Antitumor effect of recombinant *Mycobacterium smegmatis* expressing MAGEA3 and SSX2 fusion proteins

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Abstract. Mycobacterium smegmatis (M. smegmatis), which is a nonpathogenic and fast-growing mycobacterium, is a potential vaccine vector capable of expressing heterologous antigens. Spontaneous humoral and cellular immune responses have been demonstrated against cancer/testis antigens (CTA), including melanoma-associated antigen A (MAGEA) and SSX. In the present study, recombinant plasmids expressing MAGEA3 and SSX2 were constructed. The recombinant plasmids were transferred into M. smegmatis to generate the novel antitumor DNA vaccine. As MAGEA3 and SSX2 were in different ligation sequences, the two DNA vaccines were recombinant M. smegmatis MAGEA3-SSX2 (rM.S-MS) and recombinant M. smegmatis SSX2-MAGEA3 (rM.S-SM), respectively. The expression levels of Fusion proteins were assessed by western blotting. BALB/c mice were immunized with rM.S and western blot analysis was used to determine whether antibodies against MAGEA3 or SSX2 were produced in immunized mice. EC9706 cells were inoculated into BALB/c nude mice and the mice were maintained until an obvious visible tumor appeared on the back. Subsequently, the blood from the rM.S immunized BALB/c mice was injected into the BALB/c nude mice via the tail vein. In order to evaluate the antitumor effect of the vaccines, tumor volume and weight were measured 5 to 21 days after injection. Mice were euthanized on day 21 of tumor growth, and the tumor was dissected and weighed. The two fusion proteins were expressed in the rM.S and the specific fusion protein antibodies

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were expressed in the blood of immunized BALB/c mice. The tumor volumes and weight in the recombinant *M. smegmatis* MAGEA3 (rM.S-M) and recombinant *M. smegmatis* SSX2 (rM.S-S) groups were significantly reduced compared with the control group. Furthermore, the decrease in tumor volumes and weight in the rM.S-MS and rM.S-SM groups was more severe than in the rM.S-M or rM.S-S groups. There was no significant difference in the antitumor effect of the rM.S-MS and rM.S-SM groups. The present findings suggest that this rM.S may be a potential candidate therapeutic vaccine for the treatment of cancer.

Introduction

Immunotherapy is a new avenue of cancer treatment for a range of different cancer types. It is now understood that the immune system is capable of recognizing and eliminating cancer cells, but tumors evade and suppress host immune responses and therefore persist and spread (1-3). During the past few decades, anticancer immunotherapy has evolved from a promising therapeutic option to a robust clinical reality. Many immunotherapeutic regimens are now approved for use in cancer patients, and many others are being investigated as standalone therapeutic interventions or combined with conventional treatments in clinical studies.

Mycobacterium smegmatis (*M. smegmatis*) is a fast-growing saprophytic environmental bacterium, which is a non-pathogenic and commensal genus (4,5). *M. smegmatis* also has a number of properties such as growth rapidily and can be transformed effectively with many genes, that renders it an ideal vaccine vector. Further more, *M. smegmatis* is reported to activate dendritic cells and trigger CD8-mediated immune responses, and immunization with rM.S can generate more durable memory T cells than intramuscular DNA vaccination (6,7). These findings indicate the potential role of mycobacteria as recombinant vaccine delivery vector.

Immunogenic target antigen is another crucial element for developing a successful vaccine. The melanoma-associated antigen A3 (MAGEA3) is a member of the large cancer/testis antigens (CTA), which are frequently aberrantly expressed in a wide range of cancer (8-12). MAGEA gene family is regarded as a promising target of specific immunotherapy because MAGEA is expressed mainly in cancers that have acquired maliganat phenotypes and contribute towards malignancy (13). MAGEA3 is an tumor antigenic nonapeptide that is identified in various tumors and associated with a broad set of HLA (human MHC locus) molecules (14). Consequently, MAGEA3 antigen is a genuinely selective target for tumor-specific active immunotherapy.

It is well known that novel and effective adjuvants can elicit stronger cellular and humoral adaptive immune responses to antigenic targets. The expression of a particular CTA is limited to only a subset of patients with a particular tumor type; therefore, for human application, this is too weak to induce a substantial response against difficult antigens. In order to expand the number of patients and tumor types that can be treated, it is necessary to expand the repertoire of antigens by this approach. We developed another CTA, SSX2 (synovial sarcoma X breakpoint 2), which is the primary member of the SSX family expressed in different kinds of cancers inculding prostate, lung, breast and multiple myeloma and pancreatic cancer (15-19). SSX2 gene encodes for the human tumor-specific antigen HOM-MEL-40, which is an immunogenic protein known to trigger spontaneous antibody responses (20). The SSX2 protein can induce spontaneous immune responses. Therefore, the development of vectors expressing SSX2 opens up a wide array of possibilities in the immunotherapy of cancer.

In this study, we designed two fusion proteins with different ligation sequences, MAGEA3-SSX2 and SSX2-MAGEA3, from *M. smegmatis* for tumor immunotherapy and detected their tumor therapeutic effect by mice tumor-burdened experiments.

Materials and methods

Bacterial strains and growth conditions. The *M. smegmatis* strain MC²155 was supplied by Yinlan Bo's Laboratory at the Fourth Military Medical University (Xi'an, China). *M. smegmatis* cultures were grown in 7H10 solid medium (7H10 solid medium contained 3 ml/l glycerin, 0.5 g/l Tween-80, 100 ml/l OADC and 19/l middle brook 7H10 agar powder) and incubated at 37°C for 2-3 days; the medium was supplemented with hygromycin (50 ng/ml) when selecting for the recombinant plasmid. *Escherichia coli* cultures were grown in Luriae-Bertani (LB) broth or plates (LB broth contained 10 g/l trypeptone; 15 g/l NaCl; 5 g/l yeast extract; and 15 g/l agar powder) and incubated at 37°C overnight; the media were supplemented with ampicillin (100 μ g/ml) when selecting for the recombinant plasmid.

Plasmid and strain construction. The pDE22 vector was supplied by Yinlan Bo's laboratory at the Fourth Military Medical University. The *E. coli* strain DH5- α was purchased from MBI Fermentas (Vilnius, Lithuania). The pUC57 vector was purchased from Tiangen (Beijing, China). Taq DNA polymerase and Pst I endonuclease were obtained from Takara Biotechnology Co., Ltd. (Dalian, China). BamHI endonuclease, ClaI endonuclease, EcoRV endonuclease and T4 DNA ligase were obtained from MBI Fermentas (Burlington,

ON, Canada). All other media components and chemicals used were of the highest purity grade available commercially from Beijing Chemical Plant, China.

Splicing overlap extension polymerase chain reaction (SOE-PCR) primers were synthesized by Shanghai Bioengineering Company (Shanghai, China). The MAGEA3 gene was cloned from DNA of EC9706 cell via PCR using the primer pair: Sense primer 5'-GCCGATATCATGCCT CTTGAGCAGAGGAGTC-3' and antisense primer 5'-GCT GCCGCCGCCGCCGCTGCC-3'. The SSX2 gene was cloned from DNA of EC9706 cell via PCR using the primer pair: Sense primer 5'-GCCGATATCATGAACGGAGAC GACGCCTTTC-3' and antisense primer 5'-GCTGCCGCC GCCGCCGCTGCC-3'. The cloned genes MAGEA3 and SSX2 were constructed from two kinds of different connection sequence gene fragments, MAGEA3-SSX2 (MS) and SSX2-MAGEA3 (SM). The MAGEA3-SSX2 fragment was amplified using the primer pair: Sense primer 5'-CGGCGG CGGCGGCAGCATGCCTCTTGAGCAGAG-3' and antisense primer 5'-CCATCGATTTACTCGTCATCTTCCTCA GGG-3', and the SSX2-MAGEA3 fragment was amplified using the primer pair: Sense primer 5'-CGGCGGCGGCGG CAGCATGAACGGAGACGACG-3' and antisense primer 5'-CCATCGATTCACTCTTCCCCCTCTCTCAAA-3'. The MAGEA3-SSX2 and SSX2-MAGEA3 fusion expression cassettes were generated using the gap repair method as above, and a linker designed was used to maintain the correct biological activity of both MAGEA3 and SSX2. A verified clone with the correct sequence (AuGCT Biotechnology, Beijing, China) was transferred into a pDE22 cloning vector, then cut with the appropriate restriction endonucleases and inserted in the E. coli-mycobacterium shuttle plasmid pDE22 construct. Plasmid DNA was introduced into M. smegmatis by electroporation using standard techniques (21) to generate the rM.S strain expressing the two kinds of fusion protein MAGEA3-SSX2 and SSX2-MAGEA3.

Western blot analysis. To monitor the expression of the M. smegmatis MAGEA3 and SSX2 transgenes, the rMS strains were grown in 7H10/ADC until mid-log phase and blocked with 10% bovine serum albumin. The lysate of grown rM.S was fractionated on 20% SDS-polyacrylamide gels and blotted onto nitrocellulose filters (Invitrogen; Thermo Fisher Scientific, Inc., Waltham, MA, USA). The membranes were blocked with 5% non-fat milk and incubated with a rabbit anti-human MAGEA3 antibody (Abgent, Inc., San Diego, CA, USA) at a dilution of 1:100 and a rabbit anti-human SSX2 antibody (Abcam, Cambridge, MA, USA) at a dilution of 1:200 or a mouse anti-\beta-actin monoclonal antibody at a dilution of 1:2,000 (Sigma-Aldrich; Merck KGaA, Darmstadt, Germany). The membranes were subsequently incubated with a goat anti-mouse or anti-rabbit horseradish peroxidase secondary antibody (Sigma-Aldrich; Merck KGaA). The protein complexes were detected using enhanced chemiluminescence reagents (Pierce; Thermo Fisher Scientific, Inc.).

The production of antibodies against MAGEA3 and SSX2 in the blood of immunized mice was determined using the purified MAGEA3 protein (Abnova, Walnut, CA, USA) or SSX2 protein (Abnova) separated by SDS-PAGE. All experiments were carried out at least three times. Immunization of mice. Seven-week-old and specific pathogen-free male BALB/c mice provided by the laboratory animal center of the Fourth Military Medical University were used for immunogenicity studies. All animal protocols were reviewed and approved by the Institutional Animal Care and Use Committee of the Fourth Military Medical University (ID11013). Mice were randomly divided into six groups (6 per group) to receive subcutaneous injections as follows: Normal control group (NC) received 0.2 ml saline/mouse, M. smegmatis group infected with the M. smegmatis strain and received 1x10⁶ CFU empty pDE22 vector/mouse via the tail vein, recombinant M. smegmatis MAGEA3 (rM.S-M) infected with the M. smegmatis strain and transfected with pDE22-MAGEA3 at a dose of 1x10⁶ CFU/mouse, recombinant M. smegmatis SSX2 (rM.S-S) infected with the M. smegmatis strain and transfected with pDE22-SSX2 at a dose of 1x10⁶ CFU/mouse, recombinant M. smegmatis MAGEA3-SSX2 (rM.S-MS) infected with the M. smegmatis strain and transfected with pDE22-MAGEA3-SSX2 at a dose of 1x10⁶ CFU/mouse, recombinant M. smegmatis SSX2-MAGEA3 (rM.S-SM) infected with the *M. smegmatis* strain transfected with pDE22-SSX2-MAGEA3 at a dose of 1x10⁶ CFU/mouse. Mice were immunized once every 5 days with rM.S for a total of three times.

Immunotherapy in the tumor-bearing mice. Seven-week-old, specific pathogen-free male BALB/c nude mice provided by the laboratory animal center of the Fourth Military Medical University were housed and monitored in a specific pathogen-free environment with sterile food and water in our animal facility. The human esophageal EC9706 cancer cell line, which was MAGEA3 and SSX2 double-positive cancer cell, was maintained in culture and prepared for injection as previously described (22). EC9706 tumor cells were cultured and inoculated subcutaneously into one site on the back surface of each BALB/c nude mouse at a concentration of 1x10⁶ cells. Mice were cultured and observed until an obvious visible tumor appeared on the mouse back. Tumor-bearing mice were randomly divided into six groups with 6 mice each. The mice received the following different treatment: Normal control group (NC) receiving 100 µl 0.9% saline/mouse, M. smegmatis group, rM.S-M group, rM.S-S group, rM.S-MS group and rM.S-SM group infected with the blood of the immunized mice from the same groups as the above via the tail vein at a dose of 100 μ l/mouse, respectively.

In the present study, all data collection was completed from 5 to 21 days after injection. The sizes of tumors were measured using a digital caliper in three dimensions (L x W x H). The height of the tumors was determined by physically grasping the tumor by its base. Tumor volume was calculated using the following equation: Tumor Volume=(π ·H (H2 + 3·(L+W/2)))/6. Mice were euthanized by inhalation of CO₂ gas on day 21 of tumor growth. Tumors were dissected and weighed.

Statistical analysis. Data are presented as the mean \pm standard deviation (SD) from at least three independent experiments. Statistical analysis was performed using SPSS16.0 (SPSS, Inc., Chicago, IL, USA). Student's t-test and χ^2 test were used to analyze the difference between different groups. The



Figure 1. The protein expression of MAGEA3 and SSX2 in EC9706 cancer cells. Western blot assay was performed to detect the MAGEA3 and SSX2 protein expression in human esophageal EC9706 cancer cell line. β -actin is shown as a loading control. MAGEA3, melanoma-associated antigen A3.

comparison of multiple groups was carried out using one-way ANOVA followed by Tukey's post hoc test. P<0.05 was considered to indicate a statistically significant difference.

Results

MAGEA3 and SSX2 were double-positive expressed in human esophageal EC9706 cancer cell line. Firstly, we detected the protein expression level of MAGEA3 and SSX2 in EC9706 cells, and the results showed that both MAGEA3 and SSX2 were highly expressed (Fig. 1).

Construction of rM.S strains and the expression of two fusion proteins. The optimized MAGEA3-SSX2 and SSX2-MAGEA3 fusion segments were synthesized in the pUC57 vector by Shanghai Generay Biotechnology Co., Ltd. (Shanghai, China) in *Bam*HI and *Eco*RV restriction sites at the 5'and 3'ends. Then, the generated MAGEA3-SSX2 and SSX2-MAGEA3 fusion segments, along with the vector, were digested with *Pst*I and *Cla*I restriction enzymes and ligated into pDE22 in the corresponding enzyme site at the 5'and 3' ends. Agarose gel electrophoresis of the *Pst*I and *Cla*I digested plasmid showed the expected bands of 1,596 bp, representing the insert and pDE22, respectively (Fig. 2A). Also, sequencing showed that the target DNA was inserted correctly into the multi-cloning site of pDE22.

The pDE22-based constructs were electroporated into the fast-growing, non-pathogenic *M. smegmatis* strain to obtain a rM.S strain that could be easily manipulated in most laboratories. Following induction, MAGEA3-SSX2 and SSX2-MAGEA3 were expressed and corresponded to their predicted molecular masses of 67 kDa (MAGEA3 approximately 46 kDa and SSX2 approximately 21 kDa) on an SDS-PAGE gel (Fig. 2B). Western blotting showed that the MAGEA3-SSX2 and SSX2-MAGEA3 were recognized by the purchased rabbit anti-human SSX2 polyclonal antibody and anti-human MAGEA3 polyclonal antibody (Fig. 2C).

Antibody production in immunized mice. Mice were immunized once every 5 days with rM.S for a total of three times, as described in Methods. The expression of fusion protein-specific antibody levels after final vaccination is shown in Fig. 3. Immunized mice blood fusion protein-specific antibodies levels were detected by Western blot. Significantly specific specific antibodies levels were observed in mice of the groups vaccinated with the rM.S compared to the control group. In contrast, no specific antibodies were expressed between the control group and the *M. smegmatis* group.

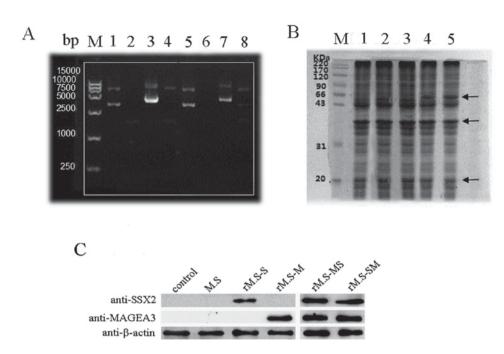


Figure 2. Identification of successfully constructed rM.S vaccine and detection the expression of MAGEA3-SSX2 (MS) and SSX2-MAGEA3 (SM) fusion proteins in the rM.S. (A) Agarose gel electrophoresis for the verification of recombinant pDE22 vector. Lane M, DNA marker; line 1, the empty vector of pDE22; line 2, rM.S; line 3, pDE22-MS; line 4, pDE22-MS digested with *Eco*RV and *Cla*1; line 5, the empty vector of pDE22; line 6, rM.S-SM; line 7, pDE22-SM; line 8, pDE22-SM digested with *Eco*RV and *Cla*1. (B) SDS-PAGE showed the expression of the MAGEA3 and SSX2 transgenes in the rM.S. Lane M, protein marker; line 1, *M. smegmatis*; line 2, MAGEA3-*M. smegmatis*; line 3, SSX2-*M. smegmatis*; line 4, MAGEA3-SSX2-rM.S; line 5, SSX2-MAGEA3-rM.S. MAGEA3-SSX2 and SSX2-MAGEA3 predicted molecular masses of 67 kDa (MAGEA3 approximately 46 kDa and SSX2 approximately 21 kDa). (C) Western blot analysis of MAGEA3 and SSX2 transgenes expression in the rM.S. MAGEA3, melanoma-associated antigen A3. rM.S, recombinant M. smegmatis; rM.S-MS, recombinant M. smegmatis-SSX2; rM.S-SM, recombinant

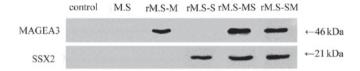


Figure 3. Western blot analysis of specific fusion protein antibody in immunized mouse blood. Mice were immunized once every 5 days with rM.S for a total of three times. The expression of blood fusion protein-specific antibody levels after final vaccination was determined using western blot analysis. MAGEA3, melanoma-associated antigen A3; rM.S, recombinant *M. smegmatis*; rM.S-MS, recombinant *M. smegmatis* MAGEA3-SSX2; rM.S-SM, recombinant *M. smegmatis* SSX2-MAGEA3; rM.S-S, recombinant *M. smegmatis*-SSX2; rM.S-M, recombinant *M. smegmatis*-MAGEA3.

The antitumor effect of the rM.S-MS and rM.S-SM was better than that of the single rM.S-M or rM.S-S. To establish the tumor-bearing mouse model, mice were injected using 1x10⁶ EC9706 cells into one site of the ventral surface. Approximately 5 days later, the back surface of every mouse formed a macroscopic and palpable tumor. Following the detection of a palpable tumor, mice were treated daily by rM.S injection with 100 µl/mouse or saline control. The rM.S-treated tumor volumes were lower compared with the control group (Fig. 4A). As shown in Fig. 3, rM.S-M and rM.S-S-stimulated mice group showed a slight decrease in tumor volume compared to the decrease in tumor volume of the rM.S-MS- or rM.S-SM-stimulated mice groups. There was no significant difference in tumor volume between rM.S-MS or rM.S-SM mice. This was accompanied by a significant reduction in the final tumor weight in the rM.S-treated mice. Tumors were dissected and weighed on day 21 of tumor growth (Fig. 4B). As shown in Fig. 3, thte rM.S-M- and rM.S-S-stimulated mice groups showed a slight decrease in tumor weight compared to the rM.S-MS- or rM.S-SM-stimulated mice groups. There was no significant difference in tumor weight between rM.S-MS or rM.S-SM mice. The result shows that MAGEA3 and SSX2 rM.S are better than the single MAGEA3 *M. smegmatis* or single SSX2 *M. smegmatis* for the demonstration of an anti-tumor effect. In addition, there was no significant difference in the antitumor effect of rM.S-MS or rM.S-SM.

Discussion

Mycobacterium smegmatis is a nonpathogenic species of the genus Mycobacterium which is easily manipulated to produce recombinant bacteria. Consequently, it is widely used as a live vaccine against cancer (23,24). rM.S has been used to express various proteins in studies. A particular rM.S vaccine expressing a fusion protein containing ESAT-6 and CFP10 induced higher humoral and cellular immunity than the M. bovis BCG vaccination in a mouse model (25). Similarly, Xu et al reported that fusion protein Ag85A-IL17A was expressed in rM.S vaccine which attenuated allergic airwat inflammation (26). Here, we demonstrated two kinds of novel vaccine vectors: rM.S-MS and rM.S-SM. rM.S-MS is a recombinant non-pathogenic M. smegmatis MC²155 expressing the MAGEA3-SSX2 fusion protein; rM.S-SM is a rM.S expressing the SSX2-MAGEA3 fusion protein. Moreover, the *M. smegmatis* MC²155 strain has been shown to be nonpathogenic following intravenous infections of SCID mice (27); we also verified the treatment effect

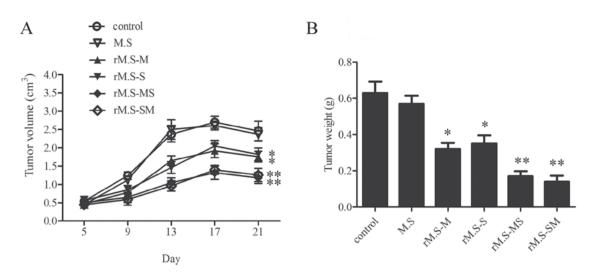


Figure 4. rM.S inhibits the esophageal tumor growth in an esophageal cancer mouse model. (A) Tumor volume was calculated for the 5 different time points and the rM.S treated tumor volumes were lower than those of the control group. (B) Tumors were dissected and weighed on day 21 of tumor growth. Data presented are mean \pm standard deviation. *P<0.05 vs. control or MS; **P<0.05 vs. rM.S-M or rM.S-S. rM.S, recombinant *M. smegmatis*; rM.S-MS, recombinant *M. smegmatis*; rM.S-MS, recombinant *M. smegmatis* SSX2-MAGEA3; rM.S-S, recombinant *M. smegmatis*-SSX2; rM.S-M, recombinant *M. smegmatis*-MAGEA3.

of tumors in the tumor-bearing mouse model through vaccine with recombinant strains.

We have shown that our recombinant vaccines rM.S-MS and rM.S-SM are capable of inducing a specific immune response in two different vaccination schemes (Fig. 2). This response probably reflects the immunogenicity of the recombinant fusion protein when used in this vaccine vector. The results of our group and others have confirmed that MAGEA3 and SSX2 tumor antigens with immunogenic and antigenic properties are represented by the epitopes expressed in the recombinant fusion protein (28-32). Both MAGEA3 and SSX2 belong to the group of cancer-testis antigens (CTA); CTA are immunogenic antigens with an expression that is largely restricted to testicular germ cells and a variety of malignancies, making them attractive targets for cancer immunotherapy (11,33). The cancer-testis-X genes have been the principal targets of developing immunotherapies (33). MAGEA3 and SSX2 are all immunogenic CTA that have been shown to elicit coordinated humoral and cell-mediated immune responses (33,34).

MAGEA3 is one of the best-characterized tumor antigens. Due to its tumor-restricted expression pattern and its recognition by both cytotoxic and helper T cells, it constitutes a promising tumor antigen for anticancer immunotherapy. Roeder et al demonstrated that MAGEA3 is a frequent tumor antigen of metastasized melanoma (29). Vansteenkiste et al demonstrated that MAGEA3 cancer immunotherapy is an active immunotherapy that has been evaluated in NSCLC (35). SSX gene products are expressed in tumors of different histological types and can be recognized by tumor-reactive CTLs from cancer patients. The immunogenicity of SSX-2 has been previously corroborated by detection of specific humoral and CD8+ T cell responses in cancer patients (36). Ayyoub et al also demonstrated that an SSX2-derived immunodominant T cell epitope is recognized by CD4+ T cells from melanoma patients in association with HLA-DR (37,38). This is because MAGEA3 and SSX2 are able to trigger an immune response in the tumor. The aim of our study is to build a fusion protein composed of MAGEA3 and SSX2 and verify whether the fusion proteins' antitumor immune responses are enhanced by either of the two. In the present study, two fusion proteins were amplified in M. smegmatis. ELISA was used to detect whether anti-fusion protein antibodies are produced or not in vivo in immunized mice after the injection of rM.S vaccine. In Fig. 2, rM.S-MS group mice blood and the rM.S-SM group mice blood are able to produce both anti-MAGEA3 and anti-SSX2 antibodies. There was no significant difference in antibody concentration between the two groups. However, the antibody concentration of the rM.S-MS and rM.S-SM groups was significantly increased compared with the rM.S-M group and rM.S-S group, which only produce single protein antibodies. The results are in line with our expectations. Multiple-antigen combination tests may be more useful for the development of diagnostic antibody tests because of many antigens inducing serological responses.

The BALB/c mouse is the animal most commonly used as an *in vivo* model for *M. smegmatis* infection and constructed tumor-bearing mice model. The human esophageal EC9706 cancer cell line is a MAGEA3 and SSX2 double-positive tumor. In this study, BALB/c mice were immunized with rM.S-M, rM.S-S, rM.S-MS and rM.S-SM to provide experimental data to evaluate the effect of these proteins in M. *smegmatis*; blood was obtained containing the corresponding antibody to the treatment of tumor-bearing mice. According to our data after immunization, we extracted blood from the mice with the best antibody concentration for the following experiment. Treatment with the rM.S inhibits esophageal tumor growth *in vivo* in mice. The rM.S treated tumor volumes and weight were significantly reduced compared with the control group (Fig. 3).

In conclusion, in the current study we constructed rM.S vaccines, rM.S-MS and rM.S-SM, and demonstrated the immunogenicity of the vaccine. In addition, we demonstrated that the rM.S vaccine can activate the immune system and enhance

the antitumor effect. The antitumor effect of the rM.S-MS and rM.S-SM is better than that of the single rM.S-M or rM.S-S.

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Availability of data and materials

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

Authors' contributions

WJ and YX conceived and designed the experiments. WJ, XL, JK, YL and YB performed the experiments. WJ and XL analyzed the data. WJ, XL and JK contributed reagents/materials/analysis tools. WJ and YX wrote the paper.

Ethics approval and consent to participate

All animal protocols were reviewed and approved by the Institutional Animal Care and Use Committee of the Fourth Military Medical University (ID11013).

Patient consent for publication

Not applicable.

Competing interests

The authors declare that they have no competing interests.

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