CORRESPONDENCE Characterization of TGFβ-specific CD4⁺T cells through the modulation of TGFβ expression in malignant myeloid cells

Rasmus Erik Johansson Mortensen¹, Morten Orebo Holmström ^{1,2} and Mads Hald Andersen^{1,2^{IM}}

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Transforming growth factor beta (TGFB) is a pleiotropic cytokine with functions related to angiogenesis, tumor suppression, and immune tolerance [1]. TGF β is involved in tumor immune escape and is a target for cancer immune therapy [1]. We have reported that healthy donors and cancer patients harbor CD8⁺ cytotoxic T lymphocytes (CTLs) that are specific to TGF_β-derived epitopes and that these CD8⁺ CTLs kill malignant myeloid cells in a TGFβ-dependent manner [2]. We also found that T cells specific to the TGF\beta-derived epitope $\text{TGF}\beta_{141-160}$ (REAVPEPVLLSRAELRLLRL, or TGF_β-15) that were expanded from a patient with cancer contained both CD4⁺ and CD8⁺ TGF β -15-specific T cells. The presence of CD4⁺ subsets, such as Th1 and Th9 cells, is important to antitumor immunity and has been shown to increase the efficacy of PD-L1/PD-1 immune checkpoint therapy [3, 4]. Hence, we set out to characterize CD4⁺ T cells that were specific to TGFβ-derived epitopes.

Using magnetically activated cell sorting, we enriched CD4⁺ T cells from the TGFβ-15-specific culture described above, thereby establishing CD4+ T cell cultures that were specific to TGFβ-15. These T cells were characterized by fluorescenceactivated cell sorting (FACS) and intracellular cytokine staining (ICS) as described previously [2]. TGF β -15-specific CD4⁺ T cells produced high amounts of interferon-gamma (IFN-y) and tumor necrosis factor alpha (TNF-α) upon stimulation with TGFβ-15 (Fig. 1A). A pan-HLA class II-blocking assay showed that the response was HLA-II dependent (Fig. 1B), and blocking HLA-DR, DQ, and DP (Fig. 1B) showed that the response was restricted to HLA-DR. Donor genotyping identified the HLA-DRB1*15:01:01 G allele. As we wanted to investigate the ability of T cells to recognize TGFβ-expressing cells, we searched the TRON Cell Line Portal (http://celllines.tron-mainz.de/) for HLA-DR-compatible cancer cell lines and found the TGF_β-producing malignant myeloid cell line THP-1, which expresses HLA-DRB*15.

We have previously shown that stimulation of THP-1 cells with the cytokines IL-4 and TGF β enhances their expression of TGF β [2]. Hence, we investigated whether increased amounts of TGF β produced by THP-1 cells could increase their recognition by TGF β -15-specific CD4⁺ T cells. Importantly, the active form of the cytokine TGF β does not include the TGF β -15 epitope, which is found in the N-terminal latency-associated peptide of TGF β . Accordingly, any increased recognition of THP-1 cells was not due to the processing and presentation of the TGF β used for stimulation. Using western blotting, we confirmed that treatment of THP-1 cells for 72 h with 200 U/ml IL-4 and 5 ng/ml TGFB increased the expression of TGFB (Fig. 1C). THP-1 cells that were treated with these cytokines for 72 h were then used as target cells in an ICS experiment with an effector:target (E:T) ratio of 3:1. The results showed increased expression of TNF-a, IFN-y, and CD107a in T cells that were stimulated with cytokine-treated THP-1 cells compared with unstimulated THP-1 cells (Fig. 1D, E). An increase in TGFB expression by THP-1 cells was already evident after 48 h of stimulation with cytokines [2], and CD4⁺ T cells stimulated with THP-1 cells that were treated with cytokines for 48 h also responded with increased cytokine (data not shown). The experiment was repeated with 8 replicates using another TGF β -15-specific CD4⁺ T cell culture. The number of replicates allowed us to perform statistical analysis. Again, we identified significant increases in cytokine release and CD107a expression in T cells stimulated with cytokine-treated THP-1 cells (Fig. 1F). Next, we asked whether the increased T cell activation was caused by increased HLA-II expression on cytokine-treated THP-1 cells. Using FACS analysis with Fc blocking and an appropriate isotype control, we demonstrated that this was not the case (Fig. 1G). In fact, cytokine-treated THP-1 cells expressed less HLA-II than untreated THP-1 cells.

To demonstrate that the decreased expression of TGF β induced by THP-1 cells resulted in decreased activation of TGF β -15-specific CD4⁺ T cells, we transfected THP-1 cells with TGF β siRNA [2]. THP-1 cells that were transfected with TGF β siRNA had lower expression of TGF β than mock-transfected THP-1 cells (Fig. 1H) and comparable levels of HLA-II expression (Fig. 1I). Accordingly, we found a difference in the activation of T cells stimulated with either mock- or siRNA-transfected THP-1 cells (Fig. 1J). The experiment was repeated with 8 replicates and showed a statistically significant decrease in the activation of T cells stimulated with siRNA-transfected THP-1 cells (Fig. 1K).

There is mounting evidence pointing to the existence of a class of self-reactive T cells, known as anti-regulatory T cells, that are not deleted in the thymus [5]. These cells can recognize and kill regulatory immune cells and cancer cells that produce immunosuppressive molecules such as indoleamine 2,3-dioxygenase 1 (IDO) [6] and programmed death ligand-1 (PD-L1) [7]. Activation of these anti-regulatory T cells through vaccination with epitopes from these immunosuppressive molecules induces an immune response against cancer cells and regulatory cells in the tumor.

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¹National Center for Cancer Immune Therapy, Department of Oncology, Copenhagen University Hospital, Herlev, Denmark. ²Institute for Immunology and Microbiology, Copenhagen University, Copenhagen, Denmark. ^{Sem}email: mads.hald.andersen@regionh.dk

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Fig. 1 Characterization of TGF β epitope-specific CD4⁺ T cells. **A** Intracellular cytokine staining (ICS) of the TGF β -15-specific CD4⁺ T cell culture stimulated with TGF β -15 (left) and of the unstimulated cells (right). **B** ICS of TGF β -15-stimulated cells treated with a pan-HLA-II-blocking antibody (left) and HLA-DR-, DQ-, and DP-blocking antibodies (right). **C** Western blot analysis of lysates from THP-1 cells that were unstimulated or treated with IL-4 and TGF β 1 for 72 h (left). Relative expression of TGF β in lysates were analyzed using ImageJ software (right). **D** ICS of T cells stimulated with THP-1 cells that were either untreated or treated with IL-4 and TGF β 1 for 72 h with an effector:target ratio of 3:1. **E** Expression of CD107a on T cells from the experiment shown in **D**. **F** Experiment performed as in **D**, **E** with 8 replicates, showing the ICS results (left) and CD107a expression (right). Error bars indicate the standard error of the mean. Statistical analysis was performed using the Mann–Whitney *U*-test. **G** FACS analysis of the expression of HLA-II by THP-1 cells that were either mock transfected or transfected with TGF β siRNA (left). Relative expression of TGF β in lysates were analyzed using ImageJ software (right). **I** FACS analysis of the expression of TGF β in lysates were analyzed using ImageJ software (right). **I** FACS analysis of the expression of TGF β in lysates were analyzed using ImageJ software (right). **I** FACS analysis of the expression of TGF β is INNA (left). Relative expression of TGF β is Ilysates were analyzed using ImageJ software (right). **I** FACS analysis of the expression of TGF β is Isinulated with TGF β siRNA 48 h prior to analysis. **J** ICS performed on TGF β -15-specific T cells stimulated with THP-1 cells that were either mock or TGF β 1 siRNA transfected 96 h before ICS. The experiment was performed using different conditions: 2 h of stimulation and an E:T of 3:1 (left); 4 h of stimulation and an E:T of 3:1 (middle); or 5 h of stimulation a

This response can decrease local immune suppression, enhancing the tumor-specific immune response. Therapeutic cancer vaccines against IDO and PD-L1 to enhance the anti-regulatory T-cell response have shown clinical potential in melanoma and nonsmall cell lung cancer [8, 9]. Our discovery of CD4⁺ and CD8⁺ T cells specific for TGFB is noteworthy, as vaccination with TGFB-derived epitopes could enhance the TGFβ-specific T cell response in cancer patients. Using the HLA-DR-compatible malignant myeloid cell line THP-1 and modulating these cells to increase or decrease intracellular TGFB, we showed that the recognition of these cells by our TGFβ-specific CD4⁺ T cells was TGFβ dependent. Systemic CD4⁺ T cell responses to tumor antigens are important in enhancing cytotoxic CD8⁺ T cell immunity in patients [4]. Our discovery of TGF β -specific CD4⁺ T cells is of great importance, as CD4⁺ TGF β -specific T cell responses in patients could be used to enhance the responses of the TGFβ-specific CD8⁺ CTLs we described previously.

TGF β is produced not only by cancer cells but also by several regulatory cells in the tumor microenvironment, including Tregs, tumor-associated macrophages, and cancer-associated fibroblasts [1]. Hence, the induction of a synergistic CD4⁺ and CD8⁺ TGF β -specific T cell response by therapeutic cancer vaccination with one or several TGF β -derived epitopes might result in the specific recognition of cancer cells, cancer-associated fibroblasts, and other regulatory immune cells in the tumor microenvironment. This recognition could revert local immune suppression, induce a proinflammatory environment and push the balance toward antitumor immunity, thereby improving responses to other therapies.

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AUTHOR CONTRIBUTIONS

REJM performed experiments, interpreted the data, and wrote the manuscript; MOH performed experiments, interpreted the data, and wrote the manuscript; and MHA conceived the project, interpreted the data, and wrote the manuscript.

COMPETING INTERESTS

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MHA has developed an invention based on the use of TGF β for vaccinations. The rights of the invention have been transferred to Copenhagen University Hospital Herlev, according to the Danish Law of Public Inventions at Public Research Institutions. The capital region has licensed the rights to the company IO Biotech ApS. The patent application was filed by IO Biotech ApS. MHA is board member, consultant and shareholder in IO Biotech. The other authors declare no competing financial interests.

ADDITIONAL INFORMATION

Correspondence and requests for materials should be addressed to Mads Hald Andersen.

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