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Altered Effector CD4⁺ T Cell Function in IL-21R^{-/-} CD4⁺ T Cell-Mediated Graft-Versus-Host Disease

Iekuni Oh,* Katsutoshi Ozaki,* Akiko Meguro,* Keiko Hatanaka,* Masanori Kadowaki,[†] Haruko Matsu,* Raine Tatara,* Kazuya Sato,* Yoichiro Iwakura,[‡] Susumu Nakae,[§] Katsuko Sudo,[¶] Takanori Teshima,[†] Warren J. Leonard,^{||} and Keiya Ozawa*

We previously showed that transplantation with *IL-21R* gene-deficient splenocytes resulted in less severe graft-versus-host disease (GVHD) than was observed with wild type splenocytes. In this study, we sought to find mechanism(s) explaining this observation. Recipients of donor CD4⁺ T cells lacking IL-21R exhibited diminished GVHD symptoms, with reduced inflammatory cell infiltration into the liver and intestine, leading to prolonged survival. After transplantation, CD4⁺ T cell numbers in the spleen were reduced, and MLR and cytokine production by CD4⁺ T cells were impaired. These results suggest that IL-21 might promote GVHD through enhanced production of effector CD4⁺ T cells. Moreover, we found that CD25 depletion altered neither the impaired MLR *in vitro* nor the ameliorated GVHD symptoms *in vivo*. Thus, the attenuated GVHD might be caused by an impairment of effector T cell differentiation itself, rather than by an increase in regulatory T cells and suppression of effector T cells. *The Journal of Immunology*, 2010, 185: 1920–1926.

Interleukin-21 was discovered as a costimulatory cytokine for T cell proliferation and NK cell expansion *in vitro* (1, 2). IL-21 is produced by activated CD4 T cells (1), and its receptor is expressed on T, B, and NK cells (1, 3). It was also reported that IL-21 suppresses dendritic cell function (4) and increases hematopoietic progenitor cells (5). IL-21 is known to play critical roles in Ig production (6), whereas reports have differed regarding its contributions to Th1-, Th2-, and Th17-mediated effects and differentiation (6–15). IL-21 contributes to Th17 differentiation, but it may not be required for this process (7, 9, 14, 15). A relationship between IL-21 and autoimmune disease has been established. Overexpression of IL-21 induces inflammation, and in a systemic lupus erythematosus model mouse (the BXSb.6-Yaa^{+/J}) with high serum levels of IL-21 (16), the development of disease is abrogated when these mice are crossed to IL-21R knockout (KO) mice (17). In addition, autoimmune NOD mice do not develop diabetes in the absence of IL-21 signaling (18–20).

Graft-versus-host disease (GVHD) is a major complication following hematopoietic stem cell transplantation (21), sometimes with a fatal outcome. Previously, we showed that transplantation with *IL-21R* gene-disrupted splenocytes resulted in less severe

GVHD than was seen with wild type (WT) splenocytes (22). We sought to elucidate the mechanism(s) for this observation; in this article, we demonstrate dysregulated effector function of activated CD4⁺ T cells in IL-21R^{-/-} mice.

Materials and Methods

Mice

IL-21R^{-/-} and IL-17^{-/-} mice were generated previously (6, 23). Both were on a C57BL/6 background. Male and female mice were used as donors. C57BL/6-DBA2-F1 male mice were purchased from Clea Japan (Tokyo, Japan). All mice used in experiments were 6–12 wk old. All mice were housed in a Jichi Medical University mouse facility, which is regulated by an intramural small animal committee, and were treated in accordance with university guidelines.

In vitro T cell stimulation and MLR

Cells were cultured in RPMI 1640 (Invitrogen, Carlsbad, CA) supplemented with 10% FCS (Sigma-Aldrich, St. Louis, MO), 2 mM L-glutamine (Invitrogen), 50 μM 2-ME (Sigma-Aldrich), 0.1 mg/ml streptomycin, and 100 U/ml penicillin G (Invitrogen). Nonspecific pan T cell stimulation was performed using anti-CD3/CD28 beads for 3 d, according to the manufacturer's instructions (DynaL Biotech, Oslo, Norway). Alloantigen-specific T cell stimulation was induced by cocultivation of CD4 T cells with 30 Gy-irradiated splenocytes from C57BL/6-DBA2-F1 mice for 4 d.

GVHD models

We used IL-21R^{-/-} bone marrow (BM) to eliminate the effects of WT T cells in BM. We compared transplantations with IL-21R^{-/-} CD4⁺ T cells versus WT CD4⁺ T cells. C57BL/6-DBA2-F1 mice were irradiated with 11 Gy and injected *i.v.* with 5 × 10⁶ IL-21R^{-/-} BM and 5 × 10⁶ purified CD4⁺ T cells from WT or IL-21R^{-/-} mice. The cells were purified using CD4 microbeads and AutoMACS (Miltenyi Biotec, Tokyo, Japan); the purity was >80–90%.

Pathological analysis

Two weeks after transplantation, mice were sacrificed; liver, skin, and intestine were subjected to formalin fixation, paraffin embedding, excision, and H&E staining. Photographs were taken with an Olympus BX51 microscope at ×400 magnification.

Flow cytometric analysis

Fc-block (BD Biosciences, San Jose, CA) was used to prevent nonspecific Ab binding to Fc receptors. Abs to CD4 (RM4-5), CD8 (Ly-2), CD25

*Division of Hematology, Department of Medicine, Jichi Medical University, Tochigi; [†]Department of Medicine and Biosystemic Science, Kyushu University Graduate School of Medical Science, Fukuoka; [‡]Center for Experimental Medicine and [§]Frontier Research Initiative, Institute of Medical Science, University of Tokyo; [¶]Animal Research Center, Tokyo Medical University, Tokyo, Japan; and ^{||}Laboratory of Molecular Immunology, National Heart, Lung, and Blood Institute, National Institutes of Health, Bethesda, MD 20892

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Address correspondence and reprint requests to Dr. Katsutoshi Ozaki, Division of Hematology, Department of Medicine, Jichi Medical University, 3311-1 Yakushiji, Shimotsuke-shi, Tochigi 329-0498, Japan. E-mail address: ozakikat@jichi.ac.jp

The online version of this article contains supplemental material.

Abbreviations used in this paper: BM, bone marrow; GVHD, graft-versus-host disease; KO, knockout; Treg, regulatory T; WT, wild type.

(7D4), H-2^b (AF6-88.5), H-2^d (SF1-1.1), IFN- γ (XMG1.2), and TNF- α (MP6-XT22) were purchased from BD Biosciences, and anti-Foxp3 (FJK-16a) was from eBioscience (San Diego, CA). Intracellular staining was performed with a Cytofix/Cytoperm kit (BD Biosciences), according to the manufacturer's instructions. Cells were stimulated with anti-mouse CD3/CD28 beads for 5 h in the presence of GolgiStop (BD Biosciences). The stimulation was omitted for Foxp3 intracellular staining. An LSR flow cytometer (BD Biosciences) was used for data collection, and data were analyzed using CellQuest software (BD Biosciences).

ELISA

ELISA kits for IL-2, IL-4, and IFN- γ were from BD Biosciences, and ELISA kits for IL-21, IL-17, TNF- α , and TGF- β 1 were from R&D Systems (Minneapolis, MN). Concentrations were determined according to the manufacturer's instructions.

CD25 depletion in vitro and in vivo

In vitro purification of CD4⁺ T cells and depletion of the CD25⁺CD4⁺ subpopulation were performed by cell sorting using a FACSaria (BD Biosciences), which yielded highly pure populations (>98%). In vivo CD25 depletion was performed by injecting anti-CD25 Ab, as described previously (24, 25). Briefly, a hybridoma producing anti-CD25 Ab (PC61; American Type Culture Collection, Manassas, VA) was cultured in serum-free medium (Protein-Free Hybridoma Medium-II from Invitrogen), and the Ab was purified from supernatant by ammonium sulfate precipitation and a PD10 column (GE Healthcare, Buckinghamshire, U.K.). The purified product was quantified using the Bradford assay (Bio-Rad, Hercules, CA) at OD595, and 1 mg was injected i.p. weekly from day 0 for 3 wk. Control rat nonspecific IgG was purchased from Invitrogen.

Quantitative RT-PCR

At day 21 after bone marrow transplantation, CD25⁻CD4⁺ T cells were purified by cell sorting from recipients of WT or IL-21R^{-/-} CD4⁺ T cells; RNA was isolated (RNeasy, Qiagen, Valencia, CA), reverse transcribed using the SuperScript First-Strand Synthesis System for RT-PCR (Invitrogen), and PCR amplified using TaqMan Gene Expression Assay's primer for mouse Foxp3 (Mm00475156) and β -actin (Mm00607939) and an ABI Prism 7700 sequence detection System (Applied Biosystems, Foster City, CA).

Statistical analysis

Kaplan–Meier plots were used to compare survival rates. The log-rank test was used to evaluate *p* values. Statistical analyses were performed using Stat Mate ver. 6 (ATMS, Tokyo, Japan). The Student *t* test was used; all error bars in this study represent SD, unless otherwise specified.

Results

Purified CD4⁺ T cell transplantation and pathological analysis

Decreased GVHD was observed when we transplanted IL-21R-deficient splenocytes compared with WT bulk splenocytes (22). Although we sought to find molecular mechanism(s) for the ameliorated GVHD, no clue was immediately evident from the transplantation experiments (22). Thus, in this study, we used purified CD4⁺ T cells instead of bulk splenocytes in an effort to augment the differences observed. We used a well-known model of CD4⁺ T cell-mediated GVHD (26), in which C57BL/6 mice were donors, and C57BL/6-DBA2-F1 mice were recipients. In this model, the difference between WT and IL-21R^{-/-} cells seemed to be greater than in the previous experiments using bulk splenocytes (22). All recipients of WT CD4⁺ T cells died within 55 d, whereas those receiving IL-21R^{-/-} CD4⁺ T cells survived during this time period (Fig. 1A). Moreover, recipients of IL-21R^{-/-} CD4⁺ T cells recovered from body weight loss by day 14, but those receiving WT CD4⁺ T cells did not recover and continued to lose weight (Fig. 1B). In recipients of IL-21R^{-/-} CD4⁺ T cells, pathological analysis showed markedly reduced infiltration into the regions surrounding bile ducts and portal veins and into the interstitial region of small intestine compared with the infiltration observed in recipients of WT CD4⁺ T cells (Fig. 2, upper and middle panels). Apoptotic bodies near the surface area of crypts in the small intestine were barely visible in recipients

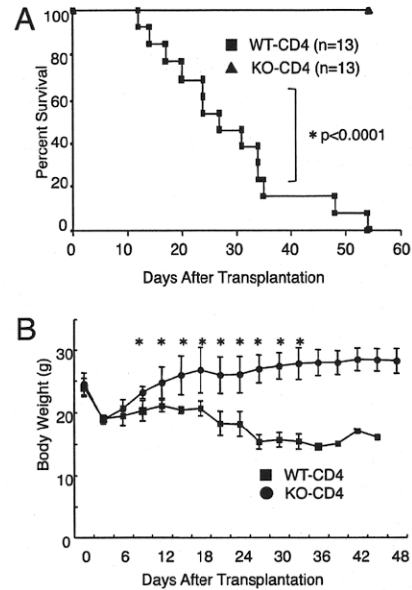


FIGURE 1. A role for IL-21 in CD4⁺ T cell-mediated GVHD. *A*, Survival of recipients of WT and IL-21R^{-/-} CD4⁺ T cells. C57BL/6-DBA2-F1 mice were irradiated with 11 Gy and received 5 × 10⁶ IL-21R^{-/-} BM with 5 × 10⁶ WT or IL-21R^{-/-} CD4⁺ T cells. Shown are combined data from two independent experiments. Thirteen recipients each for WT and IL-21R^{-/-} CD4⁺ T cells were analyzed. The log-rank test was used to calculate *p* values. *B*, Body weight after BM transplantation. Statistical significance was assessed with the Student *t* test.

of IL-21R^{-/-} CD4⁺ T cells, in contrast to recipients of WT CD4⁺ T cells, in which apoptotic bodies were evident (Fig. 2, arrowheads in middle panel). No significant difference was observed in skin

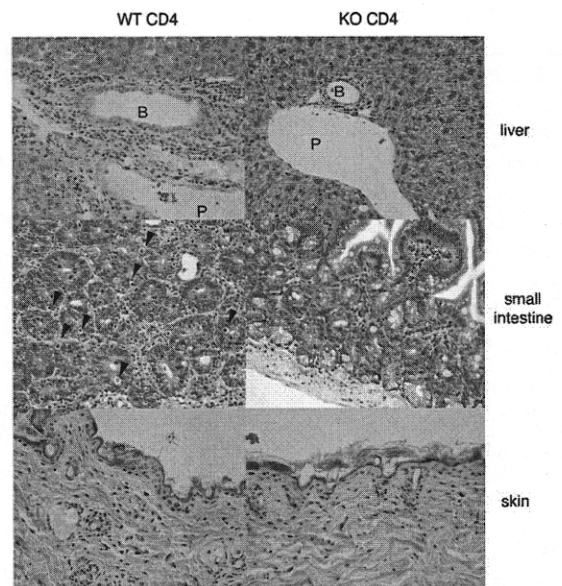


FIGURE 2. Pathological analysis of recipients. Liver, small intestine, and skin were stained with H&E (original magnification ×400). In recipients of WT CD4⁺ T cells, cell infiltration is evident around the portal vein (P) and the bile duct (B) and into the interstitial region in small intestine. Arrowheads indicate apoptotic bodies near the surface of crypts. These changes were barely visible in recipients of IL-21R^{-/-} CD4⁺ T cells. Skin did not show any significant difference between recipients of WT and IL-21R^{-/-} CD4⁺ T cells. Shown is a representative result of six mice analyzed in each group. Only one recipient of IL-21R^{-/-} CD4⁺ T cells showed apoptotic bodies in the lumens of intestine and infiltration around the bile duct and portal vein, as was observed in the recipients of WT CD4⁺ T cells.

pathology among recipients of WT CD4⁺ and IL-21R^{-/-} CD4⁺ T cells. These results suggested that IL-21 might be essential for CD4-mediated GVHD, at least in this setting.

Normal cytokine production by splenocytes after transplantation is dependent on IL-21

The above observations suggested that IL-21-mediated donor CD4⁺ T cell activation was involved in the exacerbation of GVHD. Because we could not find any significant difference in serum cytokine concentrations after transplantation (Supplemental Fig. 1), we assessed T cell differentiation by cytokine production in the presence of cellular stimulation. Interestingly, at days 14 and 21 after transplantation, bulk splenocytes from recipients of IL-21R^{-/-} CD4⁺ T cells exhibited defective cytokine production, with decreased levels of IFN- γ , TNF- α , and IL-4; in contrast, levels of IL-2, IL-17, and IL-21 were not significantly diminished (Fig. 3, left panels). Before transplantation, IL-21R^{-/-} CD4⁺ T cells did not show any significant defect in IFN- γ , IL-4, or TNF- α production (Fig. 3, right panels), suggesting that the defect was acquired after transplantation. This defect in effector T cell function might represent a mechanism for the difference in the development of GVHD by mice receiving WT versus IL-21R^{-/-} CD4⁺ T cells.

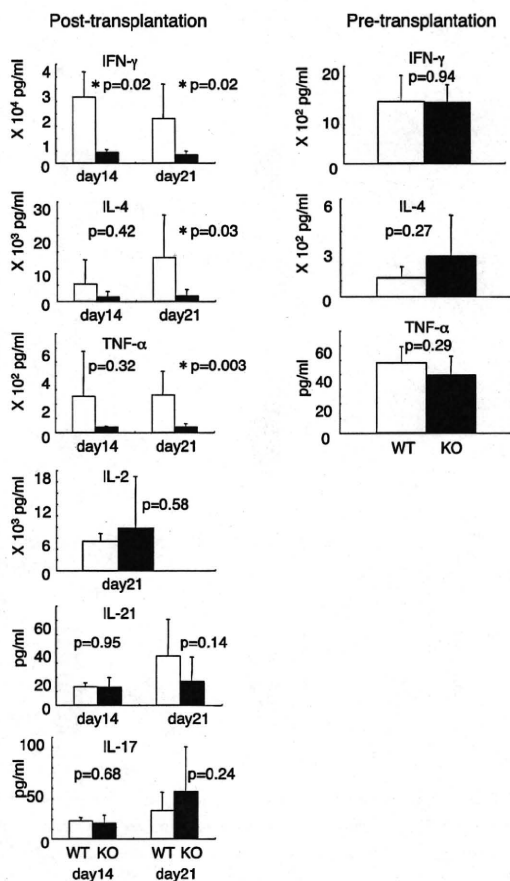


FIGURE 3. Cytokine production by bulk splenocytes before and after CD4⁺ T cell transplantation. At days 14 and 21 after transplantation, splenocytes (5×10^5) were taken and stimulated with anti-CD3/CD28 Abs for 18 h. Concentrations of cytokines in the supernatants were determined by ELISA. Twelve or 13 recipients of WT CD4⁺ T cells and 10 recipients of IL-21R^{-/-} CD4⁺ T cells were analyzed. Prior to transplantation, five WT and eight IL-21R^{-/-} mice were analyzed. At days 14–21 after transplantation, the proportion of donor cells in the spleen was >95%. The Student *t* test was used to calculate *p* values. *Statistical significance ($p < 0.05$).

CD4⁺ T cells were responsible for the low production of cytokines

To elucidate the basis for diminished cytokine production, we examined the number of donor CD4⁺ T cells in the spleen at days 14–21 after transplantation. The number of donor H-2K^d-CD4⁺ T cells was significantly lower in recipients of IL-21R^{-/-} CD4⁺ T cells than in recipients of WT CD4⁺ T cells (Fig. 4A; $p = 0.03$, Welch *t* test; $n = 15$ versus 12), although the ranges overlapped. Because it is thought that donor T cells proliferate in secondary lymphoid organs, such as the spleen, and then infiltrate into target organs (27), the reduced number of CD4⁺ T cells in the spleen is consistent with the reduced infiltration into the liver and small intestine, as shown above (Fig. 2). To identify the cells responsible for defective cytokine production, we performed intracellular staining and ELISA with purified CD4⁺ T cells. After anti-CD3/CD28 stimulation, the proportion of IFN- γ ⁺ and TNF- α ⁺ cells in splenic CD4⁺ T cells was lower in recipients of IL-21R^{-/-} CD4⁺ T cells than in those receiving WT CD4⁺ T cells (Fig. 4B). Moreover, posttransplantation, the levels

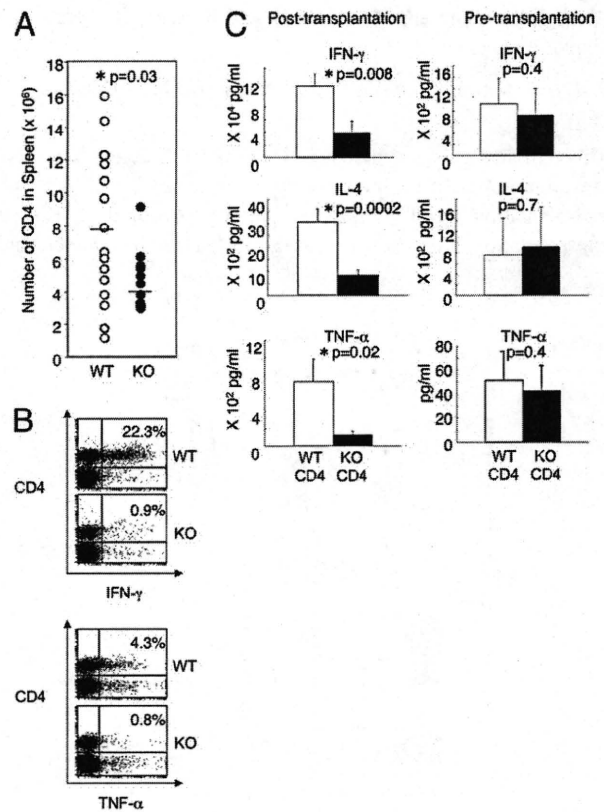


FIGURE 4. Cytokine production by splenic CD4⁺ T cells before and after transplantation. **A**, Absolute number of donor H-2K^d-CD4⁺ T cells in the spleen. The number of donor CD4⁺ T cells was determined by multiplying the number of splenocytes by the percentage of H-2K^d-CD4⁺ T cells. Each dot depicts the number of donor CD4⁺ T cells in a mouse. Horizontal lines indicate the average. Fifteen recipients of WT CD4⁺ T cells and 12 recipients of IL-21R^{-/-} CD4⁺ T cells were assessed. **B**, Intracellular staining of splenocytes after anti-CD3/CD28 stimulation. Splenocytes (1×10^6) were stimulated with anti-CD3/CD28 Abs for 5–6 h and stained with anti-IFN- γ or anti-TNF- α Ab in combination with anti-CD4 Ab. A total of three recipients in each group were analyzed, and a representative result is shown. **C**, Cytokine production by CD4⁺ T cells in vitro. At days 14 or 21 after transplantation, splenic CD4⁺ T cells (5×10^5) were purified and stimulated with anti-CD3/CD28 Abs for 18 h. Concentrations of cytokines in the supernatants were determined by ELISA. Twelve mice were analyzed in each group after transplantation. Five or six WT and eight or nine IL-21R^{-/-} mice were analyzed before transplantation. *Statistical significance ($p < 0.05$).

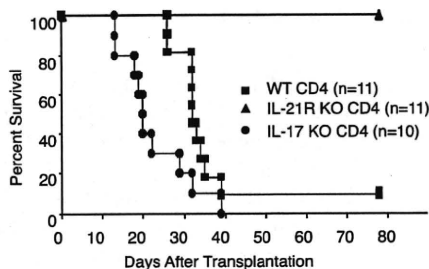


FIGURE 5. IL-17^{-/-} CD4⁺ T cells induced lethal GVHD. Survival of recipients of WT, IL-21R^{-/-} (IL-21R KO), or IL-17^{-/-} (IL-17 KO) CD4⁺ T cells. Lethally irradiated (11 Gy) C57BL/6-DBA2-F1 mice were transplanted with 5 × 10⁶ IL-21R KO BM and 5 × 10⁶ WT, IL-21R KO, or IL-17 KO CD4⁺ T cells. The data represent the combined results of two independent experiments.

of IFN-γ, TNF-α, and IL-4 production were significantly diminished with splenic-purified CD4⁺ T cells from recipients of IL-21R^{-/-} CD4⁺ T cells compared with those receiving WT CD4⁺ T cells (Fig. 4C, left panels). Before transplantation, IL-21R^{-/-} CD4⁺ T cells did not show any defect in IFN-γ, TNF-α, and IL-4 production (Fig. 4C, right panels).

IL-17 production and GVHD induced by IL-17^{-/-} CD4⁺ T cells

Although IL-21 is not essential for Th17 differentiation, IL-21 can promote it. To evaluate the effect of IL-21^{-/-} CD4⁺ T cell transplantation on IL-17 production, we measured IL-17 after transplantation. As shown in Fig 3, bottom left panel, bulk

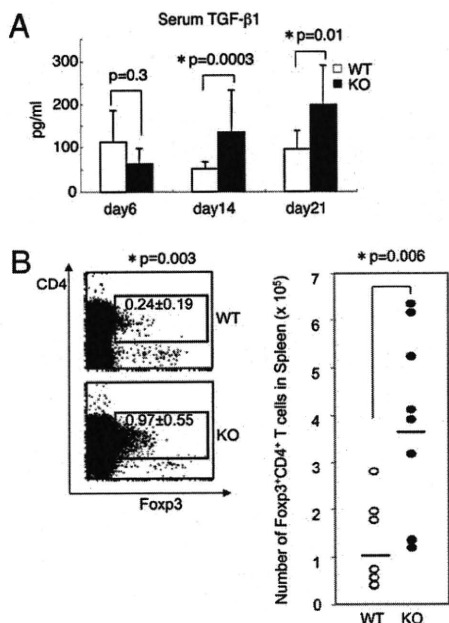


FIGURE 6. Increase in splenic Treg cells. *A*, Upregulation of serum TGF-β1. Serum TGF-β1 concentrations at the indicated day after transplantation were determined by ELISA. Three samples from recipients of WT CD4⁺ T cells and 4 samples from recipients of IL-21R^{-/-} CD4⁺ T cells at day 6, 23 samples from recipients of WT CD4⁺ T cells and 25 samples from recipients of IL-21R^{-/-} CD4⁺ T cells at day 14, and 8 samples from recipients of WT CD4⁺ T cells and 7 samples from recipients of IL-21R^{-/-} CD4⁺ T cells at day 21 were analyzed. *Statistical significance ($p < 0.05$). *B*, The percentage and absolute number of splenic Fosp3⁺CD4⁺ regulatory T cells at day 14 after transplantation. The left panel shows a representative flow cytometric result from eight or nine similar samples. The right panel indicates the number of all samples; the averages are indicated by the horizontal lines.

splenocytes from recipients of IL-21R^{-/-} CD4⁺ T cells produced comparable amounts of IL-17 at days 14 and 21 after transplantation compared with mice receiving WT CD4⁺ T cells. Moreover, we found that IL-17^{-/-} CD4⁺ T cells induced lethal GVHD analogous to WT CD4⁺ T cells (if anything, death occurred earlier), suggesting that IL-17 is dispensable for this process, in contrast to the essential role of IL-21, as reflected by the survival of mice receiving IL-21R^{-/-} CD4⁺ T cells (Fig. 5).

Regulatory T cell number in spleen

We next determined the serum concentration of the major immunosuppressive cytokine, TGF-β1, at days 6–21 after transplantation. We found an increase in TGF-β1 only after transplantation (Fig. 6A; $p = 0.0003$ at day 14; $p = 0.01$ at day 21, Student *t* test). In splenocytes from recipients of IL-21R^{-/-} CD4⁺ T cells, the production of TGF-β1 and IL-10 by in vitro T cell stimulation was not upregulated; in fact, it tended to be diminished (Supplemental Fig. 2), suggesting that the increase in serum TGF-β1 might be due to cells other than T cells. Because naive T cells can differentiate into regulatory T (Treg) cells in the presence of TGF-β1 (28), and it was reported that IL-21^{-/-} T cells were predisposed to differentiate into Treg cells (8), we also investigated whether more Treg cells were induced in recipients of IL-21R^{-/-} CD4⁺ T cells. The proportion

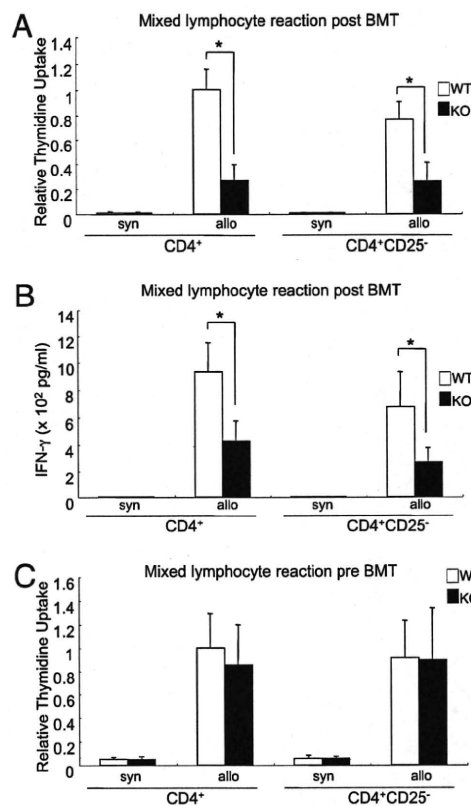


FIGURE 7. An impaired CD4 alloreaction is not dependent on CD25⁺ CD4⁺ T cells. CD4 alloreaction in vitro was impaired after transplantation, and this impairment was not restored by CD25⁺ T cell depletion. *A*, At day 14 after transplantation, 1 × 10⁵ sorter-purified splenic CD4⁺ or CD25⁺ CD4⁺ T cells (>98% purity) were cultured with 4 × 10⁵ irradiated allogeneic C57BL/6-DBA2-F1 splenocytes for 4 d. The cells were pulsed with 1 μCi of [³H]thymidine for the last 24 h. Relative thymidine uptake to the value of WT CD4⁺ T cells is depicted. *B*, Culture was the same as in *A*, but IFN-γ concentrations in the supernatants were determined by ELISA. *C*, Sorter-purified splenic CD4⁺ or CD25⁺ CD4⁺ cells from nontransplanted mice were cultured with irradiated allogeneic C57BL/6-DBA2-F1 splenocytes. Relative thymidine uptake to the number of WT CD4⁺ T cells is depicted. * $p < 0.05$.

of splenic Foxp3⁺CD4⁺ Treg phenotype cells in recipients of IL-21R^{-/-} CD4⁺ T cells was higher than in recipients of WT CD4⁺ T cells, but the total percentage was still only ~1% (Fig. 6B, left panel). The absolute number was ~4-fold higher, but the actual number was only ~4 × 10⁵ of the total number of splenocytes (~4 × 10⁷) (Fig. 6B, right panel). In contrast to posttransplantation, pretransplantation splenocytes from IL-21R^{-/-} mice did not show an increase in Foxp3⁺CD4⁺ T cells compared with cells from WT mice (Supplemental Fig. 3), suggesting that the increased Treg cell after transplantation was an induced Treg cell during GVHD reaction. For that reason, we did not deplete CD25⁺ cells prior to transplantation.

CD25 depletion did not restore the suppressed alloreaction in vitro and did not exacerbate the ameliorated GVHD

To investigate the importance of Treg cells in diminishing GVHD, we performed an MLR, which corresponds to alloreaction in vitro, with or without CD25⁺CD4⁺ T cells. Because Foxp3 is an intracellular protein, and Foxp3 staining cannot be used to purify or deplete Treg cells, anti-CD25 Ab is widely used for this purpose (9, 29–32). The impaired MLR of IL-21R^{-/-} CD4⁺ T cells after transplantation was not restored by CD25 depletion (Fig. 7A), nor was the impaired IFN-γ production by IL-21R^{-/-} CD4⁺ T cells in an MLR (Fig. 7B). Moreover, analogous to cytokine production by anti-CD3/CD28 stimulation (Fig. 3), IL-21R^{-/-} CD4⁺ T cells before transplantation were not defective for alloreaction (Fig. 7C).

Consistent with the in vitro experiments above, CD25⁺ depletion in vivo did not alter the severity of GVHD in recipients of IL-21R^{-/-} CD4⁺ T cells, without altering the body weight loss and survival (Fig. 8A, 8B). In contrast, the severity of GVHD in recipients of WT CD4⁺ T cells seemed to be slightly diminished by CD25⁺ depletion (Fig. 8A, 8B). In this condition, as previously reported (30), the depletion efficacy of CD25⁺CD4⁺ T cells was >95% and that of Foxp3⁺CD4⁺ T cells was ≥50% (Fig. 8C, upper

and lower panels). Interestingly, Foxp3 expression was higher in CD25⁻CD4⁺ T cells from recipients of IL-21R^{-/-} CD4⁺ T cells than from recipients of WT CD4⁺ T cells (Fig. 8D). Together with the results in vitro (Fig. 7), this suggests a relationship between the unresponsiveness of CD25⁻CD4⁺ T cells and greater expression of Foxp3.

Discussion

In this article, we reported evidence indicating that IL-21 is critical for the pathogenesis of CD4⁺ T cell-mediated GVHD, at least in part because of its effects on CD4 differentiation. In this study, we focused on CD4⁺ T cell-mediated GVHD; a role for IL-21 in CD8⁺ T cell-mediated GVHD remains to be investigated.

We found a profound defect in T cell effector function only after transplantation, although serum cytokine concentrations showed no obvious difference. According to these results, T cell differentiation into Th1 and Th2 cells seemed to be altered in the absence of IL-21 during GVHD. Cytokines are believed to have positive and negative roles in GVHD. For example, although T cells from IFN-γ-deficient mice resulted in more severe GVHD (33–35), T cells from Stat4 (Th1)-deficient mice resulted in less severe GVHD than did T cells from WT mice with less severe colitis (36). In contrast to IFN-γ^{-/-} T cells, T cells from IL-4-deficient mice induced less severe GVHD (34); analogously, T cells from Stat6 (Th2)-deficient mice induced less severe GVHD than did those from WT mice (36). T cells from TNF-α-deficient mice developed less severe GVHD, with less severe colitis (37). Our data suggest a strong correlation between the defect in effector function in recipients of IL-21R^{-/-} CD4⁺ T cells and the attenuated phenotype of GVHD, indicating a role for IL-21 in this process.

IL-21, as well as IL-6, induces Th17 differentiation in the presence of TGF-β, suggesting a possible involvement of IL-17 in

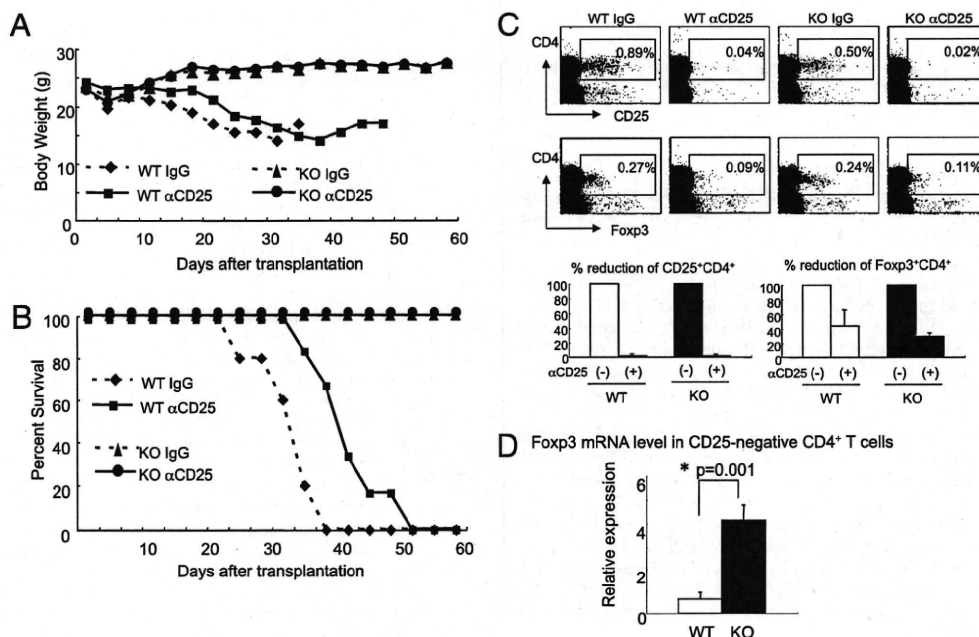


FIGURE 8. The ameliorated GVHD induced by IL-21R^{-/-} CD4⁺ T cells is not dependent on CD25⁺CD4⁺ T cells. The ameliorated GVHD induced by IL-21R^{-/-} CD4⁺ T cells was not exacerbated by depletion of CD25⁺CD4⁺ T cells. Body weight (A) and survival (B) of recipients are shown. Comparisons of recipients of WT CD4⁺ T cells and IL-21R^{-/-} CD4⁺ T cells and additional comparisons with and without anti-CD25 Ab treatment were performed. Nonspecific rat IgG was used as the control Ab. C, Splenic CD25⁺CD4⁺ T cells and splenic Foxp3⁺CD4⁺ T cells at day 14 after transplantation with or without anti-CD25 Ab treatment were analyzed by flow cytometry (upper two rows). The lower panels indicate the mean reduction in the percentage of CD25⁺CD4⁺ and Foxp3⁺CD4⁺ cells from three similar results. D, Foxp3 mRNA level in CD25⁻CD4⁺ T cells at day 21 after transplantation. Cell sorter-purified CD25⁻CD4⁺ T cells were subjected to mRNA purification, reverse-transcriptase treatment, and TaqMan quantitative PCR. Relative value to β-actin is denoted.

the phenotype we observed. However, our results with IL-17^{-/-} CD4⁺ T cells demonstrated that IL-17 was dispensable for CD4⁺ T cell-mediated GVHD, indicating that the attenuated GVHD in recipients of IL-21R^{-/-} CD4⁺ T cells was not due to an IL-17-related defect. During the preparation of this manuscript, a role for IL-17 in GVHD was reported (38–40). These reports varied, but one suggested that the lack of IL-17 promotes GVHD (38). Another report suggested that IL-17^{-/-} CD4⁺ T cells can ameliorate GVHD only at the early stages, which suggested a promoting effect for IL-17 at an early stage of GVHD (39). The third report suggested that ex vivo-differentiated Th17 cells induced skin and lung GVHD (40). Thus, the role of IL-17 may be complex and dependent on the specific experimental conditions.

Because there are reciprocal relationships between Th1/Th2 and Treg cell differentiation (41–43) and between IL-21 and Treg cell differentiation (8), we investigated the level of Treg cells in the spleens of recipients. Foxp3⁺CD4⁺ T cells were increased in percentage and absolute number but still represented only ~1% of splenocytes. Regarding the relationship between the defective effector T cell function and the increased number of Treg cells, it is possible that increased Treg cells suppress functional effector T cells. Alternatively, it is possible that effector differentiation itself is defective, and the resulting effector T cells cannot respond to alloantigen, analogous to the situation in T cell anergy, and that the increased Treg cell number is also a result of a dysregulated differentiation. Our results might be more consistent with the latter possibility, given that Treg cell depletion by anti-CD25 treatment did not alter the results in vitro and in vivo, although the efficiency of depletion of Foxp3⁺CD4⁺ T cells in vivo was incomplete. It is also conceivable that the upregulation of Foxp3 in CD25⁻CD4⁺ T cells (which would not be removed by CD25⁺ depletion) in the absence of IL-21 signaling might result in unresponsiveness or poor responsiveness of effector T cells and that more than one mechanism can contribute to the attenuated GVHD.

Disclosures

K. Ozaki and W.J.L. are inventors on patents and patent applications related to IL-21.

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False-positive GM test in myeloma patients

**High incidence of false-positive *Aspergillus* galactomannan test
in multiple myeloma**

Yasuo Mori,^{1,2} Yoji Nagasaki,^{1,2} Kenjiro Kamezaki,¹ Katsuto Takenaka,¹

Hiromi Iwasaki,² Naoki Harada,¹ Toshihiro Miyamoto,¹ Yasunobu Abe,³

Nobuyuki Shimono,¹ Koichi Akashi,^{1,2} and Takanori Teshima²

¹Medicine and Biosystemic Science, ²Center for Cellular and Molecular Medicine,
and ³Medicine and Bioregulatory Science, Kyushu University Graduate School
of Medical Sciences, Fukuoka, Japan

Corresponding author: Takanori Teshima, MD, PhD

Center for Cellular and Molecular Medicine, Kyushu University Hospital, 3-1-1
Maidashi, Higashi-ku, Fukuoka 812-8582, Japan.

Phone: +81-92-642-5947, Fax: +81-92-642-5951

E-mail: tteshima@cancer.med.kyushu-u.ac.jp

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False-positive GM test in myeloma patients

Invasive aspergillosis (IA) remains one of the most significant causes of morbidity and mortality in patients with hematological malignancies undergoing chemotherapy and hematopoietic stem cell transplantation (HSCT), mainly due to the difficulty in its early diagnosis. Monitoring of galactomannan (GM) antigen, an exoantigen of *Aspergillus*, in the blood by sandwich ELISA is a useful and non-invasive method for early diagnosis of IA. The GM test has a sensitivity of 67-100% with a specificity of 81-99% in neutropenic patients and allogeneic transplant recipients (1-3). Although it has been widely used as a diagnostic criterion for IA (4, 5), one of the major limitations of this assay is false-positivity, particularly in pediatric patients (1), patients with graft-versus-host disease (GVHD) (6, 7), and those taking dietary GM (8, 9) or fungus-derived antibiotics, such as piperacillin-tazobactam (PIPC/TAZ) (10-12).

Multiple myeloma results from malignant proliferation of a single clone of plasma cells, which produces a monoclonal immunoglobulin. Opportunistic infection is a major cause of death in patients with myeloma (13, 14). The risk for infection primarily resides during periods of chemotherapy-induced neutropenia or in the terminal stages of the disease. Therefore, monitoring of *Aspergillus* is recommended during chemotherapy-induced neutropenia.

124 patients with hematological disorders hospitalized in our institution from April 2007 to September 2009 were analyzed retrospectively. The clinical characteristics of these patients are summarized in supplementary materials (Table 1) Twenty-seven patients had plasma cell associated disorders (IgG type

False-positive GM test in myeloma patients

myeloma: 14, IgA type myeloma: 1, IgD type myeloma: 1, Bence-Jones type myeloma: 3, lymphoplasmacytic lymphoma/Waldenstrom macroglobulinemia: 3, plasma cell leukemia: 1, primary AL amyloidosis: 1, and POEMS syndrome: 2). The remaining 97 patients were diagnosed as acute leukemia (MDS/AML: 32, ALL: 15) or malignant lymphoma (n = 50). Out of the 124 patients, those receiving cytotoxic chemotherapy, autologous peripheral blood stem cell transplantation (PBSCT), and allogeneic HSCT were 81, 10, and 28, respectively. 111 patients received antifungal prophylaxis, mostly with FLCZ or ITCZ. Eight patients were administered PIPC/TAZ at sampling time points. Seventy patients were low-risk for the development of IA, while the remaining 54 patients were high-risk.

In 21 of the 124 (16.9%) patients, GM antigenemia was positive at least 2 consecutive times and their characteristics are shown in Table 1. However, only 7 of the 21 patients showed clinical features of IA and were diagnosed with probable IA in the lung (cases 1-7). Clinical features were relieved with treatment with VRCZ in these patients, confirming a diagnosis of IA. All of the 7 patients had received antifungal prophylaxis, but not antibiotics known to cause false-positive results, such as PIPC/TAZ (10-12), amoxicillin/clavulanic acid, or amoxicillin (17). Four of the 7 patients were high-risk for the developing IA and 3 were low-risk. Possible IA was diagnosed in 4 patients in the absence of positive GM results. On the other hand, no proven or probable IA was detected in 103 patients with negative GM antigenemia.

False-positive GM test in myeloma patients

Fourteen of the 21 (66.7%) patients with positive GM antigenemia did not satisfy the diagnostic criteria of proven or probable IA (cases 8-21 in Table 2); thus their episodes were considered to be false-positive. None of the 14 patients were treated with antibiotics potentially causing false-positivity of GM test. Antifungal prophylaxis had been given in all 14 episodes (FLCZ in 5, ITCZ in 5, and MCFG in 4). These patients did not show any clinical features suggestive of IA. Chest CT scans did not show any abnormal findings in 10 patients, while 4 patients showed abnormalities in the lung which were not suspicious of IA. Diagnosis of these lung lesions were history of pneumoconiosis in one patient, proven bacterial pneumonia in one, and idiopathic interstitial pneumonia in the remaining 2 patients. These lesions were not deteriorated without antifungal treatment. (1→3)-β-D-glucan was negative in all patients showing false-positive GM test. With a median follow-up of 10 months (range: 1-19 months), these patients did not develop fungal infection without treatment. On the other hand, false negative GM results were obtained in 4 patients (3.2 %). In the current study, the sensitivity, specificity, PPV, and NPV of the GM ELISA test were 63.6% (7/11), 87.6% (99/113), 33.3% (7/21), and 96.1% (99/103), respectively.

Surprisingly, 11 out of the 14 patients showing the false-positive results had diagnosis of multiple myeloma. The false-positivity of GM antigenemia was significantly higher in myeloma patients (11/22, 50%) than those with other hematological malignancies (3/102, 2.9%) ($p < 0.001$). Moreover, in myeloma patients false-positive results were exclusively detected in those with IgG subtype. Thus, rate of false-positivity was extremely high in patients with IgG

False-positive GM test in myeloma patients

myeloma (11/14, 78.6%). We could not find any difference in the characteristics of IgG myeloma patients with or without GM-false positivity, including serum levels of IgG (5083 ± 2077 mg/dl versus 4713 ± 3729 mg/dl). In addition, the GM test remained to be false-positive even after normalization of IgG levels by chemotherapy in 8 of the 11 patients. We also evaluated GM antigenemia in frozen serum samples collected prior to chemotherapy in 3 myeloma patients who showed false-positivity after chemotherapy to rule out the possibility that administration of myeloma-specific chemotherapy is associated with the false-positivity, and confirmed GM positivity prior to chemotherapy in these samples.

In a univariate logistic regression analysis, IgG myeloma and low risk category were strongly associated with false-positive GM antigenemia. Sex, type of treatment, antibiotics, corticosteroid usage, and serum levels of immunoglobulins were also significant or marginally associated with false-positivity. Multivariate analysis confirmed diagnosis of IgG myeloma as the only independent risk factor for false-positivity (odds ratio, 59.41; 95% confidence interval, 8.19 – 431.0; $p < 0.001$) (see supplementary materials, Table 2). In patients with other diseases, the GM assay showed a high sensitivity (7/11, 63.6%), specificity (96/99, 97.0%), PPV (7/10, 70%), and NPV (96/100, 96%). In contrast, for patients with IgG myeloma, specificity and PPV of the assay were very low (3/14, 21.4%, and 0/11, 0%), while NPV was 100% even in this cohort.

False-positive GM test in myeloma patients

A recent meta-analysis addressing the accuracy of a GM assay for diagnosing IA confirmed the clinical usefulness of this test with a sensitivity of 71% and a specificity of 89% (18). Although our study demonstrated similar sensitivity (7/11:64%) and specificity (103/117:88%) of the GM test, PPV (33%) was lower, compared to previous studies that demonstrate 40-60% PPV (7, 19-21). This difference is due to an unexpectedly high incidence of GM antigen false-positivity (11.3%) in our study. It should be noted, however, that screening of GM antigen was performed less frequently in this study compared to previous studies, where GM antigenemia was evaluated 2 to 3 times per week (1, 2, 22), and such a frequent monitoring is ideal to assure the optimal PPV and NPV.

Diagnosis of multiple myeloma is a major risk factor for GM false-positivity. In particular, the false-positivity was exclusively observed in patients with IgG myeloma and was not observed in patients with other types of plasma cell disorders. These observations should be confirmed in a larger study because some studies previously reported the usefulness of GM antigen assay as a diagnostic tool for IA among patients with hematological malignancies including IgG myeloma (23, 24), and only small numbers of patients with plasma cell disorders other than IgG myeloma were included in this study. Low risk category of developing IA was a risk for false positive results in a univariate, but not multivariate analysis. IgG myeloma remained a strong risk for false positivity even after the compensation by the risk categorization.

Mechanisms of high frequency of GM false-positivity in myeloma

False-positive GM test in myeloma patients

patients remain to be investigated. (1→3)-β-D-glucan, which is released from the fungal cell wall, is also widely used to support diagnosis of fungal infections and adopted as one of the microbiological criterion for probable IA in the revised EORTC/MSG definition (5). A previous study reported that high levels of immunoglobulins interfere with the measurement of (1→3)-β-D-glucan by causing precipitation of insolubilized proteins and increase the non-specific optical density levels of reaction fluid (25), although (1→3)-β-D-glucan was negative in patients showing false-positive GM test in this study. This phenomenon has not been reported in the GM assay. However, serum levels of IgG were not directly associated with the false-positivity; IgG levels did not differ between IgG myeloma patients with and without false-positivity. In addition, the GM test remained to be false-positive even after normalization of IgG levels by chemotherapy in 8 of the 11 patients.

Causative role of PIPC/TAZ, amoxicillin/clavulanic acid, and amoxicillin in GM false-positivity has been well documented (10-12, 17), therefore collection of samples prior to infusion of these antibiotics and the use of a relatively higher cut-off level (> 0.7) are recommended in patients receiving these agents (12). In this study, no patients with GM false-positivity received these antibiotics at the time of sampling. It has been hypothesized that dietary contamination by GM causes GM false-positivity by the translocation of dietary GM into the systemic circulation through the disrupted intestinal mucosal barrier, especially in patients with gastrointestinal GVHD after allogeneic HSCT (6, 7, 9). In our cohort, 1 (3.6%) of the 28 patients who underwent allogeneic HSCT showed

False-positive GM test in myeloma patients

false-positivity. This patient with lymphoma had acute GVHD involving in the skin and intestine. A previous study demonstrated that false-positive results were preferentially observed in patients with febrile neutropenic sepsis (26), although subsequent study was unable to replicate this result (22). A recent study revealed that serum GM antigen levels was significantly higher in severely neutropenic patients ($< 0.1 \times 10^9/L$) than in the other patients (27). However, we did not find such an association in this study.

In conclusion, the incidence of false-positive GM antigenemia was high in patients with IgG myeloma. Although the results should be confirmed in a prospective study including larger numbers of patients, positive results of GM antigenemia may be interpreted with caution, and intimate survey including CT scan or other microbiological markers will be recommended in myeloma patients.

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False-positive GM test in myeloma patients

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False-positive GM test in myeloma patients

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False-positive GM test in myeloma patients

Table 1. Characteristics of 21 patients with positive GM test

Case	Age /Sex	Disease	Treatment	GM (C.O.I.)	Radiological / clinical findings	Diagnosis	Times / duration of false positivity	Antibiotics	Antifungal agents*	Steroids	Neutrophils (x 10 ⁹ /L)	IgG (mg/dl)	IgA (mg/dl)	IgM (mg/dl)	Risk of IA
1	49/F	MDS/AML	CTx	1.1	nodules with halo	probable IA	-	CFPM	FLCZ	none	0.01	1419	227	50	High
2	63/M	MDS/AML	CTx	1.2	nodules	probable IA	-	MEPM, AMK	FLCZ	none	0	1253	238	62	High
3	50/F	AML	CTx	1.2	nodules	probable IA	-	MEPM, AMK	ITCZ	none	0	1301	293	235	High
4	57/F	MDS	allo-SCT	2.3	nodules	probable IA	-	CAZ, VCM	VRCZ	none	0.009	1198	256	120	High
5	63/M	ML	CTx	1.3	consolidation with pleural pain	probable IA	-	none	MCFG	DEXA	2.034	689	126	92	Low
6	57/F	ML	CTx	1.5	nodules	probable IA	-	CFPM	FLCZ	PSL	0.081	675	117	67	Low
7	59/F	ML	CTx	2.2	consolidation with pleural pain	probable IA	-	CZOP	MCFG	PSL	0.036	719	95	20	Low
8	70/M	ML	CTx	0.8	pneumoconiosis	F-P	7/7 months	none	FLCZ	PSL	5.096	1870	315	60	Low
9	52/M	ML	allo-SCT	2.5	bacterial pneumonia	F-P	2/1 month	CZOP, CPFX	ITCZ	mPSL	0.806	562	49	19	High
10	54/M	ML	CTx	0.9	negative	F-P	5/3 months	CFPM	FLCZ	none	0.395	691	46	10	Low
11	63/F	IgGk MM	CTx	0.8	negative	F-P	2/1 month	none	ITCZ	DEXA	1.497	4770	25	27	Low
12	56/F	IgGk MM	none	1.2	negative	F-P	3/2 months	CAZ	FLCZ	none	3.449	7196	5	38	Low
13	69/M	IgGk MM	CTx	0.5	negative	F-P	4/4 months	none	MCFG	DEXA	0.72	3944	13	<10	Low
14	69/M	IgGk MM	CTx	1.2	negative	F-P	6/3 months	none	FLCZ	DEXA	1.777	2385	6	28	Low
15	64/M	IgGk MM	CTx	1.6	old inflammatory change	F-P	5/2 months	none	MCFG	PSL	1.919	3269	30	<10	Low
16	51/F	IgGk MM	CTx	2.1	IP	F-P	23/15 months	none	ITCZ	PSL	3.791	8006	99	30	Low
17	64/M	IgGk MM	CTx	1.6	negative	F-P	13/8 months	none	ITCZ	DEXA	0.899	3950	6	<10	Low
18	75/M	IgGk MM	CTx	0.6	negative	F-P	2/1 month	MEPM	FLCZ	DEXA	0.823	8986	<5	<10	High
19	49/F	IgGk MM	CTx	1.1	negative	F-P	8/3 months	none	MCFG	DEXA	0.724	4563	37	15	Low
20	74/F	IgGA MM	CTx	0.7	negative	F-P	6/2 months	none	MCFG	DEXA	1.451	3924	25	18	Low
21	56/M	IgGA MM	CTx	1.1	negative	F-P	10/3 months	none	ITCZ	DEXA	1.605	4917	9	10	Low

* administered at initial positive sampling, MDS : myelodysplastic syndrome, AML : acute myelogenous leukemia, ML : malignant lymphoma, MM: multiple myeloma, CTx : chemotherapy, MEPM : meropenem, AMK : amikacin, CAZ : ceftazidime, VCM : vancomycin, CZOP : ceftazopran, F-P : false-positive, CFPM : cefepime, CPFX : ciprofloxacin, mPSL : prednisolone, PSL : piperacillin/tazobactam, CPFX : ciprofloxacin, PSL : prednisolone, mPSL : prednisolone, DEXA : dexamethasone.