A Bifunctional Fusion Membrane- Based Biocompatible Nanovaccine to

Potentiate Cancer Immunotherapy

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Supplementary Figure 1. STCM-NPs promote antigen presenting by DCs and activation of T cells, related to Figure 1. (A) $SA\beta G$ staining of B16-F10 tumor cells following senescence induction via treatment with different concentrations of cisplatin for 6 days. Scale bar, 100 µm. (B) Sketch depicting the co-culture assays. (C)

Representative flow cytometry images of mDCs (CD11c⁺ CD80⁺ and CD11c⁺ CD86⁺) co-incubated with different groups (PBS, LPS, TCMs, EMs, and STCMs) for 48 h in *vitro* (n = 3). (D) The proportion of mDCs and the levels of MHC-I and MHC-II on mDCs after being co-cultured with various formulations for 48 h *in vitro* (n = 3). (E) Representative flow cytometry images of IFN- γ^+ CD8⁺ T cells (lower) and GZMB⁺ CD8⁺ T cells (uper) after being co-cultured with BMDCs pre-treated with various formulations for 72 h in vitro. (F) Percentages of CD45⁺ T cells, CD3⁺ T cells and CD4⁺ T cells (n = 3) after being co-cultured with BMDCs pre-treated various formulations for 72 h *in vitro*. (G) Percentages of CD8⁺ T cells, GZMB⁺ T cells and IFN- γ^+ CD8⁺ T cells (gated from $CD3^+CD8^+$ cells; n = 3). (H) The SDS-PAGE image of STCM-NPs (red rectangle), and EM-NPs (black rectangle). The amounts of protein in gels were 10 ug. (I) Diameter (n = 3) of EM-NPs, STCM-NPs and FM-NPs. Data are presented as mean \pm SD. One-way ANOVA with subsequent multiple comparison tests were conducted, where ns indicates no significance, p < 0.05, p < 0.01, p < 0.01, p < 0.001 and *****p* < 0.0001.



Supplementary Figure 2. Additional analysis and characterization of Bio-HCP-

NPs. (A) Zeta size curves of Bio-HCP-NPs. (B) WCA analysis of PS, PS@PEG-NPs and Bio-HCP-NPs (C) XPS surface elements analysis of PS, PS@PEG-NPs and Bio-HCP-NPs. (D) Isothermal N2 absorption and desorption curves of Bio-HCP-NPs in total-pore BET testing. (E) Horvath-Kawazoe differential pore volume plot of Bio-HCP-NPs in total-pore BET testing. (F) BJH pore size curves of Bio-HCP-NPs in totalpore BET testing. (G) FESEM images of series Bio-HCP-NPs formed with different proportion of monomers. (H) Additional morphological and structural characterization of Bio-HCP-NPs in 90/10 wt% St/EG ratio. (I) Pore sizes distribution curves of series Bio-HCP-NPs formed with different proportion of monomers.



Supplementary Figure 3. Bio-HCP@FM-NPs activate DCs, related to Figure 3. (A)

Flow cytometry gating strategy for measuring the percentages of T cells, IFN- γ^+ T cells and GZMB⁺ T cells (n = 3). (B-C) Percentages of T cells, CD4⁺ T cells in direct coculture assay (n = 3). (D-E) Expression of CD86 and CD80 on DCs in direct co-culture assay (n = 3). (F-I) Percentages of T cells, CD4⁺ T cells, CD80⁺ DCs and CD86⁺ DCs in indirect co-culture assay (n = 3).



Supplementary Figure 4. GO and KEGG analysis of BMDCs, related to Figure 3.

(A) GO and KEGG analysis of BMDCs.



Supplementary Figure 5. Assessment of biodistribution and targeting capability of Bio-HCP@FM-NPs vaccination in *vitro* and *vivo*. (A) Quantitative analysis of fluorescence signals from FITC-labeled nanoparticles in DCs at 24 h post-injection. (B-H) The quantitative analysis of nanoparticle accumulation in major organs, tumor and lymph nodes from Bio-HCP@EM-NPs, Bio-HCP@STCM-NPs and Bio-HCP@FM-NPs injected B16-F10 cells orthotopic tumor-bearing mice *in vivo*. (I) Retention ability of Bio-HCP@EM-NPs, Bio-HCP@STCM-NPs and Bio-HCP@FM-NPs in tumor tissues of B16-F10 cells orthotopic tumor-bearing murine model quantified by flow

cytometry at different time points. Data are presented as mean \pm SD, with n = 3 per group. Statistical significance is indicated as follows: n.s. (no significance), *p < 0.05, **p < 0.01, ***p < 0.001, ***p < 0.0001.





Supplementary Figure 6. Bio-HCP@FM-NPs show good biosafety. (A-B) The

cytotoxicity of Bio-HCP@FM-NPs against B16-F10 and LLC cells after co-incubation with nanovaccines. (C) H&E staining of heart, liver, spleen, lung, and kidney in B16-F10 mouse melanoma tumor model on day 7 after treatment. The scale bar is 200 μ m. (D) Hemanalysis was performed on blood drawn from mice on day 7 after treatment. Aspartate transaminase (AST), white blood cell (WBC), alanine transaminase (ALT), red blood cell (RBC), blood urea nitrogen (BUN), and creatinine (CR) are presented as the means ± SEM. All data are mean ± SD; n = 3. **p* <0.05, ***p* < 0.01, ****p* < 0.001, and ns, not significant.



Supplementary Figure 7. Bio-HCP@FM-NPs increase tumor immune response in the TME, related to Figure 5. (A) Flow cytometry gating strategy for measuring the percentages of T cells, IFN- γ^+ and GZMB⁺T cells in tumor tissue. (B-C) Percentages

of T cells and CD4⁺T cells in tumor tissue. (D-E) Expression of IFN- γ and GZMB on CD4⁺T cells in tumor tissue (n = 3). (F) Percentages of TAMs in tumor tissue.



Supplementary Figure 8. Combinatorial efficacy of personalized Bio-HCP@FM-NPs vaccination and anti-PD-1 against postsurgical tumor recurrence, related to Figure 7.

(A) B16F10 tumor-growth curves for each mouse in different groups (n = 6). (B) Average tumor-growth curves of *C57BL/6J* mice bearing LLC tumor with different treatments as indicated (n = 6). (C) LLC tumor-growth curves for each mouse in different groups (n = 6).



Supplementary Figure 9. Validation of the immune cell depletion efficiency, related to Figure 7. (A-B) Flow cytometry analysis of changes in immune cells from peripheral blood and spleen of mice subjected to different immune cell depletion treatments as indicated (n = 6-8 per group).



Supplementary Figure 10. Both innate and adaptive immunity are required for tumor recurrence after personalized Bio-HCP@FM-NPs vaccination, related to Figure 7. (A) B16F10 tumor-growth curves for each mouse in different groups (n = 6). (B) Average tumor-growth curves in LLC-bearing mice pre-treated with different monoclonal antibodies, before the combination treatment of Bio-HCP-NPs vaccination

and anti-PD-1 (n = 6). (C) LLC tumor-growth curves for each mouse in different groups (n = 6).

Sample	C (Atom %)	O (Atom %)
PS	82.16	17.84
PS@PEG-NPs	79.02	20.98
Bio-HCP-NPs	91.58	8.42

Table S1. Elemental analysis of XPS in series polymer nanoparticles.

Table S2. Specific surface area (SBET), average pore size (DBJH) and pore volume

((Vp)) of series Bio-HCP-NPs before and after load	led drug.
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Sample	<i>S</i> _{ВЕТ} (m ² /g)	D _{BJH} (nm)	<i>V</i> _P (cm ⁻³ /g)
Bio-HCP-NPs	8.08	24.45	0.06
PS@PEG-NPs	907.05	6.65	1.02
Bio-HCP-NPs@GM-CSF	349.33	7.50	0.57

Table S3. Specific surface area (SBET), average pore size (DBJH) and pore volume

((Vp)) of series Bid	o-HCP-NPs forme	d from different	proportions of monomers.
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Sample	$S_{\rm BET}~({ m m}^2/{ m g})$	D _{BJH} (nm)	<i>V</i> _P (cm ⁻³ /g)
90/10 St/EG	907.05	6.65	1.02
80/20 St/EG	873.29	3.73	0.55
70/30 St/EG	624.26	6.31	0.40

60/40 St/EG	406.74	11.96	0.78
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