Supporting Information

Lymph node-targeting adjuvant/neoantigen-codelivery vaccines for combination glioblastoma radioimmunotherapy

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Figure S1: HPLC chromatograms and UV absorption of CpG-A1-MEB. CpG-A1-MEB showed both the characteristic peaks for DNA and MEB at 11.51 min of retention time, indicating successful MEB conjugation with CpG-A1.



Figure S2: (A) The UV–Vis absorption spectrum of MEB at different concentration and the standard curve from 550 nm to 260 nm. (B) Absorption of free MEB (1.25 or 2.5 mg/mL) and CpG-A1-MEB (after 500x dilution of its stock stolution) (For CpG-A1-MEB, A260 = 0.8988 and A550 = 0.08278). Using standard curve of MEB absorption at 550 nm vs concentration, the MEB concentration in the CpG-A1-MEB was calculated to be 1.9832 mg/mL. At this concentration, the absorption of free MEB at A260 is 0.04267. By comparing the UV absorption of CpG-A1-MEB with that of free MEB at 1.25 mg/mL or 2.5 mg/mL, the influence of MEB absorption at 260 nm on CpG-A1-MEB absorption at 260 nm is very small.



Figure S3: HPLC chromatograms of A3-Ntrk1, A3-Rtn2 and A3-Imp3.



Figure S4: Image of agarose gel electrophoresis showing the products of A3-Rtn2, A3-Ntrk1 and A3-Imp3, free mixing of A3 with peptide Rtn2, Ntrk1 and Imp3 are used as control (A3+Rtn2, A3-Ntrk1 and A3-Imp3).



Figure S5. Stability of Y-shaped DNA in PBS containing 10% FBS (Solution) at 37 °C for 24 hours by gel electrophoresis.



Figure S6: Synthesis of A3-SIINFEKL conjugate. (A) Representative HPLC chromatograms of A3 and A3-SIINFEKL. (B) Image of agarose gel electrophoresis showing the products of A3-Rtn2, A3-Ntrk1, A3-Imp3 and A3-SIINFEKL.



Figure S7. Analysis of T cell responses elicited by AAco-AlbiVax in mice. (A) Study design of AAco-AlbiVax T cell responses in C57BL/6 mice (n = 5), the same study shown in Figure 3. (B-D) Flow cytometry data showing

the peripheral T cell responses elicited by AAco-AlbiVax as above on Days 21, 42 and 56, including MFI of PD-1 expression on peripheral CD8⁺ T cells, CD8⁺ T cell percentages, percentages of cytokine⁺ T cells among CD8⁺ T cells and CD4⁺ T cells, and quantification of CD8⁺ memory T cells (Central memory: CD62L⁺CD44⁺, effector memory: CD62L⁻CD44⁺ and naive T cells: CD62L⁺CD44⁻). Data: mean ± SEM; * p < 0.05; ** p < 0.01; *** p < 0.001 (Student's t-test).



Figure S8. Flow cytometric analysis of peripheral blood from the above immunized mice (Figure S6, Figure 3) on day 35. (A) Intracellular cytokine staining results showing the percentages of cytokine⁺ CD4⁺ T cells on total PBMC CD4⁺ T cells. (B) Flow cytometry plots of CD8⁺ memory T cells (central memory: CD62L⁺CD44⁺, effector memory: CD62L⁻CD44⁺ and naive T cells: CD62L⁺CD44⁻). The quantified data are shown in Figure 3H. (C) Frequencies of CD8⁺ T cells in peripheral blood by flow cytometry at day 35. (D) Quantification of CD45⁺CD11b⁺F4/80⁺ macrophages in peripheral blood by flow cytometry at day 35.



Figure S9. Quantification of CD45⁺CD11c⁺ DCs, CD8⁺ DCs (CD11c⁺CD8⁺B220⁻) and CD45⁺CD11b⁺F4/80⁺ macrophages among PBMCs from the immunized mice as shown in Figure S6 or Figure 3 (Day 35). Data: mean \pm SEM; * p < 0.05; ** p < 0.01; *** p < 0.001 (Student's t-test).

Table S1. DNA sequences.

	Sequences (5' - 3')	
CpG	TCCATGACGTTCCTGACGTT	
A1	TTTTT GAGTCAGTCACTGTGTCGTG	
CpG-A1	TCCATGACGTTCCTGACGTT TTTTT GAGTCAGTCACTGTGTCGTG - SH	
A2	TTTTT CGAGAACCTATGACTGACTC	
A3	NH2 - TTTTT CACGACACAGTAGGTTCTCG	