TCL-SPION–Apt bioconjugates as targeted MR contrast agents for imaging of PCa needs to be further validated using in vivo models. TCL-SPION–Apt bioconjugates can be used as therapeutic carriers for the delivery of Dox, leading to selective delivery to PSMAexpressing cells without significant loss in cytotoxicity. The lack of sensitive and specific imaging agents, and effective therapeutic approaches for disseminated PCa, makes multifunctional nanoparticle technologies, such as TCL-SPION–Apt bioconju-

Figure 4. Fluorescence spectra of doxorubicin solution (12 μ g in 0.45 mL) with increasing amounts of a) TCL-SPION (from top to bottom: 5, 15, 30, 36, 52, 100, 150, 260, 360, and 520 µg) and b) TCL-SPION-Apt (from top to bottom: 4, 13, 22, 31, 44, 88, 133, 220, 311, and 440 µq).

Figure 5. MTT cell proliferation assay (LNCaP **:** PC3 **:** $* p < 0.005$, n=3).

gates, a potential approach for both the detection and the treatment of disseminated PCa. $[34-37]$ More broadly, the unique advantages of such multifunctional nanoparticles with diagnostic and therapeutic capabilities include: 1) targeted delivery of therapeutics to disease cells only, 2) observation of therapeutic delivery, and 3) detection of therapeutic response. Through the use of other disease-specific aptamers or other

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targeting molecules, as well as other strategies to conjugate therapeutic agents, similar multifunctional nanoparticles can be developed for applications in medicine.

Experimental Section

TCL-SPION–Apt bioconjugate: A solution of carboxy-TCL-SPION (50 μ L, 1.5 mg)^[29] was treated with N-(3-dimethylaminopropyl)-N'ethylcarbodiimide (EDC) (25 µL, 400 mm) and N-hydroxysuccinimide (NHS) (25 μ L, 100 mm) and gently shaken for 15 min. N-terminated Aptamer (RNA-TEC, Belgium, 1 mg in 100 µL) was then added, and the solution gently shaken for a further 4 h. Unreacted aptamer was removed using centrifugal filtration, five times (3,000 rpm, Nanosep centrifugal devices, 300 K, Pall Corp). Gel electrophoresis was carried out on 1.8% agarose gels; 0.05 mg of TCL-SPION–Apt bioconjugate and TCL-SPION were loaded. Tris-acetate-EDTA (TAE) buffer was used for the electrophoresis experiments. (Figure 1 b and Supporting Information, figure 1)

Iron (Prussian blue) stain: LNCaP and PC3 cell lines were grown in eight-well microscope chamber slides in RPMI-1640 and Ham's F-12 K medium respectively; both were treated with aqueous penicillin G (100 U mL⁻¹), streptomycin (100 μ gmL⁻¹), and 10% fetal bovine serum (FBS). Cells were grown to 70% confluency $(40000 \text{ cells cm}^{-2})$. Prior to dosing, cells were washed with PBS buffer and incubated with fresh media for 30 min. Cells were dosed with TCL-SPION–Apt bioconjugate or TCL-SPION $(0.1 \text{ mg} \text{ mL}^{-1})$ $(n=4)$ and incubated for 3-24 h at 37 °C, then washed two times with PBS and fixed with 4% formaldehyde. The cells were stained using the HT20 Iron Stain Kit (Sigma–Aldrich), and imaged with light microscopy.

Quantification of Dox loading: The amount of Dox loading onto TCL-SPION–Apt bioconjugate was calculated by fluorescence titration method. Before titration, the concentrations of TCL-SPION and TCL-SPION–Apt bioconjugate were determined from a standard curve of TCL-SPION at 310 nm (data not included).^[22] Increasing concentrations of TCL-SPION (7.46 mg mL $^{-1}$; 5, 15, 30, 36, 52, 100, 150, 260, 360 and 520 µg) or TCL-SPION-Apt bioconjugate $(6.3 \text{ mg} \text{ mL}^{-1}$; 4, 13, 22, 31, 44, 88, 133, 220, 311 and 440 μ g) were added stepwise to a fixed concentration of Dox (12 μ g in 0.45 mL). After each addition, the solution was mixed well and incubated at room temperature for 10 min. The fluorescence spectra were recorded by exciting the solution at 480 nm and recording the emission at 500–720 nm (3 mm slit) on a Shimadzu RF-PC100 spectrofluorophotometer. The maximum loading amount was defined as the concentration of nanoparticle required to give a 95% reduction in fluorescence emission compared with the spectra of an untreated solution of Dox.

 T_1 and T_2 relaxation time measurements: LNCaP and PC3 cell lines were grown in six-well plates to \sim 100% confluency. Prior to dosing, cells were washed with PBS buffer and incubated with fresh media for 30 min. Cells were dosed with TCL-SPION–Apt bioconjugate or TCL-SPION (0.1 mg mL $^{-1}$) ($n=4$) and incubated for 6 h at 37 \degree C, then washed two times with PBS, detached with trypsin, and fixed with 4% formaldehyde. Approximately one million cells of each sample were resuspended in 100 µL of Matrigel and placed inside a standard eight-well strip (Greiner Bio-One). Measurements were performed on a single-sided NMR probe at 25 °C at an operating frequency of 18.55 MHz (Profile NMR MOUSE, ACT Center for Technology, Aachen, Germany). A Minispec spectrometer (Bruker Optics, The Woodlands, TX, USA) was used for pulse sequence gen-

eration and data acquisition. A computer-controlled motion stage (Newport Corporation, Irvine, CA) positioned each well over the sensitive volume, and custom software, written in LabView (National Instruments, Austin, TX), coordinated the measurements. The acquisition time was approximately 30 s per sample. The longitudinal relaxation time T_1 was measured using a saturation recovery sequence. The signal intensity was measured with a short Carr Purcell Meiboom Gill (CPMG) echo sequence following a saturation pulse sequence, and a recovery time Δ . The echo time, TE, was 0.035 ms and 114 echoes were acquired for each time point. Seven time points were acquired per sample. The transverse relaxation time, $T₂$, was measured using a CPMG pulse sequence lasting 200 ms with an echo time of 0.035 ms. The data were averaged over 16 scans and the recovery time (TR) was 1.25 s for Sample 2 and 2.5 s for the others. The NMR sensor's static field gradient contributes to lower T_2 values and limits the maximum measurable T_2 . The measured $T₂$ values were all within the operating range of the instrument. The data were fit using a custom script running on MATLAB (The Mathworks, Natick, MA).

$$
I = I_0 \left(1 - e^{-\frac{A}{T_1}} \right) I = I_0 e^{-\frac{t}{T_2}} \tag{1}
$$

MTT cell proliferation assay: LNCaP and PC3 cell lines were grown in 48-well plates in RPMI-1640 and Ham's F-12 K medium, respectively; both were treated with aqueous penicillin G (100 U mL⁻¹), streptomycin (100 μ g mL⁻¹), and 10% FBS, at concentrations so as to allow 70% confluence in 24 h $($ -40000 cells cm⁻²). Prior to dosing, cells were washed with PBS buffer and incubated with fresh media for 30 min.

Cells were dosed with TCL-SPION–Apt(Dox) bioconjugates (0.1 mg mL $^{-1}$, 5 μ m Dox), TCL-SPION (0.1 mg mL $^{-1}$), or free Dox (5 μ m) and incubated for 3 h at 37 °C, then washed two times with PBS, and further incubated in fresh growth media for a total of 48 h. Cell viability was assessed colorimetrically with the MTT reagent (ATCC) following the standard protocol provided by the manufacturer. The absorbance was read with a microplate reader at 570 nm.

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