



Human MAIT cells are devoid of alloreactive potential: prompting their use as universal cells for adoptive immune therapy

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ABSTRACT

Background Mucosal-associated invariant T (MAIT) cells are semi-invariant T cells that recognize microbial antigens presented by the highly conserved MR1 molecule. MAIT cells are predominantly localized in the liver and barrier tissues and are potent effectors of antimicrobial defense. MAIT cells are very few at birth and accumulate gradually over a period of about 6 years during the infancy. The cytotoxic potential of MAIT cells, as well as their newly described regulatory and tissue repair functions, open the possibility of exploiting their properties in adoptive therapy. A prerequisite for their use as ‘universal’ cells would be a lack of alloreactive potential, which remains to be demonstrated.

Methods We used *ex vivo*, *in vitro* and *in vivo* models to determine if human MAIT cells contribute to allogeneic responses.

Results We show that recovery of MAIT cells after allogeneic hematopoietic stem cell transplantation recapitulates their slow physiological expansion in early childhood, independent of recovery of non-MAIT T cells. *In vitro*, signals provided by allogeneic cells and cytokines do not induce sustained MAIT cell proliferation. *In vivo*, human MAIT cells do not expand nor accumulate in tissues in a model of T-cell-mediated xenogeneic graft-versus-host disease in immunodeficient mice.

Conclusions Altogether, these results provide evidence that MAIT cells are devoid of alloreactive potential and pave the way for harnessing their translational potential in universal adoptive therapy overcoming barriers of HLA disparity.

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INTRODUCTION

Mucosal-associated invariant T (MAIT) cells are innate-like T cells expressing a semi-invariant T-cell receptor (TCR) which recognizes microbial-derived riboflavin (vitamin B₂) precursors such as 5-(2-oxopropylidene amino)-6-D-ribitylaminoouracil (5-OP-RU) presented by MR1, a monomorphic major

histocompatibility antigen (MHC) class-I like molecule.^{1–3} On TCR engagement, MAIT cells proliferate, release proinflammatory cytokines and kill target cells, supporting their role in antimicrobial defense.^{4–6} Most bacteria and yeasts, but not mammal cells, are able to synthesize riboflavin and hence provide MR1 ligands.⁷ This TCR-MR1 recognition pathway, therefore, represents a sophisticated discriminatory mechanism to target microbial antigens while sparing the host. MAIT cells can also be activated in a TCR-independent way in response to inflammatory cytokines such as interleukin 12 (IL-12) and IL-18.^{8–10} MAIT cells are predominantly localized in the liver and barrier tissues, in agreement with their expression of several chemokine receptors.^{4,11} They are also abundant in the adult blood, but very few in cord blood. We previously showed that the postnatal expansion of MAIT cells was a very slow process requiring at least 5–6 years to reach adult levels, and likely resulted from repeated encounters of a few MAIT cell clones with MR1-restricted microbial antigens.¹²

The curative effect of allogeneic hematopoietic stem cell transplantation (HSCT) in hematological malignancies is based on the capacity of donor T cells to eliminate residual tumor cells (graft-vs-leukemia effect). A serious drawback is that donor T cells may also recognize normal non-hematopoietic cells, leading to graft-versus-host disease (GVHD), which is characterized by activation, expansion and migration to target tissues of donor alloreactive T cells.^{13–17} In the first weeks after HSCT, the T cell compartment recovers through peripheral expansion of graft-derived T cells in response to increased levels of homeostatic cytokines and to host’s



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allogeneic antigens. Reconstitution of a fully diversified T-cell repertoire occurs only later by resumed thymic output of newly generated naïve T cells.¹⁸ Even under favorable conditions, it takes at least 2 months to produce naïve T cells, and a plateau of thymic output is reached only after 1–2 years.

We observed that MAIT cell recovery was delayed up to 6 years after allogeneic cord blood transplantation, mimicking the long postnatal expansion period.¹² Other studies confirmed that MAIT cell numbers failed to normalize for at least a year after HSCT^{19–21} and suggested that a low MAIT cell number was associated with an increased risk of GVHD, although the mechanisms remain unclear.^{19–21, 22} In mouse HSCT models, recipient's residual MAIT cells protected mice from acute intestinal GVHD through microbial-induced secretion of IL-17 promoting gastrointestinal tract integrity and reducing alloantigen presentation, ultimately limiting the proliferation of donor-derived alloreactive T cells.²³ Recently, transcriptomic and functional studies have revealed new tissue repair and regulatory functions of MAIT cells.^{24–28} Altogether, these results open the possibility of exploiting the effector or tissue repair/regulatory properties of MAIT cells in adoptive immunotherapy, provided that they do not cause alloreactivity. Here, we used *in vitro* and *in vivo* models to provide evidence that human MAIT cells are devoid of alloreactive potential. Our results pave the way for harnessing the translational potential of MAIT cells in universal adoptive therapy overcoming barriers of HLA disparity.

PATIENTS AND METHODS

Patients

Three independent cohorts of patients undergoing allogeneic HSCT for hematological malignancy were studied. Patient and graft characteristics are described in [table 1](#).

Cohort 1 included 40 children recipients, for whom peripheral blood mononuclear cells (PBMCs) samples were prospectively collected at Robert Debré Hospital between January 2013 and December 2015. Blood samples were collected prior to conditioning (\pm day –15 before HSCT) and at 1, 3, 6, 12 and 24 months after HSCT as the standard of care for assessment of immunologic recovery. Patients who died before day 180 were not included in the analysis.

Cohort 2 included 64 additional HSCT children in stable remission for whom PBMCs samples were collected at time of a routine visit at Robert Debré Hospital 2–16 years after HSCT.

All children from cohorts 1 and 2 received unmanipulated bone marrow transplant from HLA-matched sibling donor (MSD) or unrelated donor (MUD). Myeloablative conditioning was provided with VP16 and total body irradiation (TBI), or with cyclophosphamide and busulfan. *In vivo* T-cell depletion by ATG was given in the majority of MUD recipients. Primary prophylaxis of GVHD consisted

of a calcineurin inhibitor alone (MSD recipients) or with methotrexate (MUD recipients).

Cohort 3 included 49 adult donor/recipient pairs for whom frozen annotated PBMCs were provided by the CRYOSTEM consortium (<https://doi.org/10.25718/cryostem-collection/2018>) and Société Francophone de Greffe de Moelle-Thérapie Cellulaire. Patients received bone marrow or peripheral blood stem cells from a MSD and were given myeloablative or non-myeloablative conditioning. Blood samples were collected before conditioning, and at 3 and 12 months post-HSCT in the absence of acute GVHD (aGVHD). In case of aGVHD, samples were collected at the time of diagnosis before any treatment, 1 month later (ie, toward 3 months post-HSCT) and at 12 months.

Cells and reagents

PBMCs were isolated and used immediately or frozen. 5-OP-RU was synthesized as described in.^{29–31} Human MAIT cells were expanded for 6 days in human T-cell culture medium (RPMI-1640, Invitrogen, Life Technologies) containing 10% human AB serum (EuroBio), IL-2 (100 U/mL, Miltenyi) and 300 nM 5-OP-RU.

Flow cytometry

MAIT cells were analyzed in fresh whole blood, or in isolated PBMCs where indicated. Multiparametric 14-color flow cytometry analyses were performed as described in online supplemental methods. MAIT cells were defined as CD3⁺CD4⁺CD161^{high}V α 7.2⁺ cells in the first part of the study (HSCT patients). This population fully overlapped with the population labeled by MR1:5-OP-RU tetramers (³¹ and [figure 1A](#)). Thereafter, we used the specific MR1:5-OP-RU tetramer when it became available from the National Institutes of Health (NIH) tetramer core facility.

In vitro stimulations

Human carboxyfluorescein succinimidyl ester (CFSE)-labeled (5 μ M) PBMCs were cultured in RPMI-1640 supplemented with IL-2 (100 U/mL), IL-15 (50 ng/mL), IL-7 (50 ng/mL, all from Miltenyi Biotec), or IL-12/IL-18 (50 ng/mL each, R&D Systems) and/or 300 nM 5-OP-RU. For mixed lymphocyte reactions (MLR), CFSE-labeled PBMCs used as responders (1×10^6 /mL) were incubated with γ -irradiated allogeneic stimulator PBMCs (1:1 ratio) in 96-well round-bottom plates. Cells were harvested at day 6 and stained before flow cytometry analysis.

Adoptive transfer of xenogeneic cells

NOD-Scid-IL-2R γ^{null} (NSG) mice (Jackson laboratory, Bar Harbor, Michigan, USA) were bred in the animal facility of St-Louis Research Institute and housed under specific pathogen-free conditions. Eight- to 10-week-old female mice were irradiated (1.3 Gy) 24 hours prior to injection of 5×10^6 human PBMCs (huPBMCs) in the caudal vein. Development of GVHD was monitored three times per week based on weight loss, hunching posture, reduced mobility and hair loss. Human chimerism in peripheral

Table 1 Characteristics of HSCT recipients

	Children longitudinal cohort	Children cross-sectional cohort	Adults retrospective cohort
Patients (n)	40	64	49
Sex, n (%)			
Female	17 (42.5)	23 (36)	18 (36.7)
Male	20 (57.5)	41 (64)	31 (63.3)
Age, mean years (range)	9.4 (0.75–16)	7 (0.9–17.8)	44.4 (17–65.8)
Primary disease, n (%)			
ALL	23 (57.5)	25 (39.1)	6 (12.2)
AML	11 (27.5)	11 (17.2)	19 (38.8)
JMML	3 (7.5)	2 (3.1)	–
NHL	1 (2.5)	1 (1.6)	13 (26.5)
MDS	1 (2.5)	2 (3.1)	5 (10.2)
CML	1 (2.5)	3 (4.7)	2 (4.1)
Others	–	20 (31.2)	4 (8.2)
Donor, n (%)			
MSD	19 (47.5)	32 (50)	49 (100%)
MUD	21 (52.5)	32 (50)	–
Stem cell sources, n (%)			
Peripheral blood	–	–	30 (61.2)
Bone marrow	40 (100)	64 (100)	19 (38.8)
Conditioning, n (%)			
MA	40 (100)	64 (100)	24 (49)
NMA	–	–	25 (51)
In vivo T-cell depletion, n (%)	12 (30)	18 (28)	20 (40.8)
Acute GVHD			
Grades 1–2	24 (60)	NA	
Grades 3–4	9 (22.5)	NA	4 (8.2)

ALL, acute lymphoblastic leukemia; AML, acute myeloid leukemia; CLL, chronic lymphoblastic leukemia; CML, chronic myeloid leukemia; JMML, juvenile myelomonocytic leukemia; MDS, myelodysplastic syndrome; MSD, matched sibling donor; MUD, matched unrelated donor; NHL, non-Hodgkin's lymphoma.

blood (per cent of human CD45⁺ cells) was assessed weekly. When indicated, mice received 1 nmol 5-OP-RU and/or 100 ng human IL-15 and/or 10 µg CpG Oligonucleotide (class B ODN, Invivogen) intraperitoneally (i.p) every 2 days from the day of PBMCs infusion. Mice were sacrificed at the indicated time, or when weight loss was >15%. Peripheral blood, spleen, liver, lungs and intestine were harvested, and cells were isolated as described in online supplemental methods.

Statistics

Differences between groups were analyzed using non-parametric tests for paired (Wilcoxon) and unpaired (Mann-Whitney or Kruskal-Wallis) groups, or two-way analysis of variance. Correlations were assessed using the Spearman's rank correlation. Two-sided values of $p < 0.05$ were considered significant. Analyses were performed using Prism software V.9 (GraphPad).

All data including outliers were included with one predetermined exception: flow cytometry cell-subset percentages were considered non-evaluable if the parent subset contained <100 events.

RESULTS

MAIT cell reconstitution is delayed for several years after HSCT

Our previous findings have shown that it takes up to 6 years to recover normal MAIT cell values after cord blood transplantation, suggesting that MAIT cells do not proliferate in response to allogeneic stimulation.¹² However, MAIT cells in cord blood are naïve, and their number is 1–2 logs lower than in adult blood, which could contribute to this slow recovery. We, therefore, analyzed the kinetics of reconstitution of MAIT cells and non-MAIT CD3⁺ T

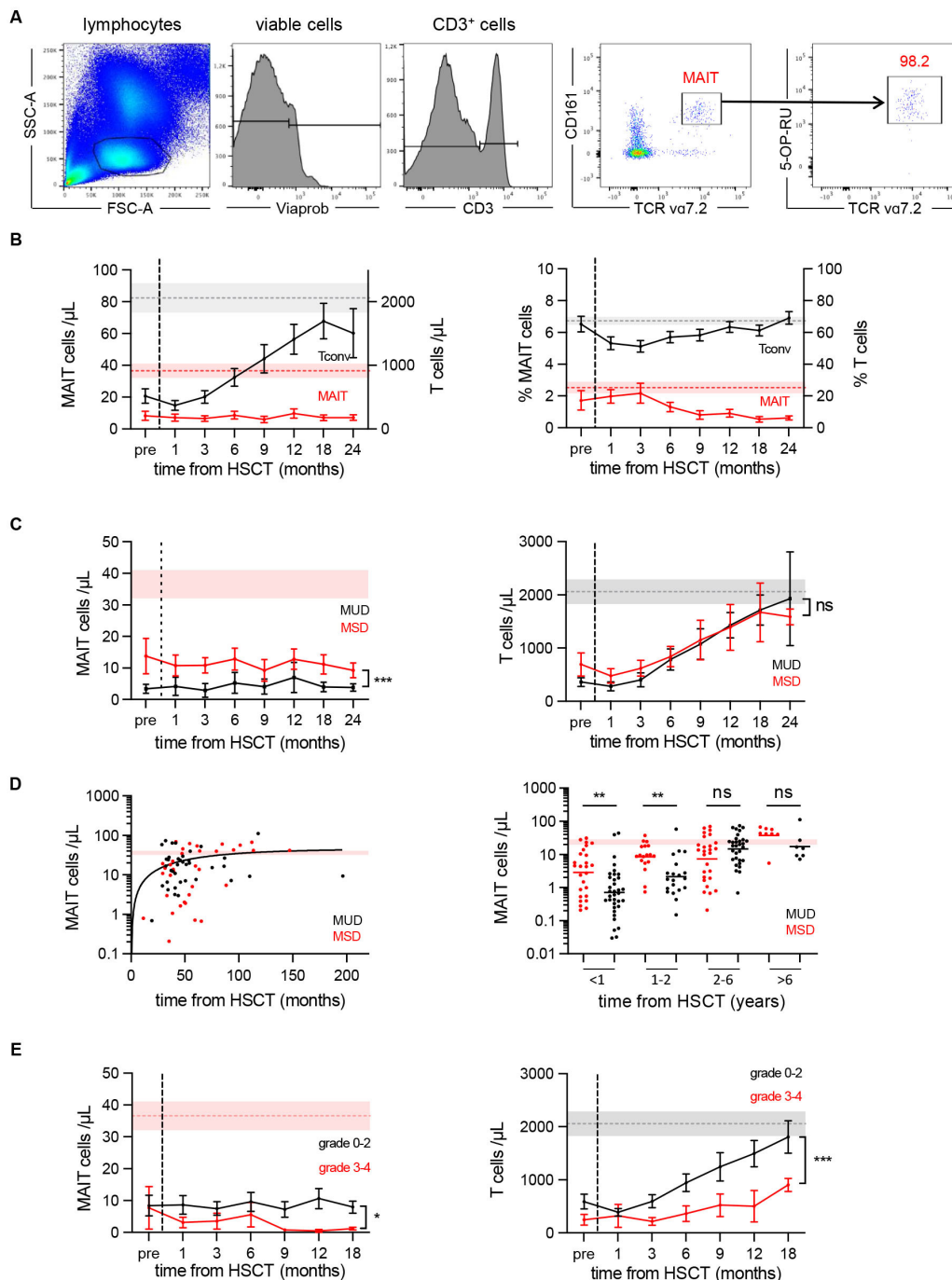


Figure 1 MAIT cell recovery takes several years after HSCT in children. (A) Gating strategy and representative staining of $V\alpha 7.2^+$ $CD161^{\text{high}}$ MAIT cells in the $CD3^+$ $CD4^-$ T cell gate in the peripheral blood. This population fully overlaps with the population labeled by MR1:5-OP-RU tetramers, as shown in the right panels. The frequencies of MAIT cells are expressed as a percentage of $CD3^+$ lymphocytes. (B) Absolute counts (left panel) and percentages (right panel) of MAIT and non-MAIT T cells detected in the peripheral blood of 40 children before conditioning (pre) and from month 1 to month 24 after HSCT. Data are shown as mean \pm SEM. (C) Recovery dynamics of MAIT cells (left panel) and T cells (right panel) in transplant recipients of matched sibling donors (MSD, red line) and unrelated donors (MUD, black line). Results show the mean absolute numbers \pm SEM during the first 24 months after HSCT. Two-way ANOVA, *** $p=0.0005$; ns: not significant. (D) MAIT cell recovery takes at least 6 years to reach normal values. Left panel: Relationship between log₁₀-transformed MAIT cell absolute numbers and time from transplantation up to 16 years after HSCT in MSD (red dots) and MUD (black dots) recipients. Right panel: Recovery is slower in MUD than MSD recipients for at least to 2 years after HSCT. Mann-Whitney U test ** $p\leq 0.002$; ns: not significant. (E) Comparison of MAIT (left panel) and T (right panel) cell recovery in patients without or with mild (grade 0–2) or severe (grade 3–4) aGVHD. Results show the mean absolute numbers \pm SEM 2-way ANOVA, * $p=0.04$, *** $p=0.0006$. Light gray and red areas show the reference physiological T cell and MAIT cell count interval, respectively, obtained from 36 age-matched healthy donors. aGVHD, acute graft-versus-host disease; ANOVA, analysis of variance; HSCT, hematopoietic stem cell transplantation; MAIT, mucosal-associated invariant T.

cells (hereafter referred to as T cells, which contain other unconventional T cell subsets not excluded from our analysis) in other HSCT settings.

Forty children who received an unmanipulated bone marrow transplant after myeloablative conditioning were studied longitudinally up to 24 months after HSCT. While the number of T cells gradually increased from 1 month after transplantation and returned almost to normal after 1 year, there was virtually no increase in the number of MAIT cells during the study period (figure 1A,B). Two years after HSCT, MAIT cell values remained five times lower than in age-matched donors (figure 1B). The number of and MAIT cells before and up to 3 months after HSCT was lower in MUD than in MSD recipients, likely due to a longer time to transplant and more frequent use of in vivo T-cell depletion in MUD recipients. However, while T cells reached comparable values after 3 months regardless of the donor type, MAIT cell numbers remained around two times lower in MUD recipients than in MSD recipients during the 24 months follow-up (figure 1C).

To extend our findings to a longer follow-up period, we analyzed 64 additional children at time of a routine visit 2–16 years after transplantation. As observed after cord blood transplantation,¹² the number of MAIT cells increased very slowly (even more slowly in MUD than MSD recipients), to reach a plateau approximately 6 years after HSCT (figure 1D). This slow recovery was not related to the underlying malignancy, gender or age of the recipient, pre-HSCT conditioning (with or without TBI), severe microbial infection during the first months after HSCT, or duration of immunosuppressive treatment (data not shown).

T-cell reconstitution is impaired in patients with aGVHD, at least in part because of defective thymic production of HSC-derived T cells.¹⁸ We previously observed that thymus-derived naïve MAIT cells appeared in cord blood recipients 6 months after transplantation.¹² Here, we found that the number of MAIT cells after 6 months was lower in patients with severe (grades 3–4) aGVHD compared with those without or mild (grades 0–2) aGVHD, as also observed for non-MAIT T cells (figure 1E). These data suggest that the alteration of thymic function associated with aGVHD participates in the defective recovery of MAIT cells after HSCT.

The reconstitution of MAIT cells after HSCT is affected by their number in the donor and by the existence of aGVHD

To further explore the potential link between MAIT cell numbers aGVHD and MAIT cell numbers, we retrospectively analyzed 49 adult HSCT donor/recipient pairs from the national Cryostem consortium, including 25 patients with aGVHD. As observed in children, MAIT cells did not significantly expand during the 1-year study period, while the number of T cells increased sharply. Moreover, absolute numbers of MAIT and T cells at 3 months were lower in patients with severe aGVHD than in those without or mild aGVHD, whereas subsequent reconstitution was not

affected (figure 2A). The proportion of Ki67⁺ cells at the time of aGVHD, which could represent alloreactive proliferating cells, was higher in T cells than in MAIT cells. One year after HSCT, Ki67 expression was hardly detectable in both cell types (figure 2B and online supplemental figure 1). Expression of activation (CD69, CD25) or maturation (CD45RO) markers was not modified at time of aGVHD (data not shown).

MAIT cell recovery was not influenced by the conditioning regimen (myeloablative or non-myeloablative) or by the source of HSC (peripheral blood or bone marrow) (data not shown).

Notably, the number of MAIT cells in the donor was significantly correlated to their number in the recipient after HSCT, whereas no correlation was observed for T cells (figure 2C). However, there was no association between the number of MAIT cells in the donor and the presence of severe aGVHD (figure 2D).

Altogether, these data indicate that the number of MAIT cells in the donor influences the early reconstitution of MAIT cells in the recipient, but do not provide evidence for a role of MAIT cells in aGVHD.

MAIT cells do not proliferate in response to alloantigen stimulation, except in the presence of MR1 ligand

Graft-derived conventional T cells expand through lymphopenia-induced homeostatic proliferation and response(s) to host's allogeneic antigens. However, not all T cells respond with the same efficiency to these proliferative cues.³²

We first analyzed the in vitro responsiveness of MAIT cells to homeostatic cytokines. CFSE-labeled PBMCs were treated with IL-7 or IL-15, or IL-2 as control, alone or in combination with the microbial-derived MR1 ligand, 5-OP-RU, and the proliferation of MR1-tetramer⁺ MAIT cells was monitored according to CFSE dilution (figure 3A,B). In the absence of cytokine, MAIT cells did not respond to 5-OP-RU alone, as reported.^{24,26} Most MAIT cells proliferated in response to IL-15 or IL-7 (but not to IL-2) alone, but the number of divisions was significantly higher when 5-OP-RU was added to IL-15 or IL-2.

Chemotherapy and radiation induce the secretion of IL-12 and IL-18, which may trigger early TCR-independent activation of MAIT cells transferred with the graft. We observed a low proliferative response of MAIT cells to IL-12/IL-18 combination, which increased in the presence of 5-OP-RU but remained lower than proliferation induced by IL-15 (figure 3A,B).

We next explored the capacity of MAIT cells to respond to allogeneic cells in a MLR. Unlike non-MAIT T cells, MAIT cells barely proliferated in response to allogeneic stimulation (figure 3C). However, the addition of 5-OP-RU to the MLR induced a strong proliferation of MAIT cells (figure 3D), suggesting that cytokines (IL-2 or other) produced by neighboring alloreactive T cells during the culture period allowed MAIT cells to proliferate only in response to the MR1 ligand.

Overall, these results suggest that signals provided by allogeneic cells and cytokines present in the post-HSCT

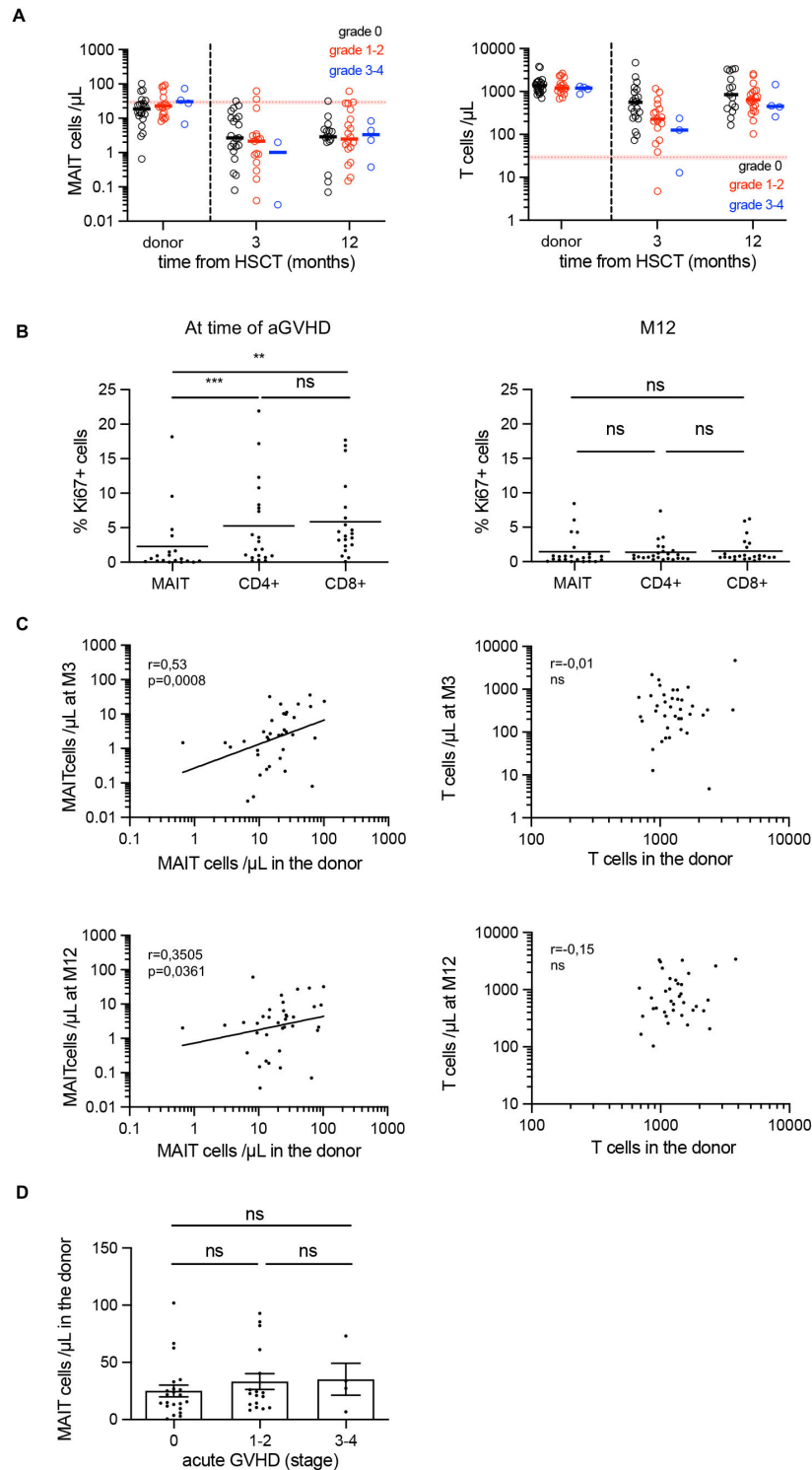


Figure 2 MAIT cell recovery in adult HSCT recipients. (A) Recovery kinetics of $V\alpha 7.2^+$ $CD161^{\text{high}}$ MAIT cells (left panel) and non-MAIT T cells (right panel) in patients from the CRYOSTEM cohort ($n=49$) according to the absence or the presence of mild (grade 1–2) or severe (grade 3–4) aGVHD. Patients with aGVHD were sampled at the time of disease diagnosis and 1 month later, that is, toward M3. Results show the individual and mean absolute numbers. Shaded areas show MAIT and T cell count intervals in the donors. Two-way ANOVA, not significant. (B) Quantification of $Ki67^+$ proliferating cells among MAIT, CD4 and CD8 T cells at time of aGVHD (left) and 12 months after HSCT (right). Scatter plots with bar showing the mean percentage of $Ki67^+$ cells. Wilcoxon non-parametric paired test, $**p=0.004$; $***p=0.0008$; ns: not significant. (C) Correlation between the number of MAIT (left) or T (right) cells in the donor and that in the recipient 3 months (top panels) and 12 months (bottom panels) after HSCT. Spearman correlation r and p values are indicated. (D) Comparison of the number of MAIT cells in the donor according to the absence or presence of mild (grade 1–2) or severe (grade 3–4) aGVHD. Kruskal-Wallis non-parametric paired test, ns: not significant. aGVHD, acute graft-versus-host disease; ANOVA, analysis of variance; HSCT, hematopoietic stem cell transplantation; MAIT, mucosal-associated invariant T.

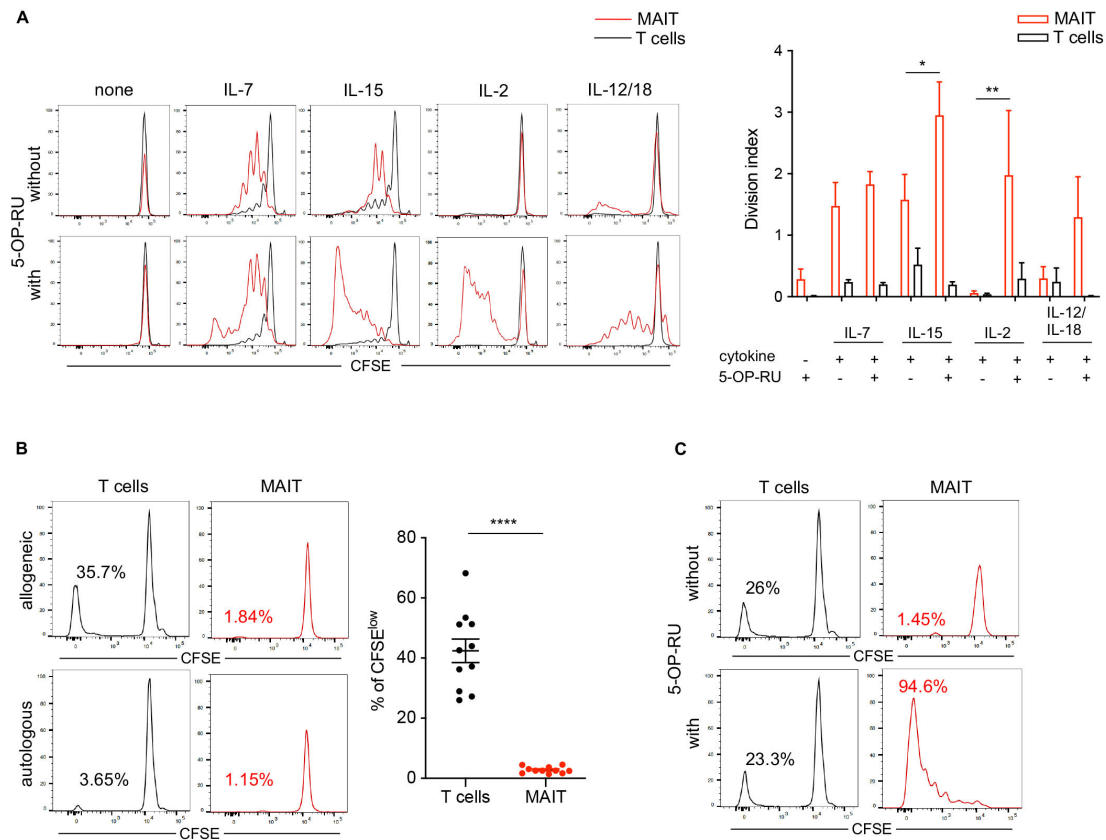


Figure 3 MAIT cell proliferative response to cytokines and allogeneic cells in vitro. Carboxyfluorescein succinimidyl ester (CFSE)-labeled human PBMCs (10^6 /mL) were cultured for 6 days in the indicated conditions. Proliferation of MR1:5-OP-RU tetramer⁺ MAIT cells (red lines) and non-MAIT T cells (black lines) was quantified by CFSE dilution at the end of the culture. (A, B) Proliferative response to stimulation by IL-7 (50 ng/mL), IL-15 (50 ng/mL), IL-2 (100 U/mL) and IL-12/18 (50 ng/mL each) in the presence or absence of the synthetic MR1 ligand, 5-OP-RU (300 nM). (A) Representative CFSE staining gated on MAIT and T cells after a 6 day culture. (B) Division index (mean number of divisions in the total population) of MAIT and T cells. Mean \pm SEM of 3 independent experiments. Paired t test, * $p=0.045$; ** $p=0.004$. (C) Mixed lymphocyte reaction: CFSE-labeled responder PBMCs were cultured with irradiated allogeneic PBMCs at 1:1 ratio. Responder T cells were identified at the end of the 6 days culture by gating on CD3⁺ T cells following exclusion of dead cells. Left panel: Representative CFSE staining gated on MAIT (red) and T (black) cells after 6 days culture in the presence of allogeneic (upper panel) or autologous (lower panel) PBMCs. Numbers in the quadrants indicate the percentage of proliferating (CFSE^{low}) cells among each population. Right panel: Individual values and means \pm SEM of proliferating T and MAIT cells ($n=11$ experiments using different recipient/donor pairs). Paired t test, **** $p<0.0001$. (D) Mixed lymphocyte reaction was performed in the absence (upper panel) or presence (lower panel) of 300 nM 5-OP-RU. Representative CFSE staining gated on MAIT (red) and T (black) cells after a 6 day culture. IL-7, interleukin 7; MAIT, mucosal-associated invariant T; PBMCs, peripheral blood mononuclear cells.

period are not sufficient to induce sustained expansion of MAIT cells.

Human MAIT cells do not expand in immunodeficient mice and do not cause xenogeneic GVHD

To further explore the potential of human MAIT cells to expand in vivo and participate to GVHD tissue lesions, we used a model of xenogeneic GVHD (xeno-GVHD) in which low doses of huPBMCs are injected into irradiated immunodeficient NSG mice. In this model, human T cells consistently expand in the mouse and mediate an acute GVHD-like syndrome with extensive T-cell tissue infiltration and damage of mouse skin, liver, intestine and lungs, resulting in death by 30–50 days.^{33 34}

Mice were injected with 5×10^6 huPBMCs, among which MAIT cells represented around 3% of T cells. The presence of CD45⁺ huPBMCs was determined at different times in

tissues of recipient mice, including those where MAIT cells are known to preferentially reside. Four weeks after transfer, variable proportions of CD45⁺ cells were found in peripheral blood and tissues, almost all of which were non-MAIT T cells and less than 0.05% were MAIT cells (figure 4A,B).

Next, mice injected with huPBMCs were monitored to evaluate aGVHD progression and euthanized when weight loss was >15% (± 45 days after transfer). A massive accumulation of T cells was observed in tissues, in particular in the spleen, lungs and liver. However, MAIT cells were very few in all compartments (figure 4C).

Human MAIT cells efficiently recognize the murine MR1 molecule,³⁵ ruling out a defective presentation of MR1 ligands to human MAIT cells in NSG mice. However, since MAIT cells do not proliferate significantly in vitro in the absence of 5-OP-RU, the availability of MR1 ligand could be a factor

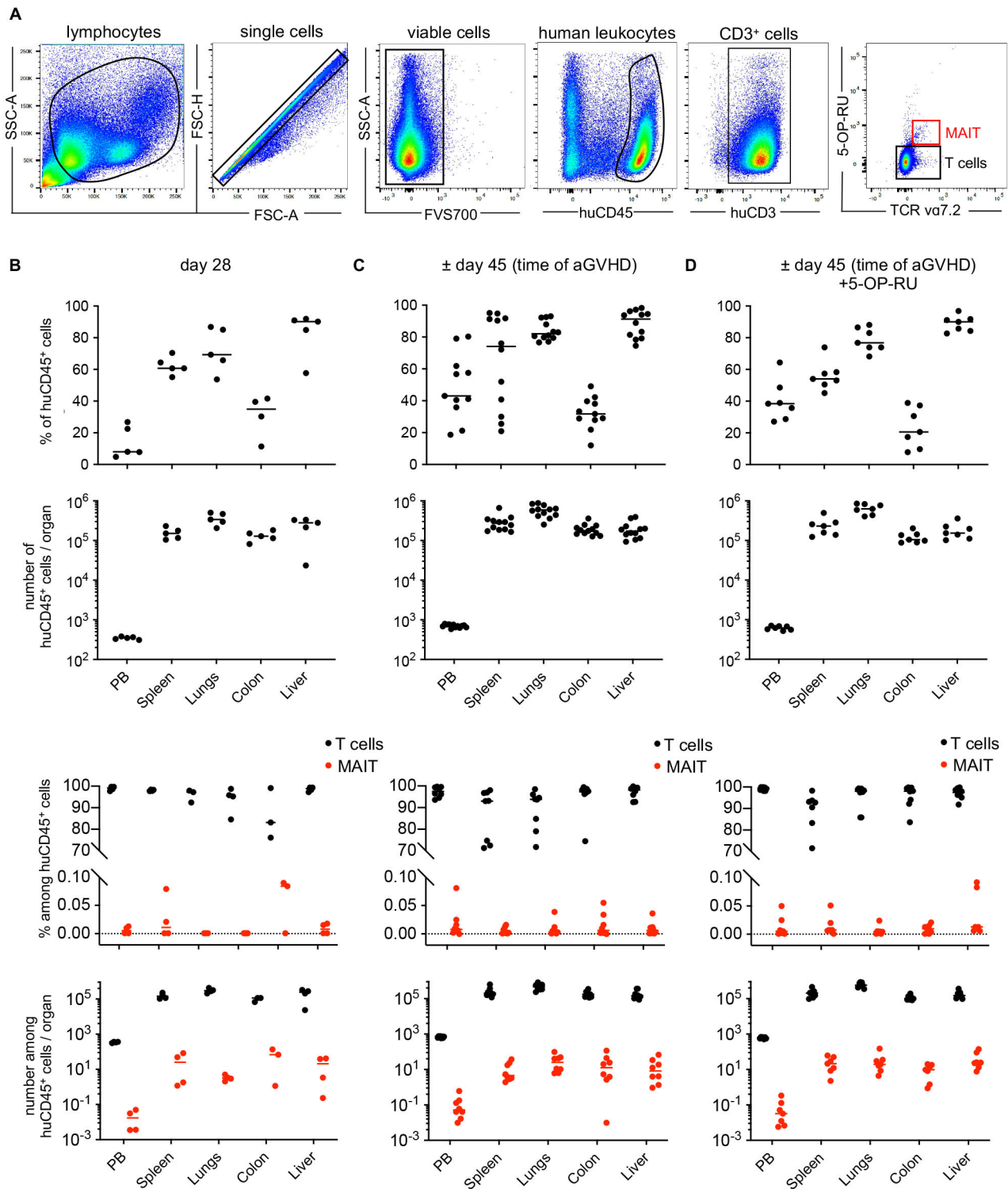


Figure 4 Absence of MAIT cell expansion in a xenogeneic GVHD model. Irradiated (1.3Gy) NSG mice injected with 5×10^6 huPBMCs were monitored to evaluate aGVHD progression and euthanized at the indicated time. Peripheral blood, spleen, liver, lungs and colon were harvested, and cells were isolated. (A) Gating strategy and representative staining of V α 7.2 tetramer⁺ MAIT cells and non-MAIT cells among viable human CD45⁺ CD3⁺ lymphocytes in the spleen. (B–D) The proportion of CD45⁺ leukocytes among viable cells (ie, human chimerism, upper panels), and the proportion of non-MAIT T cells (black dots) and MR1:5-OP-RU tetramer⁺ MAIT cells (red dots) among CD45⁺ cells (lower panels) were determined by flow cytometry. Results show individual values and median (horizontal bar). (B) Mice were sacrificed at day 28 after injection (n=5 mice). (C) Mice were euthanized when they exhibited signs of aGVHD (\pm day 45, n=11 diseased mice). (D) Mice received 5-OP-RU every 2 days (1 nmol i.p.) from the day of huPBMC injection and were euthanized at time of aGVHD (n=7 diseased mice). aGVHD, acute graft-versus-host disease; huPBMC, human peripheral blood mononuclear cell; MAIT, mucosal-associated invariant T; NSG, NOD-Scid-IL-2R γ^{null} .

limiting their expansion or survival in mice housed under specific-pathogen-free conditions. Mice were thus injected with 1 nmol 5-OP-RU intraperitoneally every 2 days from the day of huPBMC injection, a dose previously shown to activate endogenous MAIT cells.³⁶ This did not result in any increase in MAIT cells in peripheral blood or tissues (figure 4D).

It is not clear whether mouse cytokines can sustain homeostatic proliferation and survival of human MAIT cells in NSG mice due to a species barrier between human lymphoid cells and the murine recipient microenvironment.^{37–39} This is a key question for IL-15, as it is mostly mouse-derived in the xeno-GVHD model given the low human myeloid chimerism. Indeed, we found that human MAIT cells cultured with mouse IL-15 did not proliferate at all *in vitro* (figure 5A). Proliferation was increased when 5-OP-RU was added to mouse IL-15, although it remained lower than with human IL-15. Murine and human IL-7 had similar effects on MAIT cell proliferation, regardless of the presence of 5-OP-RU (figure 5A). Thus, IL-15 availability may be a factor limiting the proliferation of MAIT cells in NSG mice.

We, therefore, sought to determine the division rate of MAIT cells after transfer of CFSE-labeled huPBMCs into NSG mice. One week after transfer, substantial proportions of both T cells and MAIT cells from the liver and spleen displayed low CFSE fluorescence, indicating that they were able to divide (figure 5B). Previous reports have indicated that *in vivo* expansion of mouse-derived MAIT cells requires the combination of a TCR signal and a danger signal.^{40,41} This was confirmed in our model, as administration of the Toll-like-receptor 9 agonist CpG, alone or in combination with 5-OP-RU, caused a robust proliferation of human MAIT cells and their accumulation in the liver (figure 5B) and spleen (online supplemental figure 2). Notably, these proportions further increased when the mice were treated with human IL-15 from the day of transfer.

We, therefore, investigated whether human IL-15 would be necessary for MAIT cell accumulation in target tissues during xeno-GVHD. NSG mice were treated with human IL-15 three times per week from the day of huPBMC transfer and euthanized when they developed severe signs of aGVHD (25 days after transfer, ie, earlier than control mice). While a massive infiltration of T cells was observed in all compartments, MAIT cells remained barely detectable (figure 5C).

Altogether, these results indicate that human MAIT cells, although capable of responding to combined TCR and danger signals in NSG mice, do not expand or participate in immune-mediated tissue lesions during xeno-GVHD.

DISCUSSION

The newly described tissue repair and immunoregulatory functions of MAIT cells open fascinating perspectives for their use in adoptive therapy to control immune-mediated damage in tissues where these cells are known to accumulate. However, such strategy can be considered in unrelated recipients only if MAIT cells are devoid of GVH potential. As anticipated by the very limited diversity of their TCR

that recognizes microbial antigens presented by the highly conserved MR1 molecule, our results demonstrate that MAIT cells do not proliferate in response to allogeneic signals and do not participate to aGVHD lesions. These results support our previous observation that MAIT cells were undetectable in intestinal biopsies taken at time of aGVHD, suggesting that their defect in the circulation was not related to their recruitment to target tissues.¹²

Human MAIT cells are very few at birth and accumulate gradually during infancy, with about a 30-fold expansion to reach a plateau around 6 years of age.¹² Several pieces of evidence suggest that the drivers of this peripheral expansion are related to successive encounters with microbes leading to an accumulation of MAIT cell clonotypes that will constitute the future MAIT cell pool.⁴² Indeed, MAIT cells are absent in germ-free mice and are very few in laboratory mice, but dramatically expand following challenge with riboflavin-producing microbes.^{35,36,43} Moreover, their development in mice depends on early-life exposure to defined microbes that synthesize riboflavin-derived antigens.²⁷ In human subjects with controlled infection, MAIT cells show evidence of expansion of select MAIT cell clonotypes,⁴⁴ but longitudinal analyses of MAIT cell numbers in relation to the composition of the microbial environment are still lacking.

Here, we show that the reconstitution of MAIT cells after HSCT recapitulates their physiological expansion in the infancy and takes place over a period of at least 6 years, regardless of recipient-related or donor-related factors such as age, underlying disease, donor type, stem cell source or conditioning regimen. The number of MAIT cells in the early post-transplant period is at least in part dependent on the number of MAIT cells transferred with the graft. Subsequent MAIT cell recovery relies on thymus-derived naïve MAIT cells. The low absolute MAIT cell counts in patients with grades 3–4 aGVHD, which associate with low absolute counts of non-MAIT T cells, may be a consequence of a defective thymic function. Indeed, aGVHD-associated thymic damage results in the loss of the large population of double-positive (DP) thymocytes.⁴⁵ This may decrease the thymic output of naïve MAIT cells, which undergo positive selection by recognizing MR1 at the surface of these DP thymocytes.⁴⁶ In addition, loss of diversity and increased bacterial domination early after HSCT, in particular by *Enterococcus*, has been associated with increased risk of aGVHD.^{47–49} One might speculate that blooming of *Enterococcus* (a strain unable to synthesize riboflavin) could prevent from early expansion of donor-derived MAIT cells due to lack of MR1-ligands. So far, the relationship between abundance and diversity of gut microbiota and post-transplant MAIT cell reconstitution remains controversial.^{19,22} Investigating the gut metagenome or metatranscriptome of HSCT recipients for the presence of riboflavin biosynthesis genes should answer this question.

In mouse HSCT models, conditioning-resistant host residual MAIT cells promote gastrointestinal tract integrity and limit the proliferation of donor-derived alloreactive T-cells, thus protecting from aGVHD.²³ Although these

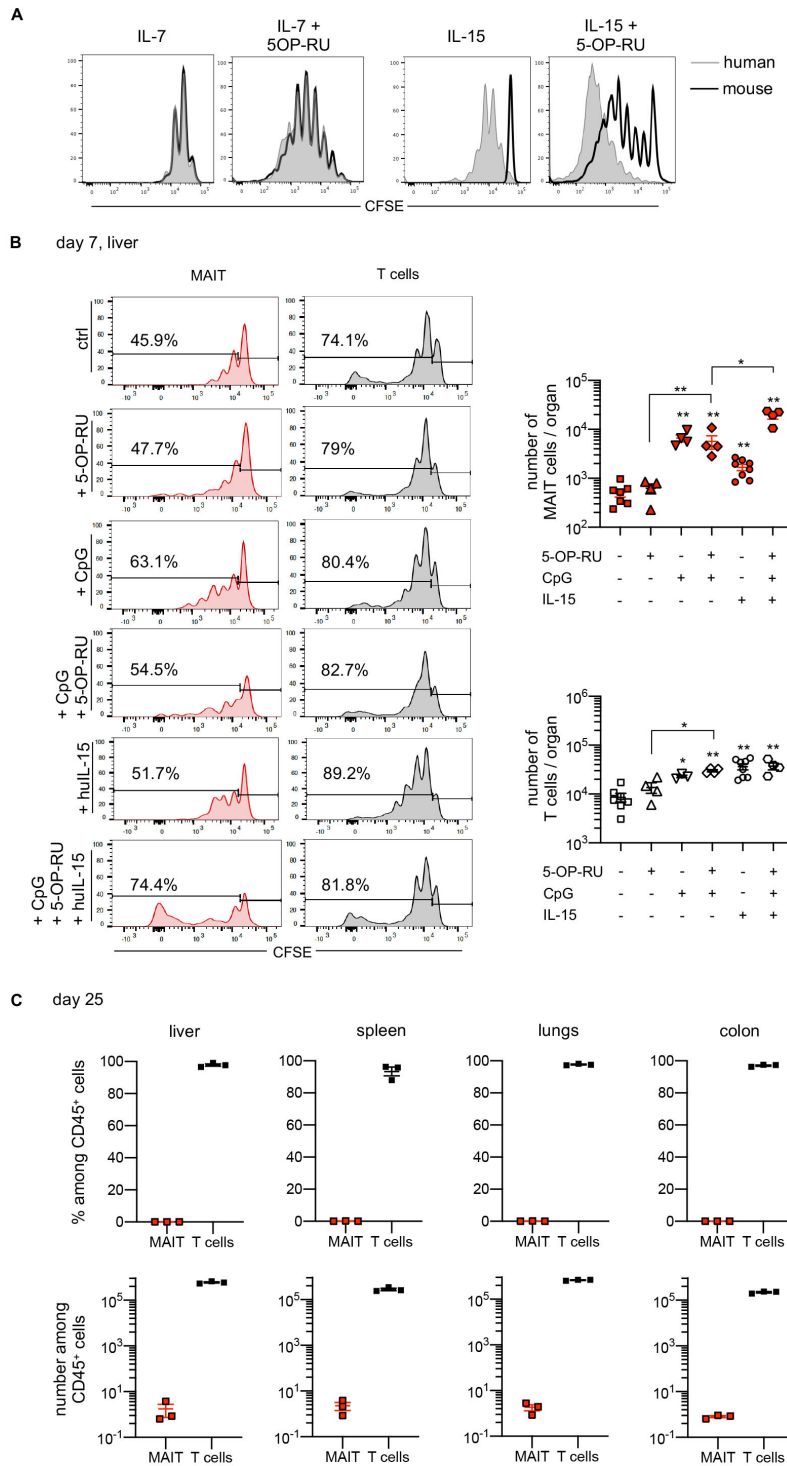


Figure 5 Human IL-15 does not increase MAIT cell accumulation in the target tissues in the xeno-GVHD model. (A) CFSE-labeled human PBMCs were cultured for 6 days with mouse (open black histograms) or human (shaded gray histograms) IL-15 (50 ng/mL) or IL-7 (10 ng/mL) in the absence or presence of 5-OP-RU (300 nM). Representative experiment showing CFSE^{low} proliferating MR1:5-OP-RU tetramer⁺ MAIT cells (gated on CD3 T cells). (B) NSG mice were injected with 20×10^6 CFSE-labeled human PBMCs without or with human IL-15 (100 ng) and/or 5-OP-RU (1 nmol), and/or CpG (10 μ g), i.p. every 2 days from the day of transfer. The liver was harvested 7 days later, and cells were analyzed by flow cytometry for CFSE expression (left panel) and numbers of MR1:5-OP-RU tetramer⁺ MAIT cells and non-MAIT T cells (right panel). Wilcoxon-Mann-Witney U tests were performed, significant differences are indicated compared with the control group (no 5-OP-RU/CpG/huIL-15), or between specified groups (* $p < 0.05$, ** $p < 0.01$). (C, D) Mice were treated with huIL-15 from the day of transfer of 5×10^6 huPBMCs and euthanized at the time of aGVHD (day 25, $n = 3$ diseased mice). Results show the proportion of human CD45⁺ leukocytes among viable cells (C), and absolute counts of T and MAIT cells (D) in the indicated compartments. Individual values and median (horizontal bar) are shown. aGVHD, acute graft-versus-host disease; CFSE, carboxyfluorescein succinimidyl ester; IL-15, interleukin 15; MAIT, mucosal-associated invariant T; huPBMCs, human peripheral blood mononuclear cells.

findings are not directly translatable to human aGVHD due to specificities of HSCT models in mice and differences between mouse and human MAIT cells,^{23 50 51} they support our observations that human MAIT cells do not contribute to aGVHD. Using a classical in vitro model of alloreactivity, we show that MAIT cells do not proliferate in response to allogeneic cells, unless combined MRI ligand and cytokine signals are present. Moreover, MAIT cells do not participate in the development of xeno-GVHD in NSG mice infused with huPBMCs. It is likely that the xeno-GVHD is caused by a fraction of T cells having a low frequency in donor PBMCs, which subsequently expand in mouse organs on recognition of murine MHC.⁵² One cannot exclude that the NSG host with an ablated immune compartment may be less likely to provide conditions for presentation of MRI-ligands to MAIT cells or the delivery of costimulatory signals. However, MRI is highly conserved across various species, with 90% of sequence similarity between mice and humans, so that murine MRI can present 5-OP-RU to human MAIT cells as efficiently as human MRI (35). That xenoreactive donor T cells may outcompete MAIT cells by limiting the availability of IL-15 seems unlikely, as demonstrated by the failure of exogenous human IL-15 to allow MAIT cell accumulation in tissues at time of xeno-GVHD. Whether MAIT cells fail to traffic or find their niche in the host due to species barriers between chemokine receptors and their ligands is also unlikely since MAIT cells were found in the spleen and liver 1 week after transfer. Moreover, administration of CpG at the time of transfer led to a strong expansion and tissue accumulation of MAIT cells, in line with other in vivo models with mouse MAIT cells.^{40 41} Altogether, our results indicate that, in contrast with T cells, MAIT cells do not find the conditions required for their activation and therefore, do not participate to tissue lesions during GVHD.

Due to ubiquitous expression of MRI and abundance of human MAIT cells in tissues, MAIT cell functions need to be tightly regulated to control the balance between healthy and pathological processes. Following HSCT, homeostatic cytokines may provide early but limited proliferation signals to graft-derived MAIT cells, at least ensuring their survival in the recipient in the early post-transplant period. However, sustained expansion of mature and/or thymus-derived naïve MAIT cells will only occur if MRI ligands are present together with inflammatory or danger signals. Our data showing the lack of alloreactive potential of MAIT cells pave the way for safely harnessing their functions in adoptive immune therapy. MAIT cells are excellent candidates for engineering universal chimeric antigen receptor cells (CAR-MAIT), as they are easy to expand in vitro, can traffic to tissues, have potent effector functions and are unlikely to cause GVHD. In addition, the novel tissue repair and regulatory functions of MAIT cells open up new avenues for exploiting these cells for clinical benefit.

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Contributors MT, NT-B, ML, GBY, MFC, AB, FM and SA designed research studies, conducted experiments, analyzed data, and/or provided patient samples. TY produced synthetic 5-OP-RU. OL and J-HD analyzed data and wrote the manuscript. SC-Z designed research studies, supervised the study, analyzed data, and wrote the manuscript with the help of other coauthors. All authors provided input, edited and approved the final version of the manuscript.

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