

Cancer Derived Exosomes Induce Macrophages Immunosuppressive Polarization to Promote Bladder Cancer Progression

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Abstract

Background: Exosomes mediated crosstalk between tumor cells and other stromal cells including tumor associated macrophages (TAM) plays an essential role in reprogramming tumor microenvironment (TME) to facilitate tumor progression. However, the mechanism of tumor derived exosomes promotes bladder cancer progression have not been defined.

Methods: Exosomes were extracted from bladder cancer cells MB49 conditioned medium by ultracentrifugation. The effects of MB49-derived exosomes on macrophages polarization were analyzed by qPCR, flow cytometry, and Western blot. The immunosuppressive phenotype and function of MB49-derived exosomes stimulated macrophages were verified by tumor xenograft assays and T cell co-culture experiments. Exosomal miRNAs were analyzed by microarray to identify potential targets regulating macrophage polarization.

Results: MB49-derived exosomes could be ingested by macrophages, consequently promoting macrophages immunosuppressive polarization. Mechanically, the MB49-derived exosomes induced macrophage M2 polarization was mediated by down-regulation of PTEN and activation of AKT/STAT3/6 signaling. Moreover, hindrance of the generation or secretion of exosomes by GW4869 inhibited macrophages differentiation into immunosuppressive phenotype and function, thereby suppressed tumor growth in a mouse subcutaneous tumor model.

Conclusion: Our study confirmed the contribution of bladder cancer derived exosomes on the establishment of immunosuppressive TME and provided a potential therapeutic target for bladder cancer treatment.

Background

Bladder cancer is one of the most common type of cancers, with more than 357,000 new cases and 130,000 deaths worldwide each year[1]. About 75% bladder cancer is non-muscle-invasive at diagnosis, and the remainder present as muscle invasion with/out metastasis. Even following effective treatment like transurethral resection of bladder cancer and intravesical chemotherapy, part of non-muscle-invasive bladder cancer can progress to muscle invasion[2]. The muscle invasive bladder cancer (MIBC) without metastasis is managed by radical cystectomy with/out neoadjuvant chemotherapy, however it has high recurrence and metastasis rate. The five-year survival rate of MIBC is less than 70%, even worse for metastatic form[3, 4]. Therefore, it is of particular importance to clarify the mechanisms of bladder cancer progression and identify new targets for bladder cancer therapy.

The tumor microenvironment (TME) refers to the environment around a tumor, consisting of tumor cells, immune cells, fibroblasts, the surrounding blood vessels and the extracellular matrix, which plays an essential role in cancer development, progression and control[5]. Since the immune cells are important determinants of TME, the interplay between host immune system and tumor cells has been intensively investigated in the past decade. Recently, immunotherapy has been used for a variety of cancers, but still

far from meeting clinical needs[6]. Tumor associated macrophages (TAM), a major component of immune cells, have been suggested to play a central role in tumor progression[7–9]. TAM is generally considered to be immunosuppressive with high *CD206/CD163* expression[10, 11]. Increasing evidence indicated that TAM could increase tumor invasion and metastasis by taming host adaptive immunity[12, 13]. It is believed that re-educating TAM to tumor killing function may potentially contribute to bladder cancer therapy. Therefore, it is urgent need to understand how TAM polarization in bladder TME during tumor progression.

Emerging evidence has suggested that the communication between tumor cells and macrophages play a key role in mediating TAM polarization[14, 15]. Signals originated from tumor cells such as transforming growth factor beta (TGF- β), could promote differentiation of non-activated macrophages into a TAM-like immunosuppressive phenotype characterized as increased anti-inflammatory cytokine and decreased pro-inflammatory cytokine expression[11, 16]. Exosomes, small vesicles derived from cells, are wrapped in a lipid bilayer that carry various biological molecules, including proteins, polysaccharides, lipids, DNA, RNA, metabolites, and mediate the information transmission and exchange of material between cells[17, 18]. Therefore, exosomes contribute to information and material exchange between tumor cells and other stromal or immune cells in TME[19]. It has been reported that exosomes originated from TAM could promote metastasis of tumor cells[20] as well as resistance to chemotherapy[21]. Reciprocally, exosomal miR-21 secreted by bladder cancer cell T24 cells could polarize THP-1 cell-derived macrophages into the M2 phenotype by activating PI3K-AKT-STAT3 signaling pathway[22]. However, the underlying mechanism of tumor cells derived exosomes mediating macrophage polarization is still not fully understood.

In current study, we aimed to investigate how bladder tumor cells derived exosomes induce macrophage polarization and the effects on tumor growth. We first investigate the effect of mouse bladder cell MB49-derived exosomes on bone marrow-derived macrophages (BMDM) polarization, as well as the influence of MB49-derived exosomes stimulated macrophages on T cell proliferation. Moreover, we also studied the potential therapeutic function of GW4869 on bladder cancer growth by inhibiting the generation or release of exosomes.

Materials And Methods

Cell culture

The mouse bladder cancer cell line MB49 was obtained from the Chinese Academy of Sciences. MB49 cells were cultured in DMEM medium (GE Healthcare Life Sciences, Pittsburgh, PA, USA), supplemented with 10% fetal bovine serum (FBS; Gibco; Thermo Fisher Science, Massachusetts, USA).

BMDM isolation

Bone marrow-derived macrophages (BMDM) were isolated from the femur and tibia of adult male mice according to our previous publication[23]. Briefly, mouse femoral tissue was isolated, and the bone marrow was flushed with 3 ml of normal saline (NS). After red blood cell lysis, bone marrow cells were

suspended at $2-4 \times 10^6$ cells/ml in RPMI-1640 medium, supplemented with 10% FBS and 30 ng/ml GM-CSF (PEPROTECH Inc., Rocky Hill, USA). BMDM were cultured in a humidified incubator at 37°C and 5% CO₂.

Isolation and identification of exosomes

MB49 cells were washed twice with PBS when its confluence reach to 90%, following cultured with serum-free RPMI 1640 medium for 24 h. The culture supernatant was collected and centrifuged at 300 g and 15,000 g for 20 min respectively, which is to remove the suspended cells and cell debris. Then the collected supernatant was subjected to ultracentrifugation at 100,000g for 70 min at 4°C (HITACHI ultracentrifuge, CS150FNX). The obtained exosome pellet was washed with PBS and concentrated after ultracentrifugation, which is to improve the purification of exosomes. The MB49-derived exosomes were verified by transmission electron microscopy (HT7800, HITACHI) and the expression of exosome specific markers CD63 and HSP90 by Western blot analysis.

RNA extraction and qRT-PCR

The total RNA of exosomes stimulated BMDM was extracted with Trizol® reagent (Invitrogen, Carlsbad, CA, USA), and reverse transcription was performed with PrimeScript™ RT Master Mix (Takara Co. Ltd.). Real-time PCR was performed on the ABI PRISM 7300HT sequence detection system (Applied Biosystems, Foster, CA, USA) using SYBR Green PCR Master Mix (Application Takara, Otsu). The relative expression level of mRNA was calculated by the $2^{-\Delta\Delta Ct}$ method. The sequences of all primers are shown in the supplementary table 1. Each experiment was repeated three times.

Western blot

BMDM were lysed with RIPA buffer (Beyotime institute of Biotechnology, Shanghai, China). The bicinchoninic acid protein assay (BCA) kit (Beijing Leagene Biotech co. Ltd) was used to determine the protein concentration. Total 20 µg protein were separated on SDS-PAGE gel and transferred to 0.45 µm PVDF membrane. The membrane was blocked with 5% skimmed milk at room temperature for 1 h. The primary antibodies were incubated overnight at 4°C. The peroxidase-conjugated secondary antibody was incubated for 60 min at room temperature. The protein bands were then visualized using an enhanced chemiluminescence kit (Beyotime institute of Biotechnology) and scanned by an imaging system (Bio-Rad Laboratories, Inc. Hercules, CA, USA). The ImageJ software v1.8.0 (National Institutes of Health) was used to quantify the density measurement. The primary and secondary antibodies used in this study were listed in supplemental table 2.

Exosomes tracing

To detect the uptake of MB49-derived exosomes by BMDM, the fluorescent dye azide cyanine-Cy5.5 (0.2 µM, Fanbo BioChemical, Beijing, China) was added to the exosomes at 37°C for 30 min, following the Cy5.5 labeled exosomes were centrifuged at 100,000 g, 4°C for 70 min to remove residual dye, and then incubated with BMDM for 3 h in the dark at 37°C. The nuclei were stained with DAPI (Beyotime institute of Biotechnology, Shanghai, China) at room temperature for 5 min, and then observed with a laser scanning

confocal microscope or analyzed by flow cytometry analysis. For tracking the uptake of circulating MB49-derived exosomes in vivo, we first injected Cy5.5 labeled exosomes (1 µg in 100 µl PBS) into the mouse via the tail vein. After 12 h, the bladder was harvested and digested with 0.1 mg/ml collagenase D / DNase I (100 U/ml) (Sigma-Aldrich, Inc. St. Louis, MO, USA) solution. The Cy5.5 positive cells in CD45⁺F4/80⁺CD11b⁺ macrophages were analyzed by flow cytometry.

T cell co-culture experiments

MB49 cells were treated with/without 10 µM GW4869 (MCE, HY-19363) for 24 h. Afterwards, the exosomes were collected from culture supernatant by ultracentrifugation. After stimulating BMDM with exosomes for 48 h, Carboxyfluorescein succinimidyl ester (CFSE, abcam, 113853)-labeled splenic T cells were added and co-cultured for another 3 d. Then the percentage of CD4⁺/CD8⁺ T cells was detected by flow cytometry.

Flow Cytometry

To detect the effects of MB49 conditional medium (CM) and MB49-derived exosomes on BMDM, the freshly isolated BMDM were stimulated with MB49 CM or MB49-derived exosomes for 48 h. Then BMDM were collected and stained with CD45, F4/80, CD206 and CD11b. For the tumor associated immune cells identification, the MB49 derived subcutaneous tumor tissues were taken and digested with 0.1 mg/ml type D collagenase (Sigma-Aldrich, Inc. St. Louis, MO, USA) for 30 min at 37°C, and the single cell suspension were filtered and stained with macrophage markers (CD45, F4/80, CD206 and CD11b). The macrophage population was detected by using flow cytometer (BD Canto, Franklin Lakes, USA), and data were analyzed by FlowJo X (Tree Star, Ashland, OR, USA) software. Each experiment was repeated three times. The used antibodies were listed in supplementary table 2.

Microarray analysis of exosomal miRNAs

To determine the miRNAs contained in MB49-derived exosomes, a microarray analysis using Agilent mouse miRNA microarray kit (Agilent Technologies) was performed (OE Biotech Company, Shanghai, China). The sample preparation, miRNAs labeling, microarray hybridization and washing processes were carried out according to the manufacturer's protocols. Data extraction and visualization were performed using Feature extraction software (version 10.7.1.1, Agilent Technologies).

Tumor xenografts

To evaluate the influence of exosomes to macrophage polarization in vivo, we established a mouse subcutaneous tumor model. Briefly, total 1×10^6 MB49 cells were transplanted subcutaneously into the right flank of a male 8–10 week-old mouse. From day 3, GW4869 (2.5 µg/g in DMSO/saline) was intraperitoneally injected in every 3 days for total 4 times. A sham control group was injected with DMSO/saline alone. Tumor volumes were calculated as $1/2 \times \text{length} \times \text{width}^2$. The mice were sacrificed on day 15 and the tumor/body weight ratio was measured. The animal experiment was approved by the Ethical Review Committee of the First Affiliated Hospital of Zhengzhou University.

Statistical analysis

The statistical analysis was performed using GraphPad Prism 8 software. Comparisons between two groups were made using *t-test* as appropriate. For multiple comparisons, one-way ANOVA was applied. Statistically significant differences are indicated as follows: *P < 0.05; **P < 0.01; ***P < 0.001.

Results

MB49 cells conditioned medium induces macrophages differentiation into M2 phenotype

Previous studies have suggested that secreted factors from tumor cells could mediate the polarization of macrophages[24, 25]. In order to investigate the effect of bladder cancer cells secreted factors on macrophage polarization, mouse BMDM were treated with bladder cancer cell MB49 cells culture supernatant and gene expression was analyzed by RT-qPCR. The expressions of M2 related genes, including interleukin-10 (*Il-10*), *Cd206* and transforming growth factor beta (*Tgfb*) were increased, whilst M1 gene *Inos* (Fig. 1a) was decreased. The MB49 cells culture supernatant mediated macrophages M2 polarization was further verified by showing the increased F4/80⁺ CD206⁺ macrophage population (Fig. 1b). These results indicated that MB49 cells conditioned medium (CM) can promote the polarization of BMDM to M2 phenotype.

Identification of exosomes derived from MB49 cells and ingested of exosomes by macrophages in vitro and in vivo.

We further asked if exosomes in MB49 cells CM could mediate macrophages M2 polarization. Exosomes were extracted by ultracentrifugation and the morphology was observed through an electron microscope (Fig. 2a). In addition, the expression of exosome markers HSP90 and CD63 was also detected by Western blot (Fig. 2b). To interrogate whether macrophage can uptake MB49-derived exosomes, BMDM were co-cultured with the Cy5.5 labeled exosomes. Immunofluorescence and flow cytometry analysis demonstrated that MB49-derived exosomes can be ingested by macrophages (Fig. 2c, d). In order to further study the possibility of exosomes enrichment and absorption in bladder macrophages, we injected Cy5.5-labeled MB49-derived exosomes into mice via the tail vein. The results of flow cytometry showed that the average fluorescence intensity of Cy5.5 in bladder F4/80⁺ CD11b⁺ macrophages was much higher than that of the control group (Fig. 5e). Based on the above results, we confirmed that exosomes originated from MB49 cells can be ingested by macrophages both in vitro and in vivo.

MB49-derived exosomes mediated the polarization of macrophages into immunosuppressive phenotype

To further identify the effect of MB49-derived exosomes on BMDM phenotype and function, we first detected the gene expression level of M1/M2 macrophage markers in MB49-derived exosomes treated BMDM. RT-qPCR results showed that the expression of *Il10*, *Cd206* and *Tgfb* in MB49-derived exosomes stimulated BMDM was significantly increased, whilst the expression of M1 macrophage gene *Inos* was reduced, indicating MB49-derived exosomes could polarize macrophage to M2 phenotype (Fig. 3a). To

further confirm the polarization of M2 macrophages was exosomes dependent, we utilize GW4869, an exosome inhibitor, to detect the expression of M1/M2 macrophage markers. As expected, GW4869 treatment significantly reduced exosomes release, as showing reduced CD63 and HSP90 expression in MB49-derived exosomes by Western blot (Fig. 3b). Moreover, GW4869 treatment could inhibit MB49-derived exosomes mediated M2 macrophage gene *I110* and *Cd206* upregulation and increase M1 macrophages gene *Inos* expression (Fig. 3c). Additionally, the proportion of F4/80⁺CD206⁺ macrophages was significantly decreased in GW4869 treatment group comparing to MB49-derived exosomes group (Fig. 3d). To further investigate the immunosuppressive function of MB49-derived exosomes polarized macrophages, we co-cultured the MB49-derived exosomes stimulated BMDM with CFSE-labeled splenic T cells for 3 days. MB49-derived exosomes polarized BMDM significantly reduced the proliferation of CD4⁺/CD8⁺ T cells, whilst GW4869 treatment reversed the reduction of T cell proliferation (Fig. 3e). Collectively, these results indicate exosomes in MB49 CM play a dominant role in polarizing macrophages into M2 phenotype and immunosuppressive function.

MB49-derived exosomes promote bladder tumor cells growth in a mouse subcutaneous tumor model.

We next ask if the MB49-derived exosomes induced immunosuppressive macrophages could influence bladder tumor cells growth in vivo. In an established mouse subcutaneous tumor model, the volume/weight and growth rate of xenograft tumors were significantly decreased in GW4869 treatment group comparing to DMSO treatment group (Fig. 4a-c). Moreover, the majority of macrophages in tumors are M2 (F4/80⁺ CD206⁺) phenotype, and GW4869 treatment significantly reduced the percentage of M2 macrophages (Fig. 4d). Furthermore, the proportion of CD4⁺ and CD8⁺ T cells were significantly higher in GW4869 treatment (Fig. 4e). Taken together, these results indicate that MB49-derived exosomes can promote the growth of tumor cells by exosomes mediated immunosuppressive TME.

Exosomal miRNAs promote the polarization of macrophages in a PTEN/p-AKT/STAT3/6 dependent manner.

A growing body of evidence suggests that exosomal miRNA has multiple functions such as facilitating tumor growth, invasion, modulating TME[26]. To further elucidate miRNA profile in MB49-derived exosomes, microarray analysis was conducted. By analyzing two databases, 246 miRNAs were detected and 42666 target genes were predicted (Fig. 5a). Based on MB49-derived exosomal miRNAs target genes, a variety of cancer related signaling pathways like PI3K-AKT pathway were identified by KEEG analysis (Fig. 5b). Since PTEN is a tumor suppressor and negatively regulates PI3K-AKT signaling[27], two miRNAs, miR92a-3p and miR1231-5p were identified which can target *Pten* mRNA sites by using the online miRNA-mRNA matching tool Target Scan (Fig. 5c). In addition, Western blot results corroborate that the expression of PTEN was reduced in MB49-derived exosomes stimulated BMDM (Fig. 5c). Consequently, p-PI3K and p-AKT were significantly upregulated (Fig. 5d). Of note, p-STAT3/6 were also activated at 30 min and 1 h after MB49-derived exosomes stimulation, however the underlying mechanism needs further investigation. In summary, we identified MB49-derived exosomal miRNAs could

induce macrophage polarization to immunosuppressive phenotype by activating PTEN/AKT/STAT3/6 pathways, thus contributing to the establishment of immunosuppressive TME to facilitate tumor growth.

Discussion

Dynamic and mutual crosstalk between heterogeneous tumor cells and stromal cells including immune cells plays a vital role in tumor formation and progression. In TME, tumor cells manipulate other stromal cells by a variety of signaling networks and molecules to facilitate their growth, invasion and metastasis[5]. Macrophages are heterogeneous and their phenotype and functions are regulated by the surrounding micro-environment[28]. As the major immune cells in TME, TAM have reported to have distinct functions during different stages of tumor progression, including tumor killing, angiogenesis, promoting tumor invasion, metastasis and mediating chemoresistance[29–31]. Therefore, it is of particular importance to investigate the mechanism of tumor cells regulating macrophage polarization. In the present study, we identified that MB49-derived exosomes contribute to immunosuppressive phenotype and function of TAM polarization by activating PTEN/AKT/STAT3 signaling pathways, thus promoting bladder cancer growth. Targeting MB49-derived exosomes release could inhibit macrophage M2 differentiation and increase T cells proliferation to impede tumor cell growth, therefore this study sheds new light on bladder cancer treatment.

It is well documented that tumor cells utilized soluble molecules like cytokines or metabolites to induce macrophage polarization in TME[14]. Recently, exosomes carrying genetic information including proteins, miRNAs have been suggested to play an essential role in affecting the functions of target cells by mediating the information transmission and exchange of material[18]. Therefore, tumor cells derived exosomes may contribute to the recruitment and reprogramming of TAM. A recent study reported that exosomes–transmitted long non-coding RNA PTENP1 suppresses bladder cancer progression by suppression PTEN expression[32]. Similarly, in present study, we found MB49-derived exosomes also suppress PTEN expression and induce BMDM M2 differentiation, thus contributing to immunosuppressive TME and facilitating tumor growth. Additionally, inhibition of exosomes generation or release by GW4869 significantly reduces the percentage of M2 macrophage as well as tumor growth in mice subcutaneous tumor model.

Exosomal miRNAs have been demonstrated to interfere with tumor immunity and the microenvironment to regulate cancer progression[26]. As described in our data, 246 miRNAs were identified in MB49-derived exosomes and associated with a variety of cancer related signaling pathways, like PI3K-AKT pathway. Two miRNAs, miR-92a-3p and miR1231-5p, were found to inhibit *Pten* expression and AKT activation. It has been reported miR-92a-3p could promotes the proliferation, migration and invasion of esophageal squamous cell cancer (ESCC) by regulating PTEN[33], but its function on macrophage polarization is still not investigated. In addition, the effect of miR1231-5p in PTEN inhibition and macrophage function needs further investigation. Since AKT activation is required for macrophage M2 polarization[34, 35], in current study we found MB49-derived exosomes could activate both AKT and STAT3/6 pathways to induce macrophage M2 polarization. However, the mechanism of activation of STAT3/6 pathway needs

further exploration. Similarly, it has been demonstrated that bladder cancer cell line T24 derived exosomal miR-21 could also induce THP-1 cell M2 differentiation by activation AKT-STAT3 pathway[22].

An increasing body of evidence has demonstrated that TAM mediated immunosuppressive TME contributes to dampen adaptive anti-tumor immune responses[11]. In current study, we found MB49-derived exosomes could induce upregulation of immunosuppressive cytokines *Il10*, and *Tgfb* in BMDM. Moreover, TAM could also cause metabolic starvation of T cells by the activity of arginase and/or by production of immunosuppressive metabolites by indoleamine 2,3-dioxygenase 1/2 (IDO1/2) pathway[11, 36]. Indeed, we also observed that MB49-derived exosomes stimulated BMDM could hinder both CD4 + and CD8 + T cell proliferation, which can be partially rescued by GW4869 treatment. Furthermore, GW4869 treatment could also increase the percentage of CD4⁺ and CD8⁺ T cells and impede tumor growth in mice subcutaneous tumor model. These results indicate that MB49-derived exosomes reprogram macrophage into immunosuppressive phenotype and function, thus contributing to immunosuppressive TME and facilitating tumor progression.

Conclusion

In summary, this study further confirmed bladder cancer cells derived exosomes contribute to the establishment of immunosuppressive TME and facilitate tumor progression. The exosomal miR-92a-3p and miR-1231-5p may play a vital role in activation of AKT/STAT3/6 signaling pathway and inducing macrophages immunosuppressive differentiation. In addition, hindrance of exosomes generation or release by GW4869 inhibited bladder tumor cell growth by reversing the immunosuppressive TME.

Abbreviations

TAM: Tumor associated macrophages; TME: Tumor microenvironment; PTEN: Phosphate and tension homology deleted on chromosome ten; STAT3: Signal transducer and activator of transcription 3; STAT6: Signal transducer and activator of transcription 6; MIBC: muscle invasive bladder cancer; CD206: Cluster of differentiation 206; CD163: Cluster of differentiation 163; TGF- β : tumor growth factor beta; miR21: microRNA 21; PI3K: Phosphatidylinositol 3-kinase; BMDM: Bone marrow-derived macrophages; IL-10: Interleukin-10; Inos: Inducible nitric oxide synthase.

Declarations

Acknowledgements

Not applicable

Authors' contributions

ZZ and JY conceived this study, generated hypotheses, and designed the experiments. YZ, ZJ, and YZ performed the experiments. YZ, ZJ, and YZ analyzed the data. YZ and ZZ wrote, reviewed, and edited the

manuscript. JY supervised the project. All authors read and approved the final manuscript.

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Availability of data and materials

All data generated or analyzed during this study are included in this article.

Ethics approval and consent to participate

All procedures performed in studies involving animals were reviewed and granted by the Ethical Review Committee of the First Affiliated Hospital of Zhengzhou University.

Consent for publication

All authors have consented to publication of the results presented in this manuscript.

Competing interests

The authors disclose no potential conflicts of interest.

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Figures

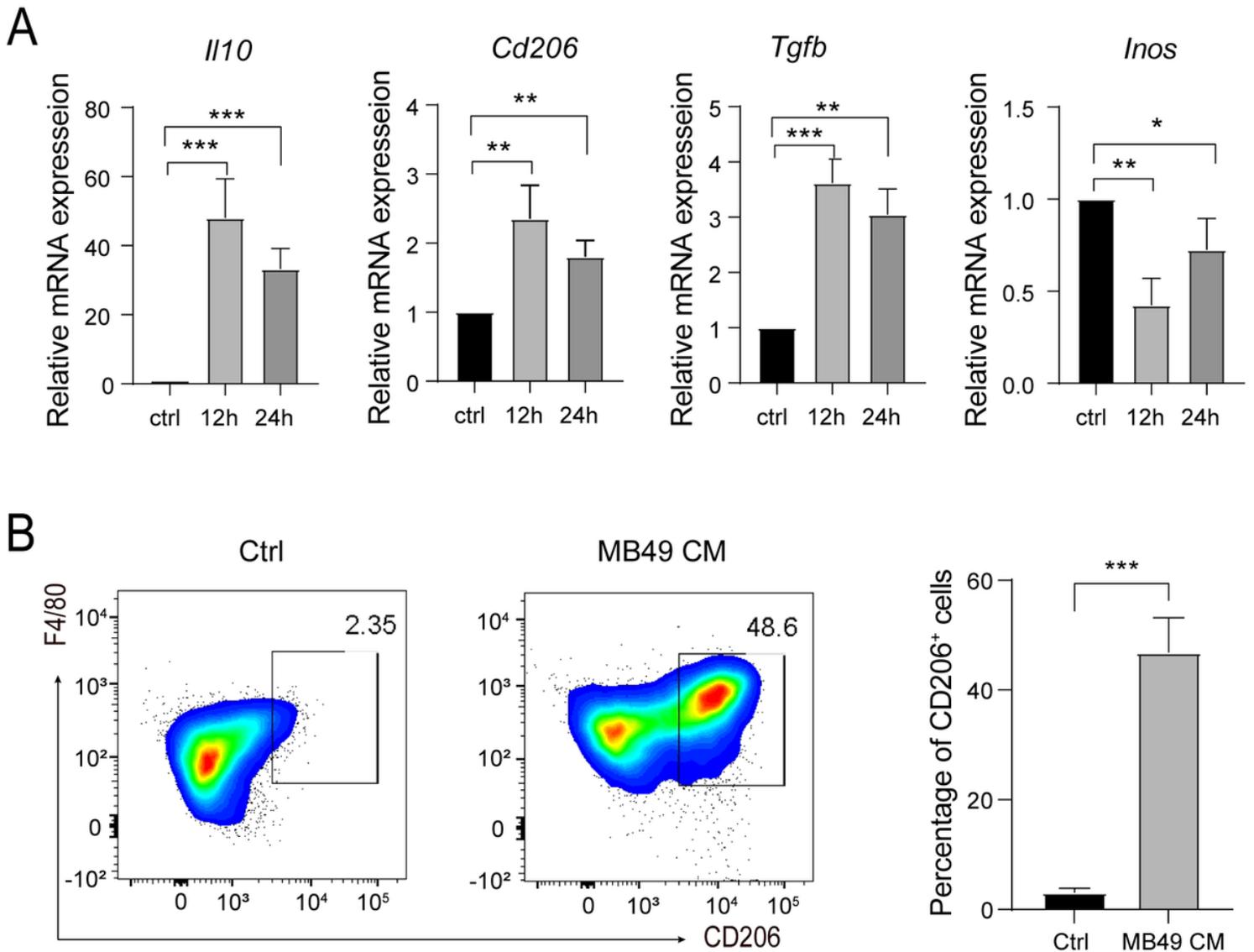


Figure 1

MB49 cells conditioned medium could promote the transformation of BMDM to M2 phenotype. a The expression of *Il-10*, *Cd206*, *Tgfb* and *Inos* was quantified by qRT-PCR in macrophages stimulated with MB49 cells conditioned medium for 12 and 24 h, respectively. Results are present as the means \pm SD of 3 independent experiments. Data were analyzed by one-way ANOVA with multiple comparisons test ($*p < 0.05$, $**p < 0.01$, $***p < 0.0001$). b BMDM were stimulated with MB49 conditioned medium for 48 h, and the proportion of F4/80⁺CD206⁺ cells was detected by flow cytometry. Representative images of each sample are shown. Data were analyzed by t-test (Mean \pm SD, $n=3$, $***P < 0.001$).

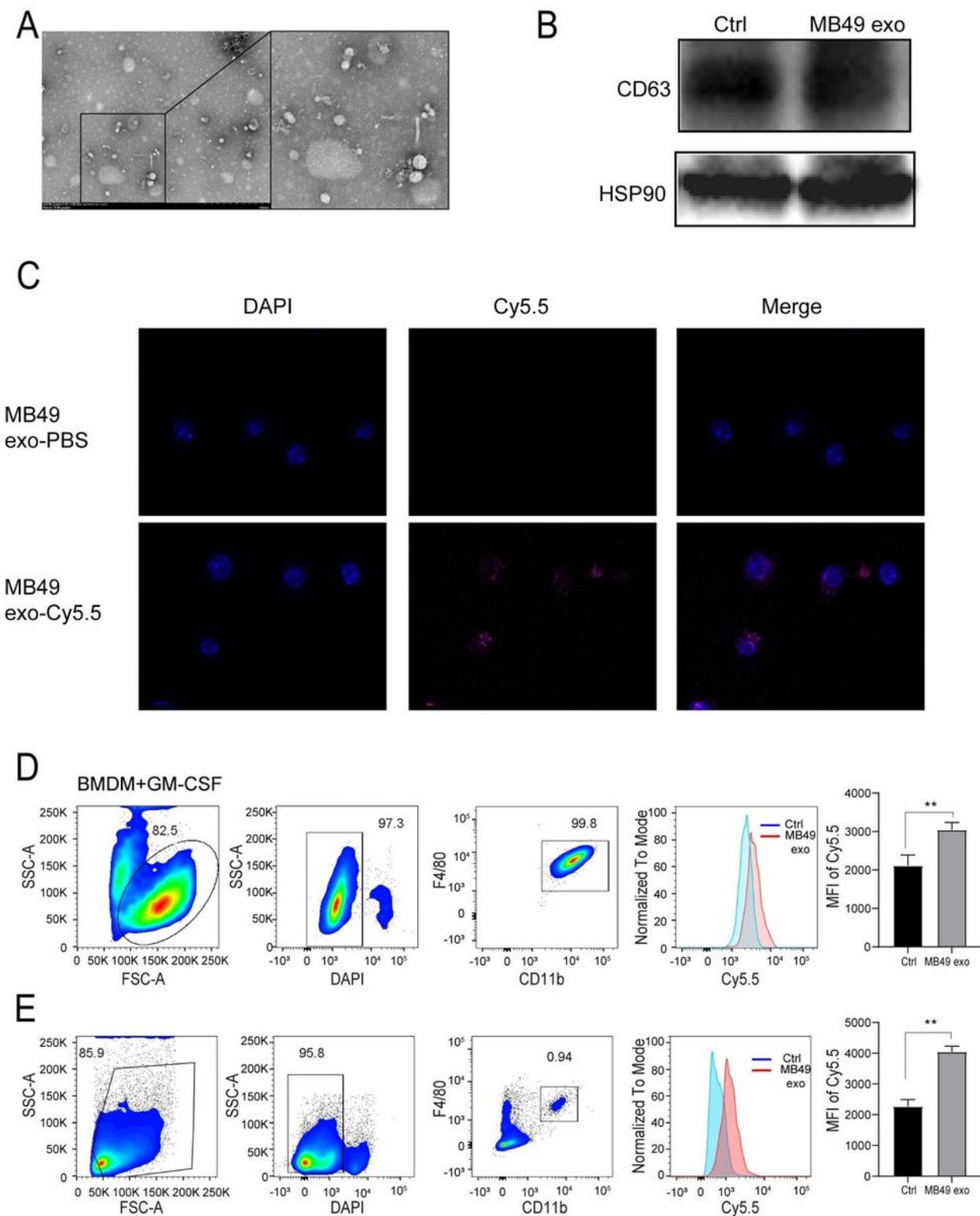


Figure 2

Identification of MB49-derived exosomes and their uptake in macrophages. a The morphology of exosomes isolated from MB49 cells were observed by electron microscope. Shown in the box are typical exosome particles. Scale bar, 500 nm. b The expression of CD63 and HSP90 were detected by Western blot analysis in MB49 exosomes. c Cy5.5-labeled exosomes were co-cultured with BMDM for 3 h, and the uptake of exosomes was observed under a confocal microscope. The representative images were shown.

Scale bar, 10 μ m. d The mean fluorescence intensity (MFI) of Cy5.5-labeled exosomes treated BMDM was measured by flow cytometry, and the control group was treated with PBS-exosomes. e Cy5.5-labeled MB49-derived exosomes or PBS MB49-exosomes were injected into mice via the tail vein. And the MFI of Cy5.5 labeled exosomes in CD45⁺F4/80⁺CD11b⁺ bladder macrophages was analyzed by flow cytometry. Each experiment was repeated three times, and the data were expressed as mean \pm SD, t-test, **P<0.01

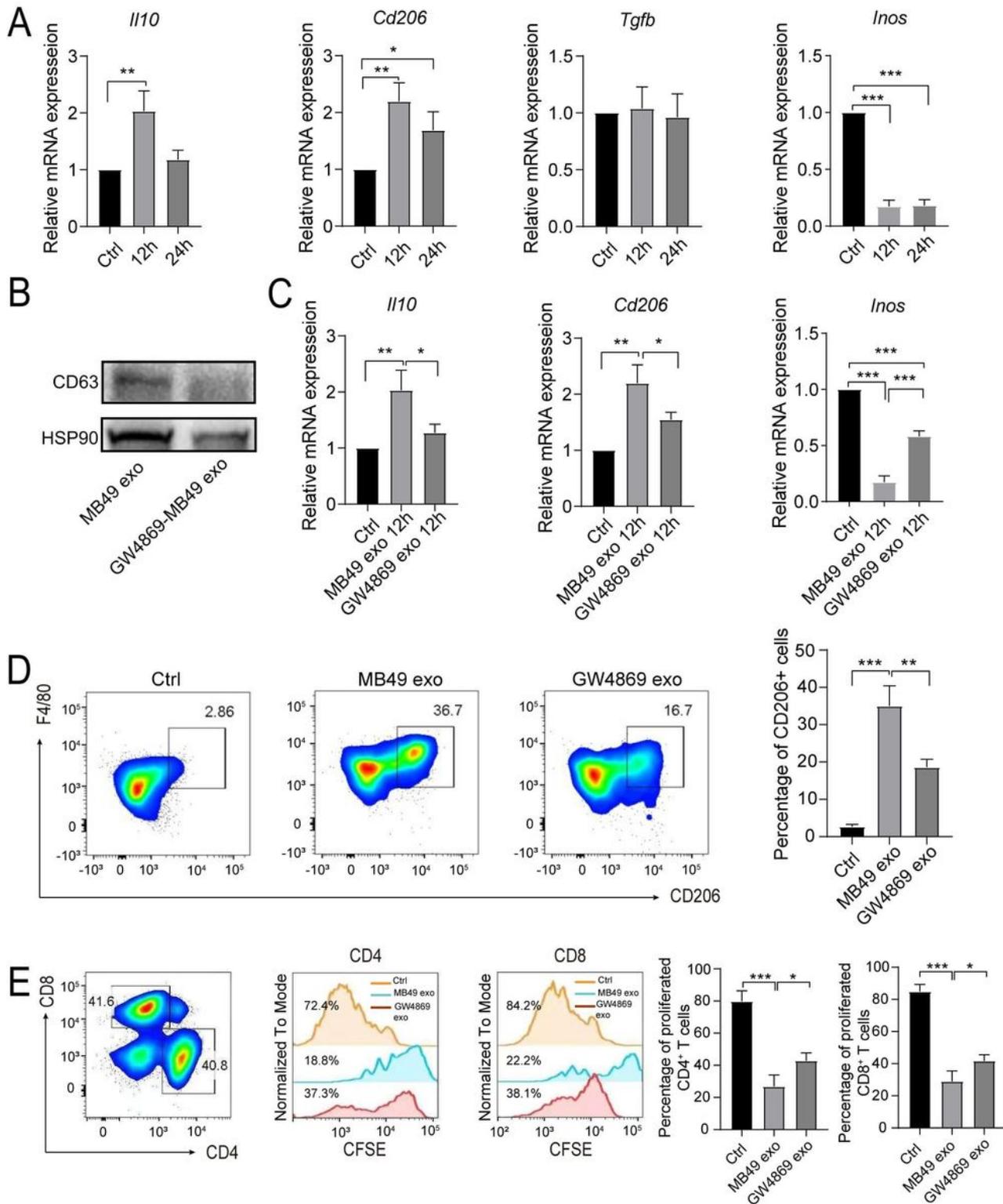


Figure 3

MB49-derived exosomes play a major role in inducing the immunosuppressive function of macrophages. BMDM were treated with MB49-derived exosomes, which were extracted from MB49 CM \pm GW4869, for 12 h or 24 h respectively. a The expression of Il10, Cd206, Tgfb and Inos qRT-PCR was measured by qRT-PCR. b The expression of CD63 and Hsp90 was analyzed by Western blot. c The expression of Il10, Cd206 and Inos was quantified by qRT-PCR. d BMDM were stimulated with exosomes for 48 h and flow cytometry were used to detect the percentage of F4/80+CD206+ macrophages, the representative plot was shown. e The exosome stimulated BMDM were further co-cultured with CFSE labeled splenic T cells for 3 d, the percentage of CFSE+ CD4+ or CD8+ T cells was measured by flow cytometry. The experiments were repeated for 3 times. Data are present as mean \pm SD, one-way ANOVA, *P<0.05; **P<0.01; ***P<0.001.

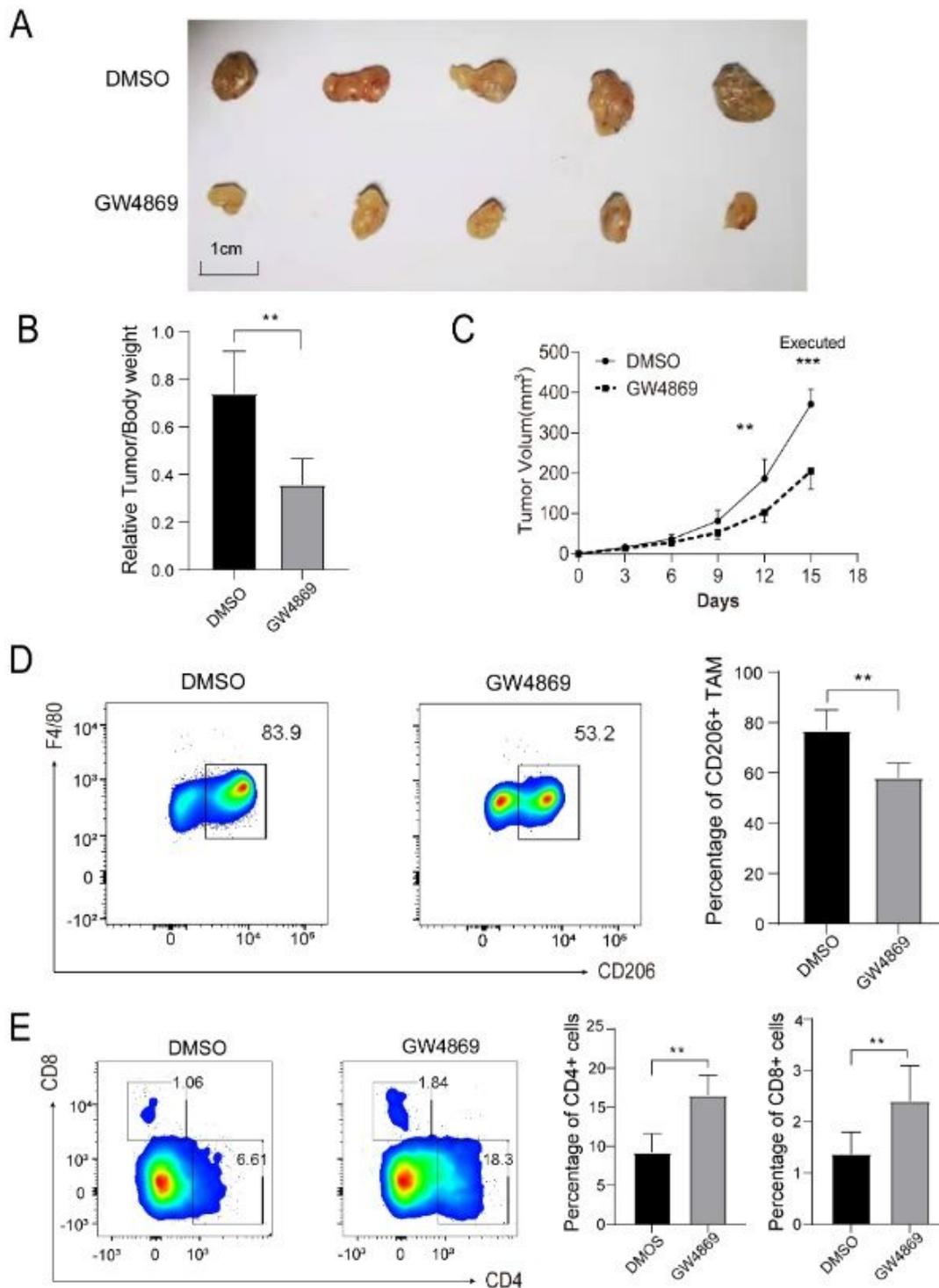


Figure 4

MB49-derived exosomes promote the growth of tumor cells. a-c Total 1×10^6 MB49 cells were subcutaneously injected to mice, DMSO or GW4869 (2.5 $\mu\text{g/g}$) were intraperitoneally injected every three days ($n=10$, 5 mice in each group). a, b Mice were sacrificed at 15 d, and the ration of tumor/body weight were measured. c The tumor volume was measured every three days and the growth curve was shown. d,

e The proportion of CD45+F4/80+CD206+ macrophage and CD4+/CD8+ T cells were measured by flow cytometry in tumors (n=5). T-test; Mean \pm SD, *P<0.05; **P<0.01; ***P<0.001.

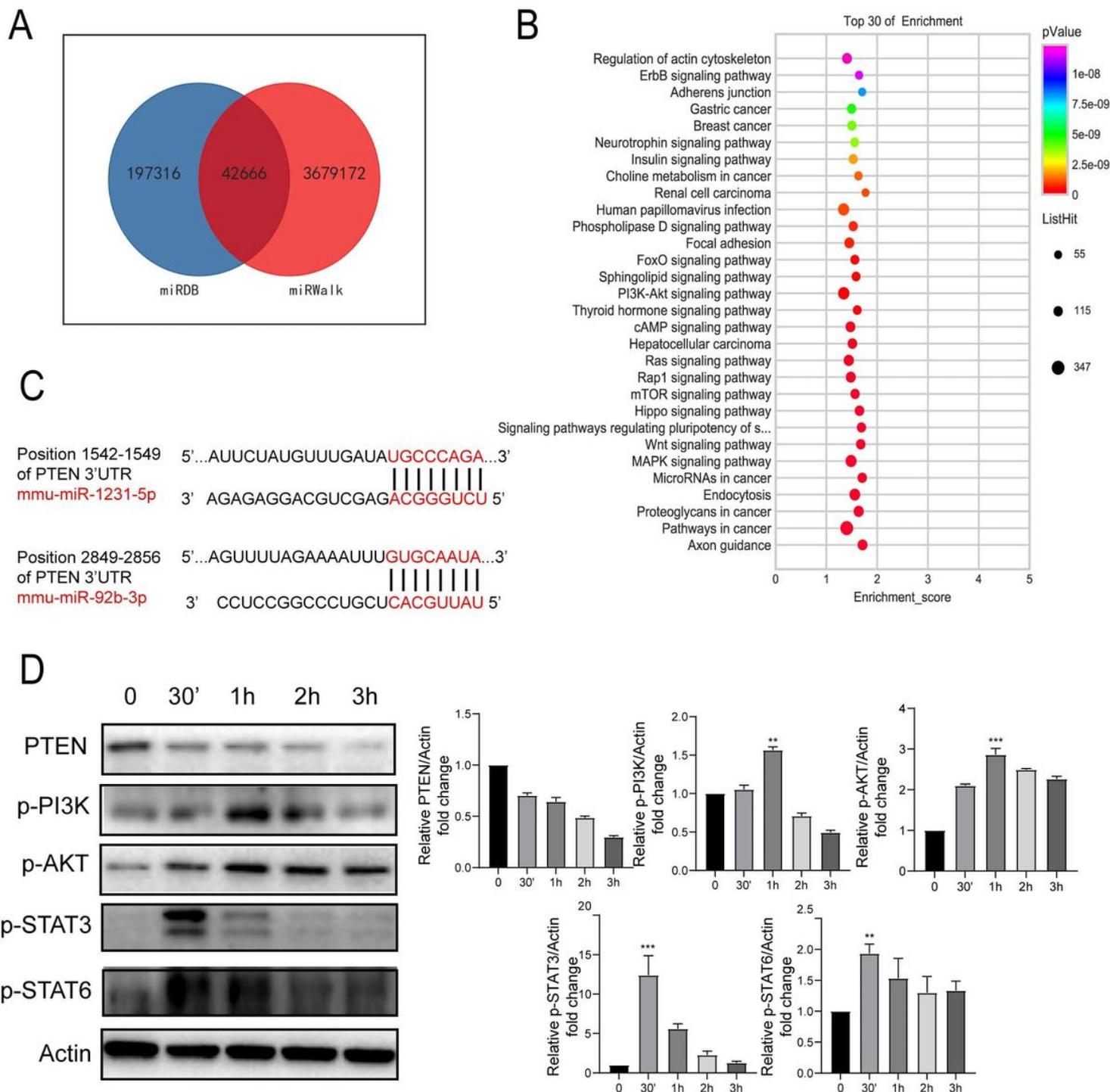


Figure 5

MB49 exosomes activate PTEN/AKT/STAT3/6 signaling pathway in BMDM. a 42666 miRNAs were identified by microarray analysis in MB49-derived exosomes. b Bubble plot of GO enrichment analysis of target genes of MB4- exosomal miRNAs. c The binding sites between miRNAs and Pten were shown. d BMDM were treated with MB49-derived exosomes for indicated time points, and the expression of p-PTEN, p-PI3K, p-AKT, p-STAT3 and p-STAT6 was analyzed by Western blot. The experiment was

performed at least three independent times and results were presented as mean \pm SD. one-way ANOVA, **P<0.01; ***P<0.001.

Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [SupplementaryTable1.docx](#)
- [SupplementaryTable2.docx](#)