#### **RESEARCH ARTICLE**



# Biosynthetic neoantigen displayed on bacteria derived vesicles elicit systemic antitumour immunity

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#### 1 | Introduction

Abstract

Neoantigens derived from mutant proteins in tumour cells could elicit potent personalized anti-tumour immunity. Nevertheless, the layout of vaccine vehicle and synthesis of neoantigen are pivotal for stimulating robust response. The power of synthetic biology enables genetic programming bacteria to produce therapeutic agents under contol of the gene circuits. Herein, we genetically engineered bacteria to synthesize fusion neoantigens, and prepared bacteria derived vesicles (BDVs) presenting the neoantigens (BDVs-Neo) as personalized therapeutic vaccine to drive systemic antitumour response. BDVs-Neo and granulocyte-macrophage colony-stimulating factor (GM-CSF) were inoculated subcutaneously within hydrogel (Gel), whereas sustaining release of BDVs-Lipopolysaccharide (LPS) and GM-CSF recruited the dendritic cells (DCs). Virtually, Gel-BDVs-Neo combined with the programmed cell death protein 1 (PD-1) antibody intensively enhanced proliferation and activation of tumour-infiltrated T cells, as well as memory T cell clone expansion. Consequently, BDVs-Neo combining with checkpoint blockade therapy effectively prevented tumour relapse and metastasis.

#### KEYWORDS

bacteria derived vesicles (BDVs), cancer vaccines, checkpoint blockade, immunotherapy, neoantigen

Cancer vaccines are expected to exert the protective or therapeutic efficacy as that of pathogenic microorganisms (Melero et al., 2014; Palena et al., 2006; Sahin & Tureci, 2018). However, the dearth of potent anti-tumour response of the tumour-associated antigens (TAAs), derived from self-antigens that are abnormally expressed in tumour cells, leads to the termination of most TAAs-vaccines-associated clinical trials in the past decades (Blass & Ott, 2021; Hollingsworth & Jansen, 2019; Kreiter et al., 2015;

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Saxena et al., 2021; Schumacher & Schreiber, 2015). Virtually, the antitumour immune responses from TAAs are basically unsatisfactory in the clinical, which is probably ascribed to the fact that TAA-specific T cells suffer central and/or peripheral tolerance (Blass & Ott, 2021; Hollingsworth & Jansen, 2019). Recently, the mutant antigens, referred as neoantigens, have been identified from individual tumour tissue by exome sequencing (Kreiter et al., 2015). Neoantigen is an abnormal protein exclusively produced by cancer cells through 'non-synonymous mutation' (Kreiter et al., 2015; Yang et al., 2019). Hence, T cells enable eradicate cancer cells presenting neoantigens, which could drive potent anti-tumour response (Hu et al., 2021; Keskin et al., 2019; Schumacher & Schreiber, 2015; Yang et al., 2019). Neoantigen-based vaccines have been shown to specifically induce tumour antigen-specific CD4<sup>+</sup> and CD8<sup>+</sup> T cell responses, which could improve patient survival combining with immune checkpoint inhibition (ICI) therapy (Saxena et al., 2021). Intriguingly, neoantigen-specific T cells can avoid 'off-target' damage to normal tissues, since neoantigens circumvent T cell central tolerance to self-epitopes (Blass & Ott, 2021). Currently, neoantigen vaccines developed with poly-neo-epitope RNA or peptide sequestered in liposome enable boost tumour-specific immune responses (Kreiter et al., 2015; Sahin et al., 2020). However, spontaneous immune recognition of neo-epitope is inefficient if no proper immunologic adjuvant is utilized. Hence, it is critical to devise an adjuvant vehicle for the unmet demand of neoantigen vaccines.

Lipid-based nanoparticles (NPs) such as liposomes, exosomes, tumour cell membrane vesicles and bacterial outer membrane vesicles (OMVs), have been intensively explored as cancer antigen vehicles to elicit systemic antitumour immunity (Cheng et al., 2021; Valentine et al., 2016; Ye et al., 2019). Intriguingly, OMVs released from Gram-negative bacteria contain foreign covmponents and thus could potentiate immune response (Carvalho et al., 2019; Kuehn & Kesty, 2005; Lee et al., 2008; Rosenthal et al., 2014; Zanella et al., 2021). As a major component of OMVs, lipopolysaccharide (LPS) is a potent activator of immune cells, such as monocytes/macrophages (Kulp & Kuehn, 2010; Sartorio et al., 2021). However, this inflammatory activation can also result in high vaccine reactogenicity. Exception of LPS, OMVs also contain pathogen-associated antigens and lipoproteins, which could modulate and activate innate immunity by the Toll-like receptor 2 (TLR2) signal pathway (Bachmann & Jennings, 2010; Kovacs-Simon et al., 2011; Mancini et al., 2020). Based on the characteristics of OMVs, it has inspired researchers to explore their application in tumour immunity (Kim et al., 2017; Huang et al., 2020; Li, Zhou et al., 2020; Li, Zhao et al., 2020; Peng et al., 2020). Recently, in order to enhance the immunization efficacy of the antigens, amounts of studies have focused on integrating antigens into OMVs. For example, Salmonella OMVs which presenting an antigen of Mycobacterium tuberculosis named Ag85B were constructed by Luirink et al., the study resulting in the the engineered OMVs could activate the bone marrow-derived dendritic cells (BMDCs), and the antigen are subsequently presented to the navie T cells (Daleke-Schermerhorn et al., 2014). E. coli OMVs displaying Omp22 could protect mice from Acinetobacter baumannii-induced lethal challenge and trigger a robust immune response after immunization (Huang et al., 2016). Moreover, OMVs carrying Cytolysin A fusion proteins have been investigated for their potential as anti-tumour agents (Murase, 2022). In addition, the design of hybrid biomimetic nanovesicles has been explored by fusion of eukaryotic and prokaryotic cell derived membrane vesicles (Chen, Huang et al., 2020; Park et al., 2021). Alternatively, OMVs have been developed as a novel type of drug delivery vehicles for enhanced enhancing cancer immunotherapy (Chen, Bai et al., 2020; Gujrati et al., 2014; Huang et al., 2020; Patel et al., 2019).

Synthetic biology, a promising discipline, aims to harness living cells including bacteria to produce chemicals, materials and biological therapeutic agents under the control of gene circuits (Chowdhury et al., 2019; Cubillos-Ruiz et al., 2021). Currently, synthetic gene circuits are devised to program bacteria to generate therapeutic agents such as antibodies, immunologic factors, and even L-arginine in tumour microenvironment to eradicate cancer (Canale et al., 2021; Gurbatri et al., 2020; McNerney et al., 2021). The exploitation of natural or bioengineered bacteria membrane derived vesicle as therapeutic bacteria vaccines has shown considerable prospects. Herein, we consulted the nucleotide sequence of mutations identified by exome sequencing (Kreiter et al., 2015) and utilized the non-pathogenic *Escherichia coli* (*E. coli*) BL21(DE3)plysS strain to engineer bacteria derived vesicles (BDVs) displaying the neoantigen peptide. BDVs-Neo and granulocyte-macrophage colony-stimulating factor (GM-CSF) were loaded into a thermosensitive hydrogel as antigen delivery vehicle, then was accompanied with PD-1 antibody to improve the anticancer efficacy and long-term memory immunity (Scheme 1). PD-1 antibody protected CD8<sup>+</sup> T cells from exhaustion by blockade of PD-1/PD-L1 axis, that facilitated cytotoxic T cells (CTLs) to recognize and eliminate the surgical residual tumour cells. Collectively, the unique features of BDVs contribute to boosting ICI therapy and reactivating the T cell antitumour immunity (Chen & Han, 2015; Gong et al., 2018; Wei et al., 2021; Zou et al., 2016).

#### 2 | MATERIALS AND METHODS

#### 2.1 | Plasmid construction

The nucleotide sequences of neoantigens were based on the next-generation sequencing (Kreiter et al., 2015). DNA fragments encoding for the Lpp-OmpA-Green Fluorescent Proteins (GFP)-M33-M47 (neoantigen) protein were synthesized commercially by Yingmao, Inc. (Chongqing, China). Using gene recombinant techniques, wild type epitopes (M33-Valine; M47-Alanine) and the mutated epitopes (M33-Asparticacid; M47-Glycine) gene segments were linked into pET21a vector between *NdeI* and *Xhol I* 



sites, respectively (Table S1). The recombinant plasmids were transformed into competent *E. coli* BL21(DE3)plysS cells (TransGen Biotech).

# 2.2 | Expression of GFP-neoantigen and the preparation of BDVs

The recombinant cells were grown in LB medium until the OD<sub>600</sub> value reached 0.6, and then the expression of GFP-neoantigen was induced by addition of 1 mM isopropyl  $\beta$ -d-1-thiogalactopyranoside (IPTG) at 16°C for 6 h. After induction, recombinant cells stably expressing GFP-neoantigen were collected by centrifugation at 5000  $\times$  g for 10 min. The cells were resuspended in cold phosphate-buffered saline (PBS) and washed three times. Then we suspended the cells with cold HM lysis buffer (0.25 M sucrose, 1 mM Ethylene Diamine Tetraacetic Acid (EDTA), 20 mM HEPES-NaOH, pH7.4, and 1x protease inhibitor cocktail (Beyotime, 100x)). Afterwards, the mixed solution was transferred to the ultrasonic cell disruptor in ice bath for ultrasonic decomposition (power: 30%, time: 2 min). The entire solution was centrifuged at  $1000 \times g$  for 5 min in order to eliminate cells. Next, the supernatant was centrifuged at  $3000 \times g$  for 5 min. Carefully aspirate the supernatant to a new tube and then centrifuged at 15,000  $\times$  g for 0.5 h to collected the sediment. The sediment was resuspended in cold HM buffer, passing through 0.8  $\mu$ m and  $0.22 \,\mu$ m filters at least 10 times in turn to get the BDVs. The BCA protein assay kit (Beyotime) was used to quantitatively calculate the total protein concentration of BDVs. At last, The BDVs were stored in HM buffer at -80°C. Fluorescence images were taken on the confocal microscope (Zeiss) to verify the expression of GFP. The preparation of thermosensitive HP hydrogel (Hyaluronic Acid, 2 wt%, Macklin; Pluronic F-127, 25 wt%, Sigma Aldrich) was performed using a cold method as described by Ye et al (Ye et al., 2019). In brief, HA and Pluronic F-127 were mixed in cold distilled water and then stored in 4°C refrigerator until the polymer had completely dissolved. Murine GM-CSF (20 ng/ml, PeproTech) and BDVs ( $25 \mu g$ ) were then added in the 150  $\mu$ l HP hydrogel, and the mixture was stored at 4°C before use. In 37°C incubator, thermosensitive HP hydrogel could rapidly form the hydrogel and act as an embedding carrier for long-term drug release.

## 2.3 | Separation of outer and inner membrane

Two liters of recombinant cells were harvested by centrifugation at 4°C. The cells were washed with 10 mM Tris-(hydroxymethyl)aminomethane-hydrochloride (Tris-HCL buffer, pH 8.0) and resuspended in 20 ml of 20% (wt/vol) sucrose (50  $\mu$ g/ml Deoxyribonuclease I). Add 2 ml of 1.5 mg/ml lysozyme (final concentration of 150  $\mu$ g/ml) and 200  $\mu$ l 100x protease inhibitor cocktail to the resuspension. Afterwards, the mixed solution was transferred to the ultrasonic cell disruptor in ice bath for ultrasonic decomposition (power: 30%, time: 2 min). Cell debris was then removed by centrifugation at 2,000 x g for 10 min, and the supernatant was decanted to a new centrifuge tube. Eight milliliters were then layered onto a sucrose step gradient containing 8 ml of 70% (wt/vol) sucrose and 10 ml of 50% (wt/vol) sucrose in Tris-HCL buffer, and then the tubes were centrifuged at 184,500 × g in a Beckman Type 70i (70 Ti) rotor for 18 h. The two bands were observed and then remove the upper-brown inner membrane (IM) layer using the pipette by suction from above. Repeat the pipetting procedure for the outer membrane (OM) fraction. The individual samples were diluted in distilled water and collected by centrifuged at 184,500 × g in a Beckman 70 Ti rotor. This procedure is a modification of the method described by Hancock (Hancock & Nikaido, 1978) and Cian (Cian et al., 2020).

# 2.4 | Morphology and characterization

The recombinant *E. coli* cells were washed with PBS and fixed with 4% paraformaldehyde for 15 min. Then cells were stained with 4',6-diamidino-2-phenylindole (DAPI, Beyotime) for 15 min at room temperature and washed with PBS buffer. Fluorescence of GFP and DAPI were observed by confocal microscope (Zeiss) with 40× objective. The morphology of *E. coli* cells and BDVs, negatively stained with 1% uranyl acetate, were observed using transmission electron microscopy (TEM; Shiyanjia Lab (www. shiyanjia.com) for the TEM image). In addition, the size distribution of BDVs were measured by the dynamic light scattering (DLS, Nano-ZS, model MPT2;). The zeta potential of the BDVs were measured by Malvern Zetasizer (Malvern Instruments).

### 2.5 | Western blot

The recombinant protein of *E. coli* cells, BDVs, OM and IM were boiled with 1x loading buffer and separated on a 10% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) gel. Then the protein was transferred onto the polyvinylidene fluoride (PVDF) membrane, which was blocked with 5% skimmed milk (tris buffer solution tween (TBST, Tween 20, 0.1%) buffer) for 2 h at room temperature. The membrane was hybridized with EGFP (Abmart) or ompF (Thermo Fisher PA5-121442)



antibodies (1:1000) overnight at 4°C. Then the membrane was washed three times for about 15 min in TBST buffer and incubated with secondary antibody for 1 h at room temperature. After three washes for 5 min each with TBST, the imaging was performed using enhanced chemiluminescence detection (ECL, FDbio-Pico).

#### 2.6 | In vitro BMDCs isolation and uptake

BMDCs were isolated and collected from the bone marrow which was flushed from the tibias and femurs of female C57BL/6 mice (6-week-old). The harvested cells were resuspended in red blood cell (RBC) Lysis Buffer (Sigma-Aldrich) for 10 min to eliminate RBCs. The remaining cells were cultured in Roswell Park Memorial Institute (RPMI) 1640 supplemented with 10% fetal bovine serum (FBS), 1% penicillin-streptomycin, 10 ng/ml interleukin 4 (IL-4, PeproTech), and 20 ng/ml GM-CSF in 6-well plates at a concentration of  $1 \times 10^6$  cells. All of the floating cells were removed and replaced with fresh medium after 24 h. The nonadherent cells under the plate were recognized as bone marrow-derived dendritic cells. Engineered BDVs (25  $\mu$ g) presenting GFP were added into the plate to verify the phagocytosis by BMDCs. The cells were incubated with BDVs for 24 h, subsequently stained with wheat germ agglutinin Alexa Fluor 647 conjugate (WGA AF647, Invitrogen, 1833658) and DAPI and analyzed by confocal microscope (Zeiss, LSN880).

To study the activation of DCs, the cells were divided into four groups: PBS (G1), blank-BDVs (G2), Normal-M33-M47 BDVs (G3), Mutation-M33-M47 BDVs (G4). The BMDCs ( $1 \times 10^6$  cells/ml) were incubated with the different formulations (with same mass of BDVs, 25  $\mu$ g) for 12 h and then collected, washed three times with PBS. The cells were divided equally into three parts and stained with anti-CD11c-Allophycocyanin (APC), anti-CD40-Phycoerythrin (PE), anti-MHC II-PE, anti-CD80-fluorescein isothiocyanate (FITC) and anti-CD86-PE antibodies (Biolegend) according to the manufacturer's protocols. The stained cells were sorted using Beckman CytoFlex flow cytometer and analyzed with FlowJo software.

#### 2.7 | In vivo activation

Six-week-old female mice were randomly separated into five groups (n = 5) and subjected to different treatments: Gel (150  $\mu$ l, #1), Gel-GM-CSF (150  $\mu$ l, 20 ng/ml GM-CSF, #2), Gel-blank BDVs + GM-CSF (150  $\mu$ l, 25  $\mu$ g BDVs and 20 ng/ml GM-CSF, #3), Gel-Normal-M33-M47 BDVs + GM-CSF (150  $\mu$ l, 25  $\mu$ g BDVs, 20 ng/ml GM-CSF, #4), Gel-Mutation-M33-M47 BDVs + GM-CSF (150  $\mu$ l, 25  $\mu$ g BDVs and 20 ng/ml GM-CSF, #5). Different types of Gel-formulas were injected on the left side of the back via subcutaneous injection. To assess the recruitment and maturation of dendritic cells, the skin of the injection area and inguinal lymph nodes were respectively harvested after treatment for 72 h. The skin of mice was harvested and snap frozen in optimal cutting temperature (OCT) compound. The immunofluorescent assay of skin was carried out according to the description below.

On the other hand, the inguinal lymph nodes were collected from each group, and monocellular suspensions were prepared to assess the maturation of dendritic cell. Cells were stained with anti-CD11c-APC, anti-CD40-PE, anti-CD80-FITC and anti-CD86-PE antibodies (Biolegend). All samples were sorted using Beckman CytoFlex flow cytometer and analyzed with FlowJo software.

#### 2.8 | In vivo tumour model

Animal experiments were performed under an approved protocol by the Administrative Committee of Animal Research in the Sun Yat-Sen University (SYSU-IACUC-2021-000874). In order to study the anti-tumour therapeutic efficacy of BDVs in vivo, 6week-old female C57BL/6 mice in SPF grade were purchased from Guangdong Medical Laboratory Animal Center (Guangdong, China) and kept in SPF level Laboratory Animal Center of Sun Yat-sen University. Mice were cared and performed according to the instructions and approval of the Institutional Animal Care and Use Committee of Sun Yat-sen University. Mice were challenged with  $1 \times 10^{6}$  B16F10-luc cells on the right side of the back on day 0. After 7 days, approximately 1 mm<sup>3</sup> in volume of the tumour was remained to simulate the presence of the residual microtumour in the clinical operation. The wound was stitched up skillfully with three surgical suture nails. Metastasis mouse model were established by intravenous injection of 5 ×  $10^5$  B16F10-luc via the tail vein. The mice were separated randomly into six groups (n = 7 or 10): Gel-PBS (150  $\mu$ l, 20 ng/ml GM-CSF, #1), Gel-blank BDVs (150  $\mu$ l, 25  $\mu$ g BDVs and 20 ng/ml GM-CSF, #2), Gel-Normal-M33-M47 BDVs (150  $\mu$ l, 25  $\mu$ g BDVs, 20 ng/ml GM-CSF, #3), Gel-Mutation-M33-M47 BDVs (150  $\mu$ l, 25  $\mu$ g BDVs and 20 ng/ml GM-CSF, #4), aPD-1 (anti-PD-1 antibody, GoInVivo Purified anti-mouse CD279 (PD-1), Biolegend, 114115, 2.5 mg/kg, #5) and Gel-Mutation-M33-M47 BDVs + aPD-1 (150  $\mu$ l, 25  $\mu$ g BDVs and 20 ng/ml GM-CSF, 2.5 mg/kg, #6). Gel-formulations were injected on the left side of the back via subcutaneous injection, and aPD-1 was injected via the tail-vein. Mice were anesthetized with isoflurane and bioluminescence images of melanomas were captured by IVIS fluorescence imaging system based on XenoLight D-Luciferin-K<sup>+</sup> Salt Bioluminescent Substrate. The mice weight and tumour size were measured every other day. The tumour size was measured



by digital caliper, and the tumour volume (mm<sup>3</sup>) was calculated as follows: width<sup>2</sup> × length /2. When the tumour volume of tumour-bearing mouse exceeds 1500 mm<sup>3</sup>, the suffering mice should be humanely euthanized and the tumours and organs were collected. The numbers of lung metastatic nodules in different groups were counted carefully. The organs (heart, liver, spleen, kidney and lung) of surgical mice and lung tissue of lung tumours mice were stained with hematoxylin-eosin staining (H&E).

#### 2.9 | Tissue immunofluorescent assay

The analysis of tissue immunofluorescent was performed following the instructions. First of all, skin and tumours were collected and frozen in optimal cutting temperature (OCT) compound (SAKURA). The proper thickness section (8  $\mu$ m) was obtained through rotary freezing microtome. Secondly, the slices from different groups were washed with PBS buffer in order to clean off the embedding medium. Then 4% paraformaldehyde was used to fix the slices, whereafter, 3% bovine serum albumin (BSA, containing 0.2% Triton X-100) was applied to block the slices. Thirdly, the slices of skin were then incubated with CD11c primary antibody (1:200 in 1.5% BSA, Abcam, ab254183) overnight at 4°C. Whereas, the slices of tumour were incubated with CD4 (1:100 in 1.5% BSA, Abcam, ab183685) and CD8 primary antibody (1:100 in 1.5% BSA, Abcam, ab217344) under the same conditions. After washing 3 times with PBS buffer, the secondary antibodies (1:500 in 1.5% BSA, ThermoFisher) were incubated with skin or tumour specimens for about 2 h in the darkness. Finally, DAPI staining was used to locate the position of cell nucleus and ensure the depth of shooting field. Fluorescence images were observed on the confocal microscope (Zeiss, LSN880).

#### 2.10 | Statistical analysis

The data values were presented as the mean  $\pm$  s.d. or the mean  $\pm$  s.e.m.. Comparisons between the experimental groups were performed using the one-way analysis of variance (ANOVA) followed by the Tukey post-hoc tests. Comparisons of survival curves were performed using the Long-Rank (Mantel-Cox) test. All statistical analyses were performed using GraphPad Prism 8 software (\*P < 0.05, \*\*P < 0.01, and \*\*\*P < 0.001). Unless otherwise stated, all experiments were performed in biological replicates.

#### 3 | RESULTS

#### 3.1 | Preparation and characterization of Mutation-M33-M47 BDVs

OMVs are naturally released from the outer membrane of Gram-negative bacteria (Carvalho et al., 2019; Kuehn & Kesty, 2005; Lee et al., 2008; Rosenthal et al., 2014; Zanella et al., 2021). To synthesize fusion neoantigens and display the neoantigens on the outer membrane of the bacteria, we subcloned pET-21a vector containing two neoantigen nucleotide sequences of M33 and M47 that had been identified from B16F10 cells (Kreiter et al., 2015). The Lpp-OmpA-based display system behind the IPTG promoter were employed to make the fusion proteins localize on the outer membrane, while GFP was fused on the N-terminal of the neoantigen as tracking tag (Scheme 1). Then the plasmid was transformed into competent BL21(DE3)plysS cells. The transfected BL21(DE3)plysS cells were then induced by IPTG for 16 h, and were observed by confocal microscope. It showed that the Lpp-OmpA-GFP-M33-M47 fusion protein was successfully expressed and localized on the outer membrane in engineered *E. coli* (Figure 1a, b). The morphology of GFP-Mutation-M33-M47 *E. coli* was characterized by TEM (Figure 1c). The BDVs were prepared by ultrasonication and hypothermal differential centrifugation and then were filtered through  $0.8\,\mu\text{m}$  and  $0.22\,\mu\text{m}$  poresized hydrophilic membrane. Hereafter, we characterized the morphology of the negatively stained GFP-Mutation-M33-M47 E. coli BDVs by TEM and confirmed that the BDVs akin to OMVs (Figure 1d). We further validated that GFP-Mutation-M33-M47 fusion protein had been presented on BDVs by the green fluorescence signals of GFP (Figure 1e). In addition, the size distribution of BDVs were measured by the dynamic light scattering (DLS), which indicated that the average diameter of BDVs was ~120 nm (Figure 1f). The zeta potential of different formulation of BDVs was around -12 mV using Malvern Zetasizer Nano ZS ZETA (Figure 1g). The characterizations of BDVs were same as the cell membrane vesicles and natural OMVs (Li et al., 2021; Park et al., 2021; Xue et al., 2022). And on top of that, we also successfully extracted the inner and outer membrane of E. Coli (Figure S1). Analysis of the OM and IM proteins by SDS-polyacrylamide gel electrophoresis showed a little bit different (Figure S2). Ultimately, western blot analysis showed that the Lpp-OmpA-GFP-M33-M47 fusion protein and outer membrane protein F (ompF) were efficiently expressed in E. coli BL21(DE3)plysS cells, BDVs and OM, respectively (Figure 1h, i).





**FIGURE 1 Characterizations of GFP-Mutation-M33-M47** *E. coli* **BDVs**. (a) Establishment of *E. coli* stably expressing GFP (Scale bar:  $2 \mu m$ ). (b) Schematic diagram illustration the presentation of neoantigens using Lpp-OmpA system on *E. coli* outer membrane. (c) The TEM images shown the size and morphology of GFP-Mutation-M33-M47 *E. coli* (Scale bar:  $2 \mu m$ ). (d) The TEM images shown the size and morphology of GFP-Mutation-M33-M47 *E. coli* (Scale bar:  $2 \mu m$ ). (d) The TEM images shown the size and morphology of GFP-Mutation-M33-M47 *E. coli* BDVs (Scale bar: 100 nm). (e) The confocal image of GFP-Mutation-M33-M47 *E. coli* BDVs (Scale bar:  $5 \mu m$ ). (f) The size distribution of GFP-Mutation-M33-M47 *E. coli* BDVs measured by dynamic light scattering (DLS). (g) The zeta potential of the *E. coli* BDVs. Blank (BL21(DE3)plysS): non-heat shock transformation control, worked as a negative control. (h) Western blot analysis indicated the expression of GFP on *E. coli* and BDVs. (i) Western blot analysis indicated the expression of GFP and ompF on *E. coli*, BDVs, OM and IM

#### 3.2 | In vitro and in vivo immune response of mutation-M33-M47 BDVs

Dendritic cells (DCs) enable internalizing tumour antigen and priming T cells to generate tumour antigen-specific T cells (Banchereau & Palucka, 2005). As the well-equipped professional antigen-presenting cells, DCs play crucial roles in the adaptive immune response. BMDCs from female C57BL/6 mice were isolated to evaluate the endocytosis of the BDVs by antigen-presenting cells. To test whether the BDVs could be internalized by BMDCs in vitro, GFP-Mutation-M33-M47 BDVs were incubated with BMDCs at 37°C for 24 h. As shown in Figure 2a and Figure S3, BMDCs could efficiently internalize BDVs. Furthermore, the expression of costimulatory markers CD11c, CD40, MHC-II, CD80 and CD86 on BMDCs were employed to assess whether the BDVs could promote the maturation of BMDCs by flow cytometry. The cells received different treatments were collected after co-incubation for 24 h. Compared with PBS (G1), the cells received blank BDVs (G2), Normal-M33-M47 BDVs (G3), or Mutation-M33-M47 BDVs (G4) all showed a higher proportion of mature DCs, demonstrating that BDVs could effectively promote the maturation of DCs (Figure 2b, c and Figures S4, S5).

In order to assess the immune stimulation capacity of BDVs vaccines, the mice were subcutaneously inoculated with Gel (#1), Gel-GM-CSF (#2), Gel-blank-BDVs + GM-CSF (#3), Gel-Normal-M33-M47 BDVs + GM-CSF (#4), or Gel-Mutation-M33-M47 BDVs + GM-CSF (#5), respectively. After inoculation, the activation extent of the immune system was investigated with skin and inguinal lymph nodes. Flow cytometry analysis data showed that neither Gel (#1) nor Gel-GM-CSF (#2) could induce the maturation and activation of DCs effectively. On the contrary, BDVs could significantly induce the maturation and activation of DCs. The activation markers of DCs including CD11c<sup>+</sup> CD40<sup>+</sup> and CD80<sup>+</sup> CD86<sup>+</sup> were intensively increased in the mature



7 of 14

**FIGURE 2** Dendritic cell activity and maturation in vitro and in vivo. (a) Confocal 3D image of BMDCs uptake of GFP-Mutation-M33-M47 E. coli BDVs in vitro (Scale bar: 10  $\mu$ m). Green: BDVs. Red: cell membrane. Blue: cell nucleus. (b, c) Quantitative analysis of CD11c<sup>+</sup> CD40<sup>+</sup> cells (b) and mature DCs (c) from different treatment groups in vitro (gated on CD11c<sup>+</sup> cells, n = 5). Error bar, mean  $\pm$  s.d.. PBS (G1), Blank BDVs (G2), Normal-M33-M47 BDVs (G3), Mutation-M33-M47 BDVs (G4). (d, e) Representative flow cytometry data (d) and statistical data (e) to show CD11c<sup>+</sup> CD40<sup>+</sup> cells induced by different formulations of BDVs in vivo (in lymph nodes) on day 3 post- injection (gated on CD11c<sup>+</sup> cells, n = 5). Cells were stained with anti-CD11c-APC, anti-CD40-PE antibodies (Biolegend). (f, g) Representative flow cytometry data (f) and statistical data (g) to show DC maturation induced by different formulations of BDVs in vivo (in lymph nodes) on day 3 post-injection (gated on CD11c<sup>+</sup> cells, n = 5). Cells were stained with anti-CD10c-APC, anti-CD40-PE antibodies (Biolegend). (f, g) Representative flow cytometry data (f) and statistical data (g) to show DC maturation induced by different formulations of BDVs in vivo (in lymph nodes) on day 3 post-injection (gated on CD11c<sup>+</sup> cells, n = 5). Cells were stained with anti-CD10c-APC, anti-CD80-FITC, anti-CD80-PE antibodies (Biolegend). Error bar, mean  $\pm$  s.d.. (h) Immunofluorescence analysis of CD11c<sup>+</sup> DCs in the skin after the treatments (Scale bar: 50  $\mu$ m). Gel (#1), Gel-GM-CSF (#2), Gel-Blank BDVs + GM-CSF (#3), Gel-Normal-M33-M47 BDVs + GM-CSF (#4), Mutation-M33-M47 BDVs + GM-CSF (#5). mean  $\pm$  s.d., n = 5. NS: no significant, \**P* < 0.05, \*\**P* < 0.01, \*\*\**P* < 0.001. One-way ANOVA with Tukey post-hoc tests (b, c, e, g)

DCs in lymph nodes, indicating BDVs could promote the maturation of DCs efficiently in vivo (Figure 2d–g). Moreover, the mice inoculated with Gel-Mutation-M33-M47 BDVs plus GM-CSF significantly enhanced the maturation of DCs compared to those receiving Gel-Normal-M33-M47 BDVs plus GM-CSF. These data indicated that BDVs with neoantigens could promote the antigen recognition and maturation of DCs, while GM-CSF facilitated the recruitment and stimulation of the clone expansion of DCs. Next, to assess the recruitment of DCs, the skin of the injected area in experimental group was harvested and analyzed by immunofluorescence. Similar to the results of FACS analysis, the density of CD11c<sup>+</sup> cells with immunofluorescence staining in skin was obviously higher in the Gel-BDVs treatment groups than in the Gel-PBS group (Figure 2h). Taken together, these data indicated that injection of BDVs with GM-CSF could trigger a robust immune response both in vitro and in vivo.

# 3.3 | In vivo therapeutic efficacy of mutation-M33-M47 BDVs combined with PD-1 blockade for anti-tumour relapse

SEV

8 of 14

Neoantigens were identified by exome sequencing of the individual surgical tumour tissue (Kreiter et al., 2015). Thus, the tumour vaccines leveraging neoantigen could be potent for protecting individual from tumour relapes and tumour metastasis (Blass & Ott, 2021). To evaluate whether BDVs-neoantigen vaccine could prevent tumour relapse after the surgery, B16F10-tumour <del>xenograft</del> mouse model was employed in the <del>present</del> experiments. Majority of the tumour (~90%) was surgically removed on day 8, and the mice were randomly assigned into 6 different treatment groups: Gel-PBS (#1), Gel-blank-BDVs (#2), Gel-Normal-M33-M47 BDVs (#3), Gel-Mutation-M33-M47 BDVs (#4), aPD-1 (2.5 mg/kg, #5), Gel-Mutation-M33-M47 BDVs + aPD-1 (2.5 mg/kg, #6) (Figure 3a). Tumour growth were monitored by bioluminescence imaging and tumour volume (Figure 3b, c). It was worth noting that mice treated with Gel-Mutation-M33-M47 BDVs combined aPD-1 (#6) showed the best inhibition of tumour growth compared with the mice received other treatments. Meanwhile, mice treated with Gel-Mutation-M33-M47 BDVs (#4) and aPD-1 (#5) also exhibited better ability of antitumour response than Gel-PBS (#1), Gel-blank-BDVs (#2) and Gel-Normal-M33-M47 BDVs (#3), indicating that the neoanitgens and PD-1 antibody both contributed to therapeutic effect in repressing tumour growth. According to Figure 3d-f, mice treated with Gel-Mutation-M33-M47 BDVs combined aPD-1 (#6) not only exhibited the strongest anti-tumour relapse efficacy, but also benifited higher survival rate. In contrast, mice in other groups all died within 36 days postchallenge (Figure 3f). Additionally, the mice had no obvious weight changes throughout the treatment (Figure S6). Meanwhile, the major organs of the mice, including heart, liver, spleen, kidney and lung were harvested and performed the H&E staining, and no abnormal phenomenon were observed, indicating that the BDVs formulations had no toxic to mice (Figure S7). These data indicated that BDVs-neoantigen vaccines combined with PD-1 antibody intensively inreased the number and activity of CD8<sup>+</sup> CTLs, and thus improved the efficacy of cancer immunotherapy.

To further explore the mechanism and synergistic therapeutic effect of the Gel-Mutation-M33-M47 BDVs in combination with aPD-1, immunofluorescence and flow cytometry were used to evaluate the situation of tumour-infiltrating lymphocytes (TILs) in the relapsed tumours. T lymphocyte is the main effector cell of anti-tumour immunity (Farhood et al., 2019; McLane et al., 2019), thus the relapsed tumour infiltrated CD4<sup>+</sup> and CD8<sup>+</sup> T cells infiltrating into the relapsed tumours were analysed by flow cytometry. Notably, the number of CD4<sup>+</sup> and CD8<sup>+</sup> T cells were significantly increased in mice that were treated with the Gel-Mutation-M33-M47 BDVs combined with aPD-1 (Figure 4a, b and Figure 58). Moreover, the proliferation of CD8<sup>+</sup> T cells in the relapsed tumours of mice treated with Gel-Mutation-M33-M47 BDVs combined with aPD-1 (#6) were significantly increased (Figure 4c, d). Gel-Mutation-M33-M47 BDVs combined with aPD-1 treatment could activate the native CD8<sup>+</sup> T cells in vivo. CD69 is one of the markers of T lymphocytes activation, therefore the number of active CD8<sup>+</sup> CD69<sup>+</sup> T cells were significantly increased in the FACS detection (Figure 4e, f). In addition, to further assess the mechanism of the anti-tumour effect of Gel-BDV formulations, we also examined the central memory CD8<sup>+</sup> T ( $T_{CM}$ ) cells. Virtually, the population of CD44<sup>+</sup> CD62L<sup>+</sup>  $T_{CM}$  cells showed a intensive increase in the Gel-Mutation-M33-M47 BDVs with aPD-1-treated mice compared with the PBS control (Figure 4g, h), indicating BDVs combination therapy could generate a strong immune memory effect to protect mice from tumour recurrence. Importantly, we found the proliferation of CD4<sup>+</sup> T cells in relapsed tumours of mice treated with Gel-Mutation-M33-M47 BDVs with aPD-1 was also significantly increased (Figure S9a, b). Nonetheless, the number of regulatory T cells Tregs (CD4<sup>+</sup> CD25<sup>+</sup> Foxp3<sup>+</sup> T cells) was decreased (Figure S9c, d).

Cytotoxic proteins such as granzyme B (Gzm B) and perforin can directly kill tumour cells by activating caspase enzymes mediated apoptosis (Russell & Ley, 2002; Voskoboinik et al., 2015). Thus, we detected the expressions of GzmB in the T cells after the treatment. Flow cytometry data showed that there was a intensive increase of Gzm B<sup>+</sup> CD8<sup>+</sup> T cells and perforin<sup>+</sup> CD8<sup>+</sup> T cells in the mice treated with Gel-Mutation-M33-M47 BDVs combined with aPD-1 (Figure S10a–d). Eventually, immunostaining of the CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the tumour tissue section showed similar results compared with FACS analysis, and the number of CD8<sup>+</sup> T cells significantly increased in the margin of the tumour section (Figure 4i). Collectively, these results indicated that the Gel-Mutation-M33-M47 BDVs combined with aPD-1 blockade could trigger anti-tumour immune response and effectively prevent tumour relapse.

#### 3.4 | In vivo therapeutic efficacy of engineered BDVs for metastatic tumours

Subsequently, we investigated whether the BDVs-neoantigen vaccine could prevent tumour metastasis. We established a metastatic mice mouse model by *i.v.* injection of B16F10-luc cells ( $\sim$ 1 × 10<sup>6</sup>) to mimic the metastasis of cancer cells. The mice were randomly devided into six experimental groups and received the same treatment as described above (Figure 5a). The growth and spread of tumours were monitored by bioluminescence of B16F10-luc cells. As shown in Figure 5b, mice treated with PBS showed obvious cancer metastasis and all of them died within 16 days. On the contrary, Gel-Mutation-M33-M47 BDVs combined with aPD-1 showed the relatively weak bioluminescence signal and exhibited much better anti-tumour ability. Notably, the typical lung metastatic tumour images showed that mice received Gel-Mutation-M33-M47 BDVs combined with aPD-1 treatment had



9 of 14

**FIGURE 3** Antitumour recurrence effect of BDVs-triggered cancer immunotherapy in the surgical bed of B16F10-luc model. (a) Schematic illustration of BDVs-triggered cancer vaccine in an incomplete-surgery B16F10-luc tumour model. (b) In vivo bioluminescence images of the B16F10-luc tumour-bearing mice (n = 5). (c) Single tumour volume of each mouse in different groups. (d) Tumour growth curves in B16F10-luc model (n = 7), Error bar, mean  $\pm$  s.e.m.. (e) Tumour weights in different treatment groups (n = 7), Error bar, mean  $\pm$  s.d.. (f) Survival curves in different groups (n = 10) (#1) Gel-PBS, (#2) Gel-Blank BDVs, (#3) Gel-Normal-M33-M47 BDVs, (#4) Gel-Mutation-M33-M47 BDVs, (#5) aPD-1 (anti-PD-1 antibody), (#6) Gel-Mutation-M33-M47 BDVs + aPD-1. NS: no significant, \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001. One-way ANOVA with Tukey post-hoc tests (d, e) or the Long-Rank (Mantel-Cox) test (f)

few metastatic tumour nodules and achieved promising anti-tumour effect in the lung metastatic model (Figure 5c). Hereafter, all lungs harvested from the mice received treatments were fixed with 4% paraformaldehyde, and numbers of metastatic nodules were counted at the end time. The results revealed that the mice treated with Gel-Mutation-M33-M47 BDVs, aPD-1 or Gel-Mutation-M33-M47 BDVs combined with aPD-1 had few metastatic nodules compared to those received other formulations treatment (Figure 5d). Moreover, the mice treated with Gel-Mutation-M33-M47 BDVs combined with aPD-1 had few metastatic nodules compared to those received other formulations treatment (Figure 5d). Moreover, the mice treated with Gel-Mutation-M33-M47 BDVs combined with aPD-1 also had a higher proportion (~70%) of survical rate after tumour injection (Figure 5e). Virtually, H&E staining results showed that mice with



FIGURE 4 BDV-neoantigen vaccine induced a robust antitumour immune response. (a, b) Representative flow cytometry plots (a) and ratios (b) of different groups of T cells in the residual tumours from different groups (gated on  $CD3^{+}$  T cells, n = 5). Cells were stained with anti-CD3-FITC, anti-CD4-Brilliant Violet 421, anti-CD8a-APC/Fire750 antibodies (Biolegend). Error bar, mean ± s.d., (c, d) Representative flow cytometry plots (c) and ratios (d) of different groups of CD8<sup>+</sup> Ki67<sup>+</sup> T cells in tumours (gated on CD3<sup>+</sup> CD8<sup>+</sup> T cells, n = 5). Cells were stained with anti-CD3-FITC, anti-CD8-APC/Fire750, anti-Ki67-Alexa Fluor<sup>®</sup> 647 antibodies (Biolegend). Error bar, mean  $\pm$  s.d.. (e, f) Representative flow cytometry plots (e) and ratios (f) of different groups of CD8<sup>+</sup> CD69<sup>+</sup> T cells in tumours (Gated on CD3<sup>+</sup> CD8<sup>+</sup> T cells, *n* = 5). Cells were stained with anti-CD3-FITC, anti-CD8-APC/Fire750, anti-CD69-APC antibodies (Biolegend). Error bar, mean ± s.d., (g, h) Representative flow cytometry plots (g) and ratios (h) of different groups of CD44<sup>+</sup> CD62L<sup>+</sup> T cells infiltrating in tumours (Gated on CD3<sup>+</sup> CD8<sup>+</sup> T cells, n = 5). Cells were stained with anti-CD3-FITC, anti-CD8-APC/Fire750, anti-CD44-PE, anti-CD62L-Alexa Fluor<sup>®</sup> 700 antibodies (Biolegend). Error bar, mean  $\pm$  s.d.. (i) The immunofluorescence images of CD4<sup>+</sup> CD8<sup>+</sup> T cells infiltrating in tumours (Scale bar: 50 μm). (#1) Gel-PBS, (#2) Gel-Blank BDVs, (#3) Gel-Normal-M33-M47 BDVs, (#4) Gel-Mutation-M33-M47 BDVs, (#5) aPD-1, (#6) Gel-Mutation-M33-M47 BDVs + aPD-1. Cells were stained with anti-CD11c-APC, anti-CD40-PE, anti-CD80-FITC and anti-CD86-PE antibodies (Biolegend). All samples were sorted using Beckman CytoFlex flow cytometer and analyzed with FlowJo software. NS: no significant, \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001. One-way ANOVA with Tukey post-hoc tests (b, d, f, h)



11 of 14

**FIGURE 5** *In vivo* suppression of antimetastatic tumour effect of BDV-neoantigen vaccine. (a) Schematic illustration of the therapy of BDV-neoantigen vaccine in B16F10-luc model of metastasis. (b) In vivo bioluminescence images of the B16F10-luc lung metastasis in different groups (n = 5). (c) Representative images of the lung metastatic nodules (Scale bar: 5 mm). (d) Numbers of lung metastatic nodules in different groups (n = 5), Error bar, mean  $\pm$  s.d.. (e) Survival curves of different groups (n = 10). Day 0 is the day of tumour cells injection via tail vein. (f) H&E-stained lung slices (Scale bar: 100  $\mu$ m). (#1) Gel-PBS, (#2) Gel-Blank BDVs, (#3) Gel-Normal-M33-M47 BDVs, (#4) Gel-Mutation-M33-M47 BDVs, (#5) aPD-1, (#6) Gel-Mutation-M33-M47 BDVs + aPD-1. NS: no significant, \*P < 0.05, \*\*P < 0.01, \*\*\*P < 0.001. One-way ANOVA with Tukey post-hoc tests (d) or the Long-Rank (Mantel-Cox) test (e)

Gel-Mutation-M33-M47 BDVs, aPD-1 and Gel-Mutation-M33-M47 BDVs combined with aPD-1 treatments had less lung metastases, indicating both Gel-Mutation-M33-M47 BDVs and aPD-1 had conspicuously better effect in delaying the development and progress of lung metastasis (Figure 5f).

### 4 | DISSCUSSION

In summary, we genetically programed bacteria to synthesize fusion neoantigens, and prepared BDVs displaying neoantigen (BDVs-Neo) as personalized therapeutic vaccine to elicit systemic anti-tumour immunity. BDVs-Neo and GM-CSF loaded within thermosensitive hydrogel could recruit and stimulate the maturature of DCs within subcutaneous. Notably, BDVs also



**SCHEME 1** Schematic of preparation of Gel-GFP-Mutation-M33-M47 BDVs combined with PD-1 antibody for immunotherapy of melanoma. (a) Preparation process of Gel-BDVs. (b) Effector T cell were actived by Gel-GFP-Mutation-M33-M47 BDVs to against melanoma

exhibited spherical structures the same as exosome and showed similar diameter as natural OMVs (Chen, Bai et al., 2020; Li et al., 2022). Moreover, BDVs-Neo not only could be easily internalized into BMDCs in vitro, but also stimulated DC maturation efficiently in vivo. BDVs-Neo vaccine formulation could improve the anticancer efficacy and long-term memory immunity through combination treatment with PD-1 antibody. CTLs could recognize and eliminate residual tumour cells through cytotoxic effects and the secretion of cytokines, such as IFN- $\gamma$ . Flow cytometry and immunofluorescence analysis of tumour tissue showed that BDVs-Neo vaccine could improve CD8<sup>+</sup> T cell activity. And beyond that, BDVs-Neo vaccine also increased the number of central memory T cells, which protected the mice from tumour relapse and metasitasis. BDVs-Neo vaccine formulation improved the anti-cancer efficacy and long-term memory immunity through combination treatment. Collectively, BDVs-Neo is a promising personalized cancer vaccine formulation that could induce robust anti-tumour response.

### **AUTHOR CONTRIBUTIONS**

X.Z. and X.L. contributed conception and design of the study. F.M., L.L., Z.Z., Z. L. and J.Z. performed the experiments. F.M., X.S., T.X. and C.X. interpreted the data. Z. L. assisted with some parts of experiments and data analysis. F.M. wrote the first draft of the manuscript. The manuscript was written through contributions of all authors. All authors have given approval to the final version of the manuscript.

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#### CONFLICT OF INTERESTS

The authors declare that no competing interests exist.

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- 14 of 14 | 0 | SEV \_\_\_\_\_
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#### SUPPORTING INFORMATION

Additional supporting information can be found online in the Supporting Information section at the end of this article.

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