Supplementary Information

Cell microparticles loaded with tumor antigen and resiquimod reprogram tumor-associated macrophages and promote stem-like CD8⁺T cells to boost anti-PD-1 therapy

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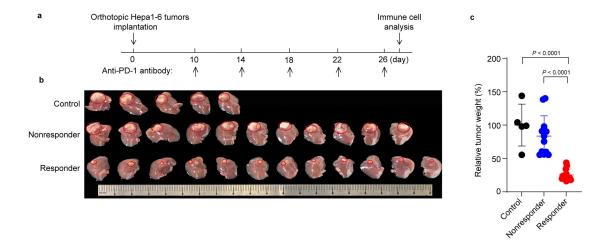
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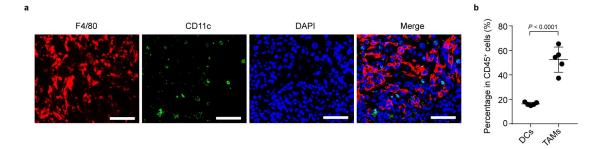
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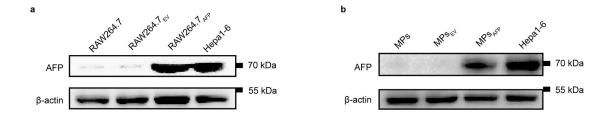
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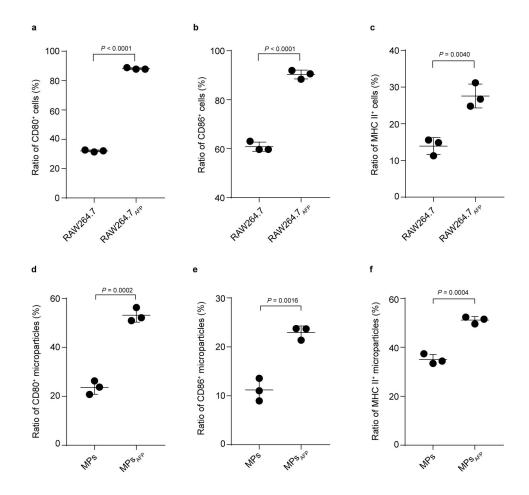
Supplementary Fig. 1 Anticancer activity of anti-PD-1 antibody in orthotopic Hepa1-6 tumor-bearing mice. (a) Schematic schedule for tumor inhibition and immune cell analysis experiment in orthotopic Hepa1-6 tumor-bearing mice after intraperitoneal injection of anti-PD-1 antibody at the dosage of 5 mg kg⁻¹. (b,c) Tumor image (b) and tumor weight (c) of PBS-, anti-PD-1 antibody-responsive and anti-PD-1 antibody-nonresponsive mice after treatment indicated in (a). Data are presented as means \pm s.d. (n = 5 mice for PBS-treated group, n = 11 mice for nonresponsive group, n = 12 mice for responsive group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



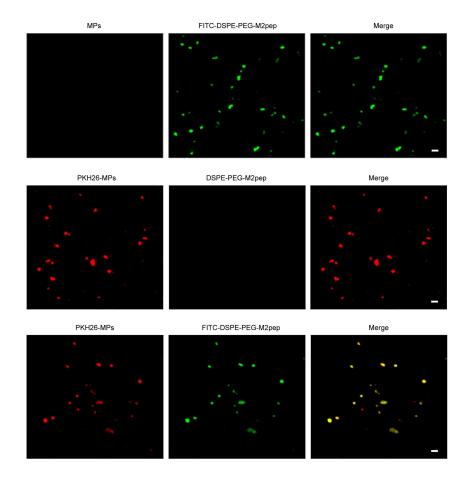
Supplementary Fig. 2 Distribution of DCs and TAMs in tumors of orthotopic Hepa1-6 tumor-bearing mice. (a) Immunofluorescence images of F4/80 (TAMs marker) and CD11c (DCs marker) in the tumor tissues of orthotopic Hepa1-6 tumor-bearing mice at 27 days after tumor inoculation. Scale bars: 50 μ m. Images are representative of three independent samples. (b) The percentages of DCs (gated as CD45+F4/80-CD11c+) and TAMs (gated as CD45+CD11b+F4/80+) in CD45+ cells in the tumor tissues of orthotopic Hepa1-6 tumor-bearing mice at 27 days after tumor inoculation by flow cytometry. Data are presented as means \pm s.d. (n = 5 mice per group; unpaired two-tailed Student's *t*-test). Source data are provided as a Source Data file.



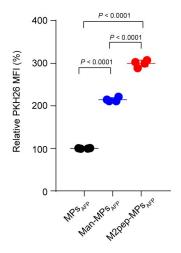
Supplementary Fig. 3 AFP expression in RAW264.7 cells and MPs_{AFP}. (a) AFP expression in RAW264.7 cells, RAW264.7 cells stably transfected with empty vector (RAW264.7_{EV}) and RAW264.7_{AFP} cells by western blotting. (b) AFP expression in MPs, MPs_{EV} and MPs_{AFP} by western blotting. Hepa1-6 cells were used as a positive control. Images are representative of three independent samples.



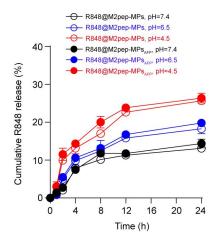
Supplementary Fig. 4 Effects of AFP overexpression on RAW264.7 cells. (a-c) Percentages of CD80⁺ (a), CD86⁺ (b) and MHC II⁺ (c) cells in RAW264.7 and RAW264.7_{AFP} cells. Data are presented as means \pm s.d. (n = 3 biological independent samples; unpaired two-tailed Student's *t*-test). (**d-f**) Percentages of CD80⁺ (**d**), CD86⁺ (e) and MHC II⁺ (f) MPs derived from RAW264.7 and RAW264.7_{AFP} cells. Data are presented as means \pm s.d. (n = 3 biological independent samples; unpaired two-tailed Student's *t*-test). Source data are provided as a Source Data file.



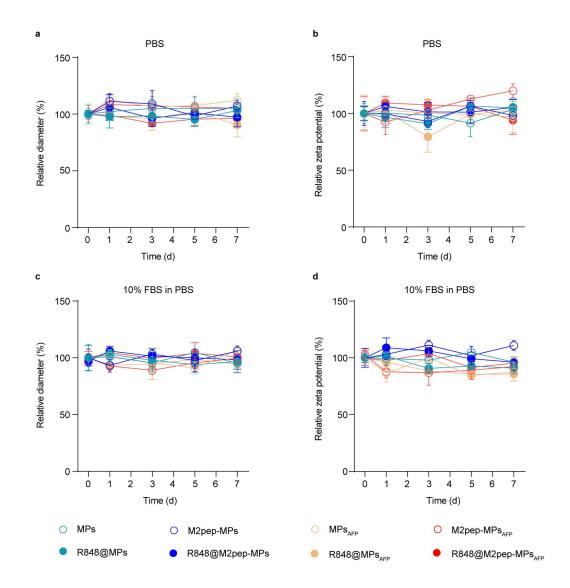
Supplementary Fig. 5 Colocalization of M2pep and MPs in M2pep-MPs by confocal microscopy. M2pep-MPs were prepared by incubating PKH26-labeled MPs with FITC-labeled DSPE-PEG-M2pep at a mass ratio of 50:1 at 4 °C for 24 h and then observed by confocal microcopy. Scale bars: 1 μm. Images are representative of three independent samples.



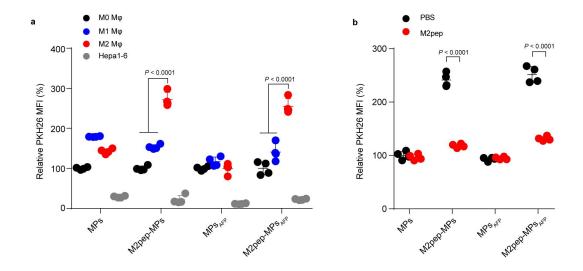
Supplementary Fig. 6 Targeting capacity of Man-MPs_{AFP} and M2pep-MPs_{AFP} to M2-like macrophages. IL-4-stimulated RAW264.7 cells (M2-like macrophages) were treated with PKH26-labeled MPs_{AFP}, Man-MPs_{AFP} or M2pep-MPs_{AFP} at the concentration of 10 μ g protein mL⁻¹ for 4 h, and PKH26 MFI was determined by flow cytometry. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 7 In vitro pH-dependent drug release of R848@M2pep-MPs_{AFP}. R848@M2pep-MPs_{AFP} and R848@M2pep-MPs_{AFP} were put in PBS at different pH values for different time intervals, and the released R848 content was determined by HPLC. Data are presented as means \pm s.d. (n = 3 independent samples). Source data are provided as a Source Data file.

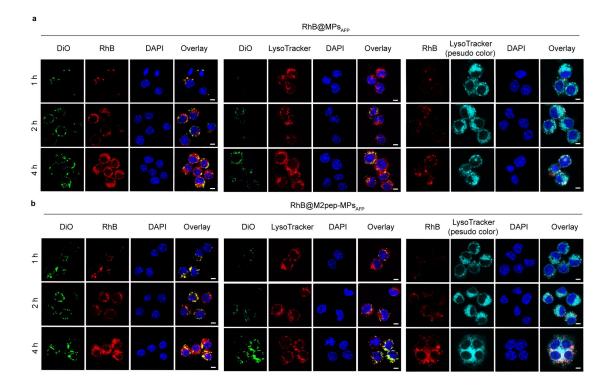


Supplementary Fig. 8 Stability of R848@M2pep-MPs_{AFP}. (a,b) Relative diameter (a) and zeta potential (b) of MPs, M2pep-MPs, MPs_{AFP} and M2pep-MPs_{AFP} with or without R848 after incubation in PBS for different time intervals. Data are presented as means \pm s.d. (n = 3 independent samples). (c,d) Relative diameter (c) and zeta potential (d) of MPs, M2pep-MPs, MPs_{AFP} or M2pep-MPs_{AFP} with or without R848 after incubation in PBS containing 10% FBS for different time intervals. Data are presented as means \pm s.d. (n = 3 independent samples). Source data are provided as a Source Data file.

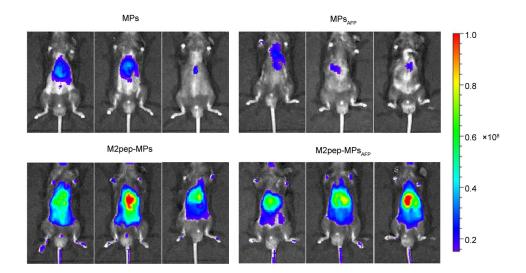


Supplementary Fig. 9 M2-like macrophage-targeting capacity of M2pep-MPs_{AFP}.

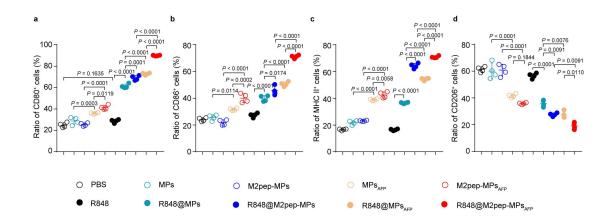
(a) Relative PKH26 MFI in RAW264.7 cells (M0 macrophages), LPS- and IFN γ -stimulated RAW264.7 cells (M1-like macrophages), IL-4-stimulated RAW264.7 cells (M2-like macrophages) and Hepa1-6 cells after treatment with PKH26-labeled MPs, M2pep-MPs, MPs_{AFP} or M2pep-MPs_{AFP} at the concentration of 10 µg protein mL⁻¹ for 4 h by flow cytometry. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (b) Relative PKH26 MFI in M2-like macrophages after treatment with PKH26-labeled MPs, M2pep-MPs, MPs_{AFP} or M2pep-MPs_{AFP} at the concentration of 10 µg protein mL⁻¹ in the presence or absence of 5 µg mL⁻¹ M2pep for 4 h. Data are presented as means \pm s.d. (n = 4 biological independent samples; unpaired two-tailed Student's *t*-test). Source data are provided as a Source Data file.



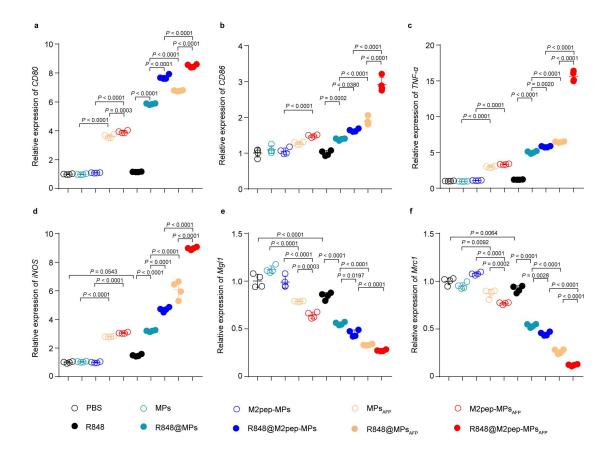
Supplementary Fig. 10 Intracellular trafficking of RhB@MPs_{AFP} and RhB@M2pep-MPs_{AFP} in M2-like macrophages. (a,b) Confocal microscopic images of M2-like macrophages after treatment with RhB-loaded DiO-labeled MPs_{AFP} (a) or M2pep-MPs_{AFP} (b) at the concentration of 10 μg protein mL⁻¹ for different time intervals and then labeling with 75 nM LysoTracker® Deep Red (lysosomes labeling dye) or 1 μg mL⁻¹ DAPI (nucleus labeling dye), respectively. Scale bars: 5 μm. Images are representative of three independent samples.



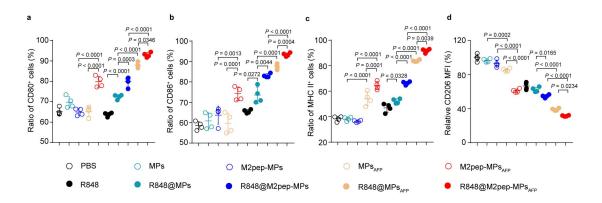
Supplementary Fig. 11 In vivo biodistribution of M2pep-MPs_{AFP}. Hepa1-6 tumor-bearing mice were intravenously injected with IR780-labeled MPs, M2pep-MPs, MPs_{AFP} or M2pep-MPs_{AFP} at the dosage of 15 mg protein kg⁻¹. At 24 h after injection, the in vivo imaging of mice was performed. (n = 3 mice per group).



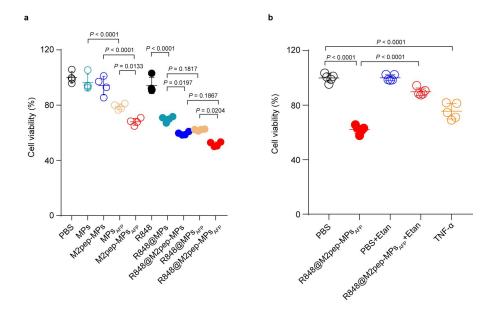
Supplementary Fig. 12 Reprogramming M2-like macrophages into M1-like phenotype by R848@M2pep-MPs_{AFP}. (a-d) Ratios of CD80⁺ (a), CD86⁺ (b), MHC II⁺ (c) and CD206⁺ (d) cells in IL-4-stimulated RAW264.7 cells after treatment with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h by flow cytometry. Data are presented as means ± s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



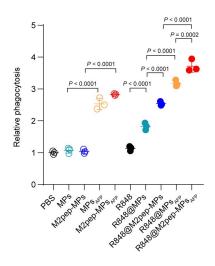
Supplementary Fig. 13 Reprogramming of M2-like macrophages to M1-like phenotype by R848@M2pep-MPs_{AFP}. (a-f) The mRNA expression levels of *CD80* (a), *CD86* (b), *TNF-\alpha* (c), *iNOS* (d), *Mgl1* (e) and *Mrc1* (f) in IL-4-stimulated RAW264.7 cells after treatment with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h by RT-qPCR. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



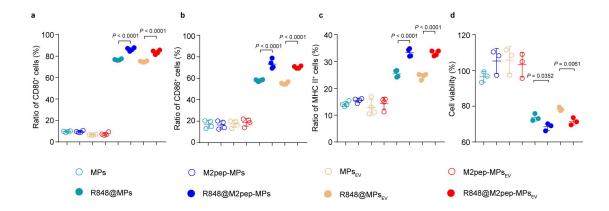
Supplementary Fig. 14 Reprogramming of M2-like BMDMs to M1-like phenotype by R848@M2pep-MPs_{AFP}. (a-d) Percentage of CD80⁺ (a), CD86⁺ (b) and MHC II⁺ (c) cells and relative CD206 MFI (d) in IL-4-stimulated BMDMs after treatment with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} derived from RAW264.7 or RAW264.7_{AFP} cells at the R848 concentration of 2 nM for 24 h by flow cytometry. Data are presented as means ± s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



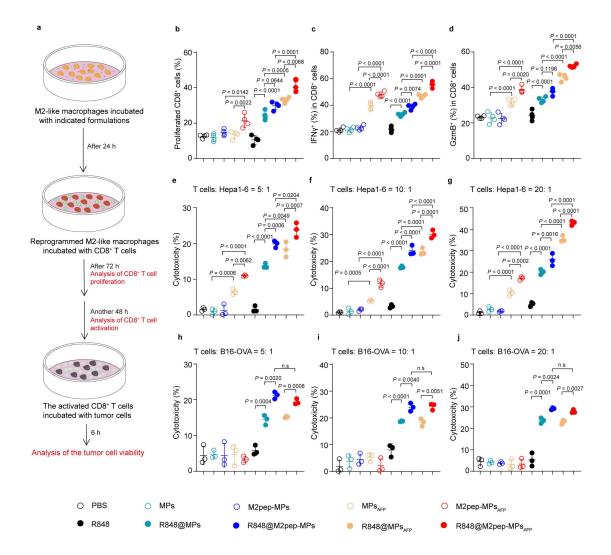
Supplementary Fig. 15 TNF-α involved the cytotoxicity R848@M2pep-MPs_{AFP}-reprogrammed M2-like macrophages against Hepa1-6 cells. (a) Cell viability of Hepa1-6 cells at 24 h after incubation with the supernatants of IL-4-stimulated RAW264.7 cells pretreated with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h by CCK-8 assay. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (b) Cell viability of Hepa1-6 cells at 24 h after incubation with the supernatants of IL-4-stimulated RAW264.7 cells pretreated with PBS or R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h in the presence or absence of 0.5 μg mL⁻¹ Etan. TNF- α at 0.5 μg mL⁻¹ was used as a positive control. Data are presented as means \pm s.d. (n = 5 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Phagocytosis Hepa1-6 cells Fig. 16 by R848@M2pep-MPs_{AFP}-reprogrammed M2-like macrophages. IL-4-stimulated DiD-labeled RAW264.7 cells were treated with PBS, MPs, M2pep-MPs, MPsAFP, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h, and then co-cultured with DiO-labeled Hepa1-6 cells at a ratio of 1:1 at 37 °C for 4 h. The phagocytosis of tumor cells by macrophages was determined by flow cytometry. Data are presented as means \pm s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

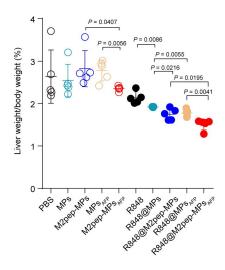


Supplementary Fig. 17 Effects of lentivirus transfection on the reprogramming of M2-like macrophages by R848@M2pep-MPs_{AFP}. (a-c) Percentage of CD80⁺ (a), CD86⁺ (b) and MHC II⁺ (c) cells in IL-4-stimulated RAW264.7 cells after treatment with MPs, M2pep-MPs, MPs_{EV}, M2pep-MPs_{EV}, R848@MPs, R848@M2pep-MPs, R848@MPs_{EV} or R848@M2pep-MPs_{EV} at the R848 concentration of 2 nM for 24 h by flow cytometry. Data are presented as means ± s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (d) Cell viability of Hepa1-6 cells at 24 h after treatment with the supernatants of IL-4-stimulated RAW264.7 cells pretreated with MPs, M2pep-MPs, MPs_{EV}, M2pep-MPs_{EV}, R848@MPs, R848@M2pep-MPs, R848@MPs_{EV} or R848@M2pep-MPs_{EV} at the R848 concentration of 2 nM for 24 h. Data are presented as means ± s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

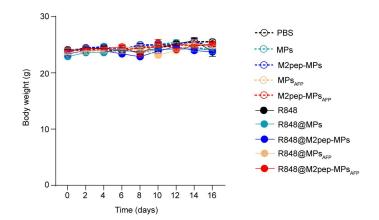


CD8⁺ 18 **Efficient Supplementary** Fig. activation cells by R848@M2pep-MPs_{AFP}-reprogrammed M2-like macrophages. (a) Schematic schedule for activating CD8⁺ T cells by R848@M2pep-MPs_{AFP}-reprogrammed M2-like macrophages. (b) Ratios of proliferated CD8⁺ T cells indicated by CFSE dilution at 3 days after co-culture with the PBS-, MPs-, M2pep-MPs-, MPs_{AFP}-, M2pep-MPs_{AFP}-, R848-, R848@MPs-, R848@M2pep-MPs-, R848@MPs_{AFP}- or R848@M2pep-MPs_{AFP}-pretreated IL-4-stimulated RAW264.7 cells (R848 concentration at 2 nM for 24 h) as indicated in (a). Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (c,d) Ratios of IFN γ^+ (c) and GzmB⁺ (d) cells in CD8⁺ T cells at 5 days after co-culture with the above-treated IL-4-stimulated RAW264.7 cells as indicated

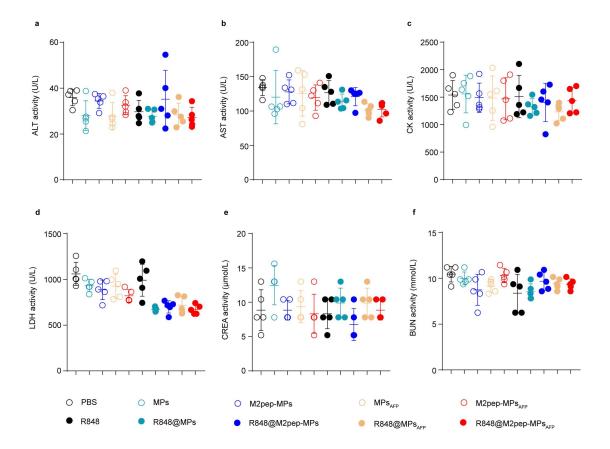
in (a). Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (e-j) Cytotoxicity of CD8⁺ T cells against Hepa1-6 or B16-OVA cells when the CD8⁺ T cells (effector cells) co-cultured with the above-treated IL-4-stimulated RAW264.7 cells for 5 days were incubated with Hepa1-6 or B16-OVA cells (target cells) at the effector/target ratio of 5:1 (e,h), 10:1 (f,i) and 20:1 (g,j) for 6 h as indicated in (a) by LDH assay. Data are presented as means \pm s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



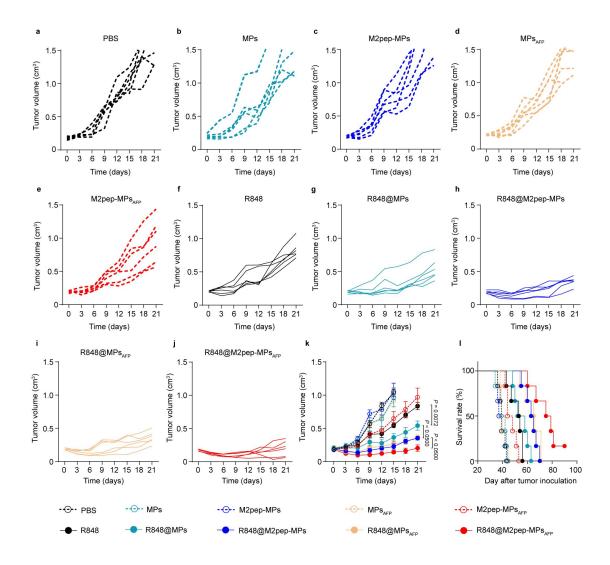
Supplementary Fig. 19 Efficient inhibition in the ratio of liver to body weight after treatment with R848@M2pep-MPs_{AFP} in orthotopic Hepa1-6 tumor-bearing mice. Orthotopic Hepa1-6 tumor-bearing mice were intravenously injected with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. At 16 days after intravenous injection, the ratio of liver to body weight was determined. Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



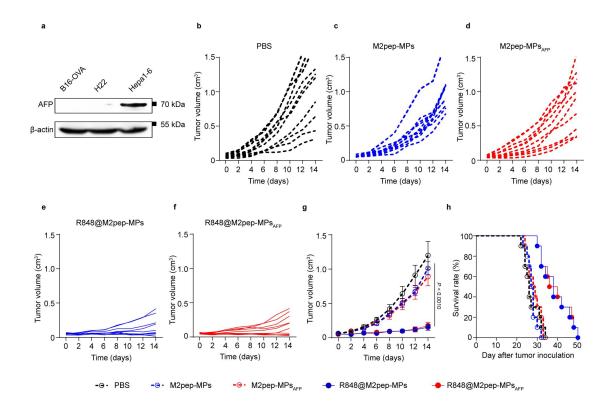
Supplementary Fig. 20 Body weight of orthotopic Hepa1-6 tumor-bearing mice after treatment with R848@M2pep-MPs_{AFP}. Orthotopic Hepa1-6 tumor-bearing mice were intravenously injected with PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@MPs_{AFP} or R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times, and the body weight was measured every other day. Data are presented as means \pm s.d. (n = 5 mice per group). Source data are provided as a Source Data file.



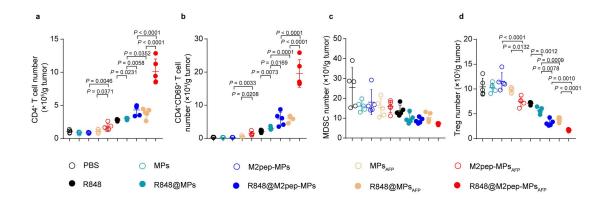
Supplementary Fig. 21 Serological analysis of orthotopic Hepa1-6 tumor-bearing mice after treatment with R848@M2pep-MPs_{AFP}. (a-e) The serological analysis of alanine aminotransferase (ALT, a), aspartate aminotransferase (AST, b), creatine kinase (CK, c), lactate dehydrogenase (LDH, d), creatinine (CREA, e) and blood urea nitrogen (BUN, f) in orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@M2pep-MPs, R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. Data are presented as means ± s.d. (n = 5 mice per group). Source data are provided as a Source Data file.



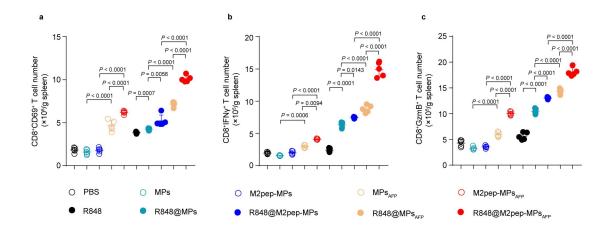
Supplementary Fig. 22 Anticancer activity of R848@M2pep-MPs_{AFP} in subcutaneous Hepa1-6 tumor-bearing mice. (a-j) Individual tumor growth curves of subcutaneous Hepa1-6 tumor-bearing mice after intravenous injection of PBS (a), MPs (b), M2pep-MPs (c), MPs_{AFP} (d), M2pep-MPs_{AFP} (e), R848 (f), R848@MPs (g), R848@M2pep-MPs (h), R848@MPs_{AFP} (i) or R848@M2pep-MPs_{AFP} (j) at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. (k) Average tumor growth curves of subcutaneous Hepa1-6 tumor-bearing mice after treatment as above. Data are presented as means ± s.e.m. (n = 6 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). (l) Kaplan–Meier survival plot of subcutaneous Hepa1-6 tumor-bearing mice after treatment as above. (n = 6 mice per group). Source data are provided as a Source Data file.



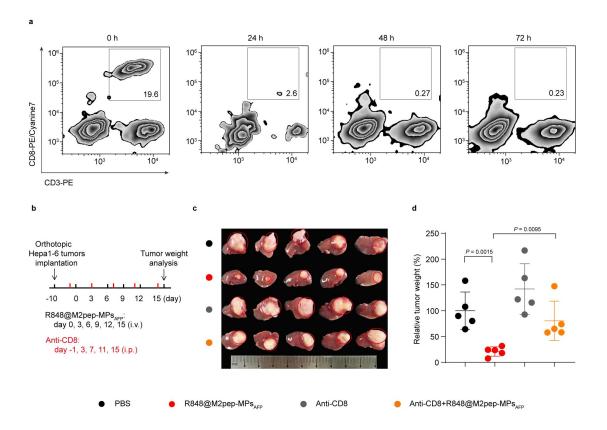
Supplementary Fig. 23 Anticancer activity of R848@M2pep-MPsAFP in subcutaneous H22 tumor-bearing mice. (a) AFP expression in B16-OVA, H22 and Hepa1-6 cells by western blotting. Images are representative of three independent samples. (b-f) Individual tumor growth curves of H22 tumor-bearing mice after injection of M2pep-MPs intravenous PBS (b), (c), $M2pep-MPs_{AFP}$ (d), R848@M2pep-MPs (e) or R848@M2pep-MPs_{AFP} (f) every three days for six times at the R848 dosage of 0.5 mg kg⁻¹. (g) Average tumor growth curves of H22 tumor-bearing mice after treatment as above. Data are presented as means \pm s.e.m. (n = 10 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). (h) Kaplan–Meier survival plot of H22 tumor-bearing mice after treatment as above. (n = 10 mice per group). Source data are provided as a Source Data file.



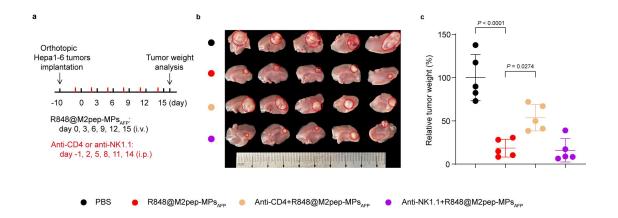
R848@M2pep-MPs_{AFP} in orthotopic Hepa1-6 tumor-bearing mice. (a-d) The numbers of CD4⁺ T cells (a), CD4⁺CD69⁺ T cells (b), MDSCs (c) and Tregs (d) in tumor tissues of orthotopic Hepa1-6 tumor-bearing mice at 16 days after intravenous injection of PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@M2pep-MPs, R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



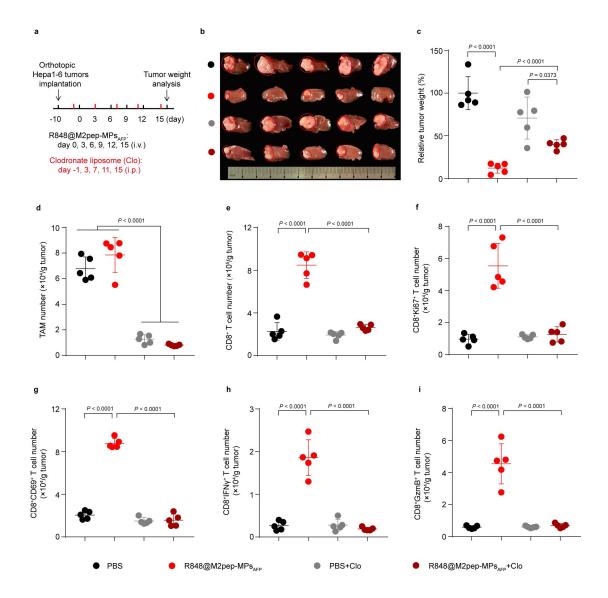
Supplementary Fig. 25 Improved systemic antitumor immunity by R848@M2pep-MPs_{AFP} in orthotopic Hepa1-6 tumor-bearing mice. (a-c) The numbers of CD8⁺CD69⁺ T (a), CD8⁺IFNγ⁺ T (b), and CD8⁺GzmB⁺ T (c) cells in spleens of orthotopic Hepa1-6 tumor-bearing mice at 16 days after intravenous injection of PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@M2pep-MPs, R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 26 CD8⁺ T cell-involved in R848@M2pep-MPs_{AFP}-induced anticancer effects in orthotopic Hepa1-6 tumor-bearing mice. (a) Percentages of CD3⁺CD8⁺ T cells in peripheral blood of orthotopic Hepa1-6 tumor-bearing mice at different time intervals after intraperitoneal injection of anti-CD8 antibody at the dosage of 100 μg per mice once. Images are representative of five independent samples. (b) Schematic schedule for the anticancer experiments in orthotopic Hepa1-6 tumor-bearing mice after CD8⁺ T cell depletion. (c,d) Tumor images (c) and tumor weight (d) of orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for 6 times in the presence or absence of intraperitoneal injection of anti-CD8 antibody at the dosage of 100 μg per mice every four days for 5 times as indicated in (b). Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

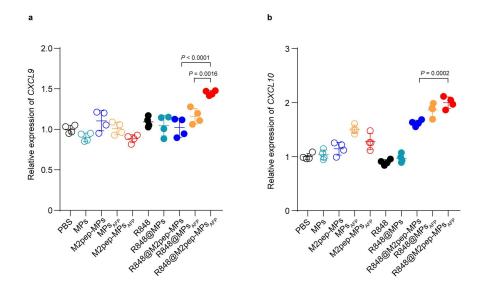


Supplementary Fig. 27 CD4⁺ T cell-involved in R848@M2pep-MPs_{AFP}-induced anticancer effects in orthotopic Hepa1-6 tumor-bearing mice. (a) Schematic schedule for the anticancer experiments in orthotopic Hepa1-6 tumor-bearing mice after CD4⁺ T or NK cell depletion. (b,c) Tumor images (b) and tumor weight (c) in orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for 6 times in the presence or absence of intraperitoneal injection of anti-CD4 or anti-NK1.1 antibody at the dosage of 100 µg per mice every three days for 6 times as indicated in (a). Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

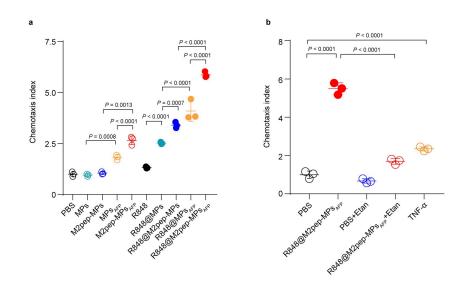


Supplementary Fig. 28 Macrophage-involved in R848@M2pep-MPs_{AFP}-induced anticancer effects and antitumor immunity in orthotopic Hepa1-6 tumor-bearing mice. (a) Schematic schedule for the anticancer experiments in orthotopic Hepa1-6 tumor-bearing mice after macrophage depletion. (b,c) Tumor images (b) and tumor weight (c) of orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for 6 times in the presence or absence of intraperitoneal injection of clodronate liposomes (Clo) at the dosage of 3.75 mg kg⁻¹ every four days for 5 times as indicated in (a). Data are

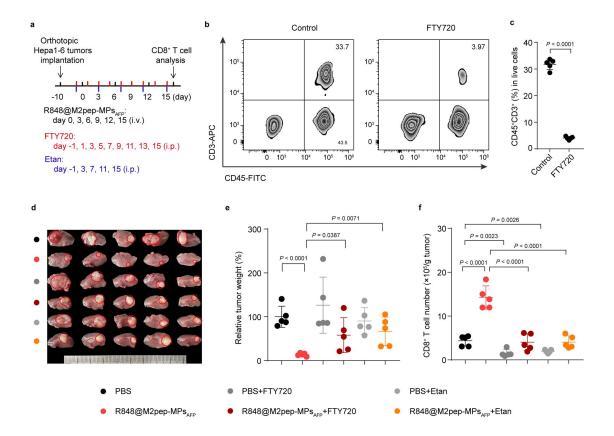
presented as means \pm s.d. for (c) (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). (d-i) The numbers of TAMs (d), CD8⁺ T cells (e), CD8⁺Ki67⁺ T cells (f), CD8⁺CD69⁺ T cells (g), CD8⁺IFN γ ⁺ T cells (h) and CD8⁺GzmB⁺ T cells (i) in tumor tissues of orthotopic Hepa1-6 tumor-bearing mice after treatment as indicated in (a). Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



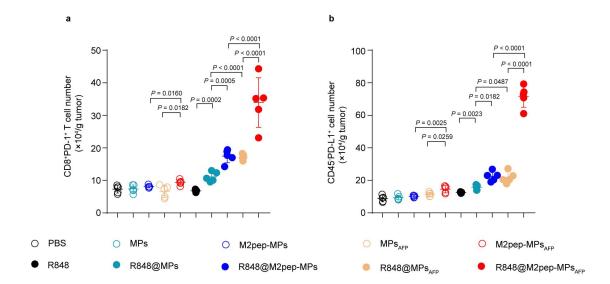
Supplementary Fig. 29 CXCL9 and CXCL10 expression in R848@M2pep-MPsafp-reprogrammed M2-like RAW264.7. (a,b) The mRNA expression levels of *CXCL9* (a) and *CXCL10* (b) in IL-4-stimulated RAW264.7 cells after treatment with PBS, MPs, M2pep-MPs, MPsafp, M2pep-MPsafp, R848, R848@MPs, R848@M2pep-MPs, R848@MPsafp or R848@M2pep-MPsafp at the R848 concentration of 2 nM for 24 h by RT-qPCR. Data are presented as means ± s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



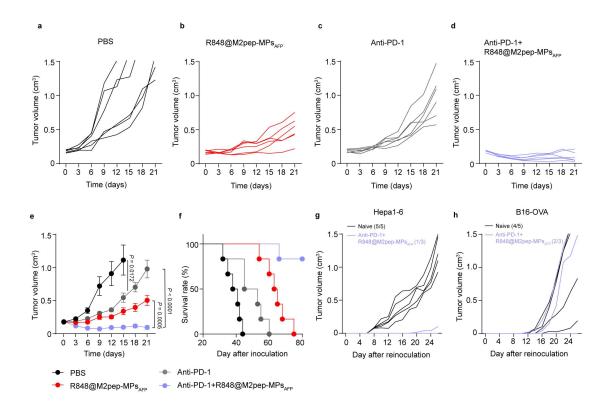
Supplementary Fig. 30 TNF-α-invovled in the recruitment of CD8⁺ T cells by R848@M2pep-MPs_{AFP}-reprogrammed macrophages. (a) Chemotaxis index of T lymphocytes seeded in the top chambers at 6 h after co-culture with the supernatants of IL-4-stimulated RAW264.7 cells pretreated with PBS, MPs, M2pep-MPs, MPs_{AFP}, R848@MPs, R848@M2pep-MPs, M2pep-MPs_{AFP}, R848, R848@MPsAFP R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h in the bottom chambers. Data are presented as means \pm s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (b) Chemotaxis index of T lymphocytes seeded in the top chambers at 6 h after co-culture with the supernatants of IL-4-stimulated RAW264.7 cells pretreated with PBS or R848@M2pep-MPs_{AFP} at the R848 concentration of 2 nM for 24 h in the presence or absence of Etan at concentration of 0.5 µg mL⁻¹ in the bottom chambers by flow cytometry. Data are presented as means \pm s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



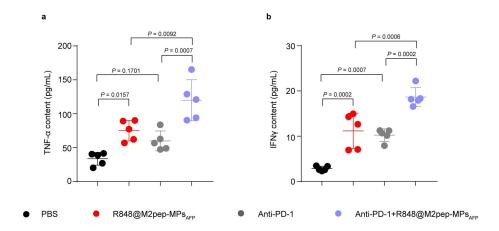
Supplementary Fig. 31 R848@M2pep-MPs_{AFP}-induced CD8⁺ T cell recruitment in orthotopic Hepa1-6 tumor-bearing mice. (a) Schematic schedule for R848@M2pep-MPs_{AFP}-induced CD8⁺ T cell recruitment experiment in orthotopic Hepa1-6 tumor-bearing mice. (b,c) Representative flow cytometric plots (b) and the percentages of CD45⁺CD3⁺ T cells in peripheral blood (c) of orthotopic Hepa1-6 tumor-bearing mice at 24 h after intraperitoneal injection of FTY720. Data are presented as means ± s.d. (n = 5 mice per group; unpaired two-tailed Student's *t*-test). (d-f) Tumor images (d), tumor weight (e) and CD8⁺ T cell numbers in tumor tissues (f) of orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of PBS or R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for 6 times in the presence or absence of intraperitoneal injection of FTY720 (1 mg kg⁻¹) or Etan (5 mg kg⁻¹) as indicated in (a). Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



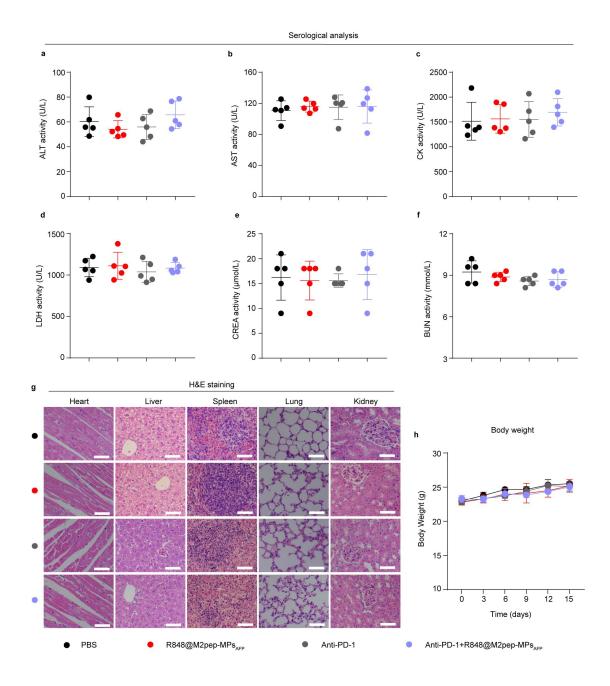
R848@M2pep-MPs_{AFP} in tumors tissues of orthotopic Hepa1-6 tumor-bearing mice. (a,b) The numbers of CD8+PD-1+ T cells (a) and CD45-PD-L1+ tumor cells (b) in tumor tissues of orthotopic Hepa1-6 tumor-bearing mice after intravenous injection of PBS, MPs, M2pep-MPs, MPs_{AFP}, M2pep-MPs_{AFP}, R848, R848@MPs, R848@M2pep-MPs, R848@M2pep-MPs, R848@M2pep-MPs_{AFP} at the R848 dosage of 0.5 mg kg⁻¹ every three days for six times. Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 33 Potent anticancer activity of combination of anti-PD-1 antibody and R848@M2pep-MPs_{AFP} in subcutaneous Hepa1-6 tumor-bearing mice. (a-d) Individual tumor growth curves of subcutaneous Hepa1-6 tumor-bearing mice after treatment with PBS (a), R848@M2pep-MPs_{AFP} (b), anti-PD-1 antibody (c) or the combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody (d) at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 6a. (e) Average tumor growth curves of subcutaneous Hepa1-6 tumor-bearing mice after treatment as above. Data are presented as means \pm s.e.m. (n = 6 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). (f) Kaplan–Meier survival plot of subcutaneous Hepa1-6 tumor-bearing mice after treatment as above. (n = 6 mice per group). (g,h) Individual tumor growth curves after rechallenge with Hepa1-6 cells (3 × 10⁶ cells, g) and B16-OVA cells (1 × 10⁶ cells, h) in naïve mice (n = 5 mice) or combination of anti-PD-1 antibody and R848@M2pep-MPs_{AFP}-cured mice (n = 3 mice). Source data are provided as a Source Data file.

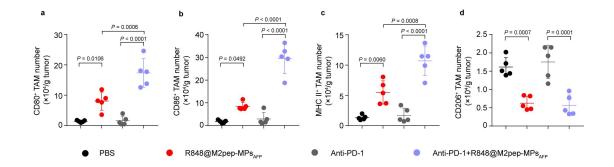


Supplementary Fig. 34 Cytokine contents in serum of orthotopic Hepa1-6 tumor-bearing mice after treatment with R848@M2pep-MPs_{AFP} and anti-PD-1 antibody. (a,b) The contents of TNF- α (a) and IFN γ (b) in serum of orthotopic Hepa1-6 tumor-bearing mice after treatment with PBS, R848@M2pep-MPs_{AFP}, anti-PD-1 antibody or a combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 6a. Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

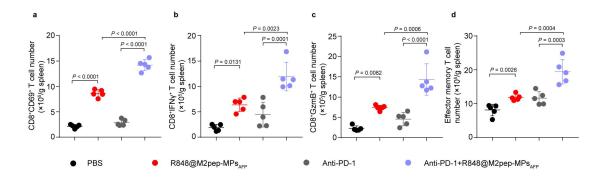


Supplementary Fig. 35 Biosafety of combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody in orthotopic Hepa1-6 tumor-bearing mice. (a-f) The activity of ALT (a), AST (b), CK (c), LDH (d), CREA (e) and BUN (f) in serum of orthotopic Hepa1-6 tumor-bearing mice after treatment with PBS, R848@M2pep-MPs_{AFP}, anti-PD-1 antibody or a combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. **6a**. Data are presented as means ± s.d. (n = 5 mice per group;

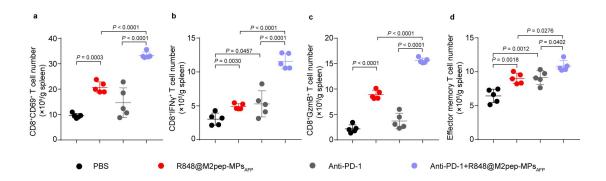
one-way ANOVA followed by Tukey's HSD post-hoc test). (g) H&E staining of major organs in orthotopic Hepa1-6 tumor-bearing mice after treatment indicated in Fig. 6a. Images are representative of 3 biologically independent mice. Scale bars: 20 μ m. (h) Body weight of orthotopic Hepa1-6 tumor-bearing mice during the treatment as indicated in Fig. 6a. Data are presented as means \pm s.d. (n = 5 mice per group). Source data are provided as a Source Data file.



Supplementary Fig. 36 Improved tumor immune microenvironment by combination of anti-PD-1 antibody and R848@M2pep-MPs_{AFP} in orthotopic Hepa1-6 tumor-bearing mice. (a-d) The numbers of CD80⁺ TAMs (a), CD86⁺ TAMs (b), MHC II⁺ TAMs (c) and CD206⁺ TAMs (d) in tumor tissues of orthotopic Hepa1-6 tumor-bearing mice after treatment with PBS, R848@M2pep-MPs_{AFP}, anti-PD-1 antibody or a combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 6a. Data are presented as means ± s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



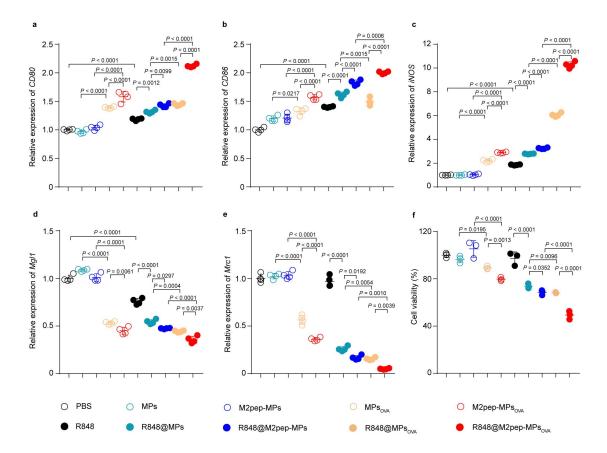
Supplementary Fig. 37 Improved systemic antitumor immunity by combination of anti-PD-1 antibody and R848@M2pep-MPs_{AFP} in orthotopic Hepa1-6 tumor-bearing mice. (a-d) The numbers of CD8+CD69+ T cells (a), CD8+IFN γ + T cells (b), CD8+GzmB+ T cells (c) and effector memory T cells (gated as CD45+CD3+CD8+CD44+CD62L-, d) in spleens of orthotopic Hepa1-6 tumor-bearing mice after treatment with PBS, R848@M2pep-MPs_{AFP}, anti-PD-1 antibody or a combination of R848@M2pep-MPs_{AFP} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 6a. Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



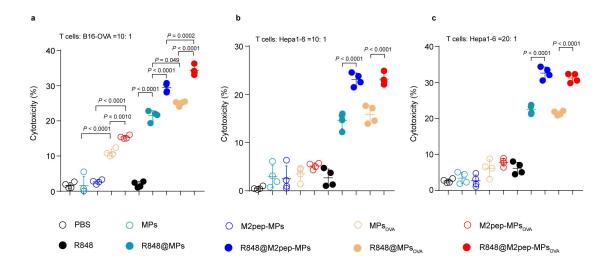
Supplementary Fig. 38 Improved systemic antitumor immunity by combination of anti-PD-1 antibody and R848@M2pep-MPsAFP in **DEN-induced** autochthonous HCC mice. (a-d) The numbers of CD8+CD69+ T cells (a), CD8⁺IFNγ⁺ T cells (b), CD8⁺GzmB⁺ T cells (c) and effector memory T cells (d) in spleens of DEN-induced autochthonous HCC mice after treatment with PBS, R848@M2pep-MPs_{AFP}, anti-PD-1 antibody combination R848@M2pep-MPs_{AFP} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 7a. Data are presented as means \pm s.d. (n = 5 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



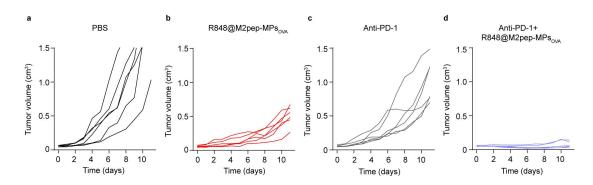
Supplementary Fig. 39 OVA expression in RAW264.7_{oVA} cells and MPsova. (a) OVA expression in RAW264.7, RAW264.7_{EV} and RAW264.7_{oVA} cells by western blotting. (b) OVA expression in MPs, MPs_{EV} and MPsova by western blotting. Images are representative of three independent samples.



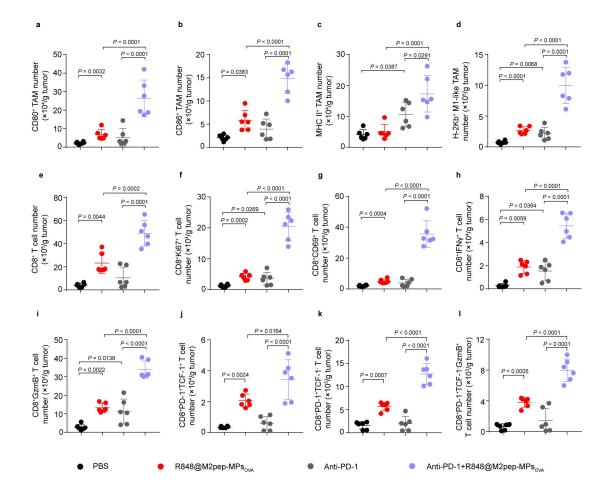
Supplementary Fig. 40 Reprogramming of M2-like macrophages to M1-like phenotype by R848@M2pep-MPsova. (a-e) The mRNA expression levels of CD80 (a), CD86 (b), iNOS (c), Mgll (d) and Mrcl (e) in IL-4-stimulated RAW264.7 cells after treatment with PBS, MPs, M2pep-MPs, MPsova, M2pep-MPsova, R848, R848@MPs, R848@M2pep-MPs, R848@MPsova or R848@M2pep-MPsova at the R848 concentration of 2 nM for 24 h by RT-qPCR. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). (f) Cell viability of B16-OVA cells at 24 h after treatment with the supernatants of the above treated IL-4-stimulated RAW264.7 cells by CCK-8 assay. Data are presented as means \pm s.d. (n = 3 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



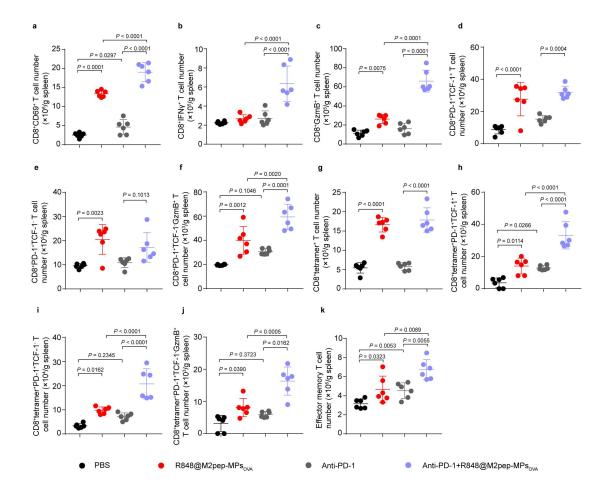
Cytotoxicity of activated **Supplementary** 41 $CD8^+$ R848@M2pep-MPsova-reprogrammed M2-like macrophages against B16-OVA or Hepa1-6 cells. (a-c) Cytotoxicity of CD8⁺ T cells against B16-OVA or Hepa1-6 cells when the CD8⁺ T cells cocultured with PBS-, MPs-, M2pep-MPs-, MPs_{OVA}-, M2pep-MPsova-, R848-, R848@MPs-, R848@M2pep-MPs-, R848@MPsova- or R848@M2pep-MPs_{OVA}-pretreated IL-4-stimulated RAW264.7 cells (R848 concentration at 2 nM for 24 h) for 5 days were incubated with B16-OVA or Hepa1-6 cells at the effector/target ratio of 10:1 (a,b) and 20:1 (c) for 6 h by LDH assay. Data are presented as means \pm s.d. (n = 4 biological independent samples; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 42 Antitumor effects of combination of R848@M2pep-MPsova and anti-PD-1 antibody in B16-OVA tumor-bearing mice. (a-d) Individual tumor growth curves of B16-OVA tumor-bearing mice after treatment with PBS (a), R848@M2pep-MPsova (b), anti-PD-1 antibody (c) or a combination of R848@M2pep-MPsova and anti-PD-1 antibody (d) at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 8i. (n = 6 mice per group). Source data are provided as a Source Data file.



Supplementary Fig. 43 Improved tumor immune microenvironment induced by combination of R848@M2pep-MPsovA and anti-PD-1 antibody in B16-OVA tumor-bearing mice. (a-l) The numbers of CD80⁺ TAMs (a), CD86⁺ TAMs (b), MHC II⁺ TAMs (c), H-2Kb⁺ M1-like TAMs (d), CD8⁺ T cells (e), CD8⁺Ki67⁺ T cells (f), CD8+CD69+ T cells (g), CD8+IFN γ + T cells (h), CD8+GzmB+ T cells (i), CD8⁺PD-1⁺TCF-1⁺ T cells (j), CD8⁺PD-1⁺TCF-1⁻ T cells (k) and CD8⁺PD-1⁺TCF-1⁻GzmB⁺ T cells (I) in tumor tissues of B16-OVA tumor-bearing mice at 11 days after treatment with PBS, R848@M2pep-MPsovA, anti-PD-1 antibody or a combination of R848@M2pep-MPs_{OVA} and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 8i. Data are presented as means \pm s.d. (n = 6 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 44 Improved immune microenvironment in spleens of B16-OVA tumor-bearing mice after treatment with R848@M2pep-MPsovA and anti-PD-1 antibody. (a-k) The numbers of CD8⁺CD69⁺ T cells (a), CD8⁺IFN γ ⁺ T cells (b), CD8+GzmB+ T cells (c), CD8+PD-1+TCF-1+ T cells (d), CD8+PD-1+TCF-1-T cells (e), CD8+PD-1+TCF-1-GzmB+ T cells (f), CD8+tetramer+ T cells (g), CD8⁺tetramer⁺PD-1⁺TCF-1⁻ T cells (h), CD8⁺tetramer⁺PD-1⁺TCF-1⁻ T cells (i), CD8+tetramer+PD-1+TCF-1-GzmB+ T cells (j) and effector memory T cells (k) in spleens of B16-OVA tumor-bearing mice at 11 days after treatment with PBS, anti-PD-1 R848@M2pep-MPsovA, antibody, combination R848@M2pep-MPsovA and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 8i. Data are presented as means \pm s.d. (n = 6 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.

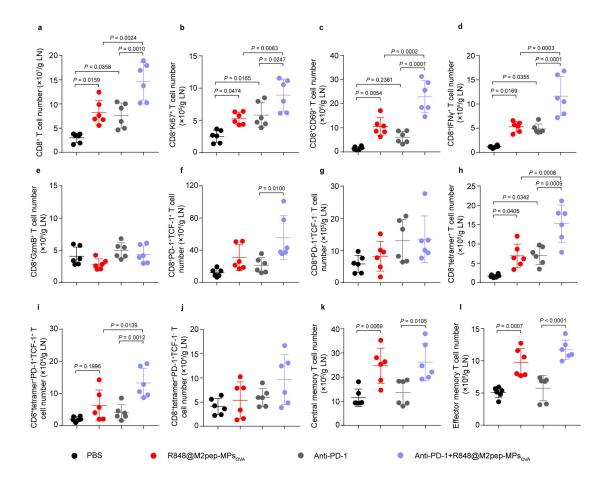
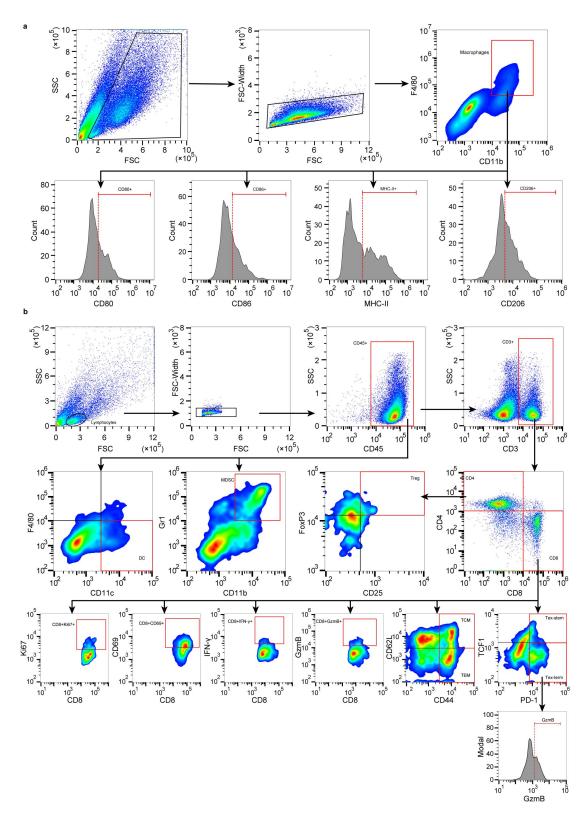


Fig. 45 Improved tumor immune microenvironment in **Supplementary** tumor-draining lymph nodes by combination of R848@M2pep-MPsovA and anti-PD-1 antibody in B16-OVA tumor-bearing mice. (a-l) The numbers of CD8⁺ T cells (a), CD8+Ki67+ T cells (b), CD8+CD69+ T cells (c), CD8+IFNγ+ T cells (d), CD8+GzmB+ T cells (e), CD8+PD-1+TCF-1+ T cells (f), CD8+PD-1+TCF-1- T cells (g), CD8⁺tetramer⁺ cells (h), CD8⁺tetramer⁺PD-1⁺TCF-1⁺ CD8+tetramer+PD-1+TCF-1- T cells (j), Tcm cells (k) and Tem cells (l) in tumor-draining lymph nodes of B16-OVA tumor-bearing mice at 11 days after treatment with PBS, R848@M2pep-MPsovA, anti-PD-1 antibody, or a combination of R848@M2pep-MPsovA and anti-PD-1 antibody at the anti-PD-1 antibody dosage of 5 mg kg⁻¹ and R848 dosage of 0.5 mg kg⁻¹ indicated in Fig. 8i. Data are presented as means \pm s.d. (n = 6 mice per group; one-way ANOVA followed by Tukey's HSD post-hoc test). Source data are provided as a Source Data file.



Supplementary Fig. 46 Gating strategy for identifying major immune cell populations in tumor tissues of tumor-bearing mice. (a) Gating strategy for identifying M1- and M2-like TAMs in tumor tissues of tumor-bearing mice presented

on Fig. 3g, h, 4b-e, 7g-j, 8l and Supplementary Fig. 28d, 36a-d and 43a-d. (b) Gating for identifying DCs (CD45+F4/80-CD11c+ cells), strategy **MDSCs** (CD45⁺CD11b⁺Gr1⁺ cells), Tregs (CD45⁺CD3⁺CD4⁺CD25⁺FoxP3⁺ cells), CD4⁺ T cells (CD45⁺CD3⁺CD4⁺ cells), CD8⁺ T cells (CD45⁺CD3⁺CD4⁺ cells), CD8⁺CD69⁺ T cells (CD45⁺CD3⁺CD8⁺CD69⁺ cells), CD8⁺Ki67⁺ T cells (CD45⁺CD3⁺CD8⁺Ki67⁺ cells), CD8⁺IFNγ⁺ T cells (CD45⁺CD3⁺CD8⁺IFNγ⁺ cells), CD8⁺GzmB⁺ T cells (CD45⁺CD3⁺CD8⁺GzmB⁺ cells), Tem cells (CD45⁺CD3⁺CD8⁺CD44⁺CD62L⁻ cells), Tcm cells (CD45⁺CD3⁺CD8⁺CD44⁺CD62L⁺ cells), stem-like CD8⁺ T cells (CD45⁺CD3⁺CD8⁺PD-1⁺TCF-1⁺ cells), terminally exhausted CD8⁺ T (CD45⁺CD3⁺CD8⁺PD-1⁺TCF-1⁻ cells) and terminally exhausted CD8⁺ T cells secreting GzmB (CD45+CD3+CD8+PD-1+TCF-1-GzmB+ T cells) in tumors, spleens or lymph nodes of tumor-bearing mice presented on Fig. 3g, 5e-l, 6e-l, 7k-r, 8l and Supplementary Fig. 24a-d, 25a-c, 28e-i, 31f, 37a-d, 38a-d, 43e-l, 44a-f, k and 45a-g, k, 1.