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Introduction

Small-interfering RNAs (siRNAs) are synthetic double-stranded RNAs capable of silencing gene expression *via* the RNAi pathway.¹ By virtue of their potential for targeting any known genes, the use of siRNAs to knock down undruggable oncogenes represents a promising anti-cancer approach.² When designed to target immune-related genes, siRNAs can also shape a tumor immune microenvironment.³ This immunogene therapy strategy has proved useful in orchestrating both the innate and adaptive immune systems to fight against cancers.^{4–6} However, due to their poor cell membrane permeability and undesired toxicity, safe and efficient delivery of siRNAs into target cells is a major barrier to advance their clinical applications.⁷ Currently, ligand conjugation⁸ and nanoparticle encapsulation⁹ are two prevailing ways to address siRNA delivery challenges. Although *N*-acetylgalactosamine conjugates and lipid nanoparticles have been approved for RNAi-based treatment of liver disorders,¹⁰ extrahepatic (e.g., tumor) delivery of siRNAs is still an unmet demand. Furthermore, to avoid side effects resulting from nonspecific siRNA

accumulation, RNAi activity needs to be confined to tumor sites.^{3,11}

Antibody–siRNA conjugates (ARCs) are a novel class of bivalent macromolecules consisting of monoclonal antibodies as carriers and siRNAs as payloads, which can be noncovalently or covalently assembled.^{12–14} Based on the specific interaction between antibodies and cell-membrane receptors, ARCs efficiently deliver siRNAs into target cells upon internalization.¹⁵ Using antibodies against tumor-associated membrane antigens, various ARCs have been constructed to direct tumor-specific delivery of siRNAs to downregulate oncogenes.^{16–25} Notably, synergistic anti-cancer treatment is feasible using all-in-one ARCs combining therapeutic antibodies and siRNAs together.^{26–28} Despite these promising accomplishments, ARCs have been rarely developed to manipulate immunotargets for anti-cancer immunotherapy. Moreover, no smart activatable ARC has been developed to minimize on-target/off-tumor side effects. This is crucial because ARCs could also be unintendedly taken up by non-target cells^{29,30} due to the nonexclusive antigen expression, and siRNA release from ARCs generally lacks controllability.

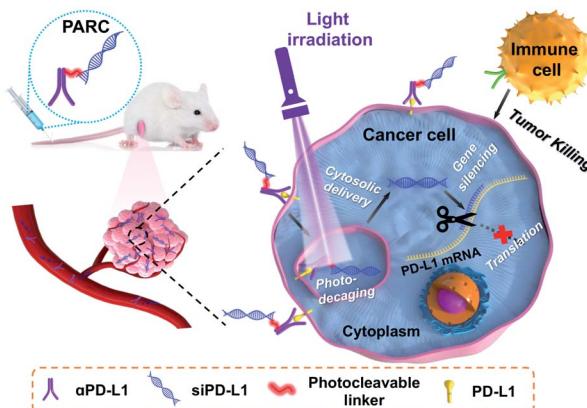
Herein, we report a novel photoresponsive antibody–siRNA conjugate (**PARC**) for remote-controllable siRNA delivery and cancer immunogene therapy (Scheme 1). As a non-invasive external stimulus, light is practical for spatiotemporal regulation of biological processes³¹ and is thereby chosen as the trigger for photoactivation. **PARC** comprises an anti-programmed death-ligand 1 antibody (α PD-L1) conjugated with a PD-L1-targeted siRNA (siPD-L1) through a photocleavable

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Scheme 1 Schematic illustration of using a photoresponsive antibody-siRNA conjugate (PARC) for tumor-specific delivery of siPD-L1. Photoirradiation on tumors liberates siPD-L1 from PARC to silence intracellular PD-L1 expression in the cytoplasm, which combines with the α PD-L1-mediated blockade of membrane PD-L1 to boost anti-cancer immunotherapy.

linker. PD-L1 is a membrane protein overexpressed on cancer cells to hamper immune cell function.³² α PD-L1 that blocks membrane PD-L1 to reinvigorate immune cell function has been approved for clinical treatment of some solid cancers.³³ Of note, α PD-L1 can get internalized into cancer cells upon interaction with membrane PD-L1,³⁴ which has been exploited for targeted delivery of cytotoxic chemo-drugs.^{35,36} On the other hand, PD-L1 is persistently produced inside cancer cells for replenishment of membrane PD-L1^{37,38} and secretion as circulating PD-L1 to exert pro-tumor effects.^{39,40} Therefore, in addition to membrane PD-L1, intracellular PD-L1 is also a pivotal target, which however is unreachable by α PD-L1. We thus integrated siPD-L1 with α PD-L1 in the PARC for not only targeted drug delivery but also synergistic inhibition of PD-L1 to amplify immunotherapeutic efficacy.

The mechanism of PARC-mediated activatable cancer immunogene therapy is proposed as follows. After systemic administration, PARC can accumulate in tumor tissues and specifically bind to PD-L1 on cancer cells, giving rise to the resurgence of immune cell activity. Following endocytosis, photoirradiation on tumors to break the cleavable linker in PARC leads to a tumor-specific release of siPD-L1. The free siPD-L1 then translocates into the cytoplasm to degrade PD-L1 mRNA, which prevents the continuous production of intracellular PD-L1 to further augment anti-cancer immunity. As a result, under light irradiation, PARC affords a synergistic blockade of PD-L1 to invoke efficacious cancer suppression.

Results and discussion

We prepared PARC by connecting siPD-L1 to α PD-L1 through a photocleavable *o*-nitrophenyl linker (Fig. 1a). While covalent ARCs have been reported before, conditional release and activation of siRNA have been barely achieved. This is due to the preferred use of a siRNA passenger strand for bioconjugation, leaving the guide strand free to execute gene silencing

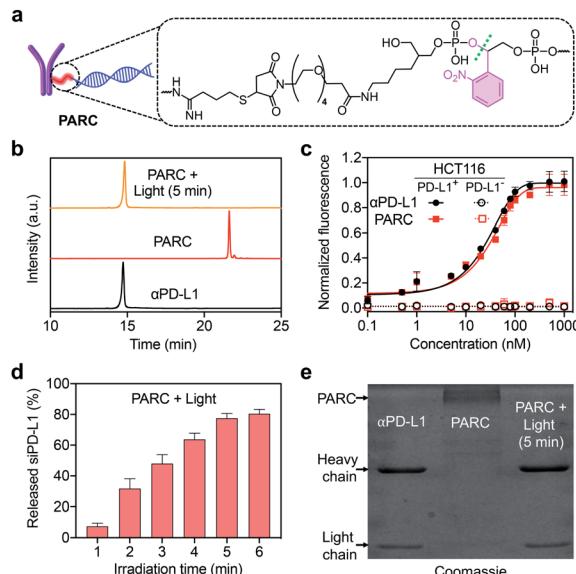


Fig. 1 (a) Chemical structure of the photocleavable linker in the PARC. (b) IEC analysis of α PD-L1, PARC, or PARC irradiated with 365 nm light (10 mW cm^{-2}) for 5 min. (c) Normalized fluorescence of PD-L1-positive or -negative HCT116 cells treated with α PD-L1 or PARC (0–1000 nM) and a fluorescent secondary antibody. (d) siPD-L1 release profiles of PARC irradiated with 365 nm light (10 mW cm^{-2}) for 1–6 min. (e) SDS-PAGE analysis of α PD-L1 and PARC without or with irradiation with 365 nm light (10 mW cm^{-2}) for 5 min. Data are shown as mean \pm SD ($n = 3$).

function.^{29,41,42} In this study, we selected the guide strand of siRNA for covalent attachment, which would allow restoration of the gene silencing function only after siRNA release. Briefly, we synthesized the siPD-L1 guide strand with an *o*-nitrophenyl moiety at its 3'-end using a standard solid-phase synthesis protocol,⁴³ followed by conversion of the terminal amine group to a maleimide group (Fig. S1 and S2†). Commercial α PD-L1 was functionalized with Traut's reagent to furnish a free sulphydryl group (Fig. S1†). After annealing with the passenger strand, siPD-L1 was coupled to α PD-L1 *via* the thiol-maleimide reaction, finally providing PARC under optimal conditions (Fig. S3†).

After ultrafiltration to remove free siPD-L1, PARC was obtained and characterized. Ion exchange chromatography (IEC) analysis revealed a significantly prolonged retention of PARC relative to α PD-L1 (Fig. 1b), indicating the successful addition of siPD-L1 to α PD-L1. We determined the average siPD-L1/ α PD-L1 ratio in PARC to be 2.2 using a previously reported method (Fig. S4†).⁴⁴ Noticeably, PARC and α PD-L1 displayed comparable binding profiles towards human colon cancer HCT116 cells that express a high level of membrane PD-L1,⁴⁵ with half maximal effective concentration (EC_{50}) values of $\sim 22 \text{ nM}$ and $\sim 19 \text{ nM}$, respectively (Fig. 1c). By contrast, such binding was undetectable when siPD-L1 was pre-transfected into HCT116 cells to knock down PD-L1 (Fig. 1c and S5†). These results suggest a negligible impact of siPD-L1 bioconjugation on the PD-L1-binding activity of α PD-L1 in PARC. We then investigated the photo-responsiveness of PARC. Upon exposure to 365 nm light,



PARC gradually unleashed siPD-L1, reaching a plateau (~78%) at 5 min (Fig. 1d). Meanwhile, sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) analysis also revealed a gradual liberation of α PD-L1 from **PARC** under photoirradiation (Fig. S6 \dagger). A complete release of α PD-L1 was observed after irradiation for 5 min (Fig. 1e), which was also confirmed by the IEC analysis (Fig. 1b). These findings could be ascribed to photocleavage of the *o*-nitrophenyl moiety in **PARC**, as judged by mass spectrometry analysis of the released siPD-L1 (Fig. S7 \dagger) and long-term stability of **PARC** in the dark (Fig. S8 \dagger).

We next explored the targeted delivery of siPD-L1 by **PARC** *in vitro*. To this end, we prepared a fluorescently labeled **PARC** with siPD-L1 and α PD-L1 tagged by Cy3 and Cy5, respectively (Fig. 2a and S1 \dagger). Cellular uptake of Cy5/Cy3-labeled **PARC** was monitored using a confocal fluorescence microscope. As shown in Fig. S9 \dagger , Cy5 and Cy3 fluorescence signals were detected both on the membrane and inside of HCT116 cells, which reached a maximum at 1 h post incubation. Notably, the obvious overlap of Cy5 and Cy3 signals indicates that **PARC** does not disassemble during internalization, warranting conditional release of siPD-L1. In comparison, we found no significant uptake of Cy5/Cy3-labeled **PARC** in PD-L1-negative HCT116 cells even after 24 h incubation (Fig. S10 \dagger). These results suggest a specific and effective cellular uptake of **PARC** mediated by the α PD-L1-PD-L1 interaction. We then irradiated Cy5/Cy3-labeled **PARC**-treated HCT116 cells with 365 nm light for 5 min at 1 h after incubation. Cy5 and Cy3 fluorescence signals were found to gradually separate (Fig. S11 \dagger). At 12 h post incubation, Cy3 signals were uniformly distributed throughout the cytoplasm, whereas Cy5 signals were mainly located on the cell membrane (Fig. 2b). Meanwhile, when co-staining the lysosome, we found that the colocalization ratio between Lysotracker and Cy3 was only ~18% (Fig. 2c). In contrast, under dark conditions, Cy5 and Cy3 signals remained overlapped, and the colocalization ratio between Lysotracker and Cy3 was ~90% (Fig. 2b and c).

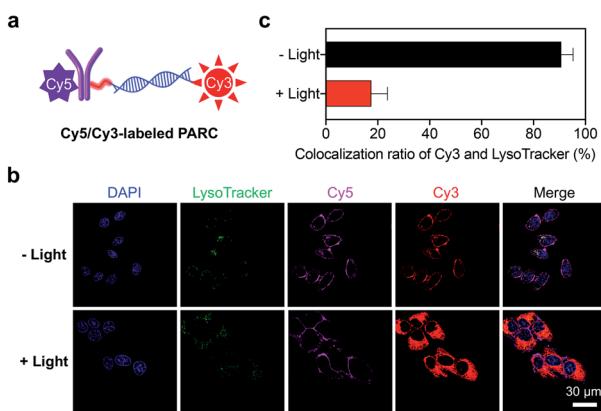


Fig. 2 (a) Schematic illustration of a Cy5/Cy3-labeled **PARC** structure. (b) Confocal fluorescence images of HCT116 cells treated with Cy5/Cy3-labeled **PARC** (150 nM) for 12 h. Cells were irradiated with 365 nm light (10 mW cm^{-2}) for 5 min at 1 h. The nucleus and lysosome were stained with DAPI and Lysotracker, respectively. (c) Colocalization ratios of fluorescence signals between Cy3 and Lysotracker. Data are shown as mean \pm SD.

These data thus demonstrate that, after cellular uptake and light irradiation, **PARC** could effectively release siPD-L1, which then successfully translocates from the lysosome into the cytoplasm. Additionally, since lysosomal trapping is a great limiting factor suffered by ARC-mediated siRNA delivery,⁴² our results also suggest that timely release of siRNAs may facilitate their lysosomal escape. In short, **PARC** allows tumor-targeted and photocontrollable cytosolic siRNA delivery, warranting further gene silencing applications.

Next, we assessed the biological activity of **PARC** *in vitro*. HCT116 cells were treated with **PARC** and irradiated with 365 nm light, followed by analysis of PD-L1 expression. As shown in Fig. S12 \dagger , **PARC** decreased PD-L1 expression in both photoirradiation time- and dose-dependent manners. At an irradiation time of 5 min and concentration of 150 nM, PD-L1 mRNA and protein levels dwindled to ~28% (Fig. 3a) and 0 (Fig. 3b) by **PARC**, respectively. In contrast, PD-L1 expression was unaffected by α PD-L1 or **PARC** without light irradiation, suggesting that the downregulation of PD-L1 expression results from the photoinduced release of siPD-L1 from **PARC**. To further confirm this, a control siRNA (si-ctrl) with a random sequence was used to replace siPD-L1 in **PARC**, giving a control ARC termed **CARC** (Fig. S1 and S2 \dagger). Comparatively, **CARC** could not regulate PD-L1 expression even under photoirradiation (Fig. 3a and b), validating the specificity of PD-L1 knockdown by **PARC**. These results thus demonstrate that **PARC** allows photoactivatable gene silencing.

It has been established that PD-L1 on cancer cells mediates immune cell tolerance through the interaction with PD-1, and its suppression can activate anti-cancer immunity.³² To test whether **PARC**-mediated PD-L1 suppression may impair PD-1 binding, a PD-1 protein was used to stain HCT116 cells treated with **PARC**. Flow cytometry analysis showed that **PARC** alone reduced binding of PD-1 to a cancer cell membrane by ~30% (Fig. 3c and d), similar to α PD-L1 and **CARC**. This result indicates that **PARC** and **CARC** retain the PD-L1-blockade activity of α PD-L1. In comparison, under light irradiation, **PARC** but not **CARC** significantly inhibited PD-1 binding by ~81%, revealing that silencing intracellular PD-L1 could dramatically prevent PD-1 binding. Consistently, PD-L1 suppression increased the sensitivity of HCT116 cells to immune cell killing,⁴⁶ as determined by co-culture with activated human peripheral blood mononuclear cells (PBMCs, Fig. 3e). PBMCs are mixed immune cells with abundant lymphocytes such as T cells and natural killer (NK) cells that are able to mediate anti-cancer immunity upon PD-1/PD-L1 blockade.^{32,47} Specifically, α PD-L1, **CARC**, and **PARC** slightly upregulated HCT116 cell apoptosis (~24%, Fig. 3f), which could be ascribed to the α PD-L1-mediated blockade of membrane PD-L1. However, upon light irradiation, **PARC** remarkably increased the apoptotic rate to ~50%, which coincides well with its inhibition of PD-L1 expression and PD-1 binding (Fig. 3a-d). Of note, treatment with a cocktail mixture of α PD-L1 and siPD-L1 complexed with a commercial transfection agent only elicited ~40% apoptosis (Fig. S13 \dagger). These results suggest a synergistic inhibition of PD-L1 by **PARC** with photoirradiation that favors anti-cancer immunity. Furthermore, we measured immunocytokines in the culture media (Fig. 3g) and S14 \dagger .



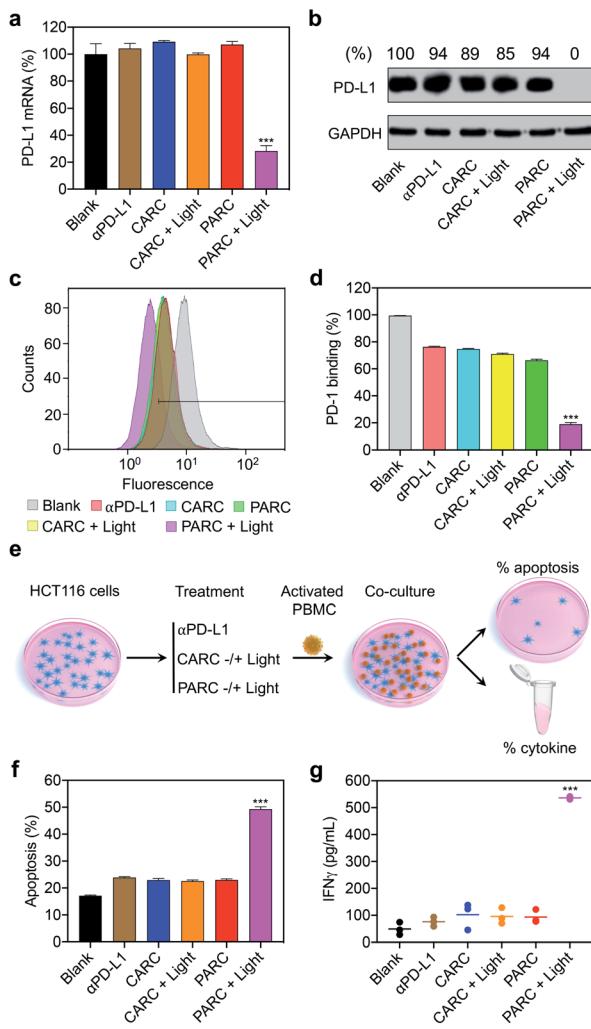


Fig. 3 (a) Reverse-transcription quantitative PCR (RT-qPCR) analysis of PD-L1 mRNA and (b) western blotting analysis of PD-L1 protein levels in HCT116 cells. GAPDH served as the internal control. Relative intensity ratios of PD-L1/GAPDH are shown at the top. (c and d) Flow cytometry analysis of immunostained PD-1 on HCT116 cells. (e) Schematic illustration of the experimental procedures for immune cell killing assay; PBMC, peripheral blood mononuclear cell. (f) Apoptosis rate of HCT116 cells determined by lactate dehydrogenase release assay. (g) IFN γ levels in the culture media. HCT116 cells were treated with α PD-L1 (150 nM), CARC (150 nM), or PARC (150 nM) for 48 h. Cells were irradiated with 365 nm light (10 mW cm^{-2}) for 5 min at 1 h. For co-culture assay, PBMCs were added to HCT116 cells at an effector-to-target ratio of 5 : 1 and incubated for another 24 h. Data are shown as mean \pm SD ($n = 3$). *** $P < 0.001$, relative to other groups.

under photoirradiation, PARC increased interferon γ (IFN γ) and tumor necrosis factor α (TNF α) levels by ~ 10.8 and ~ 7.9 fold, respectively, implying expected immune cell activation for tumor killing. Overall, these data jointly validate that PARC permits photoactivatable immunogene therapy.

Encouraged by the *in vitro* results, we further evaluated PARC *in vivo*. To find an optimal time point for light irradiation, we first performed fluorescence imaging in mice bearing subcutaneous HCT116 tumors. Cy5/Cy3-labeled PARC was used for tracing, revealing a maximum tumor accumulation at 4 h post

tail vein injection (Fig. S15†). Then, immunodeficient NCG mice bearing subcutaneous HCT116 tumors were constructed and transplanted with activated PBMCs. These mice were intravenously injected with saline, α PD-L1, CARC, or PARC at a same dose of antibody (5 mg kg $^{-1}$) and every 3 days for a total of 4 injections. For photoactivation, tumors were irradiated with 365 nm light for 5 min at 4 h post each injection. After 14 days, tumor volumes of the PARC without photoirradiation group increased from $\sim 150 \text{ mm}^3$ to $\sim 1500 \text{ mm}^3$, similar to the saline and α PD-L1 groups (Fig. 4a). The insignificant cancer suppression could be attributed to the moderate dose of α PD-L1.⁴⁸ However, with light irradiation, tumor growth of the PARC group but not the CARC group was significantly suppressed. Notably, during the treatment, body weights of all groups remained nearly constant (Fig. 4b), suggesting general biosafety. At the end of treatment, tumors of the PARC with photoirradiation group were inhibited by $\sim 62\%$ (Fig. 4c and S16†), compared to the other groups. Our results thus demonstrate that, under light irradiation, PARC can evoke potent anti-cancer effects.

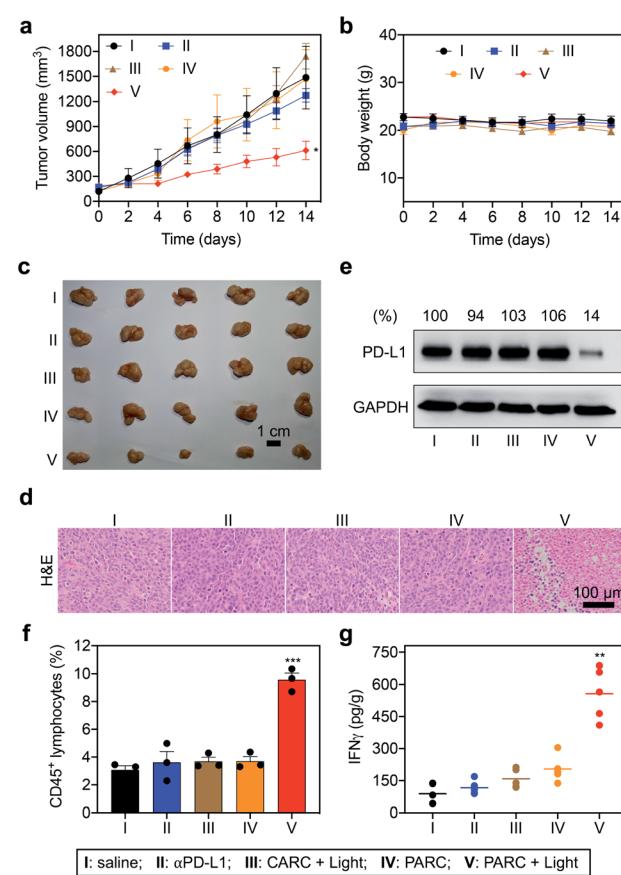


Fig. 4 (a) Tumor growth curves and (b) body weights of HCT116 tumor-bearing NCG mice that were intravenously injected with saline, α PD-L1 (0.7 nmol), CARC (0.7 nmol), or PARC (0.7 nmol) every 3 days for a total of 4 injections. Tumors were irradiated with 365 nm light (50 mW cm^{-2}) for 5 min at 4 h post each injection. (c) Optical images, (d) H&E staining, (e) western blotting analysis of PD-L1, (f) flow cytometry analysis of CD45 $^+$ lymphocytes, and (g) IFN γ levels. (e) Relative intensity ratios of PD-L1/GAPDH are shown at the top. Data are shown as mean \pm SD ($n = 3-5$). * $P < 0.05$, ** $P < 0.01$, and *** $P < 0.001$, relative to the other groups.

To validate the immunotherapeutic efficacy of **PARC**, we further analyzed tumor tissues. The histological examination with haematoxylin and eosin (H&E) staining showed obvious cell death in tumor tissues of the **PARC** with photoirradiation group compared to the other groups (Fig. 4d). TdT-mediated dUTP nick end labelling (TUNEL) staining also demonstrated tumor cell apoptosis in the **PARC** with photoirradiation group (Fig. S17†). The apoptosis was mainly detected in the area with 5 mm depth where 365 nm light can penetrate,⁴⁹ suggesting a local anti-cancer effect. However, we found no significant damage in normal tissues (heart, liver, spleen, lung, and kidney) of all groups as indicated by H&E staining (Fig. S18†), further confirming its biosafety. Western blotting analysis showed that **PARC** with photoirradiation significantly decreased PD-L1 protein levels by ~86% compared to the other groups (Fig. 4e), verifying the photoactivatable gene silencing *in vivo*. To unveil whether lymphocytes in PBMCs mediate anti-cancer immunity, we evaluated their invasion in tumor tissues. Flow cytometry analysis showed that **PARC** with photoirradiation increased the tumor infiltration of CD45⁺ lymphocytes (Fig. 4f and S19†) and CD3⁺ T cells (Fig. S20†) ~2.6 and ~3.6 fold, respectively, relative to the other groups. Meanwhile, IFN γ (Fig. 4g) and TNF α (Fig. S21†) levels in tumor tissues of the **PARC** with photoirradiation group were upregulated ~4.1 and ~3.1 fold, respectively. Taken together, these *ex vivo* data prove photoactivatable immunogene therapy by **PARC**, which also match the *in vivo* anti-cancer results.

Conclusion

In summary, we developed a photoresponsive conjugate of α PD-L1 with siPD-L1 (**PARC**), which were connected through a photocleavable linker. **PARC** could target cancer cells through α PD-L1/PD-L1 interaction, allowing not only resurgence of immune cell activity but also intracellular transportation of siPD-L1. Under light irradiation to break the photocleavable linker, siPD-L1 could get released from **PARC** and escape from the lysosome into the cytoplasm, where intracellular PD-L1 expression was subsequently silenced. Eventually, **PARC** afforded a synergistic blockade of PD-L1 to potently activate immune cells to kill cancer cells. We demonstrated the potential of **PARC** both *in vitro* and *in vivo*. Our work thus presents a novel approach to achieve tumor-targeted and photoinduced siRNA delivery, which also promises photoactivatable immunogene therapy. To further the *in vivo* applications of **PARC**, optimized or alternative photocleavable linkers⁵⁰ with long-wavelength excitation that ensure deep penetration of light as well as safe and fast release of siRNA deserve future exploration. Overall, given its high biocompatibility and versatility to assemble different antibody/siRNA combos, we believe this photoresponsive **ARC** will open new avenues for cancer treatment.

Experimental section

Preparation of **PARC** and **CARC**

α PD-L1 (2 mg mL⁻¹) was mixed with Traut's reagent (0–10 eq.) in the EDTA-PB (100 nM) phosphate buffer containing 2 mM

EDTA, pH = 6.8) buffer, and the mixture was incubated at 37 °C for 1 h. The small molecules were removed by ultrafiltration (molecular weight cut-off, 3 kDa) with EDTA-PB 5 times. The supernatant was collected and mixed with maleimide-labeled siPD-L1 or si-ctrl (10 eq.) in EDTA-PB and incubated at 37 °C for 2 h. Excess siRNAs were removed by ultrafiltration (molecular weight cut-off, 50 kDa) with EDTA-PB 5 times. To prepare Cy5/Cy3-labeled **PARC**, α PD-L1 was first labeled with Cy5 and further labeled with Cy3-modified siPD-L1. Briefly, α PD-L1 (2 mg mL⁻¹) was mixed with Cy5-NHS (0.7 mM) in PBS buffer and the mixture was incubated at 37 °C for 1 h. Small molecules were removed by ultrafiltration (molecular weight cut-off, 3 kDa) with EDTA-PB buffer 5 times. Then, Cy5-labeled α PD-L1 was further modified with maleimide/Cy3-labeled siPD-L1 using the same method as described above.

Detection of siPD-L1 release

PARC (100 nM) in PBS was irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm⁻²) for 0–6 minutes. The released siPD-L1 was collected by ultrafiltration (molecular weight cut-off, 50 kDa), and the concentration was determined using a NanoDrop 2000 (Thermo Scientific). The siPD-L1 release percentages were calculated by dividing the measured amount by the initial amount. Then, the collected siRNA was desalting, lyophilized, and characterized by mass spectrometry.

Stability test

PARC (150 nM) was incubated in PBS at 37 °C. IEC analysis was performed at 0, 6, 12, and 24 h time points to indicate the stability of **PARC**.

Cell culture

HCT116 cells were cultured in high glucose DMEM containing 10% FBS and 1% penicillin/streptomycin, and maintained in 5% CO₂ at 37 °C.

Western blotting

After reaching ~80% confluence, HCT116 cells were transfected with siPD-L1 (200 nM) or si-ctrl (200 nM) using Lipofectamine 2000 according to the manufacturer's protocol. Or, HCT116 cells were treated with α PD-L1 (150 nM), **CARC** (150 nM), or **PARC** (150 nM). For the photoirradiation group, after incubation for 1 h, HCT116 cells were further irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm⁻²) for 5 min. At 48 h, cell lysates were extracted using RIPA lysis buffer (150 mM NaCl, 10 mM Tris, pH 7.5, 1% NP40, 1% deoxycholate, 0.1% SDS, and 1× protease inhibitor cocktail). To measure the expression levels of PD-L1 in tumors, total proteins were extracted from 100 mg tumor tissues. The protein concentration was determined using a BCA protein assay kit. A total of 40 μ g of protein was denatured in 1× SDS loading buffer and resolved by 10% SDS-PAGE, followed by transferring to a PVDF membrane and block in 5% nonfat milk in TBST. The PVDF membrane was then stained with primary antibodies for PD-L1 and GAPDH. After washing, the membrane was further incubated with the



appropriate secondary antibody and finally visualized using ECL reagents.

RT-qPCR

After reaching $\sim 80\%$ confluence, HCT116 cells were transfected with siPD-L1 (200 nM) or si-ctrl (200 nM) using Lipofectamine 2000 according to the manufacturer's protocol. Or, HCT116 cells were treated with α PD-L1 (150 nM), CARC (150 nM), or PARC (150 nM). For the photoirradiation group, after incubation for 1 h, HCT116 cells were further irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm^{-2}) for 5 min. After incubation for 48 h, total RNA was isolated from HCT116 cells with TRIZOL reagent (Invitrogen). cDNA was synthesized using a SuperScript III kit (Invitrogen), and qPCR was performed with a SYBR Green PCR Master Mix (Vazyme Biotech). Primers: PD-L1: forward: 5'-TTGCTGAACGCCCTATA-3'; reverse: 5'-TGC-TTGTCCAGATGACTTCG-3'; GAPDH: forward: 5'-GGACCT-GACCT-GCCGTCTAG-3'; reverse: 5'-GTAGCCCAG-GATGCCCTTGA-3'. GAPDH served as the internal control.

Confocal fluorescence imaging

1×10^5 HCT116 cells were seeded on 35 mm glass-bottom culture dishes. When reaching $\sim 80\%$ confluence, cells were incubated with Cy5/Cy3-labeled PARC (150 nM). To explore the effect of light irradiation, at 1 h time point, cells were irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm^{-2}) for 5 min. After incubation for different time points, cells were washed with PBS three times. Then, cells were stained with DAPI and Lysotracker according to the manufacturer's instructions. The imaging acquisitions were carried out immediately using a Leica confocal microscope.

PD-1 binding analysis

After reaching $\sim 80\%$ confluence, HCT116 cells were treated with α PD-L1 (150 nM), CARC (150 nM), or PARC (150 nM). For the photoirradiation group, after incubation for 1 h, HCT116 cells were further irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm^{-2}) for 5 min. At 48 h, cells were incubated with $5 \mu\text{g mL}^{-1}$ recombinant human PD-1 FC chimera protein at room temperature for 30 min. After washing with PBS 3 times, cells were further incubated with an anti-human IgG Alexa Fluor 488 dye conjugated antibody (ThermoFisher) (1 : 200) for another 30 min. Following washing with PBS 3 times, Alexa Fluor 488-positive cells were analyzed by flow cytometry.

Immune cell killing assay

After reaching $\sim 80\%$ confluence, HCT116 cells were treated with α PD-L1 (150 nM), CARC (150 nM), or PARC (150 nM). For the photoirradiation group, after incubation for 1 h, HCT116 cells were further irradiated by using an 8 W hand-held 365 nm UV lamp (10 mW cm^{-2}) for 5 min. At 24 h, PBMCs (MT-BIO) were resuscitated and activated with an anti-CD3 antibody (100 ng mL^{-1}), an anti-CD28 antibody (100 ng mL^{-1}), and IL2 (10 ng mL^{-1}) for 24 h. Then, activated PBMCs were incubated with HCT116 cells at an effector-to-target ratio of 5 : 1 at 37°C

for another 24 h. Then, lactate dehydrogenase analysis was carried out according to the manufacturer's protocol.

In vivo and ex vivo imaging

Animal care and euthanasia were carried out with the approval of the Institutional Animal Care and Use Committee (IACUC) of Nanjing University. A total of 2×10^6 HCT116 cells were subcutaneously injected into the right back flank of 4–5 weeks old female nude mice (Nanjing Biomedical Research Institute of Nanjing University). When tumor diameters reached $\sim 1 \text{ cm}$, these tumor-bearing mice were injected *via* the tail vein with Cy5/Cy3-labeled PARC (0.7 nmol). Fluorescence imaging was then performed with an IVIS Lumina XR III small animal imaging system. To acquire fluorescence signals from siPD-L1, a Cy3 filter was used. To acquire fluorescence signals from α PD-L1, a Cy5 filter was used. Identical illumination settings (lamp voltage, filters, f/stop, field of views, and binning) were used for acquiring all images. Fluorescence images were taken at 0, 1, 2, 4 and 8 h post injection. The images were then analyzed using Living Image 3.2 software (Caliper, Hopkinton, MA). The fluorescence emission was normalized to photons per second per centimeter squared per steradian (p per s per cm^2 per sr). For *ex vivo* imaging, tumor and other major tissues were dissected and put on black paper. The *ex vivo* images were acquired immediately using an IVIS Lumina XR III system with the same illumination setting as *in vivo* imaging. The images were also processed with the same method as described above.

In vivo cancer therapy

Animal care and euthanasia were carried out with the approval of the Institutional Animal Care and Use Committee (IACUC) of Nanjing University. A total of 2×10^6 HCT116 cells were subcutaneously injected into the right back flank of 4–5 weeks old female immune-deficient NOD/Prkdc^{null}/IL2RG^{null} (NCG) mice (Nanjing Biomedical Research Institute of Nanjing University). When tumor volumes reached $\sim 100\text{--}200 \text{ mm}^3$, each mouse received 1×10^7 activated human PBMCs *via* tail vein injection to generate a PBMC-humanized mouse. IL-2 (2.75 μg per mouse) was intraperitoneally injected for 3 consecutive days. These mice were then randomly divided into five groups ($n = 5/\text{group}$) and respectively treated with saline, α PD-L1 (0.7 nmol), CARC (0.7 nmol), or PARC (0.7 nmol) through tail vein injection every 3 days for a total of 4 injections. Mice treated with CARC or PARC were subject to irradiation with a LED 365 nm UV lamp (50 mW cm^{-2}) for 5 min at 4 h after each injection. Tumor volumes and body weights were recorded every two days. Tumor volumes were calculated using the following formula: $V = (\text{width}^2 \times \text{length})/2$. After 14 days, these mice were sacrificed, and tissues were collected for further analysis.

Immune cell infiltration

Tumors were cut into small pieces and subjected to mechanical disruption and separation, followed by passing through 70 μm strainers and treatment with the erythrocyte lysis solution (BioLegend) according to the manufacturer's instructions. The single-cell suspensions were then stained with an FITC-labeled



anti-CD3 antibody and PE-labeled anti-CD45 antibody, respectively. Cells were further analyzed by flow cytometry.

Immunocytokine detection

After reaching ~80% confluence, HCT116 cells were treated with α PD-L1 (150 nM), CARC (150 nM), or PARC (150 nM). For the photoirradiation group, these treated cells were further subjected to 5 min' 365 nm light irradiation (10 mW cm^{-2}) after incubation for 1 h. At 24 h, PBMCs (MT-BIO, Shanghai) were resuscitated and activated with an anti-CD3 antibody (100 ng mL^{-1}), an anti-CD28 antibody (100 ng mL^{-1}), and IL2 (10 ng mL^{-1}) for 24 h. Activated PBMCs were incubated with HCT116 cells at an effector-to-target ratio of 5 : 1 at 37 °C for another 24 h. Then, culture media were collected for analysis. To detect the expression of TNF α and IFN γ in tumors, tumor tissues were digested. The supernatants were collected and diluted. TNF α and IFN γ levels were analyzed using ELISA kits according to the manufacturer's protocols.

Statistical analysis

Statistical comparisons between groups were evaluated by Student's *t*-test. Data are shown as mean \pm SD. **P* < 0.05, ***P* < 0.01, and ****P* < 0.001 were considered to be statistically significant.

Data availability

The data that support the findings of this study are available from the corresponding author upon reasonable request.

Author contributions

X. Wang and X. Xiao performed the experiments. Y. Feng drew the schematic figures. J. Li and Y. Zhang designed the project and wrote the manuscript.

Conflicts of interest

There are no conflicts to declare.

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References

- 1 T. C. Roberts, R. Langer and M. J. A. Wood, *Nat. Rev. Drug Discovery*, 2020, **19**, 673–694.
- 2 B. Kim, J. H. Park and M. J. Sailor, *Adv. Mater.*, 2019, **31**, e1903637.
- 3 C. Zhang and K. Pu, *Chem. Soc. Rev.*, 2020, **49**, 4234–4253.
- 4 D. W. Zheng, F. Gao, Q. Cheng, P. Bao, X. Dong, J. X. Fan, W. Song, X. Zeng, S. X. Cheng and X. Z. Zhang, *Nat. Commun.*, 2020, **11**, 1985.
- 5 H. Xiao, Y. Guo, B. Li, X. Li, Y. Wang, S. Han, D. Cheng and X. Shuai, *ACS Cent. Sci.*, 2020, **6**, 1208–1222.
- 6 K. W. Huang, F. F. Hsu, J. T. Qiu, G. J. Chern, Y. A. Lee, C. C. Chang, Y. T. Huang, Y. C. Sung, C. C. Chiang, R. L. Huang, C. C. Lin, T. K. Dinh, H. C. Huang, Y. C. Shih, D. Alson, C. Y. Lin, Y. C. Lin, P. C. Chang, S. Y. Lin and Y. Chen, *Sci. Adv.*, 2020, **6**, eaax5032.
- 7 R. L. Setten, J. J. Rossi and S. P. Han, *Nat. Rev. Drug Discovery*, 2019, **18**, 421–446.
- 8 S. Benizri, A. Gissot, A. Martin, B. Viallet, M. W. Grinstaff and P. Barthelemy, *Bioconjugate Chem.*, 2019, **30**, 366–383.
- 9 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.
- 10 J. A. Kulkarni, D. Witzigmann, S. B. Thomson, S. Chen, B. R. Leavitt, P. R. Cullis and R. van der Meel, *Nat. Nanotechnol.*, 2021, **16**, 630–643.
- 11 C. Yu, L. Li, P. Hu, Y. Yang, W. Wei, X. Deng, L. Wang, F. R. Tay and J. Ma, *Adv. Sci.*, 2021, **8**, 2100540.
- 12 J. Dugal-Tessier, S. Thirumalairajan and N. Jain, *J. Clin. Med.*, 2021, **10**, 838.
- 13 I. Dovgan, O. Koniev, S. Kolodych and A. Wagner, *Bioconjugate Chem.*, 2019, **30**, 2483–2501.
- 14 I. V. Chernikov, V. V. Vlassov and E. L. Chernolovskaya, *Front. Pharmacol.*, 2019, **10**, 444.
- 15 N. Baumer, W. E. Berdel and S. Baumer, *Mol. Pharm.*, 2017, **14**, 1339–1351.
- 16 D. Klein, S. Goldberg, C. S. Theile, R. Dambra, K. Haskell, E. Kuhar, T. Lin, R. Parmar, M. Manoharan, M. Richter, M. Wu, J. Mendrola Zarazowski, V. Jadhav, M. A. Maier, L. Sepp-Lorenzino, K. O'Neil and V. Dudkin, *Mol. Ther.*, 2021, **29**, 2053–2066.
- 17 A. R. Nanna, A. V. Kel'in, C. Theile, J. M. Pierson, Z. X. Voo, A. Garg, J. K. Nair, M. A. Maier, K. Fitzgerald and C. Rader, *Nucleic Acids Res.*, 2020, **48**, 5281–5293.
- 18 S. J. Shi, L. J. Wang, D. H. Han, J. H. Wu, D. Jiao, K. L. Zhang, J. W. Chen, Y. Li, F. Yang, J. L. Zhang, G. X. Zheng, A. G. Yang, A. Z. Zhao, W. J. Qin and W. H. Wen, *Theranostics*, 2019, **9**, 1247–1263.
- 19 N. Baeumer, N. Appel, L. Terheyden, F. Buchholz, C. Rossig, C. Mueller-Tidow, W. E. Berdel and S. Baeumer, *Nat. Protoc.*, 2016, **11**, 22–36.
- 20 K. Jiang, J. Li, J. Yin, Q. Ma, B. Yan, X. Zhang, L. Wang, L. Wang, T. Liu, Y. Zhang, Q. Fan, A. Yang, X. Qiu and B. Ma, *Biomaterials*, 2015, **59**, 77–87.



21 Y. Su, L. Yu, N. Liu, Z. Guo, G. Wang, J. Zheng, M. Wei, H. Wang, A. G. Yang, W. Qin and W. Wen, *Cancer Lett.*, 2013, **338**, 282–291.

22 Y. D. Yao, T. M. Sun, S. Y. Huang, S. Dou, L. Lin, J. N. Chen, J. B. Ruan, C. Q. Mao, F. Y. Yu, M. S. Zeng, J. Y. Zang, Q. Liu, F. X. Su, P. Zhang, J. Lieberman, J. Wang and E. Song, *Sci. Transl. Med.*, 2012, **4**, 130ra148.

23 S. Dou, Y. D. Yao, X. Z. Yang, T. M. Sun, C. Q. Mao, E. W. Song and J. Wang, *J. Controlled Release*, 2012, **161**, 875–883.

24 Y. Ma, C. M. Kowollik, P. M. Swiderski, M. Kortylewski, H. Yu, D. A. Horne, R. Jove, O. L. Caballero, A. J. Simpson, F. T. Lee, V. Pillay and A. M. Scott, *ACS Chem. Biol.*, 2011, **6**, 962–970.

25 E. Song, P. Zhu, S. K. Lee, D. Chowdhury, S. Kussman, D. M. Dykxhoorn, Y. Feng, D. Palliser, D. B. Weiner, P. Shankar, W. A. Marasco and J. Lieberman, *Nat. Biotechnol.*, 2005, **23**, 709–717.

26 W. Fu, H. Sun, Y. Zhao, M. Chen, L. Yang, X. Yang and W. Jin, *Mol. Immunol.*, 2018, **99**, 124–133.

27 Y. Lu, L. Liu, Y. Wang, F. Li, J. Zhang, M. Ye, H. Zhao, X. Zhang, M. Zhang, J. Zhao, B. Yan, A. Yang, H. Feng, R. Zhang and X. Ren, *Biomaterials*, 2016, **76**, 196–207.

28 S. Baumer, N. Baumer, N. Appel, L. Terheyden, J. Fremerey, S. Schelhaas, E. Wardelmann, F. Buchholz, W. E. Berdel and C. Muller-Tidow, *Clin. Cancer Res.*, 2015, **21**, 1383–1394.

29 O. Zavoiira, B. Brunner, P. Casteels, L. Zimmermann, M. Ozog, C. Boutton, M. W. Helms, T. Wagenaar, V. Adam, J. Peterka, C. Metz-Weidmann, P. Deschaght, S. Scheidler and K. Jahn-Hofmann, *Mol. Pharm.*, 2021, **18**, 1048–1060.

30 D. Peer, P. Zhu, C. V. Carman, J. Lieberman and M. Shimaoka, *Proc. Natl. Acad. Sci. U. S. A.*, 2007, **104**, 4095–4100.

31 J. Li, H. Kong, C. Zhu and Y. Zhang, *Chem. Sci.*, 2020, **11**, 3390–3396.

32 S. L. Topalian, J. M. Taube and D. M. Pardoll, *Science*, 2020, **367**, eaax0182.

33 J. Xin Yu, J. P. Hodge, C. Oliva, S. T. Neftelinov, V. M. Hubbard-Lucey and J. Tang, *Nat. Rev. Drug Discovery*, 2020, **19**, 163–164.

34 M. L. Burr, C. E. Sparbier, Y. C. Chan, J. C. Williamson, K. Woods, P. A. Beavis, E. Y. N. Lam, M. A. Henderson, C. C. Bell, S. Stolzenburg, O. Gilan, S. Bloor, T. Noori, D. W. Morgens, M. C. Bassik, P. J. Neeson, A. Behren, P. K. Darcy, S. J. Dawson, I. Voskoboinik, J. A. Trapani, J. Cebon, P. J. Lehner and M. A. Dawson, *Nature*, 2017, **549**, 101–105.

35 C. W. Li, S. O. Lim, E. M. Chung, Y. S. Kim, A. H. Park, J. Yao, J. H. Cha, W. Xia, L. C. Chan, T. Kim, S. S. Chang, H. H. Lee, C. K. Chou, Y. L. Liu, H. C. Yeh, E. P. Perillo, A. K. Dunn, C. W. Kuo, K. H. Khoo, J. L. Hsu, Y. Wu, J. M. Hsu, H. Yamaguchi, T. H. Huang, A. A. Sahin, G. N. Hortobagyi, S. S. Yoo and M. C. Hung, *Cancer Cell*, 2018, **33**, 187–201.

36 S. Sau, A. Petrovici, H. O. Alsaab, K. Bhise and A. K. Iyer, *Cancers*, 2019, **11**, 232.

37 Y. Wu, W. Chen, Z. P. Xu and W. Gu, *Front. Immunol.*, 2019, **10**, 2022.

38 Y. Wang, H. Wang, H. Yao, C. Li, J. Y. Fang and J. Xu, *Front. Pharmacol.*, 2018, **9**, 536.

39 G. Chen, A. C. Huang, W. Zhang, G. Zhang, M. Wu, W. Xu, Z. Yu, J. Yang, B. Wang, H. Sun, H. Xia, Q. Man, W. Zhong, L. F. Antelo, B. Wu, X. Xiong, X. Liu, L. Guan, T. Li, S. Liu, R. Yang, Y. Lu, L. Dong, S. McGettigan, R. Somasundaram, R. Radhakrishnan, G. Mills, Y. Lu, J. Kim, Y. H. Chen, H. Dong, Y. Zhao, G. C. Karakousis, T. C. Mitchell, L. M. Schuchter, M. Herlyn, E. J. Wherry, X. Xu and W. Guo, *Nature*, 2018, **560**, 382–386.

40 M. Poggio, T. Hu, C. C. Pai, B. Chu, C. D. Belair, A. Chang, E. Montabana, U. E. Lang, Q. Fu, L. Fong and R. Bleloch, *Cell*, 2019, **177**, 414–427.

41 T. Sugo, M. Terada, T. Oikawa, K. Miyata, S. Nishimura, E. Kenjo, M. Ogasawara-Shimizu, Y. Makita, S. Imaichi, S. Murata, K. Otake, K. Kikuchi, M. Teratani, Y. Masuda, T. Kamei, S. Takagahara, S. Ikeda, T. Ohtaki and H. Matsumoto, *J. Controlled Release*, 2016, **237**, 1–13.

42 T. L. Cuellar, D. Barnes, C. Nelson, J. Tanguay, S.-F. Yu, X. Wen, S. J. Scales, J. Gesch, D. Davis, A. van Brabant Smith, D. Leake, R. Vandlen and C. W. Siebel, *Nucleic Acids Res.*, 2015, **43**, 1189–1203.

43 L. Chen, Y. Sun, J. Li and Y. Zhang, *Chem. Commun.*, 2020, **56**, 627–630.

44 X. X. Wang, Y. Wang, J. Li, T. Tian, J. B. Li, Z. J. Guo and Y. Zhang, *Adv. Ther.*, 2022, **5**, 2100161.

45 Q. Liu, L. Jiang, K. Li, H. Li, G. Lv, J. Lin and L. Qiu, *Cancer Immunol. Immunother.*, 2021, **70**, 1721–1733.

46 H. Wang, H. Yao, C. Li, H. Shi, J. Lan, Z. Li, Y. Zhang, L. Liang, J. Y. Fang and J. Xu, *Nat. Chem. Biol.*, 2019, **15**, 42–50.

47 J. Hsu, J. J. Hodgins, M. Marathe, C. J. Nicolai, M. C. Bourgeois-Daigneault, T. N. Trevino, C. S. Azimi, A. K. Scheer, H. E. Randolph, T. W. Thompson, L. Zhang, A. Iannello, N. Mathur, K. E. Jardine, G. A. Kirn, J. C. Bell, M. W. McBurney, D. H. Raulet and M. Ardolino, *J. Clin. Invest.*, 2018, **128**, 4654–4668.

48 Y. Zhang, C. Fang, R. E. Wang, Y. Wang, H. Guo, C. Guo, L. Zhao, S. Li, X. Li, P. G. Schultz, Y. J. Cao and F. Wang, *Proc. Natl. Acad. Sci. U. S. A.*, 2019, **116**, 15889–15894.

49 J. Wang, Y. Liu, Y. Liu, S. Zheng, X. Wang, J. Zhao, F. Yang, G. Zhang, C. Wang and P. R. Chen, *Nature*, 2019, **569**, 509–513.

50 R. Weinstain, T. Slanina, D. Kand and P. Klan, *Chem. Rev.*, 2020, **120**, 13135–13272.

