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Macrophage-mediated tumor-targeted delivery of engineered *Salmonella typhimurium* VNP20009 in anti-PD1 therapy against melanoma



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Abstract Bacterial antitumor therapy has great application potential given its unique characteristics, including genetic manipulation, tumor targeting specificity and immune system modulation. However, the nonnegligible side effects and limited efficacy of clinical treatment limit their biomedical applications. Engineered bacteria for therapeutic applications ideally need to avoid their accumulation in normal organs and possess potent antitumor activity. Here, we show that macrophage-mediated tumor-targeted delivery of *Salmonella typhimurium* VNP20009 can effectively reduce the toxicity caused by administering VNP20009 alone in a melanoma mouse model. This benefits from tumor-induced chemotaxis for macrophages combined with their slow release of loaded strains. Inspired by changes in the tumor microenvironment, including a decrease in intratumoral dysfunctional CD8⁺ T cells and an increase in PDL1 on the tumor cell surface, macrophages were loaded with the engineered strain VNP-PD1nb, which can express and secrete anti-PD1 nanoantibodies after they are released from macrophages. This novel

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triple-combined immunotherapy significantly inhibited melanoma tumors by reactivating the tumor microenvironment by increasing immune cell infiltration, inhibiting tumor cell proliferation, remodeling TAMs to an M1-like phenotype and prominently activating CD8⁺ T cells. These data suggest that novel combination immunotherapy is expected to be a breakthrough relative to single immunotherapy.

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1. Introduction

Since Dr. William Coley treated cancer patients with heat-inactivated Gram-positive bacteria (*Streptococcus*) and Gram-negative bacteria (*Salmonella mucilago*) in the late 19th century, an increasing number of microorganisms have been developed for tumor therapy¹. These microbes, also known as oncolytic bacteria, include *Salmonella typhimurium* and can highly colonize tumors due to their facultative anaerobic properties and the anaerobic environment within the tumor^{2–4}. Because of the immunosuppressive tumor environment, the amplification of these oncolytic bacteria can be implemented rapidly in tumors followed by reactivation of tumor immunity and tumor regression^{5–7}. VNP20009 (abbreviated VNP), an attenuated strain of *S. typhimurium*, has received wide interest for its antitumor effects in preclinical models and relative safety^{8,9}. However, phase I clinical trials of VNP were terminated due to its low therapeutic efficacy⁹. Moreover, despite significantly reducing the toxicity of VNP by the deletion of *purI* and *msbB*, the strain still induced some side effects, especially splenomegaly and hepatomegaly, in mice when administered for tumor therapy by intravenous or intraperitoneal injection¹⁰. Therefore, it is critical to improve the safety of VNP as an antitumor therapy while increasing the efficacy^{7,10–12}.

In most cases, there is a strong innate immune response in the body, and invading microorganisms are quickly eliminated by antigen-presenting cells (APCs), such as macrophages and neutrophils¹³. However, *Salmonella* can produce a series of self-protective measures, including increasing bacterial antimicrobial peptide resistance gene expression¹⁴ and inhibiting intracellular lysosomal protein expression¹⁵, to survive after macrophage-mediated phagocytosis. Then, delayed release of strains is achieved because of the oncosis and rupture of macrophages due to the sustained intracellular stimulation of bacteria^{14,16,17}.

In fact, macrophages in the body are often used by *Salmonella* as natural havens to avoid being eliminated by other strong immune cells, such as neutrophils¹⁸. Immune cells, especially macrophages, are attractive natural drug carriers due to their high chemotaxis to tumors and loading capacity^{19–22}. Therefore, a hypothesis is that macrophage-mediated tumor-targeted delivery of VNP can combine the tumor-targeting characteristics of macrophages and intracellular strains that are gradually released within tumors, which also avoids the excessive antibacterial immune response induced by direct administration of VNP. Ultimately, tumor targeting and the safety of VNP as an antitumor therapy should be improved. Our previous studies showed that this strain could activate the tumor microenvironment and promote the chemotaxis of macrophages to tumors^{23,24}. This means that VNP released by macrophages within tumors could in turn promote the chemotaxis of macrophages loaded with the strain toward the tumor region.

Programmed cell death protein 1 (also known as CD279 and PD-1) and its ligand PD-1 ligand, PD-L1 (CD274), are critical

immune checkpoints that function normally to protect against autoimmunity. Their interaction is also an important strategy that many tumors use to escape immune surveillance²⁵. Despite the promising results of PD1/PDL1 blockade-based immunotherapy²⁶, a majority of patients still lack a durable response to the therapy because of the low activity of effector immune cells, specifically CD8 T cells, in the tumor^{27,28}. Therefore, increasing the response rates of cancer patients to PD1/PDL1 immune blockers is a critical challenge that needs to be overcome.

In this study, we report for the first time macrophage-mediated tumor-targeted delivery of VNP. This novel strategy combines the chemotaxis of macrophages to tumor regions, the protective and slow release of VNP by macrophages, and the colonization and tumor-killing properties of VNP. Finally, the antitumor activity of VNP was ensured, and acute organ injury caused by single VNP treatment was effectively avoided. Moreover, PDL1 levels on the surface of cancer cells were significantly upregulated after treatment with macrophages loaded with VNP, while the PD1 levels on the surface of CD8⁺ T cells in the tumor were significantly downregulated. This finding suggests that tumors in this state will be more sensitive to PD1/PDL1 blockade treatment^{26,29}. Therefore, we combined this bacteria-in-cell drug with PD1/PDL1 blockade therapy for the first time, and the therapeutic efficacy in mouse melanoma was significantly improved. We anticipate the macrophage-mediated tumor-targeted delivery of engineered attenuated *Salmonella* combined with enhanced safety and potent treatment efficacy to be a valuable reference for the research and delivery of biologically functional bacteria, and believe that stealth bacteria loaded by macrophages represent a versatile, novel and unique tool for biomedical applications. This new drug delivery strategy also offers new ideas for combining multiple cancer therapies, such as cell therapy, bacterial therapy and immunotherapy, to achieve complementary advantages.

2. Materials and methods

2.1. Cell lines and primary cells

B16F10 mouse melanoma cells, LLC mouse lung cancer cells, MC38 mouse colon cancer cells, A20 mouse B lymphoma cells, L929 mouse fibroblasts and RAW264.7 macrophages were preserved in our laboratory. Peritoneal macrophages were extracted based on the protocol provided by Choi et al³⁰. In brief, a 5% starch broth solution was prepared [1.8% nutritional broth (Solarbio, N8300, Beijing, China) and 5% soluble starch were dissolved in water], sterilized at 115 °C and stored at 4 °C. Eight-week-old female C57BL/6J mice were intraperitoneally administered 1 mL of starch broth solution, and peritoneal macrophages were harvested 2 or 3 days later. RAW264.7 cells were induced

using 100 ng/mL LPS (Beyotime, S1732, Shanghai, China) for 12 h to obtain M1-like RAW264.7 cells.

2.2. Bacterial strains, plasmid construction and transformation

VNP20009 (abbreviated VNP), VNP-RFP (transformed with a plasmid expressing RFP with a J23100 promoter), VNP-LuxCDABE (genomic insertion of LuxCDABE with a J23100 promoter), VNP-PD1nb (transformed with a plasmid expressing PD1nb with a J23100 promoter; a flag tag was added to the N-terminus of PD1nb to facilitate subsequent detection, pJ23100-flag-PD1nb), VNP-NC (transformed with an empty plasmid) and VNP-psifB-RFP (transformed with a plasmid expressing RFP with a sifB promoter; an HA tag was added to the C-terminus of RFP to facilitate subsequent detection, psifB-RFP-HA) were preserved in our laboratory. DH5 α and BL21 strains were purchased from Vazyme. All plasmids were constructed using the ClonExpress II/MultiS One Step Cloning Kit (Vazyme, C112/C113, Nanjing, China). The plasmid Pet28a-PD1nb was kindly provided by Dr. Shufeng Li (Southeast University). The construction and screening of anti-PD1 nanobodies are described in Section 2.15. The VNP strains were electrotransformed with the plasmids as described previously³¹.

2.3. Preparation of VNP-loaded macrophages [M Φ (VNP)]

M1-like RAW264.7 cells or peritoneal macrophages (PEM Φ cells) were harvested as previously described in Section 2.1, and then the macrophages and VNP were cocultured at a ratio of 1:10 in DMEM supplemented with 10% fetal bovine serum and no antibiotics at 37 °C for different times. The supernatant was discarded, and the cells were washed 3 or 4 times with PBS. The cells were incubated in DMEM supplemented with 10% fetal bovine serum and 50 μ g/mL gentamicin at 37 °C for 60 min to kill extracellular VNP but had no significant effect on the activity of strains inside the cells³². The supernatant was discarded, and the cells were washed 2 or 3 times with PBS. RAW264.7(VNP) cells were resuspended by gentle pipetting, while PEM Φ (VNP) cells were digested with 0.2% lidocaine at 4 °C for 5–8 min and resuspended. The cell suspension was centrifuged at 300 \times g and 4 °C for 5 min, and M Φ (VNP) cells (including RAW264.7(VNP) and PEM Φ (VNP) cells) were harvested from the precipitate. For immunofluorescence microscopy, RAW264.7 and PEM Φ cells were cocultured with VNP-RFP for different times. The extracellular bacteria were washed and sterilized with 50 μ g/mL gentamicin for 60 min. The cells were washed with PBS 3 times, fixed with 4% paraformaldehyde and permeabilized using 0.5% Triton X-100. After being washed with PBST 3 times, Actin-Tracker Green-488 (Beyotime, C2201S, Shanghai, China) was added and incubated for 1 h at 37 °C followed by DAPI (Beyotime, C1005, Shanghai, China) staining. Images were acquired using microscopy (Carl Zeiss, Axioplan 2, Oberkochen, Germany).

2.4. Intramacrophage VNP viability assays

Macrophages were cultured with VNP-RFP for different times [including RAW264.7(VNP) and PEM Φ (VNP) cells] and harvested as described in Section 2.3. RAW264.7(VNP) and PEM Φ (VNP) cells were adjusted to 1–10 \times 10⁴ cells/100 μ L, and the cells were transferred to a 96-well plate at 100 μ L per well. The RFP fluorescence intensity (550 nm, 585 nm) was determined by a multimode plate reader. The number of bacteria in the macrophages was

determined by using a formulated standard curve, which was generated by plotting the fluorescence intensity *versus* the concentration of a serially diluted standard VNP-RFP solution. RAW264.7(VNP) and PEM Φ (VNP) cells were incubated in 0.5% Triton X-100 (Sigma–Aldrich, 648462, St. Louis, MO, USA) lysis buffer for 10–15 min, and intracellular VNP was released from macrophages. The lysis solution was diluted and spread on LB agar plates supplemented with kanamycin. Colonies were counted after more than 12 h of incubation, and the number of live VNP in the total VNP taken up by macrophages was calculated.

2.5. Intramacrophage bacterial release assays

B16F10 cells (20,000) were added to the lower chamber of a 3.0- μ m Transwell plate (cells could not cross the pores but the bacteria could) and incubated for 6 h to adhere. Then, 40,000 RAW264.7(VNP-RFP) or PEM Φ (VNP-RFP) cells were added to the Transwell inserts, while 40,000 RAW264.7 or PEM Φ cells were added to the lower chamber [RAW264.7(VNP-RFP)-Up or PEM Φ (VNP-RFP)-Up group]. M Φ (VNP-RFP) and M Φ cells were added at opposite positions in the RAW264.7(VNP-RFP)-Low and PEM Φ (VNP-RFP)-Low groups. After 16 h of incubation, the supernatants were harvested and spread on LB agar plates with kanamycin, and the number of colonies was counted after an overnight culture at 37 °C. The adherent cells in the lower chambers were incubated in 500 μ L of DMEM supplemented with 50 μ g/mL gentamicin for an additional 60 min so that the extracellular bacteria could be killed. Cells were labeled with anti-CD11b antibodies and analyzed using flow cytometry (BD, Canto II, Franklin Lakes, NJ, USA).

To confirm the release behavior of macrophages, a VNP-psifB-RFP strain that could express RFP specifically inside cells was constructed. The psifB forward primer (5'-CTG CCC TAC CGC TAA ACA TCT-3') and psifB reverse primer (5'-CCA CAA GTG ATTATA TGATAC-3') were used to amplify the sifB promoter from the VNP genome. The strain obtained by amplification in LB medium was centrifuged at 12,000 rpm (Thermo, Fresco™17, Waltham, USA) for 10 min, and then the bacterial cells were adjusted to OD₆₀₀ = 1.0 with PBS. Subsequently, the bacteria and RAW264.7 cells were cocultured for 60 min at a MOI of 10:1, washed with PBS 3–4 times and cultured in DMEM supplemented with 50 μ g/mL gentamicin and 10% serum. After 60 min of incubation, total intracellular protein was collected for Western blot analysis. An HA-tagged antibody (CST, 3724S, Danvers, MA, USA) was used to examine the expression of RFP, and a secondary anti-rabbit IgG antibody (CST, 7074) was used. *Salmonella*-specific antibodies (Targetpharma, Nanjing, China) were used to track the strains in macrophages. RFP expression by intracellular activated bacteria was confirmed by fluorescence microscopy (Carl Zeiss).

2.6. Macrophage viability assays

The viability of VNP-loaded macrophages was examined using the trypan blue viability assay method. In brief, equal amounts of the cell suspension were mixed with 0.4% trypan blue solution (Invitrogen, T10282, Carlsbad, CA, USA), and 10 μ L was pipetted into a Countess chamber slide (Invitrogen Countess). The slide was inserted into the automated cell counter (Invitrogen Countess), and the number of viable and nonviable cells was determined.

2.7. Bacterial growth assays

The growth curves for different VNP in LB media were obtained with the Bioscreen C (OY Growth Curves Ab Ltd., Finland). In

brief, 10 μL of VNP suspension ($\text{OD}_{600} = 1.0$) was inoculated into 1 mL of LB medium, and 300 μL of solution per well was inoculated in Bioscreen C multiwell plates. The multiwell plates were incubated for 30 h at 37 °C. OD values were measured every 30 min under a brown filter with a wavelength of 600 nm.

2.8. Apoptosis assay

B16F10 cells (2.0×10^5) were plated into 12-well plates and incubated for 6–8 h until the cells adhered to the wall. Then, the collected unprocessed-VNP or released-VNP strains were cocultured with the cells at a MOI of 100 for 4 h. All cells in the plate were collected, washed and resuspended in binding buffer, stained with 1 μg of APC-conjugated annexin V protein (homemade in the laboratory) and left on ice in the dark for 30 min. All samples were added with 1 μL of PI and gently mixed before flow cytometry analysis (BD).

2.9. Macrophage chemotaxis assays

Transwell inserts (8 μm) were incubated in DMEM supplemented with 1% fetal bovine serum overnight. Approximately 3×10^4 B16F10, LLC or MC38 cells were seeded in the lower chambers, and chambers containing only DMEM served as controls. After 10 h of incubation for adherence, the medium was refreshed with 700 μL of DMEM supplemented with 10% fetal bovine serum and 50 $\mu\text{g}/\text{mL}$ gentamicin. RAW264.7, RAW264.7(VNP), $\text{PEM}\Phi$, and $\text{PEM}\Phi$ (VNP) cells were prepared, and the concentration was adjusted to 1×10^6 cells/mL using DMEM. One hundred microliters of the cell suspension was transferred to the Transwell insert and incubated at 37 °C for 16 h. Then, the Transwell inserts were removed from the plate, and the remaining cells that had not migrated from the top of the membrane were removed using a cotton-tipped applicator. After being washed 2–3 times with PBS, the Transwell inserts were placed in 4% paraformaldehyde for 20–30 min and then stained with crystal violet (KeyGen BioTech, KGA229, Nanjing, China) for 30 min at room temperature. After being washed with PBS 2–3 times, the Transwell inserts were photographed under an ortho-fluorescence microscope (Carl Zeiss).

2.10. In vitro analysis of macrophage killing of tumor cells

Approximately 60,000 B16F10, LLC or MC38 cells were plated in each well of a 0.4- μm Transwell plate (bacteria could not cross the pores, but the secreted cytokines could). After 10 h of incubation for adherence, the medium was replaced with 1200 μL of DMEM supplemented with 10% fetal bovine serum and 50 $\mu\text{g}/\text{mL}$ gentamicin. RAW264.7, RAW264.7(VNP), $\text{PEM}\Phi$, $\text{PEM}\Phi$ (VNP) and L929 cells were prepared, and the cell concentration was adjusted to 1×10^6 cells/mL. Two hundred microliters of the cell suspension was added to the Transwell inserts, and inserts containing only DMEM served as a control. The different types of cells were added to the upper chamber, and cells were not added to the lower chamber as a background control to remove interference from the upper chamber. After incubation for 12 h, the supernatant was collected. The proliferation of the tumor cells was measured using a CCK-8 assay kit (Beyotime, C0038).

2.11. qRT-PCR

qRT-PCR was performed as follows. Total RNA was extracted using TRIzol reagent (Vazyme, R401-01), and qRT-PCR was

conducted on a StepOne/StepOne Plus™ Real-Time PCR System (Applied Biosystems) using a SYBR Green PCR Master mix kit (Vazyme, Q221-01) according to the manufacturer's instructions. Primers used were as follows: CD11b forward primer (5'-ATG GAC GCT GAT GGC AAT ACC-3'); CD11b reverse primer (5'-TCC CCA TTC ACG TCT CCC A-3'); TNF- α forward primer (5'-GAC GTG GAA GTG GCA GAA GAG-3'); TNF- α reverse primer (5'-TGC CAC AAG CAG GAA TGA GA-3'); iNOS forward primer (5'-CAT TGC TGA CAG GAT GCA GAA GG-3'); iNOS reverse primer (5'-TGC TGG AAG GTG GAC AGT GAG G-3'); IL-6 forward primer (5'-CTC AAT ATT AGA GTC TCA ACC CCC A-3'); IL-6 reverse primer (5'-AAG GCG CTT GTG GAG AAG G-3'); CCR5 forward primer (5'-GTC TAC TTT CTC TTC TGG ACT CC-3'); CCR5 reverse primer (5'-CCA AGA GTC TCT GTT GCC TGC A-3'); CCR2 forward primer (5'-GCT GTG TTT GCC TCT CTA CCA G-3'); CCR2 reverse primer (5'-CAA GTA GAG GCA GGA TCA GGC T-3'); CSF1R forward primer (5'-TGG ATG CCT GTG AAT GGC TCT G-3'); CSF1R reverse primer (5'-GTG GGT GTC ATT CCA AAC CTG C-3'); 18 S rRNA forward primer (5'-GTA ACC CGT TGA ACC CCA TT-3'); 18 S rRNA reverse primer (5'-CCA TCC AAT CGG TAG TAG CG-3').

2.12. Macrophage phagocytosis assay

RAW264.7, RAW264.7(VNP), $\text{PEM}\Phi$, and $\text{PEM}\Phi$ (VNP) cells were prepared as described in Section 2.3, mixed with red fluorescence microspheres (Sigma–Aldrich, L2778) at a ratio of 1:10 (cell: microsphere) and incubated at 37 °C for 4 h. The supernatant was discarded, and free fluorescence microspheres were washed away with PBS. The cells were resuspended carefully, and phagocytosis efficiency was determined by flow cytometry.

2.13. In vivo biodistribution

RAW264.7(VNP-RFP) and $\text{PEM}\Phi$ (VNP-RFP) cells were prepared as described in Section 2.3, and 5×10^5 VNP-RFP, 2.5×10^5 RAW264.7(VNP-RFP) and 1×10^5 $\text{PEM}\Phi$ (VNP-RFP) cells in 100 μL of PBS were injected into B16F10 tumor-bearing mice *via* the tail vein. The mice were sacrificed at specific times, and tumors and other organs were dissected and lysed with a tissue pulveriser. The tissue lysate was diluted and spread on LB agar plates or analyzed using flow cytometry to determine the distribution of the strain. $\text{PEM}\Phi$ (VNP-LuxCABDE) was harvested as described previously, and the cells were incubated with the near infrared (NIR) fluorescent dye DiR (Abbkine) for 45 min. $\text{PEM}\Phi$, $\text{PEM}\Phi$ (VNP-LuxCDABE) cells (1×10^5) or VNP-LuxCDABE strains (5×10^5) in 100 μL of PBS were injected into A20 tumor-bearing mice *via* the tail vein. The fluorescence signals of LuxCDABE and DiR were detected using an *in vivo* imaging system (PerkinElmer, IVIS® Lumina III, Waltham, MA, USA).

2.14. Enzyme-linked immunosorbent assay (ELISA)

At 24 h post administration, mice were bled for sera and killed for tumor collection. The obtained tumor was added to tissue lysate (absin, abs9225, Shanghai, China) at 10 mg tumor tissue/50 μL tissue lysate and homogenized using a tissue homogenizer. The supernatant was collected by centrifugation. Both sera and tumor tissue lysates were collected for CCL2 cytokine detection by mouse CCL2 ELISA kits (absin, abs520016).

2.15. Construction and characterization of anti-PD1 nanobodies

The construction and characterization of nanobodies were performed as previously described³³. In brief, a healthy dromedary camel was immunized once a week with 0.5 mg of human PD1 protein (Sino Biological Company, Beijing, China). After 7 immunizations, 100 mL of blood from the immunized dromedary was collected. Blood lymphocytes were isolated, total RNA was extracted, and cDNA was synthesized for the following nanobody library construction. Briefly, a two-step nested PCR approach was used to amplify Nbs gene fragments. The final PCR products of 500 bp were ligated into T7Select10-3b vector arms. The ligation mixtures were packaged *in vitro*. The library size was evaluated by gradient dilution. In addition, the insertion rate was detected with PCR amplification. PD-1 protein was immobilized onto agarose beads, and the beads were incubated with phage library and then washed unbound phage particles with PBST. The bound phages on washed beads were amplified by infecting *E. coli* and cultured until cell lysis occurred. The bound phage particles were recovered and used for the next round of panning. After three rounds of biopanning, the PD-1-specific phages were enriched. Then, phage ELISA was performed, and positive plaques that gave high absorbance values toward PD-1 were selected and sequenced. By cloning in the pet-32a expression vector, VHH was equipped with an HA tag and His tag at the N-terminus and utilized for Nb purification and detection. For protein expression, the recombinant plasmid was transformed into *E. coli* BL21 (DE3) and induced with 0.1 mmol/L IPTG at 23 °C. Nanobodies were purified by nickel Sepharose affinity chromatography and dialyzed in PBS buffer. The purity of the collected proteins was checked by SDS-PAGE. The affinity constant of nanobodies for binding to PD1 was determined by ELISA. The antibody concentration resulting in 50% of the maximum absorbance value at three antigen-coating concentrations was measured and designated [Ab]₅₀. Three [Ab]₅₀ values were obtained and used for the calculation of three *K* values according to the Beatty formula. The final affinity constant is the average result of three *K* values.

2.16. PD1nb expression and secretion assay

VNP-PD1nb was incubated in 40 mL of LB medium with kanamycin until the OD₆₀₀ was between 0.6 and 0.8. Then, the solution was centrifuged at 5000 rpm (Thermo) and 4 °C for 10 min, and the supernatant and precipitate were collected. The precipitate was resuspended in 2 mL of PBS, heated in a dry bath incubator at 110 °C for 20 min to disrupt the bacteria and release the proteins, and then centrifuged at 13,000 rpm (Thermo) for 10 min, and total proteins in the bacteria were contained in the supernatant. Proteins were collected from the supernatant obtained in the first step according to the trichloroacetic acid (TCA) protein precipitation protocol. In brief, the supernatant was transferred to a 50 mL tube (Beckman Coulter, Brea, CA, USA) and centrifuged at 15,000×*g* and 4 °C for 10 min using an ultracentrifuge (Beckman Coulter). Then, the supernatant was transferred to a new 50 mL centrifuge tube, mixed with 10% TCA and incubated at 4 °C for 30 min before being centrifuged at 17,000×*g* and 4 °C for 20 min. The precipitate was resuspended in 300 μL of PBS, transferred to a 1.5-mL EP tube containing 1.2 mL of cold acetone, and centrifuged at 17,000×*g* and 4 °C for 20 min. The previous step was repeated, and the total protein secreted by VNP-PD1nb was finally collected in the precipitate and resuspended in 40 μL of PBS. The concentration of total protein in

both the precipitate and supernatant was determined by a BCA kit (Beyotime, P0012) and adjusted to the same concentration. The protein solution was mixed with 5 × SDS loading buffer (Yeasen, 20315ES20, Shanghai, China) and heated in a dry bath incubator at 95 °C. Twenty microliters of solution (approximately 15 μg of protein) was loaded for Western blot analysis. Flag-tagged antibodies (Sigma—Aldrich, F1804) were used to examine the expression and secretion of PD1nb, and secondary anti-mouse IgG (CST, 7076) was used.

2.17. Animal model

All procedures were conducted in compliance with all the relevant ethical regulations and were approved by the Nanjing University Institutional Animal Care and Use Committee. BALB/c (4–5 weeks, female) and C57BL/6J (6–8 weeks, female) mice were purchased from Changzhou Cavens Animals Corporation. B16F10 (2 × 10⁵ cells per mouse) or A20 (1 × 10⁶ cells per mouse) cells were injected subcutaneously into the right flanks of C57BL/6J mice or BALB/c mice. DTIC (dacarbazine), a recognized chemotherapeutic agent for the treatment of melanoma, was purchased from Shanghai YuanYe (S25808). The PD1nb used in this study was purified in our laboratory. When tumors grew to 80–160 mm³, DTIC (80 mg/kg) and PD1nb (5 mg/kg) were injected intraperitoneally four times every other day, while VNP (5.0 × 10⁵ cells per mouse) and cells (1.0 × 10⁵ cells per mouse), including PEMΦ, PEMΦ(VNP-NC) and PEMΦ(VNP-PD1nb), were injected through the caudal vein only one time. Tumors were measured by calipers 3–4 times weekly, and the volumes (*V*, mm³) were calculated using Eq. (1):

$$V = a^2b \times 0.52 \quad (1)$$

where *a* is the minor radius and *b* is the major radius.

For experimental lung metastasis, 1 × 10⁶ B16F10 cells in 100 μL of PBS were injected into mice *via* the tail vein. The treatment was started on Day 6, and the mice were sacrificed on Day 18. Lung metastases were determined by *ex vivo* photography, and ImageJ was applied to analyze the total surface tumor area. Blood was collected by retro-orbital puncture for routine blood examination. The serum obtained from blood was cryopreserved at –80 °C until assayed for blood biochemical indexes and ELISA. Routine blood examinations, blood biochemistry analysis, H&E staining of tumor, heart, liver, spleen, lung, and kidney sections and fluorescence immunostaining of macrophages on tumor sections were prepared by Wuhan Servicebio Corporation. To plot the survival curve, tumor-bearing mice were monitored daily and sacrificed when signs of adverse effects (pain, apathy, necrotic tumor) were observed or the humane endpoint (tumor weight equal to 10% of mouse body weight) was reached.

2.18. Flow cytometry

Tumor-bearing mice were sacrificed, and the tumors were dissected and subsequently incubated in digestion medium (10 U/mL collagenase I, 400 U/mL collagenase IV, 30 U/mL DNase I, all diluted in HBSS) for 30 min. Cell clumps were removed through a 40-μm cell strainer to obtain single-cell suspensions. Blood samples were obtained by removing the eyeball after the mice were completely anesthetized, and peripheral blood lymphocytes were obtained by a peripheral blood lymphocyte isolation kit (Solarbio, P8620). The cells were stained with fixable viability dye (BD, 564407),

incubated for 10–15 min at room temperature protected from light, and then stained using the following anti-mouse antibodies: CCR2-AF647 (clone SA203G11), CCR5-APC (clone HM-CCR5), CSF1R-APC (clone AFS98), CD45-PE-Cy7 (clone 30-F11), CD11b-APC (clone 561690), CD11b-PE (clone M1/70), F4/80-BV421 (clone T45-2342), CD86-PE (clone GL1), CD3e-FITC (clone 145-2C11), CD4-APC (clone RM4-5), CD25-BV421 (clone 3C7), CD8-APC (clone 53–6.7), PD1-APC (clone J43), PDL1-BV421 (clone MIH5), CD38-BV421 (clone 90/CD38), and CD127-PE (clone SB/199). Intracellular proteins, including CD206-APC (clone MR6F3), Ki67-BV421 (clone B56), IFN γ -PE (clone XMG1.2), and TNF α -PE (clone MP6-XT22), were stained after using a membrane-breaking fixative solution (BD, 565388). For analysis of peripheral blood immune cells in mice, blood was collected from the retro-orbital sinus of the mice. The blood samples were incubated with red blood cell lysis buffer (Solarbio, R1010) before staining with the viability dye and antibodies mentioned above. Blood was collected from the retro-orbital sinus of the mice, left at room temperature for 1 h and then centrifuged at 4 °C and 3000 rpm (Thermo) to obtain serum. A CBA kit (BD, 560485) was used for the analysis of cytokines in serum. Detection of bacterial load in tissues by flow cytometry: different organs were weighed, and the corresponding volume of 0.5% Triton X-100 (2 μ L/mg) was added according to the weight, processed by a tissue lyser and placed at 4 °C for 30 min. After cell clump removal by a cell sieve, the population of bacteria with RFP fluorescence in the lysate was detected by flow cytometry. All analyses were performed on a BD Canto II.

2.19. Statistical analysis

Data were analyzed with GraphPad Prism software version 5 (GraphPad Software). Two groups were compared using Student's *t* test. Comparisons of more than two groups were analyzed by one-way analysis of variance (ANOVA) followed by Dunnett's multiple comparisons test. Data are expressed as the mean \pm SEM for the *in vivo* mouse tumor model.

3. Results

3.1. Macrophages can function as biological carriers of VNP

In designing a valid biological carrier, we needed to determine the appropriate preparation conditions. We first constructed VNP-RFP strains that stably express red fluorescent protein (RFP) and cocultured these bacteria with the murine macrophage line RAW264.7 after the cells were induced with lipopolysaccharide (LPS) to form an M1-like phenotype (Supporting Information Fig. S1). The cells were then treated with 50 μ g/mL gentamicin for 60 min to kill extracellular bacteria (Supporting Information Fig. S2). We found that RAW264.7 cells could successfully load VNP-RFP (Fig. 1A). Fluorescence quantitative detection, which was based on the linear relationship between the fluorescence intensity and the number of bacteria (Supporting Information Fig. S3), showed that the total amount of VNP-RFP phagocytized by RAW264.7 cells positively correlated with the coculture time (Fig. 1B). Furthermore, spread plate counts were used to count the number of living VNP-RFP after macrophage phagocytosis, and the results showed that the number of live VNP-RFP loaded in RAW264.7 cells also positively correlated with the coculture time (Fig. 1C), during which the loading efficiency (live bacteria number/total bacteria number in RAW264.7) reached a peak at

60 min [220 ± 13 CFU (mean \pm SEM)/100 cells] (Fig. 1D). Although the viability of RAW264.7 cells decreased with prolonged coculture with VNP-RFP, cocultured macrophages showed acceptable cell viability at 60 min (above 90%) (Fig. 1E).

For the translatability of the approach, we also used primary peritoneal macrophages (PEM Φ cells), which have also been used as drug carriers to target tumors^{30,34}, to load VNP [PEM Φ (VNP) cells]. We first prepared high-purity peritoneal macrophages through starch broth stimulation followed by purification with adherent cultures, which ensured a high purity of the obtained macrophages (>98%) every time³⁰ (Supporting Information Fig. S4). VNP-RFP strains were cocultured with peritoneal macrophages (Fig. 1F), and unlike RAW264.7 cells, effective VNP loading by PEM Φ cells decreased with time (Fig. 1I). The number of live VNP strains loaded into PEM Φ cells was sufficient at 60 min [510 ± 10 CFU (mean \pm SEM)/100 cells] and reached a peak at 90 min [625 ± 12 CFU (mean \pm SEM)/100 cells] (Fig. 1G and H), when both the PEM Φ (VNP) cell viability (above 85%) was acceptable (Fig. 1J). For accuracy and uniformity, a coculture time of 60 min for the two kinds of macrophages and VNP with a ratio of 1:10 was chosen to obtain VNP-loaded macrophages (RAW264.7 (VNP) and PEM Φ (VNP) cells) for all subsequent experiments.

To investigate 1) whether VNP could survive for a long time, similar to wild-type *Salmonella*¹⁶, after being engulfed by macrophages, and 2) the changes in the release of intracellular strains from VNP-loaded macrophages after contact with tumor cells, we cocultured RAW264.7(VNP-RFP) or PEM Φ (VNP-RFP) cells in medium supplemented with gentamicin for 12 h and then removed gentamicin and cocultured the cells with B16F10 mouse melanoma cells *in vitro* with or without 3.0- μ m microporous membrane blocking for 16 h (Fig. 1K). When RAW264.7(VNP-RFP) or PEM Φ (VNP-RFP) cells were in indirect or direct contact with tumor cells, the VNP strains were alive and could be released from macrophages effectively (Fig. 1L and M). Moreover, when RAW264.7(VNP-RFP) and PEM Φ (VNP-RFP) cells were cocultured directly with tumor cells [RAW264.7(VNP-RFP)-Low group and PEM Φ (VNP-RFP)-Low group], more VNP strains were released into the supernatant, suggesting the acceleration of strain release by tumor cells (Fig. 1L and M left). The number of VNP strains in tumor cells, measured by flow cytometry, also showed a greater number in both the M Φ (VNP-RFP)-Low groups (Fig. 1L and M right). This contact-dependent enhancement of strain release probably results from macrophage membrane remodeling and the exocytosis of vesicles during the phagocytosis of tumor cells, which makes it easier for intracellular VNP to escape³⁵.

To further confirm the behavior of VNP release from macrophages, we constructed an attenuated *Salmonella* strain that conditionally expressed RFP inside macrophages (named VNP-psifB-RFP) based on the *sifB* promoter⁶. The *sifB* promoter, a classic *Salmonella* pathogenicity island II (SPI-II) promoter, can be activated by intracellular hypoxia, a low pH and a low phosphate environment³⁶. VNP-psifB-RFP expressed RFP only after being engulfed by macrophages (Fig. 1N). After VNP-psifB-RFP was cocultured with RAW264.7 cells for 60 min, modest protein expression by VNP-psifB-RFP was examined by Western blotting (Fig. 1O), and VNP-psifB-RFP expressing RFP was also observed inside cells (Fig. 1P). These results further indicated that the VNP-psifB-RFP strains could express RFP protein only after being activated in macrophages. Next, we prepared RAW264.7(VNP-psifB-RFP) and PEM Φ (VNP-psifB-RFP) cells and cultured them in medium supplemented with gentamicin. The appearance

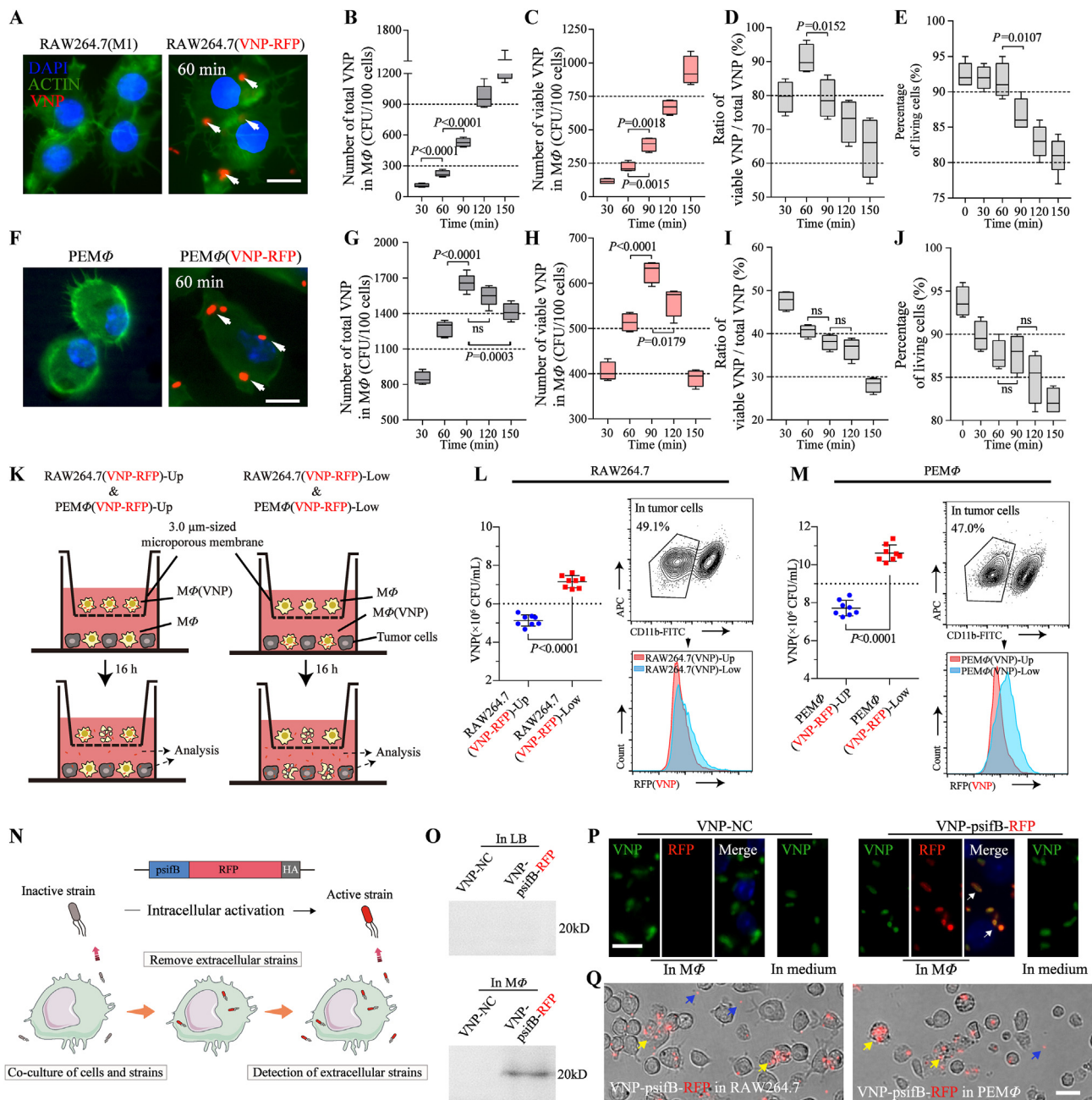


Figure 1 Preparation of Mφ(VNP) cells and the release of VNP from Mφ(VNP) cells. (A–E) RAW264.7 cells were induced with LPS (100 ng/mL, 12 h) and cocultured with VNP-RFP (1:10) for different times (30–150 min). Gentamicin (50 μg/mL) was added to kill extracellular VNP-RFP, and the quantity of the loaded bacteria in cells and cell viability were examined ($n = 5$). (A) Fluorescent images of the loaded cells. RAW264.7 cells were cocultured with VNP-RFP for 60 min and stained with DAPI and actin-FITC (Scale bar = 10 μm). (B) Changes in the total number of VNP-RFP strains phagocytosed by RAW264.7 cells. (C) Changes in the number of surviving VNP-RFP strains phagocytosed by RAW264.7 cells after different coculture times. (D) Changes in the loading efficiency (number of live bacteria/number of total bacteria) in macrophages. (E) The percentage of living RAW264.7 cells after different coculture times. (F–J) The results of repeating the experiments in (A–E) by replacing RAW264.7 cells with peritoneal macrophages (PEMφ) ($n = 5$). (K–M) Detection of VNP-RFP released from Mφ(VNP-RFP) cells. (K) RAW264.7(VNP-RFP) or PEMφ(VNP-RFP) (40,000 cells) cells were cultured in medium supplemented with 50 μg/mL gentamicin for 12 h, and then gentamicin was removed and cultured with B16F10 cells indirectly (RAW264.7(VNP-RFP)-Up & PEMφ(VNP-RFP)-Up) or directly (RAW264.7(VNP-RFP)-Low & PEMφ(VNP-RFP)-Low) by using a 3.0-μm Transwell chamber. The quantity of VNP in the supernatant or inside tumor cells was determined after 16 h. (L, M) The VNP titer in the supernatant (left) and comparison of the population of VNP inside tumor cells (right). (N) Schematic diagram of the use of VNP-ψsifB-RFP to examine bacterial release by macrophages. Due to the ψsifB promoter, VNP-ψsifB-RFP can only be activated and express RFP intracellularly, so extracellular red bacteria must have escaped from macrophages. (O) Western blot showing RFP expression of VNP-ψsifB-RFP in LB with $OD_{600} = 0.6–0.8$ or inside macrophages for 60 min. A strain carrying empty plasmid (VNP-NC) was used as a control. (P) Fluorescent images showing that VNP-ψsifB-RFP specifically expressed RFP

of VNP-*psifB*-RFP with red fluorescence outside the cells was observed 3 h later by fluorescence microscopy (Fig. 1Q). Since VNP-*psifB*-RFP activates and expresses RFP only in the intracellular environment, the extracellularly activated strain was undoubtedly released from the cells. Then, a continuous slow release of bacteria was observed for the remainder of the test period (3–15 h) (Supporting Information Fig. S5). The delayed release of the loaded VNP from cells may be due to the oncosis of macrophages and the self-escape of the strain^{14,16,17}. Based on our results, it is possible to speculate that when the external environment is suitable, such as the immunosuppressive and relatively nutrient-sufficient environment in the tumor⁷, these released strains will undergo rapid proliferation.

Since the tumor microenvironment is acidic (\sim pH 6.5–6.8)¹⁹, we prepared RAW264.7(VNP) and $\text{M}\Phi$ (VNP) cells and cultured them in medium with different pH values (7.4 and 6.7). The results showed that the acidic environment had no significant effect on the release of strains and their proliferation (Supporting Information Fig. S6). In addition, there was no significant difference between the released VNP and the unprocessed VNP in *in vitro* proliferation activity and ability to invade and kill tumor cells (Supporting Information Fig. S7). In conclusion, we confirmed that macrophages (RAW264.7 and $\text{M}\Phi$ cells) were capable of acting as natural biological carriers of VNP strains and achieved delayed slow release of intracellular bacteria without a significant effect on the proliferation activity of VNP and the invasion and killing ability of VNP on tumor cells.

3.2. Loading VNP does not weaken the phagocytosis, chemotaxis or tumoricidal activity of macrophages

It is worth confirming whether VNP loading affects some characteristics of macrophages, including phagocytosis, chemotaxis and tumor cytotoxicity. We first examined phagocytosis and quantified it with the content of fluorescent microspheres phagocytosed in cells. The results suggested that after being loaded with VNP in RAW264.7 cells, the cells were able to phagocytose more fluorescent microspheres (Fig. 2A and B). This trend was also observed in $\text{M}\Phi$ loaded with VNP, although the effect was less upregulated than that in RAW264.7 cells, which could be attributed to differences in cell origins (Fig. 2A and B). The enhanced phagocytic capacity of macrophages may be due to the upregulation of CD11b (a component of integrin CR3) expression after bacterial stimulation (Supporting Information Fig. S8), which is consistent with other reports³⁷. Considering that macrophages perform chemotactic functions through surface chemokine receptors, it is worth exploring whether phagocytosis affects the surface characterization of macrophages (*i.e.*, changes in the membrane protein). Fluorescent microspheres replaced VNP to coinubate with macrophages, and the results of flow cytometry showed that phagocytosis by macrophages did not significantly change the content of key chemokine receptor proteins on the cell surface (Supporting Information Fig. S9).

Transwell assays were utilized to evaluate the chemotactic properties of the cells. The results suggested that the chemotaxis of both RAW264.7(VNP) and $\text{M}\Phi$ (VNP) cells toward mouse melanoma tumor cells (B16F10) was not weakened, while RAW264.7(VNP) cells even showed an \sim 1.6-fold improvement compared with that of RAW264.7 cells (Fig. 2C and D). The same trend was observed for other tumor cells, including LLC (mouse lung cancer cells) and MC38 (mouse colon cancer cells) cells (Supporting Information Fig. S10A). A variety of white blood cells that are used as tumor drug delivery vehicles, including macrophages, neutrophils, and T cells^{38–40}, can cross a variety of barriers and reach tumor areas by sensing tumor-related chemokines or cytokines⁴¹. Here, we also found a significant increase in the transcript levels of key chemokine receptors, such as CCR5, CCR2 and CSF1-R⁴², in $\text{M}\Phi$ (VNP) cells (Supporting Information Fig. S11A), which also explained the enhanced tumor chemotaxis of $\text{M}\Phi$ (VNP) cells.

To evaluate the tumor cytotoxicity of $\text{M}\Phi$ (VNP) cells, a coculture experiment for tumor cells and $\text{M}\Phi$ (VNP) cells *in vitro* was performed. L929, a mouse fibroblast cell line, served as a negative control (Fig. 2E). The results revealed that both RAW264.7 and $\text{M}\Phi$ cells significantly inhibited tumor cell proliferation compared with L929 cells, while the suppression was more remarkable for macrophages after loading with VNP [RAW264.7(VNP) and $\text{M}\Phi$ (VNP-RFP) cells] (Fig. 2F). Similarly, VNP-loaded macrophages had enhanced cytotoxicity against LLC and MC38 cells (Fig. S10B). The inhibition of tumor cell proliferation in the L929 group was higher than that in the blank group, which may be due to nutritional competition between the two kinds of cells (Fig. 2F). The cytotoxic effects of macrophages without direct phagocytosis toward tumor cells could be ascribed to the significantly increased expression of proinflammatory mediators (including $\text{TNF}\alpha$, iNOS and IL6)⁸ in RAW264.7(VNP) and $\text{M}\Phi$ (VNP) cells (Fig. S11B) and the release of reactive oxygen species (ROS) and NO, which could damage tumor cells (Fig. 2G and Fig. S11C)^{43,44}. Notably, $\text{M}\Phi$ cells were more effective at phagocytosis, chemotaxis and tumor cell inhibition than RAW264.7 cells, which may be attributed to the strong activation of $\text{M}\Phi$ cells⁴⁵. This was further confirmed by comparing the levels of ROS, a well-accepted marker of activated macrophages⁴⁶ (Fig. 2G). Collectively, the phagocytosis, chemotaxis and tumor cytotoxicity of macrophages were not inhibited after VNP loading but were enhanced to some extent, which was beneficial for the antitumor activity of this bacteria-in-cell drug.

3.3. $\text{M}\Phi$ (VNP) cells increase tumor targeting and reduce the hepatosplenic toxicity of VNP

Typically, researchers have used intravenous or intraperitoneal injection to administer VNP, which is always accompanied by splenomegaly and hepatic inflammatory lesions. This is because some bacteria are off-target from tumors after administration⁸. Although intratumoral injection significantly reduces the hepatic and splenic accumulation of VNP¹⁰, this is undoubtedly not an

within 60 min after entering into macrophages, while the strain cultured in DMEM did not express RFP. The cells were stained with *Salmonella*-specific antibodies (green) and DAPI (blue). (Scale bar = 10 μm). (Q) Brightfield images showing activated VNP-*psifB*-RFP inside RAW264.7 and $\text{M}\Phi$ (yellow arrows) cells and the VNP released from macrophages (blue arrows) after 3 h of culture in medium containing 50 $\mu\text{g}/\text{mL}$ gentamicin (Scale bar = 20 μm). The data in (L) and (M) are reported as the mean \pm SEM. Boxplot representations of the spot counts. The median, interquartile range, and minimum and maximum identifiers are shown. All data are representative of three independent experiments. Statistics were calculated using the two-tailed, unpaired Student's *t* test with Welch's correction.

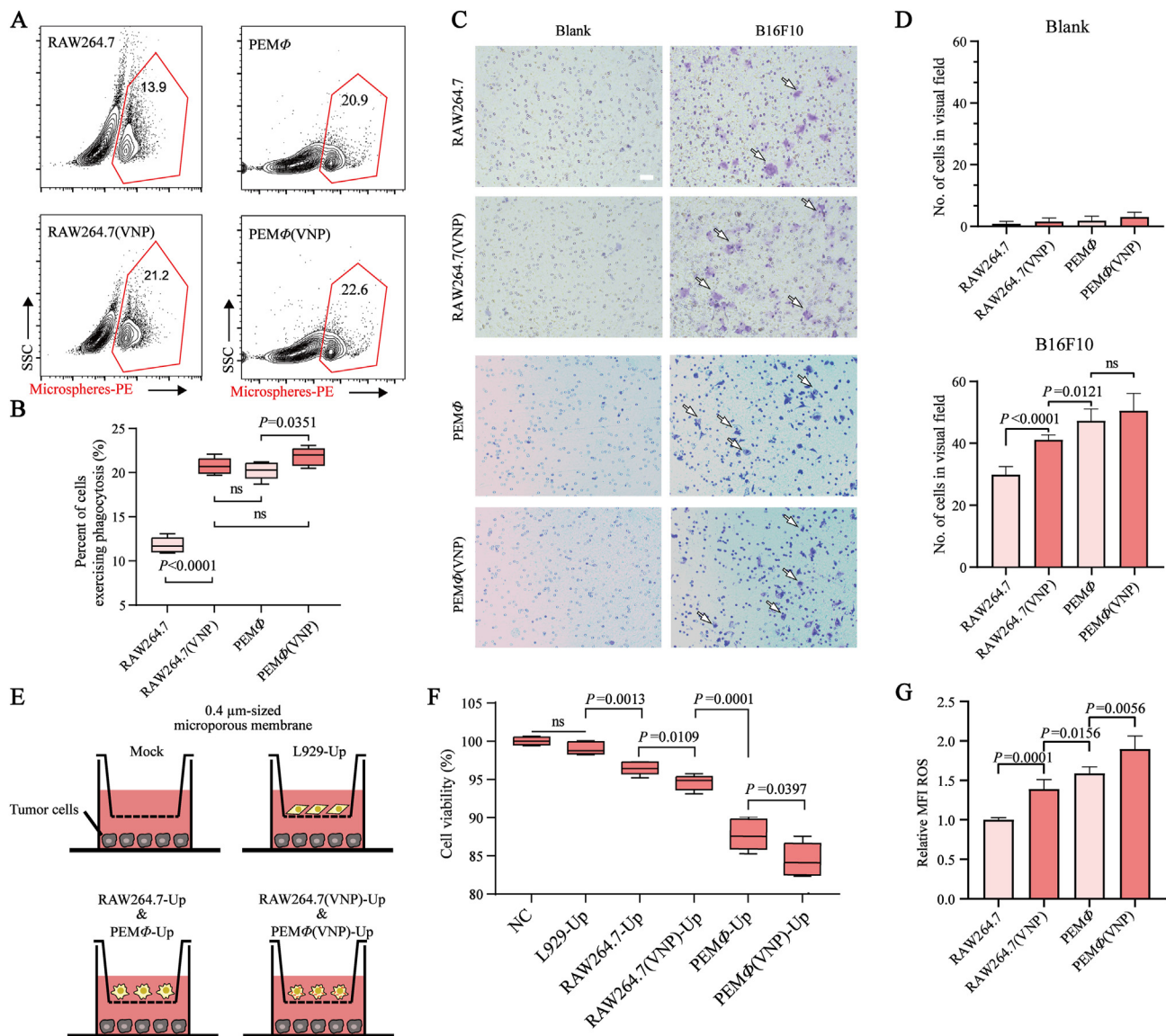


Figure 2 M Φ (VNP) cells achieve increased phagocytosis, chemotaxis and tumoricidal activity. (A, B) RAW264.7/RAW264.7 (VNP) and PEM Φ /PEM Φ (VNP) cells were cocultured with fluorescence microspheres at a ratio of 1:10 for 4 h. The percentage of macrophages that had engulfed microspheres was detected by flow cytometry, and the results are shown as boxplots in (B). (C, D) Transwell migration assay of RAW264.7/RAW264.7 (VNP) and PEM Φ /PEM Φ (VNP) cells toward B16F10 cells (coculture for 16 h with 50 μ g/mL gentamicin). Representative images of Transwell chambers (Scale bar = 100 μ m) in (C), and the number of cells crossing the pores in five random fields are visualized as boxplots in (D). (E, F) Detection of the indirect killing ability of the different groups on B16F10 tumor cells by indirect culture. (E) L929, RAW264.7, RAW264.7(VNP), PEM Φ and PEM Φ (VNP) cells were seeded in the 0.4 μ m Transwell upper chamber, and the lower chamber was seeded with B16F10 cells, which were cocultured for 12 h (supplemented with 50 μ g/mL gentamicin). (F) Detection of the proliferation of B16F10 tumor cells by CCK-8 assay in (E). (G) Detection of intracellular ROS in RAW264.7/RAW264.7 (VNP) and PEM Φ /PEM Φ (VNP) cells by flow cytometry. The relative mean fluorescence intensity (MFI) of ROS is shown ($n = 5$ in all data). Boxplot representations of the spot counts, with the median, interquartile range, and minimum and maximum identifiers, are shown. Data in (D) and (G) are reported as the mean \pm SEM. All data are representative of two independent experiments. Statistics were calculated using the two-tailed, unpaired Student's t test with Welch's correction.

optimal treatment for tumors. Herein, we loaded VNP into macrophages to camouflage the strain as a viable therapeutic strategy to avoid these side effects. According to the predetermined VNP-loading count by macrophages (see Section 3.1), RAW264.7(VNP) cells, hereinafter abbreviated as RAW(VNP) cells, and PEM Φ (VNP) cells were administered by tail vein at a dose of 2.5×10^5 and 1.0×10^5 cells per mouse, respectively, to

ensure the appropriate and same total initial viable VNP dose for B16F10 tumor-bearing mice treatment (5×10^5 CFU per mouse of VNP⁴⁷) (Fig. 3A).

To investigate whether M Φ (VNP) cells could lessen acute toxicity and the inflammatory response caused by VNP in mice, we first evaluated changes in body weight and the morphological characteristics of the liver and spleen at 24 h after different

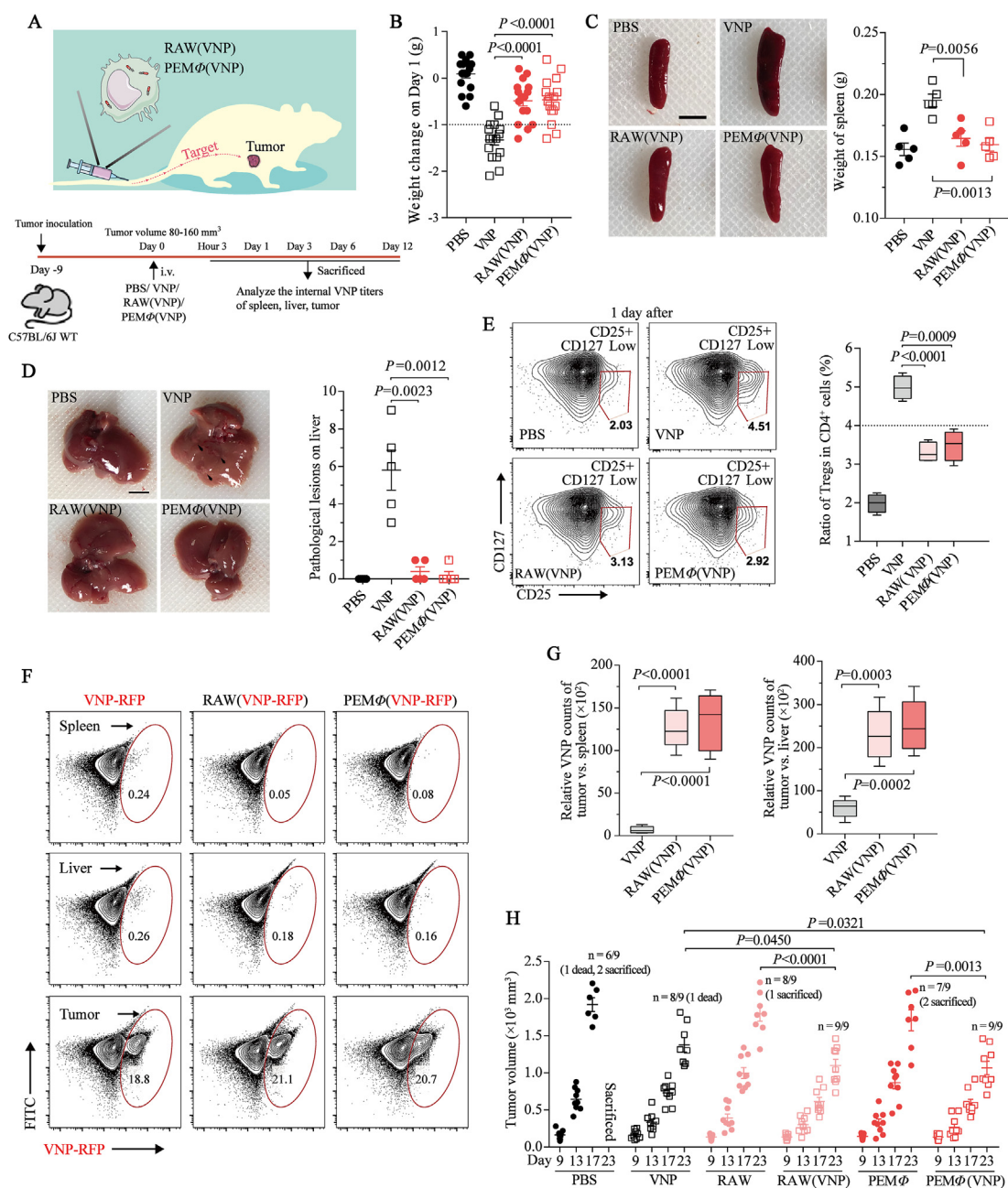


Figure 3 M ϕ (VNP) cells effectively alleviate the side effects of VNP and inhibit tumor growth. (A) B16F10 tumor-bearing C57BL/6J mice were administered VNP (5.0×10^5 CFU per mouse), RAW (VNP) (2.5×10^5 cells per mouse) and PEM ϕ (VNP) (1.0×10^5 cells per mouse) cells *via* the tail vein. The biodistribution of VNP in different treatment groups was determined at specific time points. (B) Changes in body weight 1 day after different treatments ($n = 16$ mice per group). (C, D) Comparison of splenomegaly (C) and liver pathological lesions (black arrow) (D) 1 day after different treatments ($n = 5$ mice per group, scale bar = 5 mm). (E) Percentage of Tregs among CD4⁺ T cells from the peripheral blood 1 day after different treatments ($n = 5$ mice per group). (F) Detection of the VNP population in the spleen, liver and tumor by flow cytometry on Day 6 ($n = 5$ mice per group). (G) The tumor:spleen (left) and tumor:liver (right) ratios of bacterial CFU per Gram was calculated on the basis of recovered CFU from extracted organs on Day 6 ($n = 5$ mice per group). (H) Comparison of B16F10 tumor suppression in response to different treatments ($n = 9$ mice per group). “Dead” means the mice died naturally, and “sacrificed” means the mice were humanely sacrificed when the tumors reached an ethical limit. Data in (B), (C), (D) and (H) are reported as the mean \pm SEM. Boxplot representations of the spot counts, with the median, interquartile range, and minimum and maximum identifiers, are shown. All data are representative of two independent experiments. Statistics were calculated using the two-tailed, unpaired Student’s *t* test with Welch’s correction.

treatments. The results confirmed significant alleviation of weight loss, splenomegaly and hepatic inflammatory lesions induced by VNP with this strategy (Fig. 3B–D). H&E staining and blood biochemistry analysis also revealed reductions in liver pathologic changes in the RAW (VNP) and PEM Φ (VNP) groups (Supporting Information Fig. S12). It has been reported that the content of Treg cells in the blood is related to the level of inflammation *in vivo*⁴⁸. Thus, the CD25⁺ CD127^{low} CD4⁺ Treg cell numbers in the blood after different treatments at 24 h were measured using flow cytometry. We found a significant decrease in Treg cell content after RAW(VNP) or PEM Φ (VNP) cell treatment compared with administration of VNP alone (Fig. 3E). All these results indicated that M Φ (VNP) cell treatment effectively avoided the side effects caused by a single administration of VNP.

The lower inflammatory response *in vivo* after M Φ (VNP) cell treatment is presumably because of the reduction in strain titer in normal organs. To validate this hypothesis, the colonization of VNP *in vivo* was determined by the plate count method. The results indicated that regardless of which kind of macrophage was used to load VNP, less VNP was observed in both the liver and spleen (Supporting Information Fig. S13B and S13C). The colonization of VNP in each tissue on Day 6 was selected as the representative. Flow cytometry showed VNP in the liver, spleen and tumor more intuitively (Fig. 3F). Within each group, we found no significant difference in the bacterial titers in the tumors (Fig. S13A). This is understandable because M Φ (VNP) cells need to gradually release the strains, causing a lower initial number of strains compared to administering VNP alone. Strains usually grow rapidly in tumors^{5,12}. Although M Φ (VNP) cells continuously release intracellular VNP, achieving a rapid increase in the intratumoral bacterial titer later, it may have been difficult to obtain a significant difference with administering VNP alone. This made it difficult to evaluate the efficiency of strains reaching the tumor by comparing their titers. Therefore, tumor targeting, which refers to the ratio of the bacterial titer in the tumor to that in normal organs⁴⁹, was used to assess the differences in the tumor-specific delivery effects of VNP between these groups. The tumor:spleen and tumor:liver ratios of bacterial CFU per Gram on Day 6 were increased by 7.8–48.8 and 2.6–12.2 times in the RAW(VNP) group, respectively, while those in the PEM Φ (VNP) group were increased by 6.9–51.7 and 2.1–12.9 times, respectively (Fig. 3G). These results highlight the increased tumor targeting of macrophage-mediated delivery of VNP. Moreover, the VNP titers in other organs, including the heart, lung and kidney, in the macrophage-loaded VNP group were also decreased compared to those in the single VNP group (Fig. S13D–S13F), although the H&E staining of these organ sections did not reveal noticeable histological deficiency differences between these groups, suggesting the negligible treatment-induced toxicity of the cell preparations to these organs (Fig. S13G).

We hypothesized that the lower inflammatory reaction and side effects and higher tumor targeting of VNP from M Φ (VNP) cell treatment as bacterial immunogens are camouflaged by macrophages also protect the VNP from being quickly cleared by neutrophils and then load the strain into the tumor. We found that M Φ (VNP) cells exhibited great natural migration toward tumor cells in Transwell assays (Fig. 2C and D, and Fig. S10A). Furthermore, direct injection of the mixture of macrophages and VNP into mice, instead of loading, did not reduce the toxicities of VNP and failed to stabilize the body weights of mice or reduce the accumulation of VNP in the liver and spleen (Supporting Information Fig. S14). Comprehensive hematology showed that there

were no significant changes in any of the indexes 1 day after M Φ (VNP) cell administration compared with PBS, although there was a tendency to lower neutrophil percentage than VNP (Supporting Information Fig. S15). These results all confirmed the reliability of this conjecture and the higher security of this delivery strategy.

The antitumor effects of different treatments were also evaluated, and we found that macrophages alone had comparatively effective antitumor abilities (Fig. 3H), which is easy to understand because we used activated M1-like macrophages that show a tumor-killing effect (Fig. 2). Compared with the PBS group, the VNP group showed significantly inhibited tumor progression, while the M Φ (VNP) group showed even better antitumor activity (Fig. 3H). These results indicated that this novel strategy not only avoided the side effects caused by the single administration of VNP but also effectively improved the therapeutic effect on tumors. Both RAW(VNP) and PEM Φ (VNP) cells showed lower toxicity and higher tumor repression than VNP, and we focused on PEM Φ (VNP) cells in subsequent studies due to their lower immunogenicity and higher translatability.

Macrophages are usually detected in the tumor area within 12 h after intravenous injection^{19,20}. To indicate the dynamic change in tumor accumulation, the biodistribution of PEM Φ (VNP-LuxCDABE) cells at various time points was analyzed. Given the limited number of labeled cells, dissected mice bearing A20 (B lymphoma cells) subcutaneous xenograft tumors were monitored for better visibility. *In vivo*, both DIR-labeled PEM Φ and PEM Φ (VNP-LuxCDABE) cells were observed in A20 tumors at 12 h (Fig. 4A and B), and the released strains quickly multiplied in the tumor after 12 h, as visualized by bioluminescence imaging (Fig. 4A and C). The unprocessed VNP-LuxCDABE was used as a positive control. There was no significant difference in the bacterial bioluminescence intensity between the PEM Φ (VNP-LuxCDABE) cell group and VNP-LuxCDABE group of tumors at 24 and 36 h, which was consistent with our previous results (Fig. S13A). Chemokine secretion was analyzed by ELISA. Compared to non-tumor-bearing mice, serum CCL2, the most important chemokine for the recruitment of macrophages into tumor⁵⁰, were significantly increased approximately 1.85-fold in A20 tumor-bearing mice (Fig. 4F) and 1.29-fold in B16F10 tumor-bearing mice (Supporting Information Fig. S16A). In addition, compared with PEM Φ cells, PEM Φ (VNP-LuxCDABE) cells achieved more efficient chemotaxis and enrichment in tumors (Fig. 4A, B, D and E). One hypothesis for this difference is that the released VNP strains after PEM Φ (VNP) administration promote the chemotaxis of macrophages by increasing cytokine/chemokine levels in tumors⁵¹. Compared with PEM Φ cell administration, the serum and tumor CCL2 levels of both A20 and B16F10 tumor-bearing mice were significantly increased after PEM Φ (VNP) cells administration at 36 h (Fig. 4G, Fig. S16B), further confirming our hypothesis. These results again suggest the effectiveness of macrophage-mediated tumor-targeted delivery of strains.

3.4. M Φ (VNP) cells reverse CD8⁺ T cell dysfunction but upregulate PDL1 expression on the tumor cell surface

Changes in the tumor microenvironment after PEM Φ (VNP) cells treatment are worth exploring. Herein, we used macrophages as the Trojan horse of VNP for antitumor delivery and treatment, so we first examined the population of macrophages in the tumor with a flow cytometry assay. Compared with macrophages administered

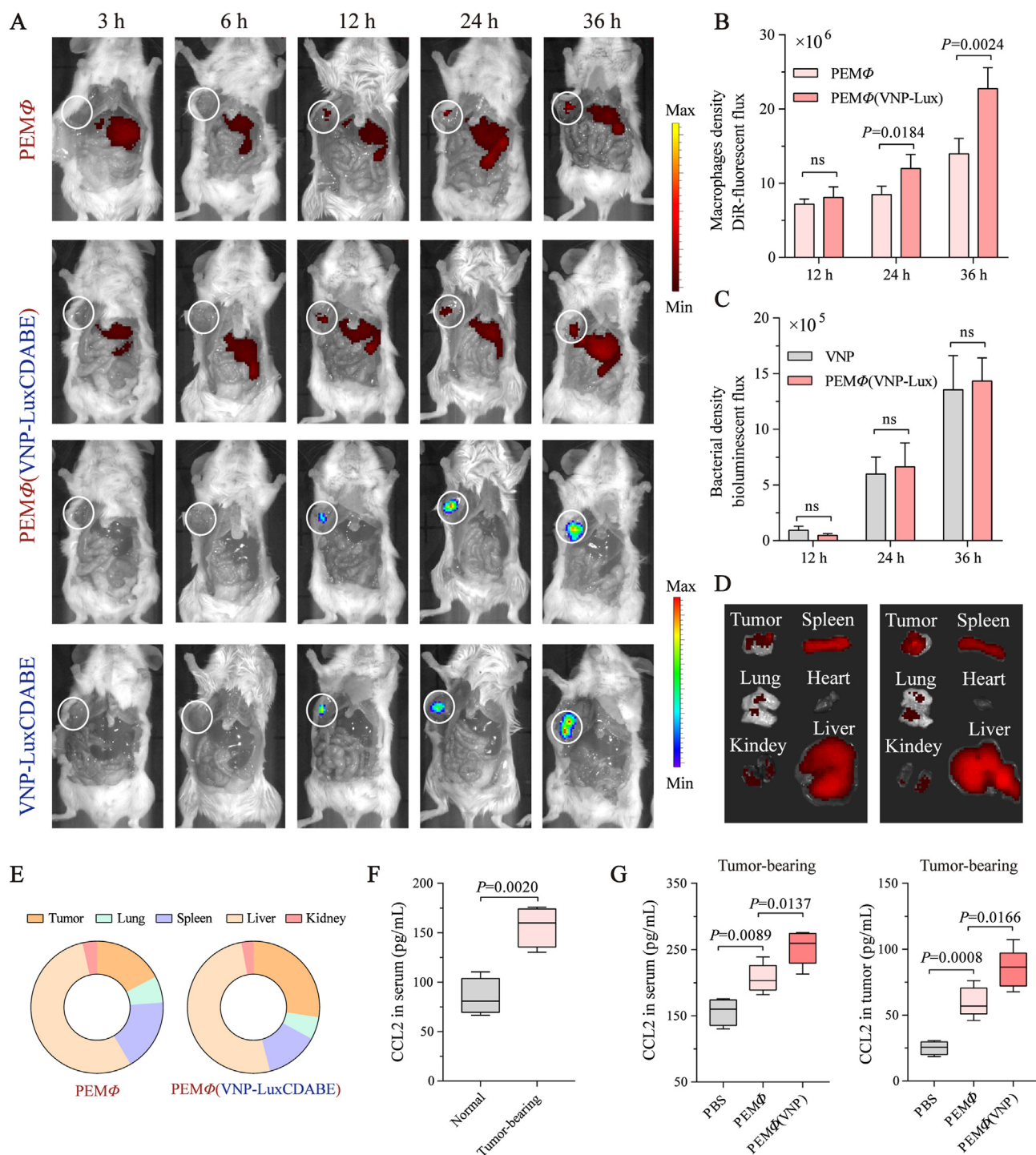


Figure 4 VNP released by macrophages rapidly proliferates in tumors and feedback to accelerate the recruitment of M Φ (VNP) cells into tumors. (A) Tumor trafficking and biodistribution of DiR-labeled PEM Φ cells (1.0×10^5 cells per mouse) or PEM Φ (VNP-LuxCDABE) cells (1.0×10^5 cells per mouse) and VNP-LuxCDABE strains (5.0×10^5 CFU per mouse) in A20 tumor xenograft models were analyzed by live-animal imaging at different time points after intravenous administration. Mice were sacrificed and dissected along the abdomen for better visualization. (B, C) Changes in macrophage density DiR-fluorescent flux (B) and bacterial density bioluminescent flux (C) in tumors 12, 24 and 36 h after administration ($n = 3$ or 4 mice per group). (D) A20 tumor-bearing mice were killed, and organs/tumors were explanted for *ex vivo* imaging 36 h after a single injection of 1×10^5 PEM Φ or PEM Φ (VNP) cells ($n = 4$ mice per group). (E) Proportion of DiR-fluorescent flux in different organs/tumors in (D). (F) Detection of CCL2 concentration in peripheral blood of normal mice and A20 tumor-bearing mice (tumor size = 100 mm^3 , $n = 4$ mice per group). (G) Detection of CCL2 concentration in peripheral blood and tumor of A20 tumor-bearing mice (tumor size = 100 mm^3) 36 h after administration ($n = 4$ or 5 mice per group). Data in (B) and (C) are reported as the mean \pm SEM. Boxplot representations of the spot counts, with the median, interquartile range, and minimum and maximum identifiers, are shown. The experiment was performed once. Statistics were calculated using the two-tailed, unpaired Student's *t* test with Welch's correction.

alone, both VNP and PEM ϕ (VNP) cells recruited more macrophages after colonization in the tumor region, while there were no differences between these two groups (Fig. 5A). It has been reported that VNP can remodel macrophages from M2-like to M1-like through LPS stimulation⁵². Here, we also found that the number of tumor-infiltrating M2 macrophages decreased, whereas the number of M1-like macrophages increased in VNP and PEM ϕ (VNP) cells on Day 3 (Fig. 5B and Supporting Information Fig. S17).

We next examined the population of CD8⁺ T cells within the tumor and found that VNP or PEM ϕ (VNP) cells had no effect on the population of CD8⁺ T cells within the tumor compared with PBS treatment (Fig. 5C and Supporting Information Fig. S18). Remarkably, VNP or VNP-loaded macrophages decreased the proportion of PD1⁺ CD38^{High} CD8⁺ T cells (Fig. 5D), which have been described as a population of dysfunctional cells that fail to respond to antigenic stimulation and do not exhibit effector functions^{53,54}. This suggested that VNP could effectively reverse CD8⁺ T cell dysfunction. The reversal may be attributed to some

inflammatory factors induced by VNP on dysfunctional CD8⁺ T cells^{10,51,54,55}.

Ideally, there should be improved CD8⁺ T cell-mediated tumor killing after PEM ϕ (VNP) cell treatment due to the reversal of dysfunctional CD8⁺ T cells. However, we found significantly increased expression of programmed cell death protein receptor 1 (PDL1) on the surface of tumor cells after the administration of VNP and PEM ϕ (VNP) cells compared with that of PBS. The upregulation of PDL1 on the surface of tumor cells was also significant when macrophages were administered alone (Fig. 5E). It is important to note that PDL1 binds to PD1 on the surface of CD8⁺ T cells and inhibits the proliferation and cytotoxicity of CD8⁺ T cells⁵⁶, ultimately impairing the antitumor efficacy of CD8⁺ T cells. This may help explain why VNP or VNP-loaded macrophages have a mediocre killing effect on tumors that was not much better than M1-like macrophage treatment alone (Fig. 3H). Similarly, the antitumor effect of PEM ϕ (VNP) cells was not much more markedly improved than that of single VNP treatment, which could also be attributed to the greater content of

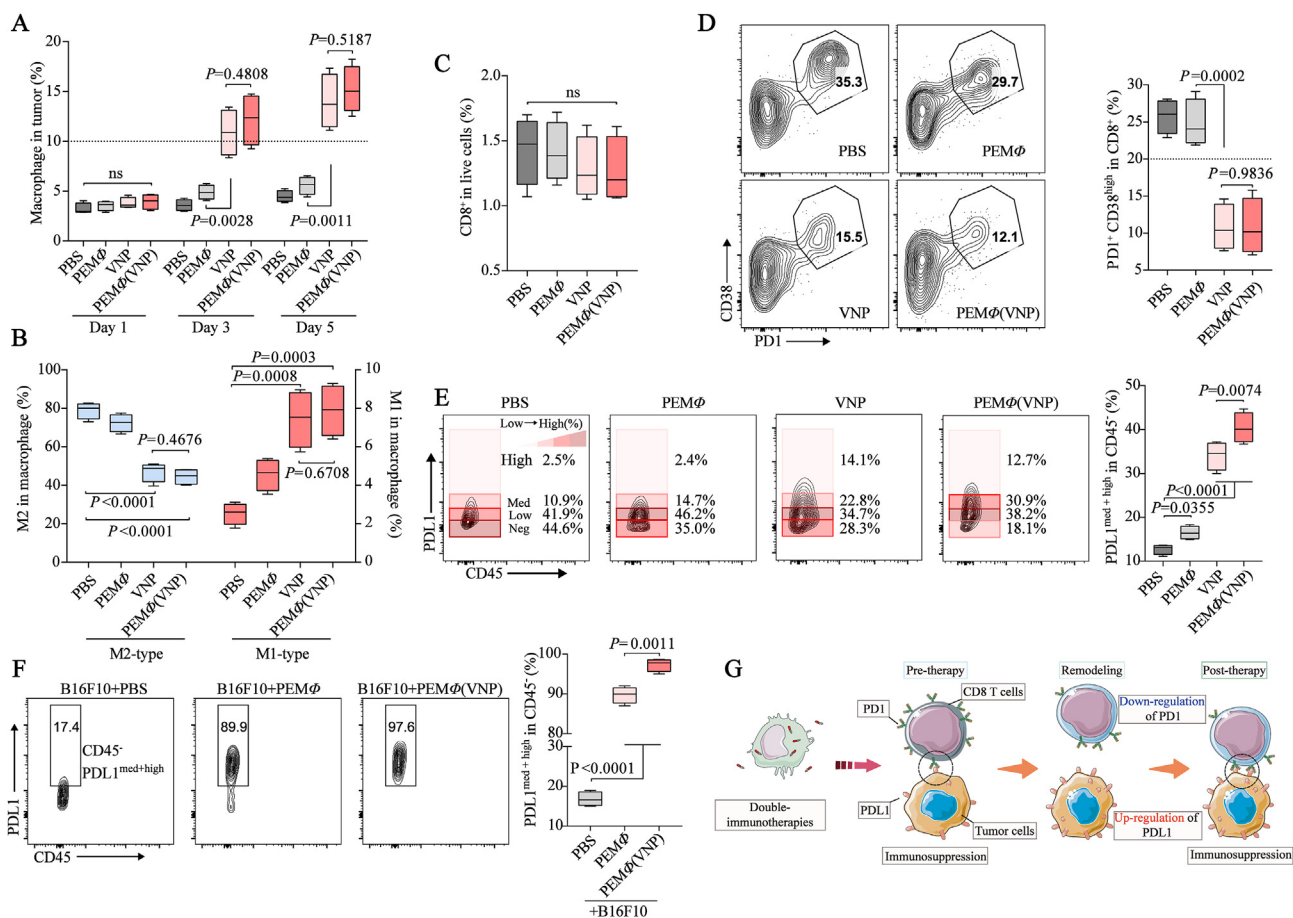


Figure 5 M ϕ (VNP) cells reverse CD8⁺ T cell dysfunction but upregulate PDL1 expression on the tumor cell surface. (A, B) Changes in the tumor-infiltrating macrophage population on Days 1, 3 and 5 (A) and phenotype (M1-like and M2-like) on Day 3 (B) after different treatments in the B16F10 tumor model ($n = 4$ mice per group). (C, D) The percentage of CD8⁺ T cells in tumors (C) and dysfunctional CD8⁺ T cells (PD1⁺ CD38^{high}) among CD8⁺ T cells after different treatments in the B16F10 tumor model (D) (Day 3, $n = 4$ mice per group). (E) Proportion of PDL1 on the surface of tumor cells after different treatments on Day 3 ($n = 4$ mice per group). (F) Changes in PDL1 on the surface of B16F10 cells 12 h after coculture with PEM ϕ and PEM ϕ (VNP) cells at a ratio of 1:1 ($n = 3$; 50 μ g/mL gentamicin was added to the medium). (G) Schematic diagram: PEM ϕ (VNP) cells activate CD8⁺ T cells but upregulate PDL1 expression on the tumor cell surface and further immunosuppress CD8⁺ T cells. Boxplot representations of the spot counts, with the median, interquartile range, and minimum and maximum identifiers, are shown. All data are representative of two independent experiments. Statistics were calculated using the two-tailed, unpaired Student's t test with Welch's correction.

PDL1 on the surface of tumor cells in the PEM ϕ (VNP) group (Fig. 5E). *In vitro* experiments further verified that PEM ϕ and PEM ϕ (VNP) cell treatments upregulated PDL1 on the surface of B16F10 cells, while the upregulation was more obvious in the PEM ϕ (VNP) group (Fig. 5F). This finding suggested that macrophages and VNP synergistically promote the upregulation of PDL1 on the tumor cell surface, which may be attributed to the

increased inflammatory cytokines secreted by intratumoral macrophages after activation⁵⁷ (Fig. S11B). Moreover, melanoma cells secrete PDL1-abundant exosomes to myeloid cells⁵⁸ and then achieve stronger immunosuppression against CD8⁺ T cells through these myeloid cells^{59,60}. Here, we also found an increase in PDL1 on the surface of tumor-related myeloid cells of the VNP and PEM ϕ (VNP) groups (Supporting Information Fig. S19),

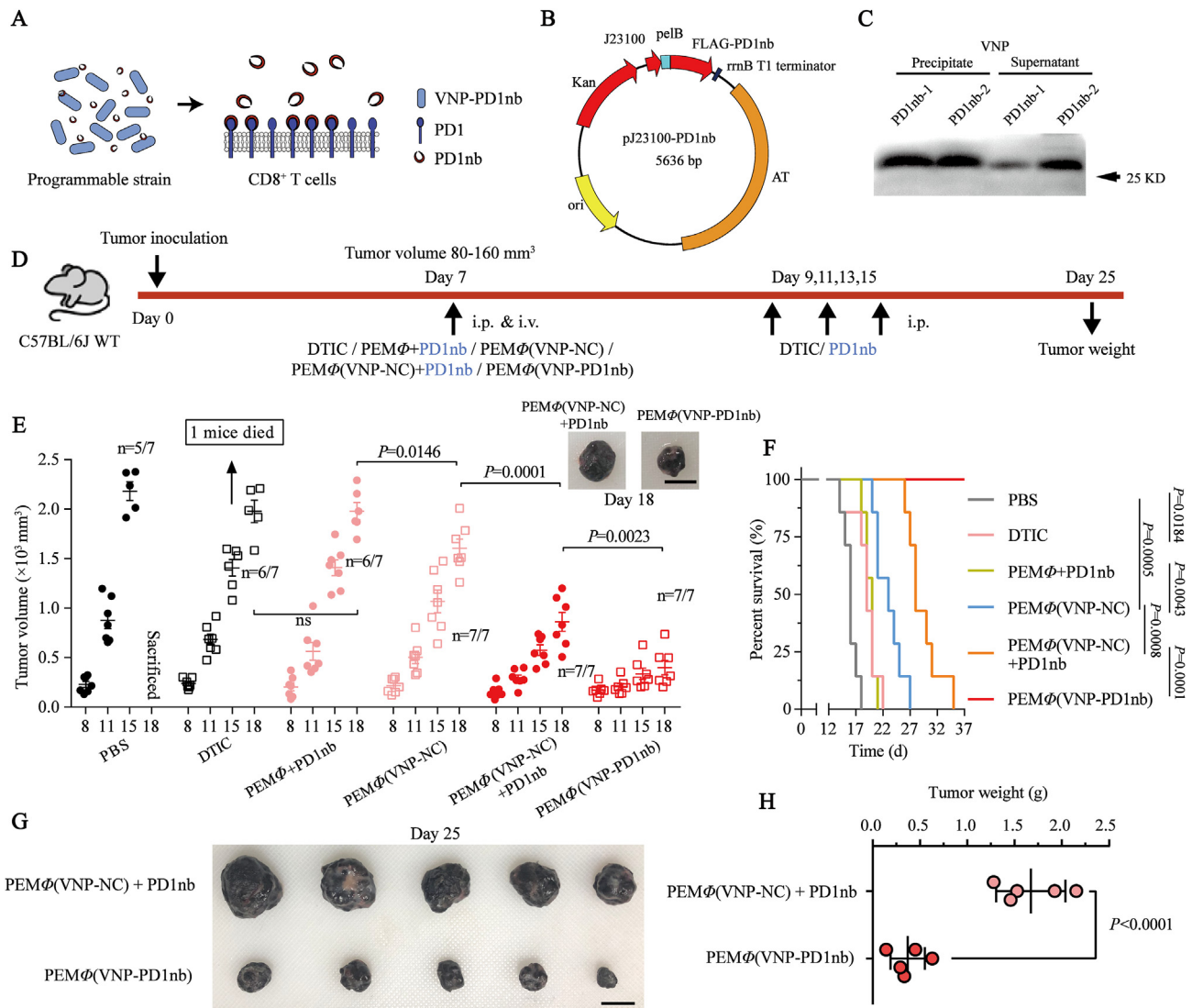


Figure 6 M ϕ (VNP-PD1nb) cell treatment achieved significant tumor inhibition in a B16F10 tumor-bearing mouse model. (A) Engineered *Salmonella typhimurium* VNP20009, with macrophage-mediated tumor-targeted delivery, constitutively expresses and secretes anti-PD1 nanoantibodies that bind to PD1 on the CD8⁺ T cell surface in the tumor. (B) Schematic diagram of the pJ23100-PD1nb plasmid. The flag tag was inserted into the N-terminus of the protein to facilitate subsequent detection, the pelB signal peptide realized the secretion of the expressed PD1nb, and the AT element prevented plasmid loss. (C) Immunoblot analysis to check the bacterial expression and secretion of PD1nb *in vitro*. Samples were separated into precipitates of total strains and supernatants of culture medium fractions. (D) Experimental scheme of the antitumor effect in the six different groups. DTIC (daphnane diterpenes, 80 mg/kg) and PD1nb (5 mg/kg) were injected intraperitoneally four times every other day, and cells (1.0×10^5 per mouse), including PEM ϕ , PEM ϕ (VNP-NC) and PEM ϕ (VNP-PD1nb) cells, were injected through the caudal vein only once. (E) Tumor growth profiles of B16F10 tumor-bearing mice after the different treatments ($n = 7$ mice per group, scale bar = 10 mm). (F) Survival curve of mice treated as described in (D). The survival of the mice was closely monitored several times per day, and the mice were killed when they reached a humane endpoint. (G, H) Tumors were photographed (G) and weighed (H) 25 days after B16F10 cell inoculation to compare the differences in tumor size between the PEM ϕ (VNP-NC) + PD1nb and PEM ϕ (VNP-PD1nb) groups ($n = 5$ mice per group, scale bar = 10 mm). “Dead” means the mice died naturally, and “sacrificed” means the mice were humanely sacrificed when the tumors reached an ethical limit. Data in (E) and (H) are reported as the mean \pm SEM. All data are representative of two independent experiments. Statistics were calculated using the two-tailed, unpaired Student’s *t* test with Welch’s correction in (E) and (H) and the Mantel–Cox test in (F).

which could further limit the effectiveness of bacteria in treating tumors.

Overall, PEM Φ (VNP) cells enhanced the fraction of M1-like macrophages and reversed dysfunctional CD8⁺ T cells in the tumor area, which exerted potentially stronger antitumor activity. However, the upregulation of PDL1 on the surface of tumor cells after PEM Φ (VNP) cell treatment inhibited the cytotoxicity of CD8⁺ T cells with PD1/PDL1 blockade. The schematic diagram in Fig. 5G shows the possible changes and interactions between CD8⁺ T cells and tumor cells during tumor therapy with VNP-loaded macrophages.

3.5. M Φ (VNP-PD1nb) cells potently and durably inhibit melanoma progression

Since M Φ (VNP) cell treatments upregulate PDL1 on the surface of tumor cells, which could bind to PD1 on the surface of CD8⁺ T cells and inhibit their cytotoxicity, it can be hypothesized that this novel dual immunotherapy [M Φ (VNP) cell treatment] will have a significantly improved antitumor effect after combination with PD1/PDL1 blockade, especially the blockade of PD1 considering its “Buckets effect” (Fig. 6A and Supporting Information Fig. S20). To test this conjecture, we first constructed and obtained a variety of anti-PD1 nanoantibodies, including VHH1 and VHH2 (Fig. S21A–S21F), and the affinity constants of nanobodies for binding to PD-1 were determined by ELISA. The results showed that VHH1 has better neutralization activity ($K_{\text{aff}} = 0.82 \times 10^7$ L/mol) than VHH2 ($K_{\text{aff}} = 1.54 \times 10^7$ L/mol) (Fig. S21G). Thus, VHH1 (subsequently called PD1nb) was used in subsequent experiments. Considering the two disadvantages of PD1 blockade therapy in clinical use, including off-target effects and the need for continuous administration, the expression of anti-PD1 antibody by programmed VNP may be a better choice. Thus, the VNP-PD1nb strain, which could stably express and secrete PD1nb, was constructed (Fig. 6B), and the VNP strain transferred with empty vector plasmids was used as a control. Western blotting verified the high expression and secretion of PD1nb by the strain *in vitro* (Fig. 6C).

It is important to note that macrophages loaded more VNP-PD1nb strains than VNP-NC strains during the same time (Supporting Information Fig. S22), and the number of intracellular VNP-PD1nb strains in PEM Φ at 30 min, which reached a level similar to that of VNP-NC at 60 min (Fig. S22C), was chosen as the coculture time in the subsequent experiments. This enhanced phagocytosis of macrophages to VNP-PD1nb strains is consistent with observations that anti-PD1 antibodies effectively improve macrophage phagocytosis⁶¹. To assess whether PEM Φ (VNP-PD1nb) cells effectively release intracellular strains and whether the released VNP-PD1nb strain can still express and secrete PD1nb, we collected the culture supernatant of PEM Φ (VNP-PD1nb) cells at different time points and counted the number of strains and PD1nb content in the supernatant (Supporting Information Fig. S23A). The results revealed that the VNP-PD1nb strains were continuously released from macrophages and multiplied over time (Fig. S23B). Western blotting confirmed that the VNP-PD1nb strains, which were released from PEM Φ (VNP-PD1nb) cells, still effectively expressed and secreted PD1nb (Fig. S23C). Moreover, increasing PD1nb content secreted by the strain was detected over time in the culture supernatant because of the higher amount of strains (Fig. S23C and S23D). It is not difficult to hypothesize that the PD1nb released by the engineered

VNP strains will further achieve antitumor immunotherapy through the well-known PD1/PDL1 immune blockade.

We confirmed that the macrophage-mediated dual-targeted delivery of VNP [M Φ (VNP)] to tumors significantly improves tumor targeting of the strains while alleviating the strains' side effects and ultimately enhancing the tumor treatment effect (Figs. 3 and 4). Encouragingly, macrophage-mediated tumor-targeted delivery of VNP-PD1nb [M Φ (VNP-PD1nb)] also demonstrated fewer side effects and stronger therapeutic effects than administration of VNP-PD1nb strains (Supporting Information Fig. S24). This effectively verifies the universal applicability of this delivery strategy. To further examine the effect of PEM Φ (VNP-PD1nb) cell treatment, prepared PEM Φ (VNP-NC) and PEM Φ (VNP-PD1nb) cells (1.0×10^5 per mouse) were administered *via* the tail vein to B16F10 tumor-bearing mice. Daphnane diterpenes (DTIC), a recognized chemotherapeutic agent for the treatment of melanoma⁶², were used as a positive control (Fig. 6D). The results showed that both PEM Φ (VNP-NC) and PEM Φ (VNP-PD1nb) cells achieved superior therapeutic outcomes, while PEM Φ (VNP-PD1nb) cells achieved better antitumor effects than PEM Φ (VNP-NC) cells (Fig. 6E). Melanoma often has a high metastatic capacity because of its high degree of malignancy⁶². Encouragingly, both PEM Φ (VNP-NC) and PEM Φ (VNP-PD1nb) cells suppressed the lung metastasis of melanoma, while mice with PEM Φ (VNP-PD1nb) cells showed fewer lung metastases (Supporting Information Fig. S25). Since engineered bacteria could express and secrete proteins as long as they were alive, we hypothesized that PEM Φ (VNP-PD1nb) cells would inevitably achieve a more significant and persistent antitumor effect than the simple mixed treatments [PEM Φ (VNP-NC) + PD1nb]. Thus, we also investigated the therapeutic effect of treatment with PEM Φ (VNP-NC) cells mixed with exogenous PD1nb by injection (5 mg/kg, which is consistent with the typical dose of PD-1 monoclonal antibodies⁶³). Compared with the mixed PEM Φ + PD1nb group, both the PEM Φ (VNP-NC) + PD1nb and PEM Φ (VNP-PD1nb) groups had significantly inhibited melanoma progression, which once again showed the high efficiency of VNP as a tumor immune activator (Fig. 6E). Impressively, PEM Φ (VNP-PD1nb) cell treatment exhibited a stronger and durable antitumor effect than the simple mixture of PEM Φ (VNP-NC) cells and PD1nb treatment, especially when continuous injection of PD1nb was stopped on Day 15 (Fig. 6F–H). After treatment with PEM Φ (VNP-PD1nb) cells, there was no mortality, and all mice were alive at the end of the study period, while all of the mice treated with PEM Φ (VNP-NC) + PD1nb died within 35 days (Fig. 6F). These results suggested that the continuous production and secretion of PD1nb by engineered VNP in tumors was a more convenient and long-lasting treatment strategy. Moreover, endogenously and continuously expressed protein with strains may enable higher intratumoral PD1nb doses, which may also be one of the reasons for its better therapeutic effect than systemic injection of PD1nb.

3.6. M Φ (VNP-PD1nb) cells inhibit melanoma progression by activating the tumor microenvironment and inhibiting tumor cell proliferation

Finally, to investigate the mechanism of the improved therapeutic effect of PEM Φ (VNP-PD1nb) cell treatment, we evaluated the changes in the tumor immune microenvironment following different treatments, including PBS, PEM Φ (VNP-NC) and PEM Φ (VNP-PD1nb) cells, in melanoma. We first detected tumor necrosis. A flow cytometry assay showed that both PEM Φ (VNP-

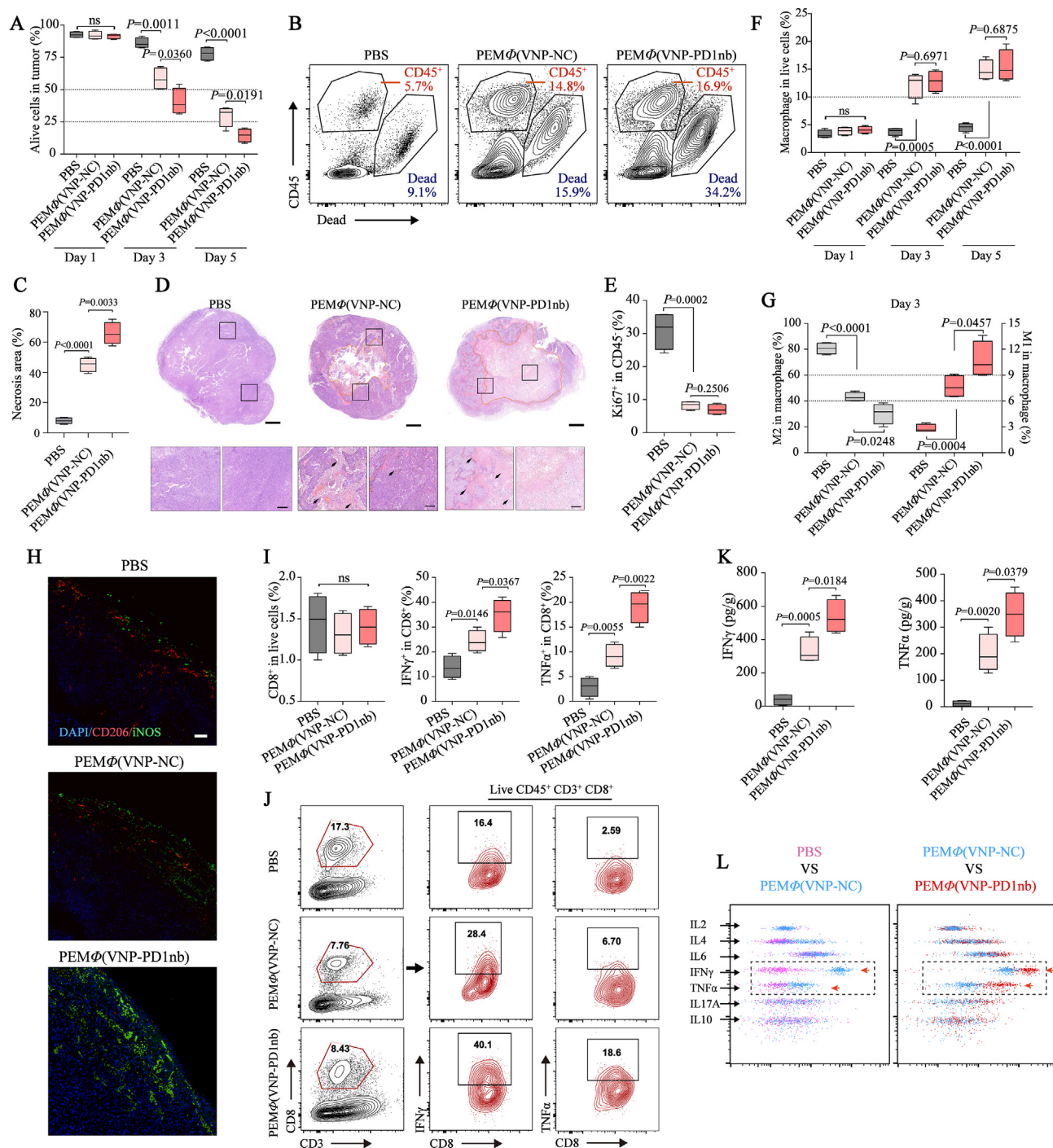


Figure 7 M ϕ (VNP-PD1nb) cells accelerate tumor cell necrosis and activate TAMs and CD8⁺ T cells. (A, B) Detection of alive cells in the tumor on Days 1, 3 and 5 by flow cytometry in (A) ($n = 4$ mice per group), and flow cytometry plots of representative data are shown in (B). (C, D) Comparison of the tumor necrosis proportion in different treatment groups by H&E staining on Day 3 in (C), and representative H&E images are shown in (D). Scale bars, 1 mm (complete tumor sections) and 200 μ m (enlarged tumor sections). (E) The proliferation of tumor cells (CD45⁺) was quantified by Ki67 staining and flow cytometry. (F, G) Changes in the tumor-infiltrating macrophage population on Days 1, 3 and 5 in (F) and phenotype (M1-like and M2-like) at Day 3 in (G) after treatment with PBS, PEM ϕ (VNP-NC) or PEM ϕ (VNP-PD1nb) cells in the B16F10 tumor model ($n = 4$ mice per group). (H) Representative immunofluorescence staining of B16F10 tumor cross sections showing cell nuclei (DAPI, blue), CD206 (marker of M2-like macrophages, red) and iNOS (marker of M1-like macrophages, green) of (G) (Scale bar = 50 μ m). (I, J) Profile of tumor-infiltrating CD8⁺ T cell and cytokine-producing CD8⁺ T cell subsets on Day 3 in the B16F10 tumor model ($n = 4$ mice per group) (I), and representative flow cytometry plots show the method of acquisition of frequency values for each T cell subset (J). (K, L) Detection of IFN γ and TNF α in peripheral blood 3 days after treatment in the B16F10 tumor model by CBA assay in (K), and flow cytometry plots of representative CBA data are shown in (L). Boxplot representations of the spot counts, with the median, interquartile range, and minimum and maximum identifiers, are shown. All data are representative of two independent experiments. Statistics were calculated using the two-tailed, unpaired Student's t test with Welch's correction.

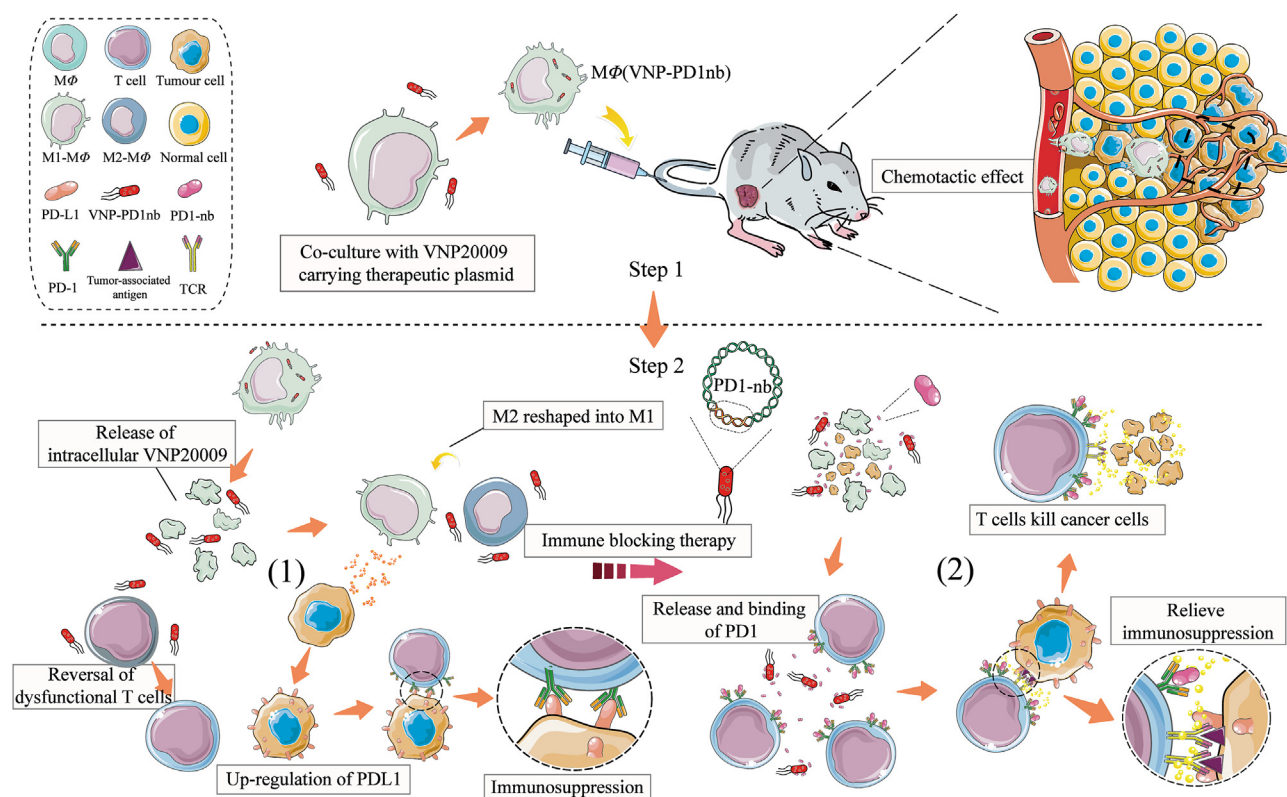


Figure 8 $M\Phi(VNP-PD1nb)$ cell treatment for tumor immunotherapy. Step 1: $M\Phi(VNP-PD1nb)$ cells act as Trojan horse engineered strains and target the tumor. Step 2: (1) After being released in the tumor, VNP-PD1nb, which is loaded in $M\Phi(VNP-PD1nb)$ cells, activates intratumoral immunity and reverses dysfunctional $CD8^+$ T cells while upregulating PDL1 in tumor cells, which results in immunosuppression. (2) PD1nb, which is released by VNP-PD1nb, can block the PD-1/PD-L1 interaction between $CD8^+$ T cells and tumor cells, thereby inhibiting immunosuppression and enhancing the tumor killing effect.

NC) and $PEM\Phi(VNP-PD1nb)$ cells caused significant intratumoral necrosis, while $PEM\Phi(VNP-PD1nb)$ cells caused more extensive necrosis than the other cells (Fig. 7A and B). H&E staining of the tumor area, as further evidence, more intuitively showed the difference in the extent of tumor necrosis after the three treatments, and the tumor necrosis areas after $PEM\Phi(VNP-PD1nb)$ cell treatment exceeded 60% within 3 days (Fig. 7C and D). Correspondingly, flow cytometry analyses revealed that both $PEM\Phi(VNP-PD1nb)$ and $PEM\Phi(VNP-NC)$ cell treatment down-regulated Ki67 (a recognized cell proliferation marker) expression in tumor cells on Day 3, indicating effective inhibition of tumor cell proliferation (Fig. 7E).

Next, we detected changes in the population and phenotype of intratumoral macrophages. The results showed significantly more infiltration of macrophages in the tumor after $PEM\Phi(VNP-NC)$ and $PEM\Phi(VNP-PD1nb)$ cell treatment than PBS treatment (Fig. 7F), which was consistent with the increased number of infiltrating lymphocytes in the tumor (Fig. 7B), yet there were no differences between the two cell treatments. We then assessed the same tumor samples using flow-based immunophenotyping for CD206 and CD86 expression to adjudicate a macrophage phenotype on Day 3. The results suggested that both cell-treated groups significantly reduced the proportion of intratumoral M2-like macrophages and increased the proportion of M1-like macrophages, while $PEM\Phi(VNP-PD1nb)$ cell treatment showed higher efficiencies (Fig. 7G), suggesting a higher increase in the tumor-killing and antigen-presentation ability of tumor macrophages. Immunofluorescence also confirmed a significant increase

in intratumoral M1-like macrophages in the $PEM\Phi(VNP-PD1nb)$ cell treatment compared with the $PEM\Phi(VNP-NC)$ cell treatment (Fig. 7H).

Cytotoxic $CD8^+$ T cells always play a role in the promotion of oncolytic therapy, especially in the treatment of PD1/PDL1 blockade^{27,28,64}. Thus, the cytotoxic $CD8^+$ T cells in tumors were detected on Day 3 after different treatments as a representative time. Flow cytometry analyses revealed that there was no significant change in the total population of $CD8^+$ T cells in the tumor among the three treatments, but markedly elevated intratumoral $IFN\gamma^+$ $CD8^+$ T cells and $TNF\alpha^+$ $CD8^+$ T cells were observed after cell treatments (Fig. 7I and J, and Supporting Information Fig. S26). More activated $CD8^+$ T cells were observed in tumors treated with $PEM\Phi(VNP-PD1nb)$ cells, which could also partially account for the stronger antitumor effect of $PEM\Phi(VNP-PD1nb)$ cells compared to $PEM\Phi(VNP-NC)$ cells.

Proinflammatory cytokines, such as $IFN-\gamma$ and $TNF-\alpha$, secreted by immune cells can directly kill tumor cells and induce strong tumor-specific immune responses⁶⁵. Thus, the sera from B16F10 tumor-bearing mice after different treatments on Day 3 were collected to measure the blood cytokine levels using a CBA kit. The results showed that the amounts of $IFN-\gamma$ and $TNF-\alpha$ induced by $PEM\Phi(VNP-NC)$ and $PEM\Phi(VNP-PD1nb)$ cell treatment were significantly higher than those induced by PBS treatment. More $IFN-\gamma$ and $TNF-\alpha$ were tested in $PEM\Phi(VNP-PD1nb)$ cell treatment (Fig. 7K and L), suggesting that $PEM\Phi(VNP-PD1nb)$ cell treatment enhanced immune activation and therapeutic efficacy. Some other proinflammatory cytokines,

including IL2, IL6 and IL17A, and anti-inflammatory cytokines, including IL4 and IL10, did not differ among the three groups (Fig. 7L). Collectively, treatment with PEM Φ (VNP-PD1nb) cells efficiently inhibited tumor cell proliferation, culminated in tumor necroptosis and activated the tumor microenvironment, including remodeling in TAMs and activation of CD8⁺ T cells, and finally prevented tumor progression.

In summary, we established a combination therapy with engineered stealth bacteria camouflaged by macrophages (Fig. 8). The three treatments, including macrophages, VNP strains and PD1nb, complement each other to form a unified system. With natural tumor-targeted macrophages as Trojan horses deliver, bacteria exhibit (1) a low inflammatory response and side effects; (2) low accumulation in normal organs; and (3) unchanged anti-tumor bioactivities. With continuously expressed and secreted PD1nb, engineered bacterial treatment exhibits (1) high immune activation, (2) significant tumor regression, and (3) a stable and durable anti-tumor effect. Given these unique advantages, we anticipate promising biomedical applications of this strategy in tumoral treatment.

4. Discussion

Specific tumor targeting and highly potent tumor suppression are two challenging targets using VNP and other oncolytic bacteria for tumor therapy. To attain improved tumor targeting, we first reported the use of macrophage-mediated tumor-targeted delivery of VNP in treating tumor-bearing mice. Although modification of attenuated *Salmonella*⁴⁹ or the tumor microenvironment²³ could also improve the tumor targeting of the bacteria in mice, all these improvements require the arrival of the bacteria at the tumor. A high titer of VNP was detected in the tumors of patients after intratumoral injection⁶⁶, and unattenuated *Salmonella* could also effectively infect the tumor area⁶⁷, which suggested that both *Salmonella* and attenuated VNP could colonize tumors and exert anti-tumor effects^{68,69}. In a phase I clinical trial of intravenous treatment for sarcoma, however, VNP did not perform well in terms of effectiveness⁹. The investigators did not detect VNP in the tumor, and there was no evidence of colonization. Thus, the accessibility of VNP in human tumors is urgently needed to avoid excessive stimulation of the immune system by bacterial immune antigens; otherwise, the bacteria will be quickly eliminated. The use of biofilms to wrap bacteria is an effective way to avoid early exposure of the bacteria in the body¹². Another effective strategy, as reported here, is to conceal VNP in macrophages to avoid premature exposure of the bacteria. This is especially valuable since the specific antigen modification of macrophages⁵⁰ can further improve the tumor targeting ability of macrophages, which is worth further exploration.

One of the major challenges of cell-mediated drug delivery is the need to inject large number of cells (10⁷–10⁸ cells in mice)³⁹, which is accompanied by sophisticated operations and high costs. The novel therapeutic strategy described here only requires a dose of 10⁵ M Φ (VNP) cells to effectively treat murine tumors, which is only 1/100–1/1000 of that required by traditional cell therapy. In fact, even if only a small amount of VNP is released into the tumor region by macrophages, a therapeutic effect can be achieved by the continuous proliferation of strains. However, the acquisition and culture of macrophages in the early stage still increases the cost of this therapy. Alternatively, low-cost allogeneic macrophages can be used to build off-the-shelf vectors that perform targeted release functions, which can then be recognized and

eliminated by the host immune system. The approach proposed here of using macrophages loaded with VNP for targeted tumor therapy is a modular approach in which macrophages can also be loaded with different recombinant oncolytic bacteria. Naturally, different types of strains may have different cell-loading efficiencies; for example, during the same processing time, macrophages loaded more VNP-PD1nb strains than VNP (Fig. S22). This study demonstrates the effectiveness and universality of macrophages as VNP delivery vectors, but other kinds of cells may also be used for this type of therapy^{39,40,70}.

For tumor suppression, we constructed an engineered VNP strain (VNP-PD1nb) that stably expressed and secreted anti-PD1 nanoantibodies and loaded the strain into macrophages for tumor treatment. This novel strategy achieved a more efficient and lasting effect than the simple combination of PD1nb and VNP-loaded macrophages for anti-tumor therapy. This strategy also provides a potential solution to the dilemma of a low response rate to PD1/PDL1 blockade in cancer patients²⁷. After immunotherapy with PD1/PDL1 blockade, a small percentage of responders suffer from a series of adverse reactions due to the accumulation of drugs in normal organs or tissues⁷¹. The bispecific delivery of macrophages and VNP theoretically blocks the side effects of anti-PD1 treatment by concentrating the production of PD1nb within the tumor. The expression and secretion of PD1nb by trace levels of VNP-PD1nb in normal organs, however, may still lead to misregulated immune activation. A feasible approach would be to induce the expression of PD1nb in a controlled manner, such as hypoxia induction³¹ or direct regulation of bacterial survival⁷. The most exciting application of immunotherapy is the ability to treat tumor metastasis⁷ and inhibit recurrence, especially when used in combination with other therapies, including chemotherapy, photothermal therapy, photodynamic therapy, etc.⁷². The combined use of macrophages, attenuated *Salmonella* and anti-PD1 nanoantibodies in this paper also achieved significant anti-metastatic effects (Fig. S25). Moreover, due to the strong loading capacity of macrophages³⁰ and the editable and carrier properties of VNP³¹, this novel immunotherapy has great room for improvement to achieve stronger effects in anti-tumor metastasis, suppress or even eliminate tumors, and inhibit recurrence. These possibilities warrant further investigation.

In conclusion, we programmed triple immunotherapies in one drug: macrophage-based immunotherapy, microbe-based immunotherapy and anti-PD1 immunotherapy using a Trojan horse strategy and delayed release. This study provides a promising future for exploring convergent immunotherapy approaches to address monotherapy challenges.

5. Conclusions

In this study, we designed, characterized and applied a novel cancer immunotherapy system. Briefly, this system allowed attenuated *S. typhimurium* VNP20009 latent in macrophages to precisely unload inside the tumor and then secrete anti-PD1 nanoantibodies. We first showed that macrophage-mediated tumor-targeted delivery of VNP20009 alleviated side effects induced by the strain and effectively suppressed melanoma in mice. Furthermore, there was a marked decrease in dysfunctional CD8⁺ T cells in tumors after this systemic treatment, which also suggests a synergistic response to PD1/PDL1 blockade. Thus, macrophages loaded with engineered VNP20009, which could express and secrete anti-PD1 nanoantibodies, were prepared and used to treat xenograft mouse melanoma models. The combination of the three immunotherapies

(macrophages, VNP20009, PD1nb) achieved much more significant and durable tumor inhibition than simple mixed treatments by activating the tumor microenvironment, including increased immune cell infiltration, TAM remodeling to an M1-like phenotype and prominent activation of CD8⁺ T cells.

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Author contributions

Zichun Hua supervised and conceived the project. Leyang Wu designed the experiments, Leyang Wu, Lin Li, Shufeng Li, Lina Liu, Wenjie Xin and Xingpeng Yin performed the experiments and analyzed the data. Leyang Wu, Lin Li and Chenyang Li wrote and Zichun Hua modified the manuscript. Xuebo Xu and Feifei Bao contributed to the discussion and provided relevant advice. All authors discussed the results and reviewed the manuscript

Conflicts of interest

The authors have applied for patents related to this study.

Appendix A. Supporting information

Supporting data to this article can be found online at <https://doi.org/10.1016/j.apsb.2022.05.006>.

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