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TITLE:

Tumor Transplantation for Assessing the Dynamics of Tumor-infiltrating CD8⁺ T Cells in Mice

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SUMMARY:

Here, we present a tumor transplantation protocol for the characterization of tumor-inherent and periphery-derived tumor-infiltrated lymphocytes in a mouse tumor model. Specific tracing of the influx of recipient-derived immune cells with flow cytometry reveals the dynamics of the phenotypic and functional changes of these cells during antitumor immune responses.

ABSTRACT:

T cell-mediated immunity plays a crucial role in immune responses against tumors, with cytotoxic T lymphocytes (CTLs) playing the leading role in eradicating cancerous cells. However, the origins and replenishment of tumor antigen-specific CD8⁺ T cells within the tumor microenvironment (TME) remain obscure. This protocol employs the B16F10-OVA melanoma cell line, which stably expresses the surrogate neoantigen, ovalbumin (OVA), and TCR transgenic OT-I mice, in which over 90% of CD8⁺ T cells specifically recognize the OVA-derived peptide OVA_{257–264} (SIINFEKL) bound to the class I major histocompatibility complex (MHC) molecule H2-K^b. These features enable the study of antigen-specific T cell responses during tumorigenesis.

Combining this model with tumor transplantation surgery, tumor tissues from donors were transplanted into tumor-matched syngeneic recipient mice to precisely trace the influx of recipient-derived immune cells into transplanted donor tissues, allowing the analysis of the immune responses of tumor-inherent and periphery-originated antigen-specific CD8⁺ T cells. A dynamic transition was found to occur between these two populations. Collectively, this experimental design has provided another approach to precisely investigate the immune responses of CD8⁺ T cells in TME, which will shed new light on tumor immunology.

INTRODUCTION:

CD8⁺ T cell-mediated immune response plays a pivotal role in controlling tumor growth. During tumorigenesis, naive CD8⁺ T cells get activated upon antigen recognition in an MHC class I-restricted manner and subsequently differentiate into effector cells and infiltrate into tumor mass^{1,2}. However, within the tumor microenvironment (TME), prolonged antigen exposure, as well as immunosuppressive factors, drive infiltrated tumor-specific CD8⁺ T cells into a hyporesponsive state known as “exhaustion”³. Exhausted T cells (Tex) are distinct from effector or memory T cells generated in acute viral infection, both transcriptionally and epigenetically. These Tex cells are mainly characterized by the sustained and elevated expression of a series of inhibitory receptors as well as the hierarchical loss of effector functions. Further, the impaired proliferative capacity of exhausted CD8⁺ T cells results in decreasing numbers of tumor-specific T cells, such that the residual CD8⁺ T cells within the TME can barely provide sufficient protective immunity against tumor progression³. Thus, the maintenance or reinforcement of intratumoral antigen-specific CD8⁺ T cells is indispensable for tumor repression.

Moreover, immune checkpoint blockade (ICB) therapy is believed to reinvigorate Tex in tumors by increasing T cell infiltration and hence, T cell numbers and rejuvenating T cell functions to boost tumor repression. The widespread application of ICB treatment has changed the cancer therapy landscape, with a substantial subset of patients experiencing durable responses^{4–6}. Nevertheless, the majority of patients and cancer types do not or only temporarily respond to ICB. Inadequate T cell infiltration in the TME has been postulated to be one of the underlying mechanisms accounting for ICB resistance^{7,8}.

Several studies have demonstrated the heterogeneity of tumor-infiltrating CD8⁺ T cells (TILs) in both patients and mouse models^{9–12}. It has been confirmed that a subset of CD8⁺ T cells expressing T cell factor-1 (TCF1) in a tumor mass exhibits stem cell-like properties, which could further give rise to terminally exhausted T cells and is responsible for the proliferation burst after ICB therapy^{12–22}. However, it has been proved that only a small proportion of

antigen-specific TCF1⁺CD8⁺ T cells exist in the TME and generate an expanded pool of differentiated progeny in response to ICB²³⁻²⁶. Whether the limited size of this population is enough to ensure the persistence of cytotoxic T lymphocytes (CTLs) to control tumor progression remains unknown, and whether there is replenishment from periphery tissues requires further investigation. Furthermore, recent research suggests the insufficient reinvigoration capacity of pre-existing tumor-specific T cells and the appearance of novel, previously non-existing clonotypes after anti-programmed cell death protein 1 treatment. This indicates that T cell response to checkpoint blockade may be due to the new influx of a distinct repertoire of T cell clones²⁷. Together with the presence of bystander non-tumor-reactive cytotoxic T cell fraction in the TME, these findings prompted the establishment of a tumor allograft model to study the role of periphery-derived CD8⁺ T cells¹¹.

Until now, several kinds of tumor implantation, as well as immune cell adoptive transfer, have been widely used in the field of tumor immunology²⁸. TILs, peripheral blood mononuclear cells, and tumor-reactive immune cells originated from other tissues can be well-characterized using these methods. However, when studying the interactions between systemic and local antitumor immunity, these models appear inadequate to examine the interactions between immune cells derived from the periphery and the TME. Here, tumor tissues were transplanted from donors into tumor-matched recipient mice to precisely trace the influx of recipient-derived immune cells and observe the donor-derived cells in the TME concomitantly.

In this study, a murine syngeneic model of melanoma was established with the B16F10-OVA melanoma cell line, which stably expresses the surrogate neoantigen ovalbumin. TCR transgenic OT-I mice, in which over 90% of CD8⁺ T cells specifically recognize the OVA-derived peptide OVA₂₅₇₋₂₆₄ (SIINFEKL) bound to the class I MHC molecule H2-K^b, enable the study of antigen-specific T cell responses developed in the B16F10-OVA tumor model. Combining this model with tumor transplantation, the immune responses of tumor-inherent and periphery-originated antigen-specific CD8⁺ T cells were compared to reveal a dynamic transition between these two populations. Collectively, this experimental design has provided another approach to precisely investigate the immune responses of CD8⁺ T cells in the TME, which sheds new light on the dynamics of tumor-specific T cell immune responses in the TME.

PROTOCOL:

All mouse experiments were performed in compliance with the guidelines of the Institutional Animal Care and Use Committees of the Third Military Medical University. Use 6–8-week-old C57BL/6 mice and naïve OT-I transgenic mice weighing 18–22 g. Use both male and female without randomization or “blinding.”

1. Preparation of medium and reagents

1.1. Prepare cell culture medium D10 as previously described²⁹ by adding 10% fetal bovine serum (FBS), 100 U/mL penicillin, 100 mg/mL streptomycin, and 2 mM L-glutamine into Dulbecco's Modified Eagle Medium.

1.2. Prepare cell culture medium R10 by supplementing RPMI-1640 with 10% FBS, 100 U/mL

penicillin, and 100 mg/mL streptomycin.

NOTE: The culture media, D10 and R10, can remain sterile and stable for at least 2 weeks when stored at 2–4 °C.

1.3. Prepare Fluorescence-Activated Cell Sorting (FACS) buffer by supplementing 1x phosphate-buffered saline (PBS) with 2% FBS and 0.01% of sodium azide.

NOTE: With the addition of sodium azide, FACS Buffer can be stored at 2–4 °C for months.

1.4. Prepare red blood cell lysis (RBL) buffer by adding 155 mM NH₄Cl, 10 mM KHCO₃, and 0.1 mM ethylenediamine tetraacetic acid (EDTA) into double-distilled water, and adjust its pH to 7.3.

NOTE: RBL buffer is stable for up to 3 months at room temperature (RT).

1.5. Prepare magnetic-activated cell sorting (MACS) buffer by supplementing PBS with 0.5% bovine serum albumin (BSA) and 2 mM EDTA.

NOTE: The solution should be passed through a 0.22 µm filter after the reagent is dissolved and preserved in asepsis.

1.6. Prepare a working solution of 2,2,2-tribromoethanol.

1.6.1. Dissolve 2.5 g of 2,2,2-tribromoethanol in 5 mL of *tert*-amyl alcohol (2-methyl-2-butanol). Stir in a vapor-bathing, constant-temperature vibrator at 180 rpm, 40 °C overnight.

1.6.2. Filter the solution through a 0.22 µm filter into a sterile container. Add double-distilled water up to a final volume of 200 mL, and mix thoroughly and continuously until the solution becomes clear and transparent.

1.6.3. Completely wrap the container with aluminum foil to exclude light and store at 4 °C.

NOTE: The final concentration of the working solution of 2,2,2-tribromoethanol is 12.5 mg/mL. A more concentrated solution is not recommended because the material is irritating at higher concentrations.

2. Preparation of B16F10-OVA cell suspension

NOTE: Cell culture should be carried out in a biosafety hood under strict aseptic conditions.

2.1. Thaw and culture a vial of B16F10-OVA cells with D10 in a cell culture incubator at 37 °C and 5% CO₂.

2.2. When the cells reach the confluency of about 80–90%, subculture the cells.

2.2.1. Remove the culture medium with a pipettor, and rinse the cells twice using PBS.

NOTE: Do not add PBS forcefully against the adherent cells in the flask or cell culture dish. Instead, pipette the PBS toward a sidewall or add it drop-wise into the flask or dish.

2.2.2. Remove the PBS, and add 1–2 mL of 0.25% trypsin-EDTA solution into the flask or dish. Rock it back and forth to cover the entire cell surface. Place the flask or dish in an incubator at 37 °C for ~1 min or at RT until the cells detach.

NOTE: An inverted microscope can be used to check whether the cells have detached.

2.2.3. Add fresh D10 to stop the trypsinization. Pipette the suspension up and down to ensure that all the cells are dissociated from the flask or dish surface.

2.2.4. Transfer the B16F10-OVA cell suspension into a 15 mL conical tube. Centrifuge the cells at $125 \times g$ for 5–7 min at RT.

2.2.5. Discard the supernatant, and resuspend the cell pellet with D10. Dispense the B16F10-OVA cell suspension into a new flask or cell culture dish containing D10 and incubate in a cell culture incubator at 37 °C and 5% CO₂.

2.3. On the day of the tumor implantation, harvest B16F10-OVA cells that are ~90% confluent as described in steps 2.2.1 to 2.2.4. Discard the supernatant, and resuspend the cell pellet with 1 mL of PBS.

2.4. Count the viable cells with a hemocytometer using 0.4% trypan blue. Adjust the cell density to 1×10^6 cells per 100 μ L by adding PBS. Keep the cells on ice.

3. Ectopic tumor implantation of B16F10-OVA cells in the inguinal region of mice

3.1. Use 6–8-week-old C57BL/6 mice weighing 18–22 g. Use both male and female without randomization or “blinding.”

3.2. Withdraw 100 μ L of the prepared B16F10-OVA cell suspension into a 1 mL tuberculin syringe. Tap the barrel to move any bubbles to the top, and gently push the plunger to remove air bubbles.

3.3. Restrain the mouse and expose its abdomen. Press the left hind leg with the little finger to tighten the skin of the left inguinal region.

3.4. Remove the mouse’s hair from its left lower abdomen with an electric shaver. Use cotton soaked in 75% ethanol to clean the posterior quadrant of the left abdomen.

3.5. Holding the syringe at a very shallow angle (0–15°) with the bevel of the needle facing upwards, insert it at the site of the left upper thigh, and advance 0.5–1 cm through the subcutaneous tissue into the inguinal region.

3.6. Pull back on the plunger prior to injection. If there is negative pressure, depress the

plunger entirely, and observe a small bolus (formation of fluid pocket) in the subcutis emerge.

NOTE: If blood is drawn back into the needle hub, withdraw and try again at another site.

3.7. Remove the needle after the injection is carried out and dispose of it appropriately. Release and place the mouse back into the cage.

3.8. Measure tumor size on days 6–8 using a vernier scale after B16F10-OVA implantation. Select mice with a ~3 mm diameter (mung bean-sized) tumor and divide them equally and randomly into two groups.

NOTE: Mice with tumors of similar size are randomly assigned as donor and recipient mice; the matched tumor tissue excised from donor mice will be transplanted into the recipient mice. Furthermore, non-operated controls and sham-operated controls should be included to evaluate the effects of surgery on adoptive cell transfer and on the general health of mice. Thus, one group of tumor-bearing mice serves as non-operated controls, receiving either CD45.1⁺CD45.2⁺ or CD45.1⁺ OT-I cells but no surgery. The other group of mice serves as sham-operated controls, receiving either CD45.1⁺CD45.2⁺ or CD45.1⁺ OT-I cells and subsequent surgery similar to the experimental group but no allograft transplantation.

4. Adoptive transfer of congenically marked OT-I T cells into tumor-bearing mice

4.1. On the day before the transfer, administer 4 mg of cyclophosphamide dissolved in 200 µL of PBS via intraperitoneal injection to each tumor-bearing mouse.

NOTE: Treatment with cyclophosphamide aims to induce lymphopenia in the host that produces “space” for transferred cells, promoting their survival and homing to lymphoid organs to function efficiently.

4.2. Use naïve OT-I transgenic mice with distinct congenic markers (6–8-week-old, 18–22 g, the same sex as the tumor-bearing mice). Use CD45.1⁺ OT-I mice and CD45.1⁺CD45.2⁺ OT-I mice to adoptively transfer OVA₂₅₇₋₂₆₄ antigen-specific T cells into tumor-bearing donor and recipient mice, respectively.

NOTE: The origin of adoptively transferred OT-I cells can be easily identified if they display distinct congenic or fluorescent markers. For instance, inject CD45.1⁺ OT-I T cells into B16F10-OVA-bearing donor mice while injecting CD45.1⁺CD45.2⁺ OT-I T cells into B16F10-OVA-bearing recipient mice. CD45.1 and CD45.2 are both isoforms of the pan-lymphocyte marker CD45 (Ly5). Other commonly used congenic markers include different isoforms of CD90 (Thy1). This protocol can be used for mice carrying different congenic markers. OT-I mice should be of the same sex as the mice receiving OT-I cell transfer to avoid rejection issues.

4.3. Isolate the lymphocytes from the spleen and lymph nodes of the OT-I mouse.

NOTE: The following procedures in this step must be performed in a biosafety cabinet to maintain strict asepsis.

4.3.1. Prepare two 60 mm × 10 mm Petri dishes. Add 3 mL of R10 medium into one dish while adding 3 mL of RBL buffer into another dish. Place a 70 µm nylon cell strainer in the dish containing RBL buffer.

4.3.2. Euthanize an OT-I mouse in an isoflurane chamber followed by cervical dislocation.

4.3.3. Harvest the spleen, inguinal (subiliac), and axillary lymph nodes of the mouse and transfer them to a 60 mm × 10 mm dish with 3 mL of R10 on ice.

NOTE: The number of OT-I mice sacrificed may be adjusted depending on the number of tumor-bearing mice to be transferred. A typical yield from a spleen and bilateral inguinal and axillary lymph nodes of OT-I CD8⁺ T cells is ~30-100 × 10⁶ cells per mouse.

4.3.4. Using the end barrel of a 1 mL syringe, macerate the spleen in 3 mL of RBL buffer through the strainer. Incubate for 3 min at RT, and terminate the reaction by adding 3 mL of cold R10 medium.

4.3.5. Mash the lymph nodes until only connective tissues remain. Rinse the filter with R10. Transfer the cell suspension into a new 15 mL conical tube. Centrifuge at 500 × *g*, 4 °C for 6 min.

4.3.6. Decant the supernatant, and resuspend the cells in 3 mL of MACS buffer. Pass the cell suspension through a new 70 µm cell strainer to remove any flocs.

4.3.7. Centrifuge the cell suspension at 500 × *g* for 5 min at 4 °C. Decant the supernatant.

4.3.8. Use a mouse CD8⁺ T cell isolation kit (see the **Table of Materials**) to purify CD8⁺ T cells by negative selection, as per the manufacturer's protocol.

NOTE: When using kits from other companies, follow the manufacturer's instructions.

4.3.9. Keep the purified cell suspension on ice. Take a small sample of cells and mix with trypan blue to count cells using a hemocytometer.

4.4. Determine the percentage of OT-I (live/dead⁻CD8⁺Vα2⁺) cells by flow cytometry.

NOTE: Simultaneous staining of congenic markers and the transgenic TCR should be performed to verify the correct phenotype of the cells prior to transfer.

4.4.1. Add 5 × 10⁴–1 × 10⁵ cells into 1 mL of FACS buffer in a 1.5 mL centrifuge tube, and centrifuge the cell suspension at 350 × *g*, 4 °C for 3 min.

4.4.2. Discard the supernatant, and disperse the cells by flicking the bottom of the tube. Place the tube on ice.

4.4.3. Prepare the following conjugated antibody mixtures (diluted in 100 µL FACS buffer): anti-CD8, 1:200; anti-TCR Vα2, 1:100; anti-CD45.1, 1:200; anti-CD45.2, 1:200; and live/dead,

1:200 (refer to the **Table of Materials**).

4.4.4. Vortex the antibody cocktail and centrifuge at $15,000 \times g$ for 3 min to pellet antibody aggregates. Store the cocktail on ice and protect it from light.

4.4.5. Resuspend the cells with 100 μL of antibody cocktail and thoroughly mix by flicking the tube. Incubate in the dark for 30 min on ice.

NOTE: Avoid disturbing the antibody aggregates at the bottom of the tube.

4.4.6. Wash the pellets twice with 1 mL of FACS buffer. Centrifuge at $350 \times g$, 4°C for 3 min. Resuspend the cells in 200 μL of FACS buffer, and transfer the cell suspension to a FACS tube.

NOTE: To maintain the viability of the OT-I cells to be transferred, test the specimen as soon as possible. If the stained OT-I cells cannot be tested immediately, keep the cells in the dark on ice or refrigerate at 4°C until analysis. Alternatively, the samples can be resuspended in 1–4% paraformaldehyde for extended storage (16 h) to prevent deterioration.

4.4.7. Run the specimen on a flow cytometer. Calculate the percentage of live/dead $^{\text{CD8}^+\text{Va2}^+}$ cells by dividing the number of live/dead $^{\text{CD8}^+\text{Va2}^+}$ cells by the number of live/dead $^+$ cells.

4.5. Determine the absolute number of OT-I cells (live/dead $^{\text{CD8}^+\text{Va2}^+}$) by multiplying the percentage of live/dead $^{\text{CD8}^+\text{Va2}^+}$ cells by the viable cell number obtained in step 4.3.9.

4.6. Adjust the concentration of OT-I cells (live/dead $^{\text{CD8}^+\text{Va2}^+}$) to $1.5 \times 10^6/\text{mL}$ with PBS.

4.7. Inject 3×10^5 distinct congenically marked OT-I cells (live/dead $^{\text{CD8}^+\text{Va2}^+}$) in 200 μL of PBS intravenously into two groups of B16F10-OVA-bearing mice (tumor-bearing mice divided into donor and recipient mice from step 3.8).

4.7.1. Withdraw 200 μL of OT-I cell (live/dead $^{\text{CD8}^+\text{Va2}^+}$) suspension into a 100 U insulin syringe (29 G), and remove bubbles as in step 3.2.

4.7.2. Place the mouse separately in a cage with an infrared lamp over the cage for 5–10 min to dilate the tail vein. Immobilize the mouse with a restraining device of appropriate size. Pull the tail to straighten it and spray with 75% ethanol to make the vein visible.

4.7.3. Hold the syringe parallel to the vein and insert it into the vein at an angle of $0\text{--}15^\circ$. Pull back the plunger slightly, and if blood enters the barrel, slowly and steadily inject the suspension at a rate of no more than 1 mL/min.

NOTE: Resistance or swelling at the injection site indicates that the needle is not inside the vein; the injection site should be moved proximally.

4.7.4. After the injection is completed, remove the syringe, and press the insertion area gently for 3–5 s to stop bleeding. Return the mouse to the cage and closely observe it for a few minutes for adverse reactions. If it has normal mobility and nasal discharge, place it back in

the company of the other mice.

5. Dissect tumor mass from tumor-bearing donor mice

NOTE: Maintain sterile conditions during surgery in sections 5 and 6. Sterilize all surgical instruments by autoclaving before and after each use. Disinfect the operating area in the biosafety cabinet with 75% ethanol followed by ultraviolet irradiation. Wear a clean gown, cap, face mask, and sterile gloves.

5.1. Eight to ten days after the adoptive transfer, select donor mice bearing comparable tumor mass of ~5 mm diameter (soybean-sized) for transplantation surgery.

5.2. Prepare a 100 mm × 20 mm dish in a biosafety cabinet, and add 10 mL of sterile ice-cold PBS.

5.3. Euthanize a tumor-bearing donor mouse in an isoflurane chamber followed by cervical dislocation. Immerse the mouse in 75% ethanol for 3–5 min and transfer to the biosafety cabinet.

NOTE: The following procedures in this step must be performed in a biosafety cabinet to maintain strict asepsis.

5.4. Place the mouse on a dissection board covered with clean absorbent paper in a supine position. Restrain the mouse limbs with dissection needles.

5.5. Cut the skin along the midline from above the urethral orifice to the xiphoid with scissors. Stretch the skin towards the left side of the mouse body with tweezers and restrain the skin with dissection needles.

5.6. Excise the tumor, keeping its capsule as intact as possible. Carefully and gently remove the connective tissue near the tumor with surgical scissors.

NOTE: To maintain the integrity of the tumor, do not peel off the tumor capsule or cut the tumor tissue into pieces.

5.7. Place the tumor tissue in a 100 mm × 20 mm dish containing 10 mL of sterile ice-cold PBS for subsequent transplantation.

6. Subcutaneous transplantation of donor-derived tumor onto the tumor-matched recipient mice

NOTE: The allograft is supposed to be implanted into the mouse's lower flank on the same side as the previously existing tumor to make two tumors drain to the identical lymph node. In the protocol presented here, as the B16F10-OVA tumor was implanted subcutaneously on the left inguinal region of the mouse (section 3), the donor-derived tumor tissue was transplanted onto the left flank of the recipient in this step. The transplantation site can be adapted to the first-implanted tumor site.

6.1. Anesthetize a tumor-matched recipient mouse with 250 mg/kg of 2,2,2-tribromoethanol via intraperitoneal injection. Pinch the footpad to assess the level of anesthesia, and wait for lack of pain reflex, which indicates the proper depth of anesthesia for performing the surgery. If vocalization or hind limb withdrawal is observed, further inject 0.01–0.03 mL of 2,2,2-tribromoethanol.

NOTE: The tumor-matched recipient mouse should be the same sex as the donor mouse that provides the allograft to avoid rejection issues.

6.2. Use veterinary ointment on eyes to prevent dryness. Shave the left flank of the mouse with an electric shaver. Apply a depilatory cream to remove the remaining hair.

NOTE: Avoid abrading the skin, which may increase the risk of contamination and infection.

6.3. Place the mouse in the biosafety cabinet. Place it in the prone position on a dissection board covered with clean absorbent paper, with the mouse's vertical axis parallel to and its head to the right side of the experimenter.

NOTE: The following procedures in this step must be performed in a biosafety cabinet to maintain strict asepsis.

6.4. Rub the skin of the shaved area with cotton soaked in povidone-iodine.

NOTE: Use povidone-iodine instead of 75% ethanol for sterilization to prevent loss of body heat.

6.5. Lift the skin at the center point between the mouse hip joints with surgical tweezers. Use the scissors to make a 5 mm-long vertical excision. Extend the cut rostrally along the dorsal midline to ~10–15 mm.

6.6. Perform a sharp dissection by inserting the closed tips of the scissors into the incision and then opening to separate the peritoneum of the left flank from the skin and soft tissue.

NOTE: To avoid causing damage to the subcutaneous tissue and peritoneum, lift the skin at the center of the incision, and then insert the closed scissors as close to the skin as possible.

6.7. Make a skin pocket at the left flank by performing sharp dissection several times. Deposit the encapsulated, intact donor-derived tumor mass into the capsule.

NOTE: Mice in the sham-operated control group receive the same surgery operation without the donor-derived tumor transplantation.

6.8. Close the incision by interrupted suture. Suture two or three stitches for each incision. Disinfect the skin around the cut with cotton soaked in povidone-iodine.

NOTE: There should be 5 mm between two consecutive stitches and a 3 mm distance from

the incision.

6.9. Place the mouse in the lateral position in a clean and warm cage. Monitor it continuously until it has regained sufficient consciousness to maintain sternal recumbency.

6.10. Administer penicillin every 8–12 h after the surgery for 3 days. Monitor the mouse's eating, drinking, moving, and the area operated on. Return the transplant recipient to the company of other animals only after it has fully recovered.

NOTE: The administration of buprenorphine is suggested to prevent post-surgical pain. The mouse typically recovers from the trauma of the surgery within 3 days. If the mouse is not back to normal feeding and mobility and shows any manifestations of infection, consult a veterinarian for interventions or euthanize it.

6.11. Sacrifice (euthanize the animals as in step 4.3.2) the mice at the indicated time points, and recover the cells of interest for flow cytometric analysis.

REPRESENTATIVE RESULTS:

The schematic of this protocol is shown in **Figure 1**. Eight days after tumor inoculation, CD45.1⁺ and CD45.1⁺CD45.2⁺ OT-I cells were injected into B16F10-OVA tumor-bearing C57BL/6 mice. The tumor was surgically dissected from CD45.1⁺ OT-I cell-implanted mice (donor) on day 8 post-transfer and transplanted into tumor-matched CD45.1⁺CD45.2⁺ OT-I cell-implanted mice (recipient) in the dorsal flank on the same side as the implanted tumor. Through flow cytometry (gating strategy shown in **Figure 2**) analysis, two populations of CD44⁺CD8⁺ tumor antigen-specific T cells can be easily identified in the TME, including CD45.1⁺ donor-derived and CD45.1⁺CD45.2⁺ recipient-derived TILs. Subsequently, the proportions of these two populations within the allografts were analyzed at indicated time points to study the dynamics of the antigen-specific CD8⁺ T cells. At day 2 post-transplantation, there were ~83% of donor-derived antigen-specific CD8⁺ T cells within the transplanted tumor, more predominant than their recipient-derived counterparts. However, the proportion of recipient-derived OT-I cells was elevated in the late stage of tumorigenesis, exceeding tumor-inherent OT-I cells derived from the donor. (**Figure 3**).

FIGURE AND TABLE LEGENDS:

Figure 1: Schematic of the experimental design. C57BL/6 mice are challenged with B16F10-OVA tumor on the inguinal area. Eight days later, different congenically marked (CD45.1⁺ or CD45.1⁺CD45.2⁺) OT-I cells are transferred into tumor-bearing mice. On day 8 post-transfer, the tumor on the CD45.1⁺ OT-I cell-implanted mice is surgically dissected and subcutaneously transplanted into tumor-matched CD45.1⁺CD45.2⁺ OT-I cell-implanted recipients in the flank on the same side as the existing tumor. Then, the mice are sacrificed, and antigen-specific T cells (OT-I cells) within the allografts are analyzed at the indicated time points. Abbreviations: CD = cluster of differentiation; i.v. = intravenous; Sac = sacrifice.

Figure 2: Gating strategy of flow cytometry analysis. Gating strategy used to identify donor-derived (CD45.1⁺) and recipient-derived (CD45.1⁺CD45.2⁺) antigen-specific CD44⁺CD8⁺ T cells within allografts. Abbreviations: SSC-A = side scattering–area; FSC-A = forward scattering–

area; FSC-W = forward scattering–width; FSC-H = forward scattering–height; SSC-W = side scattering–width; SSC-H = side scattering–height; L/D = live/dead; CD = cluster of differentiation.

Figure 3: The ratio of donor- and recipient-derived antigen-specific CD8⁺ T cells within tumor allografts. Representative flow cytometry plots showing expression of the congenic markers CD45.1 and CD45.2 used to identify donor-derived and recipient-derived OT-I cells within tumor allografts at days 2, 8, and 15 after transplantation. The numbers represent the percentages of the two subsets in the CD44⁺CD8⁺ T cell population.

DISCUSSION:

T cell-mediated immunity plays a crucial role in immune responses against tumors, with CTLs playing the leading role in eradicating cancerous cells. However, the origins of tumor antigen-specific CTLs within TME have not been elucidated³⁰. The use of this tumor transplantation protocol has provided an important clue that intratumoral antigen-specific CD8⁺ T cells may not persist for a long time, despite the existence of stem-like TCF1⁺ progenitor CD8⁺ T cells. Notably, there is a continuous influx of periphery-derived tumor-specific CD8⁺ T cells into the tumor mass.

To our knowledge, this is a relatively convenient and convincing method confirming that the maintenance of antigen-specific CD8⁺ T cells within the TME predominantly depends on the replenishment of periphery-derived tumor-specific CD8⁺ T cells instead of the self-renewal of tumor-resident TILs. Although the protocol presented here only focuses on the proportions of donor-derived and recipient-derived TILs, the phenotypic, functional, and transcriptional properties of these two populations can be readily examined with flow cytometry. Moreover, it is feasible to combine ICB antibodies to investigate the responses of a specific cell subset to ICB therapy.

In this protocol, donor-derived tumor tissue is transplanted onto the recipient mouse with an existing original tumor. Two tumors in a recipient mouse will lead to the distribution of periphery-generated T cells into two tumor masses. Moreover, the tumor burden will be nearly doubled compared to animals without transplants. In pilot experiments, we attempted to excise the original tumor on recipient mice before transplantation; however, it was technically challenging to eliminate all tumor cells by surgery thoroughly. The residual tumor cells will grow rapidly and form a new tumor tissue soon. Thus, there is a limitation for this system when comparing T cell immune responses with those in non-transplanted mice.

However, this system is still useful for the comparison of recently migrated and existing T cells within the same TME that is transplanted from donor tumor-bearing mice. Besides, there is no denying that the transplantation of tumor tissue may lead to inflammation, which might influence immune cell dynamics within the tumor. Though the impact of surgery on OT-I cell infiltration could be excluded through non-operated and sham-operated controls, we did not assess the effects of local inflammatory responses to OT-I cell dynamics.

Some considerations should be taken into account, one of which is the usage of cyclophosphamide. Cyclophosphamide³¹ is an alkylating agent widely used to treat solid organ malignancies and lymphoproliferative and autoimmune disorders. Six to eight days

after B16F10-OVA inoculation, cyclophosphamide is administered before adoptive transfer to induce the lymphodepletion of host mice and enhance the activity of the transferred OT-I cells²⁹. Although melanoma is not sensitive to this reagent, some tumor cell lines, such as EG7³², a murine thymic lymphoma cell line, respond to cyclophosphamide. Treatment of EG7-bearing mice with cyclophosphamide results in the eradication of tumors, which suggests that cyclophosphamide must be carefully used or titrated for sensitive tumor models. The recommended alternative method is a single sublethal dose of radiation (4.5–5.5 Gy) one day before the transfer, and the optimal choice depends on the characteristic of tumor cell lines.

Other steps need to be taken cautiously, including the careful selection of tumor-bearing donor mice and the delicate surgical operation during tumor transplantation. Implanted tumors would be surgically removed and transplanted into tumor-matched recipient mice 8–10 days post-transfer. Before transplantation, a comparable size of tumor mass of ~5 mm diameter is to be chosen as an allograft to reduce discrepancies between individual mice and make acquired data more reliable. Moreover, during surgery, the incision should be near the midline of the mouse back to keep the allograft at a distance from the tumor already existing in the recipient mouse. Gentle dissection is also suggested to prevent injuries on the inguinal lymph node and surrounding tissues.

The effective killing of cancerous cells requires the coordination of various components within the TME³³. The protocol presented here can be extended to the investigation of adaptive and innate immune cells such as natural killer cells, tumor-associated macrophages, and dendritic cells. Furthermore, in addition to the B16F10-OVA utilized here, this protocol can be applied to other subcutaneous tumor models. To conclude, the aforementioned tumor transplantation assay offers a new approach for the study of interactive transitions of certain types of immune cells during antitumor responses and is useful for researchers in tumor immunology.

ACKNOWLEDGMENTS:

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DISCLOSURES:

The authors have no conflicts of interest to disclose.

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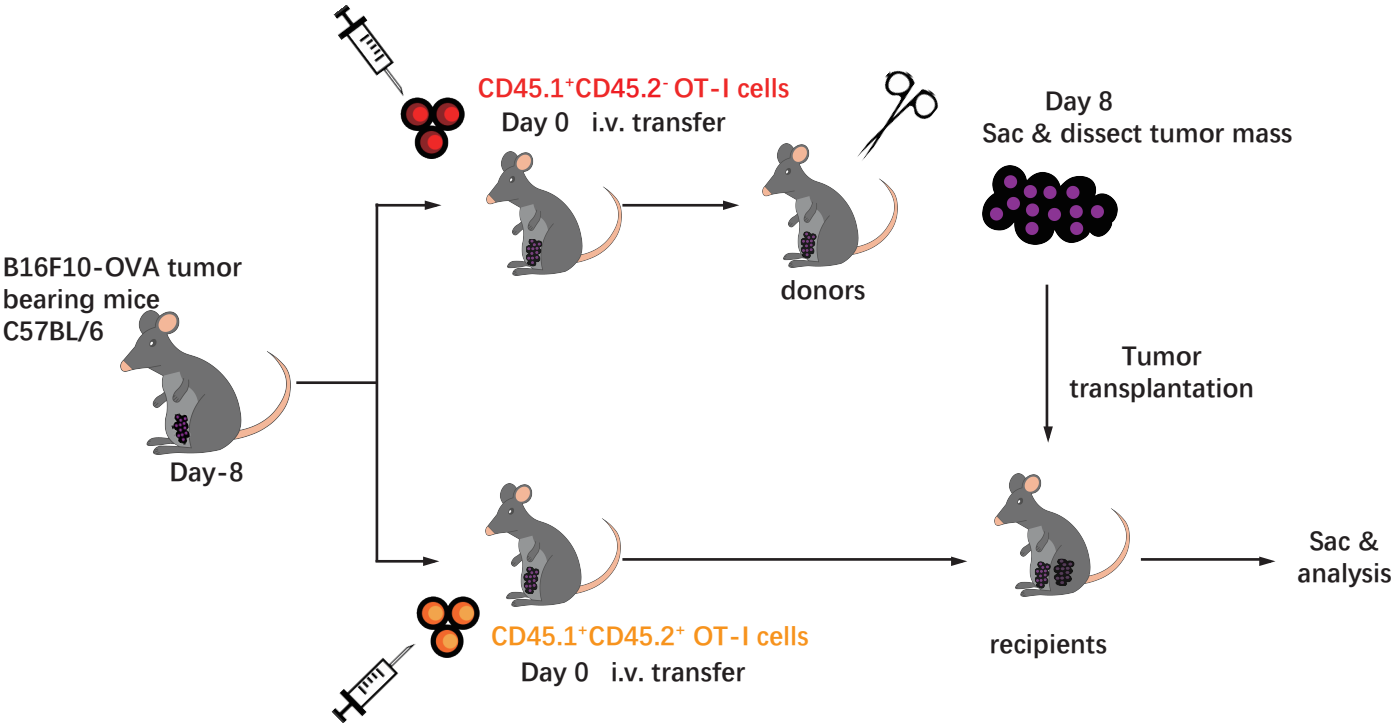
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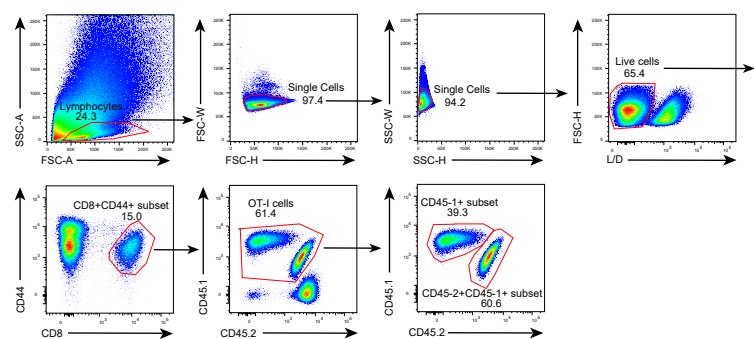
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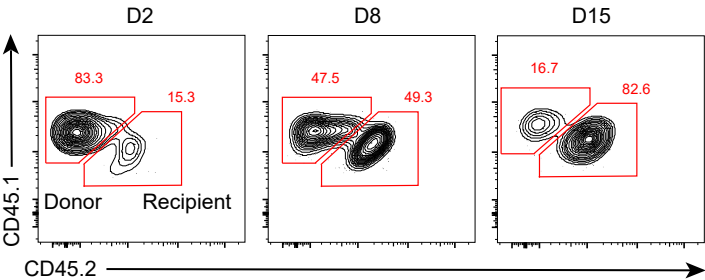
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Table of Materials

JoVE_Materials (11).xls



Editorial comments:

Please address all the comments in the attached document by tracking your changes and retain all stylistic and formatting changes.

Response:

Thanks for editors' efforts in reviewing the manuscript. We have revised the manuscript according to the comments in the attached document and the changes have been tracked.

Reviewers' comments:**Reviewer #3:**

Manuscript Summary:

The manuscript was significantly improved and is detailed and easy to follow.

Major Concerns:

I have no other concerns.

Minor Concerns:

I have no other concerns.

Response:

We are very thankful to this reviewer for supporting the publication of this manuscript in JoVE.

Reviewer #5:

In their manuscript entitled "Tumour Transplantation for Assessing Dynamics of Tumour Infiltrating CD8+ T cells in Mice", Ye and colleagues describe a method to transplant subcutaneous tumours from donor mice to tumour matched recipient mice in combination with adoptively transferred antigen specific T cells in order to dissect CD8 T cell dynamics in the tumour environment. While this is an interesting approach and I agree with reviewers 2-4 that it is worth publishing, there are a few points that need addressing before publishing of the manuscript:

3.1 The authors indicate that male and female mice can be used. They comment on the need of sex-matched OT-I T cells but will also need to clarify that tumours should only be transplanted between sex-matched mice in order to prevent rejection issues.

Response:

We thank the reviewer for pointing out this important issue. In section 6 of the new manuscript, we have emphasized that tumors should be transplanted between sex-matched mice.

4.4.7 The authors suggest that stained OT-I T cells are resuspended in 2% PFA

before FACS. This should be an optional step.

Response:

In the newly submitted manuscript, we have revised the relevant step and added a note to clarify this point.

6.1 Euthanize needs to be replaced by Anesthetize (recipient mice need to survive)

Response:

We have corrected this in the new manuscript.

6.7 The authors state that a "piece of donor derived tumor" will be placed in the generated skin capsule. They should comment on the size of the piece. Do they transplant the whole tumour or do they cut off a piece of a certain size? This might need to be standardized in order to generate reproducible data. Furthermore, the procedure how the skin pocket was generated seems a bit unclear to me. The video will certainly help to understand but maybe a figure with photos of each step of the procedure would be helpful?

Response:

We may not clearly elaborate this point in our manuscript. Virtually, in pilot experiments, we found that cutting tumor mass into pieces increased chances of the tumor necrosis and surgical site infection. Instead, transplanting a whole tumor mass with its capsule intact can largely avoid these issues. Besides, the size of allograft affects the infiltration of tumor-specific CD8⁺ T cells (OT-I cells). Hence, we selected donor mice bearing comparable tumor mass of about 5 mm in diameter (soybean-sized) before surgery and dissected the tumors with their capsule as intact as possible. Then we transplanted the encapsulated, intact tumor tissue onto the tumor-matched recipient mice. In the newly submitted manuscript, we have added related information to section 5 and section 6.

6.10. Transplanted mouse should be replaced by transplant recipient.

Response:

Thanks for the advice, and we have changed it in the revised manuscript.

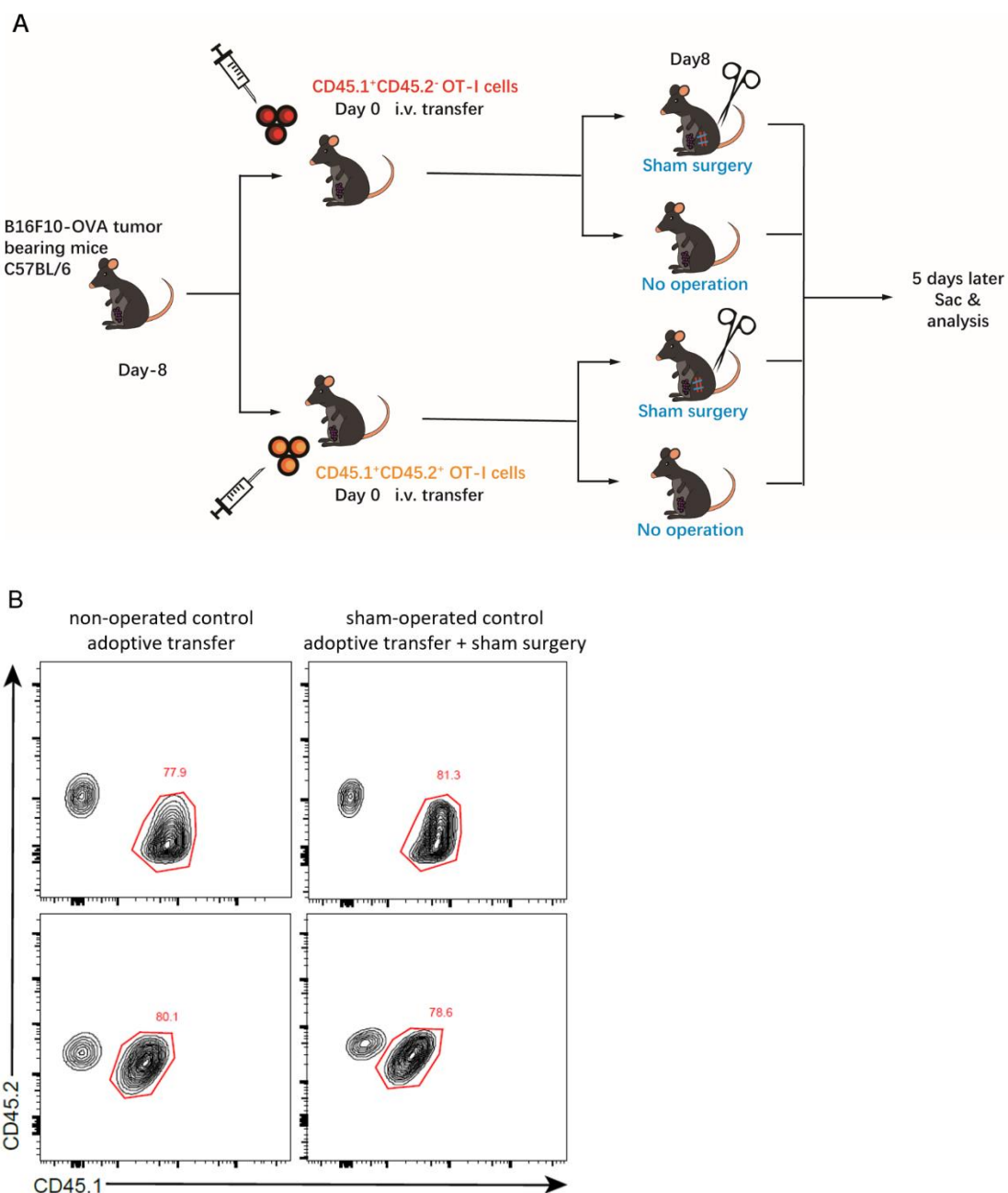
General comments/concerns:

Transplantation and surgery will lead to inflammation, which might influence T cell/immune cell dynamics within the tumor. Is there a way to control for this issue? At least, it should be discussed as a limitation of this technique.

Response:

We are thankful to the important points raised by the reviewer. To evaluate the effects

of surgery on transferred cells, we set two groups of mice including non-operated controls and sham-operated controls. The tumor-bearing mice in non-operated control group receive either CD45.1⁺CD45.2⁺ or CD45.1⁺ OT-I cells but no surgery operation, while the mice in sham-operated control group receive either CD45.1⁺CD45.2⁺ or CD45.1⁺ OT-I cells and subsequent surgery operation similar to the experimental group, but no allograft transplantation (figure A). The ratios of OT-I cells with distinct congenic marks in sham-operated mice were comparable with that in tumor-bearing mice receiving only adoptive cell transfer but no operation (figure B), confirming that surgery operations don't overtly influence the infiltration of OT-I cells. We agreed the reviewer's opinion that transplantation may lead to inflammation, which might influence immune cell dynamics within the tumor. Thus, there is a limitation for this protocol because we did not assess the effects of local inflammatory responses to OT-I cell dynamics.



(A) Schematic procedures of non-operated controls and sham-operated controls set-up. Adoptively transfer CD45.1⁺ or CD45.1⁺CD45.2⁺ OT- I cells into B16F10-OVA tumor-bearing C57BL/6 CD45.2⁺ mice, and on day8 post-transfer, perform sham surgery by cutting the dorsal skin of mice and sewing up the cut. This group of mice are set as non-operated controls. Mice that receive CD45.1⁺ or CD45.1⁺CD45.2⁺ OT- I cells but not surgery are set as negative controls. 5 days after sham operation, sacrifice mice of both sham-operated control group and non-operated control group. CD44^{hi} CD8⁺ T cells in tumor mass are recovered and analyzed by flow cytometry.

(B) Representative flow cytometry plots of proportions of CD45.1⁺ (top) or CD45.1⁺CD45.2⁺ (bottom) transferred OT- I cells in tumors of negative control mice and sham-operated mice at day5 after sham-operation. The panels are gated on CD44^{hi}CD8⁺ live cells and the subsets circled in red frame are the transferred OT- I cells.

While the transplantation of whole tumor tissue is an interesting technique, the authors should explain or at least comment on the benefits of this method compared to transfer of donor derived peripheral T cells into the recipient mouse (i.e. why do they not take donor blood, purify OT-I T cells and transfer those cells into the recipient tumor matched mouse?). This would certainly be less invasive.

Response:

Thanks to the reviewer for making this point. With this tumor transplantation assay, we are able to track dynamic phenotypic and functional changes of both newly infiltrated immune cells and already-infiltrated immune cells in TME (tumor micro-environment), which presumably differ from each other in phenotypes and functions, in the long run. However, without tumor mass transplantation, it is extremely challenging to distinguish the newly infiltrated T cells from the periphery in TME, in particular, in longitudinal assays. The transferred OT-I cells will not infiltrate to the TME as a whole, instead, a fraction of these cells will circulate in the blood, which may consistently replenish OT-I cells in TME. Therefore, we considered our protocol as a better way to investigate the temporal and dynamic changes of both peripheral and infiltrated tumor-specific T cells under different conditions.



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December 28, 2020

Journal of Visualized Experiments,

Dear editors:

Please find enclosed our manuscript entitled “Tumor Transplantation for Assessing Dynamics of Tumor Infiltrating CD8⁺ T Cells in Mice” by *Lisha et al.* for your consideration of publication as an Article in *Journal of Visualized Experiments*.

T cell-mediated immunity plays a crucial role in immune responses against tumor, with CTLs playing the leading role in eradicating cancerous cells. However, the origins and replenishment of tumor antigen specific CD8⁺ T cells within tumor micro-environment remain obscure. Within the tumor microenvironment (TME), persistence of tumor antigens as well as the immunosuppressive factors drive CD8⁺ T cells into a hypo-responsive state of exhaustion. Besides, the impaired proliferative capacity of exhausted CD8⁺ T cells results in decreasing number of tumor antigen-specific T cells and the residual CD8⁺ T cells within TME could barely provide sufficient protective immunity against tumor progression. Thus, the maintenance or reinforcement of intra-tumoral antigen-specific CD8⁺ T cells is indispensable for tumor repression. To study immune responses of antigen-specific T cell during tumorigenesis and compare the immune responses of tumor-inherent and peripheral-originated antigen-specific CD8⁺ T cells, we designed this experiment, the critical points are as follows:

- (1) We employed B16F10-OVA melanoma cell, stably expressing the surrogate antigen ovalbumin and TCR transgenic OT-I mice, enabling the study of antigen-specific T cell responses during tumorigenesis.
- (2) By transplanting tumor tissues from donor mice into tumor-matched recipient mice to specifically trace the influx of recipient-derived immune cells, it allowed us to analyze the immune responses of tumor-resident and peripheral-originated antigen-specific CD8⁺ T cells.
- (3) We found a dynamic transition between tumor-inherent and peripheral-originated antigen-specific CD8⁺ T cells.

Collectively, this experimental design has provided a new approach to precisely investigate the immune responses of CD8⁺ T cells in TME, and can be further exploited as a novel strategy in the field of anti-tumor immunity. Hence, this study

should be of broad interests to scientists in the field of anti-tumor immune response, T cell biology and tumor immunotherapy.

For potential reviewers, we recommend Dr. Bin Li (Shanghai Institute of Immunology, Shanghai JiaoTong University School of Medicine, binli@sibs.ac.cn) and Dr. Linrong Lu (Institute of Immunology, Zhejiang University School of Medicine, lu_linrong@zju.edu.cn) who are experts on the tumor immunology. Thank you for your consideration.

Sincerely yours,

A handwritten signature in black ink, appearing to be 'Lilin Ye', with a stylized, flowing script.

Lilin Ye, Ph.D.

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