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Establishment of animal models to analyze the kinetics and distribution of human tumor antigen-specific CD8$^+$ T cells

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ABSTRACT

Many patients develop tumor antigen-specific T cell responses detectable in peripheral blood mononuclear cells (PBMCs) following cancer vaccine. However, measurable tumor regression is observed in a limited number of patients receiving cancer vaccines. There is a need to re-evaluate systemically the immune responses induced by cancer vaccines. Here, we established animal models targeting two human cancer/testis antigens, NY-ESO-1 and MAGE-A4. Cytotoxic T lymphocyte (CTL) epitopes of these antigens were investigated by immunizing BALB/c mice with plasmids encoding the entire sequences of NY-ESO-1 or MAGE-A4. CD8$^+$ T cells specific for NY-ESO-1 or MAGE-A4 were able to be detected by ELISPOT assays using antigen presenting cells pulsed with overlapping peptides covering the whole protein, indicating the high immunogenicity of these antigens in mice. Truncation of these peptides revealed that NY-ESO-1-specific CD8$^+$ T cells recognized D^d-restricted 8mer peptides, NY-ESO-1$_{81\text{-}88}$. MAGE-A4-specific CD8$^+$ T cells recognized D^d-restricted 9mer peptides, MAGE-A4$_{265\text{-}273}$. MHC/peptide tetramers allowed us to analyze the kinetics and distribution of the antigen-specific immune responses, and we found that stronger antigen-specific CD8$^+$ T cell responses were required for more effective anti-tumor activity. Taken together, these animal models are valuable for evaluation of immune responses and optimization of the efficacy of cancer vaccines.

1. Introduction

A number of cancer vaccine strategies targeting tumor antigens recognized by the human immune system have been tested [1–3]. While many of these cancer vaccines elicited measurable immune responses detectable in peripheral blood mononuclear cells (PBMCs), only a subset of treated patients experienced clinical benefits, such as tumor regression [4,5]. Because of the weak clinical effectiveness of currently available cancer vaccines, not only new immunogenic antigens, effective adjuvant formulations, vectors or vaccination methods but also new methodologies to evaluate efficacy of cancer vaccines are required.

NY-ESO-1, a germ line cell protein detected by SEREX (serological identification of antigens by recombinant expression cloning) using the serum of an esophageal cancer patient, is often expressed by cancer cells, but not by normal somatic cells [3,6]. This ideal expression pattern facilitated the study of this antigen; including immuno-monitoring of cancer patients with NY-ESO-1-expressing tumors and clinical trials that focused on NY-ESO-1 [3]. While these studies have revealed that a number of different cancer vaccines, including short and overlapping peptides, protein, viral vectors and DNA, resulted in development of measurable immune responses,

Abbreviations: APC, antigen presenting cells; CTL, cytotoxic T lymphocyte; dLN, draining lymph node; ELISPOT assay, enzyme-linked immunospot assay; IFN, interferon; mAb, monoclonal antibody; PBMC, peripheral blood mononuclear cells.

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correlations between immunological and clinical responses were often week or difficult to observe [3].

MAGE-A4, another cancer/testis (CT) antigen, elicits MAGE-A4-specific $CD4^+$ and $CD8^+$ T cell responses in some patients with MAGE-A4-expressing cancers, indicating that MAGE-A4 is also an immunogenic protein [7–9]. We have recently reported a novel MAGE-A4 epitope presented by human leukocyte antigen (HLA)-A*2402 using a $CD8^+$ T cell clone 2-28 [9]. As this clone effectively killed tumor cell lines that expressed both MAGE-A4 and HLA-A*2402, this antigen could be a candidate for a cancer vaccine.

Given the poor correlation between the immune responses detected in PBMCs and clinical responses [2,3,5], it is necessary to re-evaluate existing cancer immunotherapy strategies in detail using animal models, namely reverse translational research. To this end, we developed animal models involving human tumor antigens, such as NY-ESO-1 or MAGE-A4 in this study.

2. Materials and methods

2.1. Mice

Female BALB/c mice were purchased from SLC Japan (Shizuoka, Japan) and used at 7–10 weeks of age. They were maintained at the Animal Center of Mie University Graduate School of Medicine (Mie, Japan). The experimental protocol was approved by the Ethics Review Committee for Animal Experimentation of Mie University Graduate School of Medicine.

2.2. Antibodies and reagents

Anti-H2-K^d (KD40, mouse IgG2a), anti-H2-D^d (DD98, mouse IgG2a), and anti-H2-L^d (30-5-7, mouse IgG2a) were produced and purified from each hybridoma. FITC-conjugated anti-CD8 mAb (53-6.7, rat IgG2a) and APC-conjugated anti-CD4 mAb (GK1.5, rat IgG2b) were purchased from BD Biosciences (Franklin Lakes, NJ). PE-conjugated anti-Foxp3 mAb (Fjk16s, rat IgG2a) was purchased from eBiosciences (San Diego, CA). Synthetic NY-ESO-1 and MAGE-A4 peptides (summarized in Supplementary Table 1) were obtained from Sigma Genosys (Hokkaido, Japan).

2.3. Immunization using a gene gun

Naive BALB/c mice were immunized twice at two-week intervals. Gold particles coated with 1 μg of each plasmid DNA were prepared and delivered into the shaved skin of the abdominal wall of BALB/c mice using a Helios Gene Gun System (BioRad, Hercules, CA) at a helium discharge pressure of 350–400 psi, as described previously [10,11].

2.4. Cell isolation

Spleen cell suspensions were mixed with CD8 Microbeads (Miltenyi Biotec, Bergisch Gladbach, Germany) and separated into $CD8^+$ T cells by positive selection on a MACS column. The isolated $CD8^+$ T cell populations were confirmed to contain >95% $CD8^+$ T cells.

2.5. Enzyme-linked immunospot (ELISPOT) assay

The number of IFN-γ secreting antigen-specific $CD8^+$ T cells was assessed by ELISPOT assay as described previously [10,11]. Briefly, purified $CD8^+$ T cells were cultured for 24 hours with 5×10^5 irradiated CD90-depleted splenocytes pulsed with the indicated peptides in 96-well nitrocellulose-coated microtiter plates (Millipore, Bedford, MA) coated with rat anti-mouse IFN-γ mAb (R4-6A2, BD Biosciences). Spots were developed using biotinylated anti-mouse IFN-γmAb (XMG1.2, BD Biosciences), alkaline phosphatase conjugated streptavidin (MABTECH, Sweden) and alkaline phosphatase substrate kit (BioRad), and subsequently counted.

2.6. ELISA

96-well flat-bottomed microliter plates (Immuno-NUNC) were coated with 20 ng/50 μml of NY-ESO-1 or MAGE-A4 protein, respectively, at 4 °C overnight. Wells were blocked with 1% BSA/PBS for 1 hour at room temperature and washed three times. Serum (1:100 dilution) was added and incubated at 4 °C overnight. After washing, goat anti-mouse IgG antibody conjugated with horseradish peroxidase (Promega, Madison, WI) was added (1:5000 dilution). Two hours later, color was developed with TMB substrate solution (Thermo scientific, IL) and stopped with H_2SO_4. The absorbance was measured at 450 nm and calculated after subtraction of the absorbance value of control wells without sera.

2.7. Flow cytometry and tetramer staining

Tetramer staining was performed as described previously [11]. Briefly, cells were stained with PE-labeled NY-ESO-$1_{81\text{-}88}/D^d$ or MAGE-$A4_{265\text{-}273}/D^d$ tetramers (prepared at the Ludwig Institute Core Facility, Lausanne, Switzerland) for 10 minutes at 37 °C before additional staining of surface markers for 15 minutes at 4 °C. After washing, the results were analyzed on FACSCanto (BD Biosciences) and FlowJo software (Tree Star, Ashland, OR).

2.8. Tumors

CT26 is a colon epithelial tumor derived by intrarectal injections of N-nitroso-N-methylurethane in BALB/c mice [12]. CT26 expressing NY-ESO-1 or MAGE-A4, a human cancer/testis antigen were established as described previously [11,13].

2.9. Statistical analysis

Values were expressed as mean ± SD. Differences between groups were examined for statistical significance using the Student's *t*-test. p values <0.05 were considered statistically significant.

3. Results

3.1. NY-ESO-1-specific $CD8^+$ T cells recognize D^d-restricted NY-ESO-1_{81-88} peptide

We analyzed the minimal epitope recognized by NY-ESO-1-specific $CD8^+$ T cells after immunization with NY-ESO-1. To this end, we employed a Helios Gene Gun System as we have previously detected NY-ESO-1-specific $CD8^+$ T cell responses [10,11]. To identify minimal epitopes, naive BALB/c mice were immunized twice at two-week intervals with plasmids encoding the entire sequence of NY-ESO-1. $CD8^+$ T cells were obtained from spleens and specific T cell responses were analyzed by ELISPOT assay using peptide pools shown in Supplementary Table 1. A significant number of NY-ESO-1-specific $CD8^+$ T cells was detected against peptide pool #3 (Fig. 1A). To identify the NY-ESO-1 epitope, NY-ESO-1-specific $CD8^+$ T cells were stimulated with each of these peptides. IFN-γ secretion was observed when NY-ESO-1-specific $CD8^+$ T cells were stimulated with 71–90 and 81–100 NY-ESO-1 peptides, suggesting the presence of a minimal epitope within 81–90 residues (Fig. 1B). To determine the minimal epitope, the 81–90 peptide of NY-ESO-1 was further truncated. NY-ESO-1-specific $CD8^+$ T cells recognized the 80–88 and 81–88, but not 80–87 or 82–88 peptides, thus