Mucosal-associated invariant T cells and disease

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Abstract | Mucosal-associated invariant T (MAIT) cells are unique innate-like T cells that bridge innate and adaptive immunity. They are activated by conserved bacterial ligands derived from vitamin B biosynthesis and have important roles in defence against bacterial and viral infections. However, they can also have various deleterious and protective functions in autoimmune, inflammatory and metabolic diseases. MAIT cell involvement in a large spectrum of pathological conditions makes them attractive targets for potential therapeutic approaches.

The immune system is traditionally divided into the adaptive and innate arms that together mount efficient immune responses against invading pathogens and preserve body integrity. Conventional T and B cells of the adaptive immune system induce specific responses and provide long-term memory against microorganisms, whereas the innate immune system allows immediate recognition of pathogens and moulds adaptive immune responses. Mucosal-associated invariant T (MAIT) cells are innate-like T cells and thus can be considered to span both the innate and adaptive arms. They were originally identified on the basis of expression of an invariant T cell receptor (TCR) α -chain (Va7.2–Ja33 in humans, and Va19–Ja33 in mice) associated with a limited TCR β -chain repertoire^{1–3}. In 2003, Lantz and colleagues showed that MAIT cells are restricted by the non-polymorphic MHC class I-related (MR1) molecule, which is highly conserved in mammals and expressed by most cell types³. Subsequent studies revealed the antibacterial recognition and function of MAIT cells^{4–7}. A critical advance in the biology of MAIT cells was the discovery by Rossjohn and McCluskey of the main bacterial ligands presented by the MR1 molecule to MAIT cells, namely agonist ligands derived from the microbial vitamin B₂ (riboflavin) biosynthesis pathway and modified by the bacterial and host cell metabolites glyoxal and methylglyoxal, as well as non-activating ligands derived from vitamin B₉ (folate)^{8–10} (BOX 1). The recent generation of a Va7.2-specific monoclonal antibody and MR1 tetramers has allowed MAIT cell characterization in both humans and mice. Although MAIT cells were first described as being enriched in mucosal tissues such as those of the intestinal tract and lungs, they are also abundant in the blood and are present in several peripheral tissues in physiological conditions^{11–19} (TABLE 1). The highest frequency of MAIT cells is found in the liver, whereas they are less frequent in lymphoid organs. MAIT cells are more abundant in humans (comprising

1–35% of total $\alpha\beta$ T cells) than in common laboratory mouse strains (comprising 0.01–5% of $\alpha\beta$ T cells)^{4,20}.

These innate-like T cells exhibit an effector-memory phenotype and are usually characterized in humans by high expression of the C-type lectin CD161. They produce a wide range of cytokines and molecules that induce cytotoxic responses, such as granzyme B and perforin, upon activation through their TCRs and/or cytokine receptors, and they express integrins and chemokine receptors, allowing their migration to inflamed tissues^{21–23}. Due to their localization and function, MAIT cells can act as a first line of defence against infections, but they may also contribute to autoimmune and immune-mediated diseases. Unlike existing Reviews^{23–32}, we highlight the pathophysiological roles of MAIT cells, emphasizing the common and distinct features of MAIT cells in various pathologies. We address the behaviour of MAIT cells in infectious and non-infectious pathologies and describe their homeostasis, activation, tissue migration, apoptosis and different effector functions through which they can exert protective or deleterious roles.

MAIT cells in infectious diseases

MR1-dependent MAIT cell activation in bacterial infection. The first functional studies of MAIT cells showed that they are activated by certain bacteria and yeast in an MR1-dependent manner^{7,33,34}. Subsequent analyses characterized MAIT cell agonist ligands as metabolites derived from the bacterial vitamin B₂ synthesis pathway^{8–10} that can be produced by both commensal and pathogenic strains^{3,35}. Vitamin B₂ biosynthesis pathways that are unique to yeast and bacteria allow MAIT cells to sense infection by recognition of conserved microbial metabolites. In patients with *Mycobacterium tuberculosis* infection, MAIT cells are decreased in the blood but seem to accumulate in the lungs, suggesting that they are recruited to the infected tissues and can

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Box 1 | MAIT cell ligands

The main ligands for MHC class I-related (MR1) molecules are small organic metabolites derived from two vitamin B sources: metabolites that can activate mucosal-associated invariant T (MAIT) cells are derived from riboflavin (vitamin B₂), and non-stimulatory metabolites are derived from folic acid (vitamin B₉)^{9–10}. Riboflavin is synthesized by many bacteria and yeasts, but not by mammals. The first described ligands — 7-hydroxy-6-methyl-8-D-ribityllumazine and 6,7-dimethyl-8-D-ribityllumazine — can stimulate MAIT cells when presented by MR1, due to the presence of a ribityl tail in their structure that interacts with the MAIT cell T cell receptor. Two single-ring pyrimidine compounds, 5-(2-oxopropylideneamino)-6-D-ribitylaminouracil and 5-(2-oxoethylideneamino)-6-D-ribitylaminouracil, which are formed from the spontaneous reaction of the riboflavin precursor 5-amino-6-D-ribitylaminouracil with methylglyoxal or glyoxal, respectively, and are by-products of both bacteria and host cell glycolysis, were recently described as the most potent MAIT cell-activating ligands. 6-Formylpterin and acetyl-6-formylpterin, the resulting products of folic acid photodegradation, can both bind to MR1, increasing its stability and its surface expression. These two formylpterins inhibit MAIT cell activation by competing with the activating ligands.

Due to its high structural plasticity, the MR1 pocket can present a large variety of ligands, such as drugs and drug-related molecules with diverse chemical structures¹¹⁰. As an example, MR1 can present chemical agonists such as diclofenac or 5-hydroxy diclofenac, as well as chemical inhibitors such as 3-formylsalicylic acid, 5-formylsalicylic acid and 2,4-diamino-6-formylpteridine.

Endogenous mammalian ligands have not yet been identified; however, their existence is likely. MAIT cells are selected in the thymus in an MR1-dependent manner, and this thymic education is maintained in germ-free mice^{3,35}. Several in vitro studies also support the presence of endogenous MR1 ligands leading to MAIT cell interaction with hepatic myofibroblasts or tumour cells in an MR1-dependent way in the absence of exogenous ligands^{69,148}.

contribute to immune defence^{5,7,36}. Functional in vitro studies of MAIT cells from healthy individuals show that they produce interferon- γ (IFN γ) and tumour necrosis factor (TNF) upon MR1-dependent activation by lung epithelial cells infected with *M. tuberculosis*^{5,7}, thereby inducing their death. Interestingly, IFN γ production by MAIT cells could be one mechanism by which they control *M. tuberculosis* infection, as IFN γ is known to induce antimicrobial mechanisms such as autophagy³⁷ and the activation of nitric oxide synthase 2 (NOS2)^{38,39}.

The activation of MAIT cells and their ability to kill infected cells have been confirmed in infection of epithelial cells by various bacteria, such as *Escherichia coli* and *Shigella flexneri*³³. Migration of blood MAIT cells to infected tissue is also suggested in patients who developed typhoid fever after being challenged with oral *Salmonella enterica* subsp. *enterica* serovar Typhimurium. These patients have decreased blood MAIT cell frequency compared with those who are resistant to the infection, and their remaining MAIT cells are activated and express high levels of gut-homing chemokine receptors (CCR9 and CCR6), which could allow their migration to the infected intestinal tract⁴⁰. However, the decreased blood MAIT cell frequency could also reflect their activation-induced cell death. In addition to migration, proliferation can also lead to MAIT cell expansion in infected tissue. Indeed, non-human primates vaccinated intradermally with bacillus Calmette-Guérin and challenged with pulmonary *M. tuberculosis* exhibit a higher frequency of proliferating (Ki-67⁺) MAIT cells at the site of vaccination⁴¹. A protective role for MAIT cells against bacterial infections in humans is suggested by the better outcome

of patients with bacterial sepsis who exhibit higher frequencies of circulating MAIT cells⁴².

Although studies of the frequency, phenotype and activation status of MAIT cells in patients bring some insight into their function in bacterial infections, most studies are limited to analysis of the blood. The most convincing data supporting the protective role of MAIT cells in bacterial infections have been obtained in mouse models. MAIT cells in mice are exclusively identified by mouse MR1 tetramers, owing to a lack of specific antibodies recognizing the invariant TCR Va19–Ja33 chain. Despite the low frequency of MAIT cells in mice compared with humans (TABLE 1), mouse studies showed that MAIT cells have protective roles against infection by *Mycobacterium abscessus*, *Mycobacterium bovis*, *E. coli*, *Klebsiella pneumoniae*, *Francisella tularensis* and *Legionella longbeachae*^{6,7,43–47}. For example, intraperitoneal injection of *E. coli* or *M. abscessus* in *Mr1*^{−/−} mice, which are thereby devoid of MAIT cells, results in a higher bacterial load in the spleen than in MAIT cell-containing control mice. Intranasal infections of wild-type mice with *F. tularensis*, *M. bovis*, *S. Typhimurium* or *L. longbeachae* induce an accumulation of MAIT cells in the lungs, where they produce large amounts of cytokines such as IL-17A, IFN γ and TNF, depending on the type of infection^{43,45,47,48}. MAIT cell accumulation in the lungs after *S. Typhimurium* infection is MR1 dependent and requires the presence of ligands derived from the microbial riboflavin synthesis pathway⁴⁸.

MR1-independent activation of MAIT cells. TCR-independent activation of MAIT cells has also emerged as an important mechanism against both bacterial and viral infection (FIG. 1). MAIT cells express high levels of IL-18 receptor (IL-18R) and IL-12R, and IL-18 and IL-12 have emerged as the major cytokines that activate MAIT cells upon bacterial or viral infection^{43,45,49–68} (TABLE 2). During bacterial infection, blockade or deletion of the IL-12p40 subunit impairs MAIT cell control of intracellular growth of *F. tularensis* and *M. bovis*^{43,45}. The effect of IL-18 on MAIT cells in bacterial infection is more uncertain. In vitro studies have shown that *F. tularensis* and *Enterococcus faecalis*, whether or not they produce riboflavin, are able to stimulate the production of IFN γ by MAIT cells in an IL-12- and/or IL-18-dependent manner^{49,50}. Furthermore, peripheral blood mononuclear cells or the THP1 monocytic cell line exposed to *E. coli* (riboflavin-synthesizing) or *E. faecalis* (non-riboflavin-synthesizing) produce IL-12 and IL-18 via a Toll-like receptor 8 (TLR8) pathway, and both cytokines are required for MAIT cell activation⁵⁹. However, a recent study showed that although IL-18 is required for MAIT cell activation by *F. tularensis* in vitro, this cytokine is not essential for MAIT cell IFN γ production in in vivo infection⁴⁹. Bacterial infection in vivo might induce MAIT cell activation through a more complex immune cell crosstalk, involving TLRs expressed by antigen-presenting cells (APCs)^{48,59}. Interestingly, cytokine-dependent MAIT cell activation by tissue macrophages or dendritic cells has a key role in the recruitment and differentiation of monocytes in the lungs of infected mice and thereby promotes local CD4⁺

Table 1 | Features of MAIT cells in humans and mice

Feature	Human MAIT cells	Refs	Mouse MAIT cells	Refs
T cell receptor expression	Va7.2–Ja33 Vβ2, Vβ13	2,20 2,20	Va19–Ja33 Vβ8.1, Vβ8.2, Vβ6	2,115 2,115
Percentage of total T cells	1–35%, depending on the tissue (blood 5%, oral mucosa 2%, lung 3%, spleen 1%, liver 35%, adipose tissue 2%, ileum 2%, colon 2–7%, female genital tract 2–3%)	11–19	0.01–5%, depending on the tissue and mouse strain (excluding transgenic mice) (blood 0.5%, lung 3%, spleen 0.08%, pancreatic lymph nodes 0.4%, liver 5%, adipose tissue 2%, ileum 1%, colon 1%)	70,115
Transcription factor expression	PLZF, RORγt, T-bet	4,35	PLZF, RORγt, T-bet	35,48,115
Activation and/or maturation marker expression	CD69, CD25, CD161, CD95, CD62L ^{low} , CD45RO, CD27	20,113	CD69, CD25, CD38, CD62L, CD44	69,70,115
Cytokine receptor expression	IL-7R, IL-12R, IL-15R, IL-18R, IFNAR	28,50	IL-7R, IL-12R, IL-15R, IL-18R	115
Chemokine receptor expression	CCR2, CCR5, CCR6, CCR9, CCR10, CXCR3, CX ₃ CR1, CXCR6	70,152	CCR2, CCR5, CCR6, CCR9, CXCR3, CXCR6	115
Cytokines produced	IFNγ, IFNα, TNF, IL-17A, IL-2, IL-4, IL-10, IL-22	5,16,50,73	IFNγ, TNF, IL-17A, IL-2, IL-4, IL-10, IL-13, IL-22, GM-CSF	70,115
Cytotoxic mediator expression	Granzyme A, granzyme B, granzyme K, perforin	34,70	Granzyme B, perforin	70
Natural killer cell and adhesion marker expression	PD-1, NKG2D, NKP80, NKP30, αL integrin, CD56, CD43, CD49a, CD103, CLA, α4β1 integrin, PSGL1, sLex	18,70,91,98,113,153–156	NK1.1	115

CLA, cutaneous lymphocyte-associated antigen; GM-CSF, granulocyte–macrophage colony-stimulating factor; IFN, interferon; MAIT, mucosal-associated invariant T; PD-1, programmed cell death 1; PSGL1, P selectin glycoprotein ligand 1; sLex, sialyl–Lewis X motif; TNF, tumour necrosis factor. ^aA.T., B. Kiaz and A.L., unpublished data.

T cell recruitment and activation⁴⁶. These studies highlight the ability of MAIT cells to bridge the innate and adaptive arms of an immune response.

During viral infection, MAIT cells are also activated by cytokines produced by dendritic cells or macrophages. Of note, unlike bacteria, viruses are unable to generate riboflavin-derived ligands to activate MAIT cells. In vitro, myeloid cells exposed to hepatitis C virus (HCV), dengue virus and influenza virus activate MAIT cells to produce IFNγ, TNF and granzyme B in an IL-18-dependent manner, which might act in synergy with IL-12, IL-15 and type I interferons^{62,66}. Through the production of IFNγ, TNF and granzyme B, MAIT cells inhibit viral replication and may exert cytotoxic function against virus-infected cells. In patients with acute viral infection by dengue or influenza virus, MAIT cells rapidly acquire an activated phenotype characterized by the expression of activation and exhaustion markers such as CD38, CD69, PD-1 and TIM3 (REFS^{62,63,66}). Moreover, there is a positive association between the frequency of activated blood MAIT cells and the severity of dengue disease⁶⁶, which could be due to higher viral load and/or inflammatory cytokine levels. Although patients with dengue infection have frequencies of circulating MAIT cells similar to those in healthy controls, circulating MAIT cells are reduced in patients with influenza virus infection. Accumulation of MAIT cells in the lungs of mice infected with influenza virus suggests their recruitment from the blood to infected sites, and this was associated with their increased expression of activation markers such as CD69 and CD25 and increased production of granzyme B^{62,66}. Analysis of MR1-deficient mice and transfer of MAIT cells demonstrated a protective role for MAIT cells in this infection⁶⁷.

In chronic infections by hepatitis B virus (HBV), HCV and HIV, most studies have described a decrease in MAIT cell numbers both in the blood and in tissues such as the liver. Despite their reduction in frequency and defective in vitro response to *E. coli*, the ability of MAIT cells to respond to IL-12 or IL-18 is generally preserved^{51–58,60,66,68}. Indeed, as has been observed in dengue and influenza virus infections, the remaining MAIT cells express higher levels of activation and/or exhaustion markers, such as CD38, HLA-DR, CD69 and PD-1, and produce IFNγ and granzyme B on activation.

However, some human studies have described a functional impairment of MAIT cells in chronic HBV⁶⁸ and HIV^{60,61} infection, with decreased IFNγ, TNF and IL-17A production and reduced cytotoxicity, after in vitro TCR-dependent activation, correlating with very low levels of expression of the transcription factors T-bet, EOMES, PLZF, RORγt and HELIOS⁶¹. In a small cohort of HIV-infected patients, treatment with recombinant human IL-7 restored the frequency of circulating MAIT cells, as was previously shown for conventional CD4⁺ and CD8⁺ T cells⁶⁴, and functional analysis of one treated patient showed that MAIT cell responsiveness to IL-12 and/or IL-18 or *E. coli* stimulation was also restored⁶⁵.

Overall, studies in vitro, in mouse models and in patients demonstrate the ability of MAIT cells to sense bacterial and viral infection. The protective role of MAIT cells in several infections, such as tuberculosis and influenza virus, suggests that therapeutic approaches based on the activation of MAIT cells by cytokines and/or agonist ligands could be developed. By contrast, the potential deleterious role of MAIT cells in other infections, such as dengue, warrants further investigation.

Fig. 1 | MAIT cell activation in bacterial and viral infection. Upon bacterial and viral infections, mucosal-associated invariant T (MAIT) cells can be activated in a MHC class I-related molecule (MR1)-dependent manner through their T cell receptor (TCR) recognizing MR1-presenting bacterial ligands (part **a**), and/or in an MR1-independent manner by inflammatory cytokines such as IL-12 and IL-18, produced by infected cells and sensed by MAIT cells through cytokine receptors (part **b**). MAIT cell activation induces their release of granzyme B and perforin (which kill infected cells), their proliferation and their production of cytokines (tumour necrosis factor (TNF), interferon- γ (IFN γ) and IL-17A) that inhibit viral replication and bacterial growth. Activated MAIT cells can also promote monocyte recruitment and activation. APC, antigen-presenting cell; CCL3, CC-chemokine ligand 3; GM-CSF, granulocyte-macrophage colony-stimulating factor; IL-12R, IL-12 receptor.

MAIT cells in immune-mediated diseases

MAIT cell characteristics (TABLE 1) — including activation by inflammatory cytokines, expression of chemokine receptors and integrins, and the production of large amounts of cytokines and cytotoxic molecules — suggest that MAIT cells, similarly to conventional effector memory T cells, could be involved in autoimmune and immune-mediated chronic diseases. However, in contrast to conventional T cells, circulating MAIT cells can function without specific antigen priming and following antigen-nonspecific activation by cytokines produced in an inflammatory context. In addition, inflammation increases the surface expression of MR1 protein^{69–71}, suggesting that MAIT cells could be activated through their TCRs following recognition of MR1 molecules presenting bacterial, or possibly endogenous, ligands (BOX 1). After activation, MAIT cells could act directly on target cells or could participate to the recruitment and activation of other immune cells, as in infectious diseases. Numerous reports have described MAIT cell alterations in patients with various autoimmune and immune-mediated pathologies. Despite the heterogeneity of the disease aetiologies, there are several common features, such as reduced MAIT cell frequency in

the blood, their presence in the target tissue and their increased activation and dysfunction.

Decreased circulating MAIT cell frequency. As reported in chronic infectious diseases, the number of circulating MAIT cells decreases in most autoimmune diseases and in many inflammatory and metabolic diseases^{16,18,24,69,70,72–86} (FIG. 2; TABLE 3). Blood MAIT cell frequency correlates with disease severity in several pathologies^{87–94} (TABLE 3). In multiple sclerosis (MS), the frequency of blood MAIT cells decreases during clinical relapse and progression of the disease, whereas it increases during remission^{71,95}. Similarly, after bariatric surgery, obese patients improve in their metabolic and inflammatory parameters, simultaneously increasing in their number of blood MAIT cells¹⁶. Only a few studies have characterized circulating MAIT cell subsets according to their CD4 or CD8 expression in autoimmune and inflammatory diseases. In humans, MAIT cells are mainly CD8 positive, approximately 10% are CD4 and CD8 double negative, and very few are CD4 positive. When MAIT cell subsets are characterized, the frequencies of both CD8⁺ and CD4⁺CD8[−] circulating MAIT cells are similarly decreased^{18,70,80,87,94}. The very low frequency

Table 2 | MAIT cells in infectious diseases

Infectious disease	MAIT cell frequency	MAIT cell phenotype	MAIT cell function	Refs
Bacterial infection				
Legionella	↑ in lungs	ND	↑ IFN γ and GM-CSF	47
Tuberculosis	↓ in blood; ↑ in lungs	↑ PD-1 in blood	↑ IL-17A and granzyme B	5,7,36,157
Shigellosis	↓ in blood	↑ Activation markers?	ND	33
Klebsiella	ND	ND	↓ Bacterial load in vivo	6
Viral infection				
Influenza	↓ in blood	↑ CD25 and CD69	↑ Granzyme B	62,66,67
Dengue	↓ in blood	↑ CD38 in blood	↑ Granzyme B	66
Hepatitis B virus	Marginal ↓ in blood	↑ CD38 and CD57 in blood	↑ Granzyme B and perforin; ↑ IFN γ (after <i>Escherichia coli</i> and anti-CD28 stimulation)	53,68
Hepatitis C virus	↓ in blood; ↓ in liver	↓ CCR5; ↑ CD38, PD-1, HLA-DR, TIM3, CTLA4 and CD69 in blood	↑ Granzyme B	51,52,54,56,58
HIV	↓ in blood and lymph nodes	↑ PD-1, HLA-DR, TIM3, CD38, CD57, IL-7R, CCR6, T-bet in blood	↓ Granzyme B, IFN γ and TNF	55,57,61,64,65
HTLV-1	↓ in blood	↑ CD38 and HLA-DR and ↓ PLZF in blood	↓ IFN γ (after <i>E. coli</i> stimulation)	63

CCR, CC-chemokine receptor; CTLA4, cytotoxic T lymphocyte antigen 4; GM-CSF, granulocyte–macrophage colony-stimulating factor; HTLV-1, human T cell lymphotropic virus type 1; IFN γ , interferon- γ ; MAIT, mucosal-associated invariant T; ND, not determined; PD-1, programmed cell death 1; TIM3, T cell immunoglobulin and mucin domain-containing 3; TNF, tumour necrosis factor.

of CD4⁺ MAIT cells renders their specific detection challenging, due to potential contamination by conventional T cells during flow cytometry analysis. Several mechanisms could explain the decreased frequency of blood MAIT cells observed in most, if not all, pathologies.

Presence of MAIT cells in inflamed tissues. Compared with tissues from healthy individuals or with unaffected tissues or blood from the same patients, the frequency of MAIT cells increases in the target tissues of patients with various immune-mediated diseases. TABLE 3 and FIG. 2 illustrate the diversity of MAIT cell localization^{16,18,73,77–79,82,86,89,96–100}. Mouse models allow the analysis of MAIT cells in tissues that are hard to access in humans. In non-obese diabetic (NOD) mice, the mouse model of type 1 diabetes (T1D), MAIT cell frequency continuously increases in the pancreas until disease onset. Experiments involving transfer of MAIT cells into NOD recipient mice at different stages of disease showed an increase in the recruitment of MAIT cells to the pancreas of diabetic as compared with prediabetic mice⁷⁰.

It should be noted that MAIT cell accumulation in affected tissues has not been found in all immune-mediated diseases; some diseases may instead show a redistribution of MAIT cells. For example, in the liver of patients with cirrhosis, immunohistochemical analysis shows preferential localization of MAIT cells in the inflammatory infiltrate around the portal tract and in the fibrotic septae^{69,87}, and in patients with Crohn's disease (CD), MAIT cells distribute specifically towards the injured part of ileum¹⁸. Fate-tracking studies should help to clarify whether and how MAIT cell recruitment

to inflamed tissues contributes to their local expansion, promoting their pathogenic functions.

Several studies have suggested a key role for adhesion molecules, integrins and chemokines in the migration of MAIT cells to inflamed tissues. For example, blood MAIT cells from patients with MS express higher levels of the pro-migratory molecules P-selectin glycoprotein ligand 1 and α L integrin (also known as CD11a) than do cells from healthy individuals⁷¹. Moreover, IL-18, which is elevated in the serum from patients with MS¹⁰¹, significantly upregulates the expression of α 4 β 1 integrin (also known as VLA4) by CD8⁺ MAIT cells, compared with conventional memory CD8⁺ T cells⁹⁹, and α 4 β 1 integrin mediates CD8⁺ T cell migration across the blood–brain barrier in MS¹⁰². In rheumatoid arthritis, high levels of TNF and IL-1 β in the synovial fluid can upregulate the expression of adhesion molecules, such as E-selectin, by human endothelial cells⁹¹. E-selectin promotes MAIT cell binding to endothelial cells via the sialyl–Lewis X (sLex) motif, which is elevated on synovial fluid MAIT cells compared with blood MAIT cells. In patients with rheumatoid arthritis, fewer circulating MAIT cells express the sLex motif than do circulating MAIT cells from healthy controls⁹¹, suggesting that sLex⁺ MAIT cells have migrated to the synovial fluid. TNF also upregulates the expression of CC-chemokine ligand 20 (CCL20), which might attract MAIT cells expressing CCR6 in synovial fluid⁹¹. Similarly, in children with recent-onset T1D, the frequency of blood MAIT cells is positively correlated with their expression of CCR6 (REF.⁷⁰), suggesting the involvement of CCR6 in the migration of MAIT cells to inflamed tissues. CCR6⁺ cells recognize CCL20 and β -defensins, which

Fig. 2 | MAIT cell alterations and function in immune-mediated diseases. In most autoimmune and immune-mediated diseases, mucosal-associated invariant T (MAIT) cell numbers in the blood are decreased. This may reflect two possible mechanisms: migration of MAIT cells to inflamed tissues, or increased MAIT cell death. MAIT cells express numerous chemokine receptors (such as CCR5, CCR6 and CXCR6), integrins (such as $\alpha 4\beta 1$ integrin, CD49a and CD103) and other adhesion molecules (such as P-selectin glycoprotein ligand 1 (PSGL1), αL integrin and sialyl-Lewis X motif (sLex)). In some disease settings, MAIT cells are chronically activated (showing expression of CD25, CD69 or CD40L) by MHC class I-related molecule (MR1) and/or cytokines (IL-12, IL-18, IL-6 and interferon- α (IFN α)), and this can lead to a dysfunctional or exhaustion state (characterized by the expression of PD-1 and impaired IFN γ and tumour necrosis factor (TNF) secretion) and apoptosis (marked by increased levels of FAS and intracellular active caspase 3, as well as decreased levels of BCL-2). Disease, age, body mass index, hyperglycaemia or treatment could also impact MAIT cell frequency and functions. MAIT cells have been found to have both protective and deleterious roles in various immune-mediated diseases, through multiple mechanisms involving cytokine production, direct cell contact and cellular crosstalk. MAIT cells may act directly on target cells — for example, granzyme B-mediated lysis of islet β -cells in type 1 diabetes (T1D) and MR1-dependent or cytokine-dependent activation of profibrogenic functions of liver myofibroblasts in cirrhosis and autoimmune liver disease (AILD). MAIT cells may also act through immunoregulatory pathways involving more complex cellular crosstalk such as interactions with B and T cells and with myeloid cells. Bold text is used in the middle part of the figure for tissues to which MAIT cells have been found to migrate or for which migration is strongly supported. AS, ankylosing spondylitis; B7RP1, B7-related protein 1; CLA, cutaneous lymphocyte-associated antigen; ICOS, inducible T cell co-stimulator; IL-12R, IL-12 receptor; MS, multiple sclerosis; PD-1, programmed cell death 1; T2D, type 2 diabetes; TCR, T cell receptor.

Table 3 | MAIT cells in immune-mediated diseases

Disease	MAIT cell frequency		MAIT cell phenotype		MAIT cell effect on disease	Possible mechanisms of action	Refs
	Blood	Tissues	Blood	Tissues			
Type 1 diabetes, non-obese diabetic mice	↓	↑↓↔ ^a	Activated, exhausted, apoptotic	Activated, exhausted?	Protective Deleterious	Gut mucosa integrity maintained via IL-17 and IL-22 production by MAIT cells • Islet β-cell cytotoxic death via granzyme B • Granzyme B synthesis by MAIT cells in pancreas of non-obese diabetic mice	70,124,131
Type 2 diabetes	↓	↑↓↔ ^a	Activated, exhausted	ND	Deleterious?	IL-17 and granzyme B production by MAIT cells in blood and adipose tissue	16,73
Obesity	↓ ^c	↑↓↔ ^a	Activated, exhausted	Activated, exhausted, apoptotic	Deleterious? Protective?	• IL-17 production by MAIT cells in blood and adipose tissue, insulin resistance • Granzyme B secretion in adipose tissue IL-10 production by MAIT cells in adipose tissue of lean individuals	16,73
Cardiovascular diseases	↓ ^c	ND	Apoptotic	ND	Deleterious?	Pro-inflammatory effects of MAIT cells	94
Liver cirrhosis, autoimmune liver disease, carbon tetrachloride-exposed mice, mice undergoing bile duct ligation	↓ ^c	↑↓ ^a	Activated, exhausted, apoptotic	Activated, exhausted?	Deleterious	• M1 macrophage polarization by MAIT cells • MAIT cell interaction with myofibroblasts, profibrogenic and pro-inflammatory effects • Increased circulating and hepatic IL-17 ⁺ MAIT cells	69,85,87
Alcohol-related liver disease	↓	↔ ^a	Activated, exhausted	ND	Protective	Defective antibacterial functions of circulating MAIT cells	83
Type 1 autoimmune hepatitis	↓	↑ ^a	Activated ^c , exhausted	ND	Deleterious?	Increased granzyme B ⁺ MAIT cell frequency in blood correlates with fibrosis score at diagnosis	82
Nonalcoholic fatty liver disease, mice on methionine–choline-deficient diet	↓ ^c	↑ ^a	Activated, exhausted	Activated, exhausted	Protective	Increased IL-4 and IL-10 production, decreased IFNγ production by MAIT cells and M2 macrophage polarization	93
Ulcerative colitis	↓	↑↓ ^a	Activated ^c , apoptotic	Apoptotic	Protective? Deleterious?	IL-17 and IL-22 production by MAIT cells in blood IL-17 production by MAIT cells in blood	18,79,81
Crohn's disease, 2,4,6-trinitrobenzene sulfonic acid-induced colitis mouse model	↓	↑↓ ^a	Activated, exhausted, apoptotic	ND	Protective	IL-17 production by MAIT cells in blood	18,81,125
Coeliac disease	↓	↓↔ ^a	ND	ND	ND	ND	74
X-linked lymphoproliferative syndrome type 2	↓	ND	Apoptotic	ND	ND	ND	76
Sjögren syndrome	↓	↑ ^a	Activated, exhausted	Activated	Deleterious?	IL-17 production by MAIT cells in salivary glands	78,86
Ankylosing spondylitis	↓	↑ ^b	Activated ^c , exhausted	Activated, exhausted	Deleterious?	• IL-17 and granzyme B production by MAIT cells in synovial fluid • IL-17 and IL-22 production by circulating MAIT cells	77,80,84
Rheumatoid arthritis, collagen-induced arthritis mouse model	↓ ^c	↑ ^b	Activated	Activated	Deleterious	• Reduced disease severity in <i>Mr1</i> ^{-/-} mice, enhanced disease severity in <i>Vα19</i> transgenic mice • Reduced IL-17 production in liver of <i>Mr1</i> ^{-/-} mice	89,91,117
Systemic lupus erythematosus	↓ ^c	ND	Activated ^c , exhausted, apoptotic	ND	ND	Correlation between disease activity and frequency of CD69 ⁺ circulating MAIT cells	88,89

Table 3 (cont.) | MAIT cells in immune-mediated diseases

Disease	MAIT cell frequency		MAIT cell phenotype		MAIT cell effect on disease	Possible mechanisms of action	Refs
	Blood	Tissues	Blood	Tissues			
Antineutrophil cytoplasmic antibody-associated vasculitis	↓	ND	Activated	ND	ND	ND	72,75
Multiple sclerosis, experimental autoimmune encephalomyelitis mouse model	↑↓ ^c	↑ ^a ↓ ^b	Activated	Activated	Protective, immunoregulatory	<ul style="list-style-type: none"> Increased IL-10 production by B cells and decreased pro-inflammatory molecules, due to Vα19 cell interaction with B cells via ICOS IFNγ production by T cells suppressed in a cell-contact-dependent manner via MAIT cell interaction with T cells 	71,95, 96,99,111, 118,126,158
					Deleterious?	<ul style="list-style-type: none"> Switch of MAIT cells toward T_H17-type profile Increased MR1- and cytokine-dependent activation of MAIT cells in the CNS 	
Asthma	↓ ^c	↓ ^a	Activated ^c	ND	Deleterious?	Increased circulating IL-17 ⁺ MAIT cells associated with disease severity	90,92,129,159
					Protective?	<ul style="list-style-type: none"> Lower circulating MAIT cell frequency in 1-year-old children who develop asthma at age 7 Interaction between MAIT cells and CD4⁺ T cells that induce IFNγ 	
Psoriasis	ND	↔ ^a	ND	Activated	Deleterious?	IL-17 and IL-22 production by MAIT cells in skin	98
Dermatitis herpetiformis	ND	↑ ^a	ND	ND	ND	ND	97

Exhausted? denotes a cell status based only on phenotype and not confirmed by functional analysis. Exhausted denotes cells with impaired cytokine production and function. Deleterious? denotes a deleterious effect is suggested or hypothetical and not yet demonstrated. Protective? denotes a protective effect is suggested or hypothetical and not yet demonstrated. ↑, increased frequency; ↓, decreased frequency; ↑↓, both increased and decreased frequency reported; ↔, no change; CNS, central nervous system; ICOS, inducible T cell co-stimulator; IFNγ, interferon-γ; MAIT, mucosal-associated invariant T; MR1, MHC class I-related molecule; ND, not determined; T_H17, T helper 17. ^aCompared with tissues from healthy individuals or with unaffected tissues. ^bCompared with blood. ^cCorrelation with disease severity. ^dComparison between active and inactive forms of the disease.

are chemoattractants that are increased in the pancreas and gut of diabetic patients and mouse models^{103–105}. In addition to CCR6, other skin-homing receptors, such as CCR10, CD49a, CD103 and cutaneous lymphocyte-associated antigen, are expressed by some blood MAIT cells^{19,97,98} and could participate in their migration to dermatitis herpetiformis lesions, in which MAIT cells are abundant⁹⁷. Increased expression of the chemokine receptors CXCR3 and CX₃CR1 on hepatic MAIT cells in non-infectious chronic liver diseases might reflect their activation rather than their recruitment, as MAIT cells are already abundant in the liver at the steady state¹⁵. Of note, even though MAIT cells express various chemokine receptors (TABLE 1), until now, no specific subset of MAIT cells has been identified as preferentially trafficking to distinct tissues.

Blood MAIT cells are activated and exhausted. The decreased frequency of blood MAIT cells in immune-mediated diseases could also result from their death by apoptosis subsequent to sustained activation. Accordingly, in many autoimmune and inflammatory diseases, circulating MAIT cells express higher levels of activation and/or exhaustion markers, such as CD38, CD25, CD69 and PD-1 (REFS^{16,18,69,70,73,83,85,87,89,93}). Furthermore, in some diseases MAIT cell activation correlates with disease activity score (TABLE 3). As has been observed in infectious diseases, MAIT cells could

be activated via MR1-dependent or MR1-independent pathways. In systemic lupus erythematosus (SLE), the increased activation of MAIT cells has partly been explained by the increased ability of monocytes to activate MAIT cells in a TCR-dependent manner⁸⁸. The activation of MAIT cells by MR1 in autoimmune and immune-mediated diseases is also suggested by the observation of increased MR1 expression in inflamed tissues⁷¹. In the context of T1D, in vitro exposure of a human pancreatic β-cell line to pro-inflammatory cytokines increases both *MR1* transcript levels and MR1 surface expression⁷⁰. Interestingly, gut homeostasis is altered in many autoimmune and immune-mediated diseases (BOX 2), which could lead to increased levels of bacteria-derived ligands for MR1. The observations of thymic selection of MAIT cells in germ-free mice¹⁰⁶ and TCR-dependent activation of myofibroblasts by MAIT cells in the absence of exogenous ligands⁶⁹ support the existence of endogenous ligands (BOX 1). Moreover, correlations between circulating MAIT cell frequency, CD69⁺ MAIT cell frequency and cytokine (IL-18, IL-6 and IFNα) plasma levels in patients with SLE⁸⁸, MS⁹⁹ and ulcerative colitis (UC)⁷⁹ suggest that MAIT cells can also be activated by cytokines in these diseases. The exact mechanism of MAIT cell activation is not yet well defined. As in infectious diseases, the combined triggers for MAIT cell activation of inflammatory cytokines and TCR-mediated signals might lead to

Box 2 | MAIT cells and gut alteration in immune-mediated diseases

Studying mucosal-associated invariant T (MAIT) cells in autoimmune, inflammatory and metabolic diseases can shed new light on the pathophysiology of these cells. Despite their different features, many of the pathologies listed are associated with alterations in gut homeostasis^{149–151}, including microbiota composition, intestinal permeability and gut immunity. Given the key characteristics of MAIT cells (activation by bacterial ligands, homing capacity and gut localization), they might represent sensors and/or enactors of such gut modifications. But what is the first step during autoimmune, inflammatory and metabolic diseases? Are MAIT cells first impaired by the inflammation occurring in inflamed tissues, or is microbiota dysbiosis the initial impact modifying MAIT cell function? Inflammation outside of the gut can activate and alter the function of the MAIT cells that subsequently migrate to the gut mucosa, where they are no longer able to maintain gut homeostasis, as suggested by the decreased production of IL-17 and IL-22 by intestinal MAIT cells in the non-obese diabetic (NOD) mouse model of type 1 diabetes^{24,70}. Impaired MAIT cells might also lose their ability to provide efficient antibacterial function. Together, these MAIT cell alterations could promote gut mucosa and microbiota alterations.

In a second hypothesis, initial modifications of gut homeostasis, such as increased bacterial translocation and modification of the intestinal microbiota, could alter the equilibrium between different bacteria and the level of activating and/or inhibiting ligands, thereby impacting MAIT cell functions. Identifying the kinetics and nature of MAIT cell interactions with the gut mucosa and microbiota represents a critical step for exploring MAIT cell-based treatments for autoimmune, inflammatory, metabolic and immune-mediated diseases.

MAIT cell exhaustion or dysfunction. Indeed, alterations of MAIT cell responses in vitro suggest an exhausted status. After phorbol myristate acetate–ionomycin stimulation or upon MR1-dependent activation, MAIT cells from patients with autoimmune and immune-mediated diseases are usually less activated and produce fewer T helper 1 (T_H1)-type cytokines than MAIT cells from controls^{16,18,73,77,78,82,83,85–87,89,93}. In patients with SLE, MAIT cell exhaustion has been associated with dysfunction of the Ca²⁺–calcineurin–NFAT1 signalling pathway downstream of TCR stimulation⁸⁹. Although MAIT cells from most immune-mediated diseases show defective activation and/or IFN γ and TNF production in vitro, several studies have described an increased frequency of MAIT cells producing IL-17 in the same pathologies^{16,18,70,73,77}. These observations suggest there is heterogeneity among MAIT cells, a hypothesis that is worthy of more detailed analysis.

Increased apoptosis of blood MAIT cells. MAIT cells in many immune-mediated diseases show increased apoptosis, possibly as a result of chronic activation. In children with recent-onset T1D, fewer circulating MAIT cells express the anti-apoptotic factor BCL-2, and this is positively correlated to the overall frequency of MAIT cells⁷⁰. A similar correlation is also observed in patients with liver cirrhosis⁶⁹. In patients with UC, CD and SLE, more circulating MAIT cells express activated caspases (which mediate apoptotic cell death) than in control individuals^{81,88}. X-linked lymphoproliferative syndrome type 2 (XLP2) is a rare human genetic immunodeficiency with some clinical signs similar to CD. XLP2 results from mutations in the gene encoding X-linked inhibitor of apoptosis (XIAP) and leads to a proapoptotic phenotype in invariant natural killer T (iNKT) cells and MAIT cells, which could explain the decreased frequency of such cells in the blood of these patients⁷⁶.

It would be interesting to analyse whether chronic inflammation, often associated with autoimmune diseases, dampens XIAP expression by MAIT cells.

Increased apoptosis of MAIT cells in target tissues.

As we mentioned above, MAIT cells have the propensity to migrate to inflamed tissues, and in many autoimmune and immune-mediated diseases MAIT cell frequency is increased in target tissues. However, in some tissues, such as the liver, colon and adipose tissue, where MAIT cells are abundant in the steady state, their frequency tends to decrease upon inflammation. These results suggest that increased apoptosis of MAIT cells could override potential MAIT cell recruitment in these inflamed tissues. For example, MAIT cells are less frequent in inflamed intestinal mucosa of patients with UC and CD and of children with coeliac disease, compared with healthy tissue^{74,81}. Decreased frequency of MAIT cells is also observed in adipose tissue from obese adults compared with the adipose tissue from lean adults, and in co-cultures of MAIT cells and adipose tissue from obese individuals BCL-2 expression is decreased compared with adipose tissue from lean individuals^{16,73}. Similarly, MAIT cells are less frequent in the livers of patients with cirrhosis and autoimmune liver disease (AILD) and have an activated/exhausted phenotype (CTLA4⁺ and PD-1⁺), as compared with control livers^{69,87}. Intrahepatic MAIT cell frequency is negatively correlated with the PD-1 expression of the cell. Increased MAIT cell proliferation is observed in the blood of patients with CD¹⁸ and with liver cirrhosis^{69,83}, as well as in vitro in the presence of inflamed adipose tissue from obese patients¹⁶; however, this proliferation is not sufficient to compensate for the loss of MAIT cells.

Other factors impacting MAIT cell frequency. The marked heterogeneity among patients with regard to factors — such as age^{107,108}, body mass index^{16,73}, physical activity¹⁰⁹ and ongoing treatment^{69,90,110} — that could impact MAIT cell frequency and function might explain the conflicting results in human studies^{71,111}. For example, decreased blood MAIT cell frequency is observed in adults but not in children with coeliac disease⁷⁴, and in obesity MAIT cell frequency is increased in children but decreased in adults^{16,73}. Whether the strong decrease of MAIT cell frequency reported in patients with severe asthma results from the pathophysiology or from the inhaled corticoid treatment is unclear⁹⁰. Indeed, the lower MAIT cell frequencies in the blood, sputum and endobronchial biopsies of asthmatic patients are positively associated with disease severity and negatively associated with inhaled corticosteroids. No change in MAIT cell frequency is observed in the broncho-alveolar lavage fluid from asthmatic patients, which could reflect the weak diffusion of the inhaled corticosteroids in more distal airways and the alveolar compartment⁹⁰. However, in another study, after exclusion of patients on corticoid treatment, the re-evaluated circulating MAIT cell frequency was still decreased in patients with inflammatory bowel diseases¹¹². Moreover, decreased MAIT cell frequency has been observed in other autoimmune or immune-mediated

diseases in patients without immune-modulating treatments^{16,69,70,73,74,83,93}.

An impact of hyperglycaemia on MAIT cell frequency has been proposed. In cardiovascular disease, circulating MAIT cell frequency negatively correlated to glycated haemoglobin (HbA_{1c}) levels⁹⁴, and in obese patients, decreased circulating MAIT cell frequency was greater in adults with uncontrolled HbA_{1c} levels⁷³. Moreover, exposure of MAIT cells to glucose increases their apoptosis in a dose-dependent manner⁹⁴. While no correlation was observed between HbA_{1c} level and blood MAIT cell frequency in children with T1D, granzyme B⁺ MAIT cell frequency was inversely correlated with HbA_{1c} levels at the onset of disease, but not in children with established disease⁷⁰. The potential role of granzyme B⁺ MAIT cells in T1D is further discussed below. Together, these observations suggest that decreased MAIT cell frequency in various diseases might have multifactorial origins.

Pathogenic roles for MAIT cells

Pathogenic role of MAIT cells in the liver. During chronic liver injury, signals originating from immune cells, as well as bacterial by-products, play central roles in the progression of liver fibrosis by activating the fibrogenic functions of hepatic myofibroblasts and their inflammatory properties. In the liver of patients with cirrhosis and AILD, MAIT cells accumulate in the fibrotic septa^{69,87}, making contact with hepatic myofibroblasts. Their profibrogenic functions were demonstrated in mouse models, in which deficiency of *Mr1* led to resistance to fibrosis induced by chronic carbon tetrachloride administration or bile-duct ligation, whereas expression of a Vα19TCR transgene, leading to an excess of MAIT cells, resulted in mice that were more prone to fibrosis⁶⁹. In vitro, MAIT cells enhanced the fibrogenic functions of hepatic myofibroblasts, characterized by increased proliferation and collagen secretion, and exacerbated their pro-inflammatory properties^{69,87}. Interestingly, hepatic myofibroblasts express MR1 at the cell surface and are activated by MAIT cells via an MR1-dependent pathway. By contrast, the myofibroblast pro-inflammatory properties can be induced in a contact-independent manner by TNF or IL-17A released by MAIT cells^{69,87}. In addition to directly interacting with fibrogenic cells, MAIT cells may also favour fibrosis progression by fostering a local inflammatory reaction, in that they promote polarization of macrophages towards a pro-inflammatory phenotype⁶⁹.

Pathogenic role of IL-17-producing MAIT cells. All human and most mouse MAIT cells express RORγt, a master transcription factor controlling IL-17 production^{22,113–115}. IL-17 has pathogenic roles in many autoimmune, metabolic and liver diseases¹¹⁶, suggesting that MAIT cells could be deleterious via IL-17 production. In the collagen-induced arthritis murine model, MAIT cells are pro-inflammatory¹¹⁷, in that the severity of the disease is reduced in *Mr1*^{-/-} mice and enhanced after T cell transfer from Vα19TCR-transgenic mice. Accordingly, total splenocytes from *Mr1*^{-/-} mice produce lower levels of IL-17 after anti-CD3

and anti-CD28 stimulation or IL-23 treatment. Human studies highlight a switch towards a T_H17-type profile of blood MAIT cells, with an increase in secretion of IL-17 in patients with CD¹⁸, UC^{18,79}, ankylosing spondylitis (AS)⁷⁷, obesity^{16,73}, type 2 diabetes (T2D)¹⁶ and cirrhosis⁶⁹, and of IL-22 in patients with UC¹⁸ and AS⁸⁴. Analysis of MAIT cells in patients with MS has further shown increased expression of RORγt, CCR6 and IL-7R compared with healthy controls¹¹⁸. Similarly, IL-17 production is increased in MAIT cells from the salivary glands of patients with Sjögren syndrome (SS) compared with controls^{78,86}, as well as after stimulation with IL-23 in vitro⁷⁸. In AS, the altered cytokine profile of blood MAIT cells is also observed in the synovial fluid in favour of IL-17 production^{77,84}, and IL-7 priming of MAIT cells increases IL-17 production in patients compared with healthy controls⁷⁷. Of note, IL-7 levels are elevated in the joints and gut of AS patients^{119,120}, and there is a positive correlation between IL-7 and IL-17 levels in these patients¹²¹. Interestingly, circulating MAIT cells from patients with AS have higher IL-7R expression levels than in controls⁷⁷. There is a report of IL-17 and IL-22 production by a MAIT cell clone derived from psoriatic skin lesions⁹⁸. Adipose tissue from obese individuals had an elevated frequency of MAIT cells that produce IL-17 compared with controls^{16,73}, which might induce insulin resistance in adipocytes^{122,123}. Interestingly, the pathogenic role for IL-17-producing MAIT cells is suggested in asthmatic children, in whom there is a correlation between cell frequency and disease severity (defined as the number of exacerbations)⁹². Most of these analyses in humans have shown correlations between MAIT cells, IL-17 production and disease status, but further studies are warranted to demonstrate causality.

Pathogenic role of cytotoxic MAIT cells. MAIT cells could have a pathogenic role in autoimmune and immune-mediated diseases via their cytotoxic properties. T1D is the most obvious example of this activity⁷⁰. Analysis of NOD mice has revealed an increasing level of granzyme B expression in pancreatic islets from the prediabetic stage to diabetes onset. In humans, more circulating MAIT cells produce granzyme B in children with recent onset T1D than do control MAIT cells. Interestingly, granzyme B⁺ MAIT cell frequency correlated negatively with the frequency of MAIT cells and with the age of the child at diagnosis, in agreement with the current view that T1D is more aggressive in the youngest children. These data in NOD mice and T1D patients suggest that MAIT cells could kill pancreatic β-cells, even though to date MAIT cells have not been detected in diseased islets¹²⁴. As we mentioned earlier, MR1 expression is upregulated on the human β-cell line EndoC-βH1 in the presence of pro-inflammatory cytokines, and in co-cultures MAIT cells from healthy controls can kill EndoC-βH1 cells in an MR1-dependent manner⁷⁰.

A similar pathogenic local effect of granzyme B⁺ MAIT cells is also suggested in patients with AS⁷⁷ and in obesity¹⁶. Of note, in NOD mice the presence of granzyme B⁺ MAIT cells in the intestinal mucosa increases with diabetes development, indicative of their potential

pathogenic role in this tissue before and at T1D onset (L. Beaudoin and A.L., unpublished observations). Together, these studies show that MAIT cells can mediate various pathogenic mechanisms.

A protective role for MAIT cells

Immunoregulatory role of MAIT cells. In contrast to their potentially pathogenic activities, MAIT cells also have immunoregulatory activities that may be protective in disease. MAIT cells have a protective role in the 2,4,6-trinitrobenzene sulfonic acid (TNBS)-induced colitis mouse model^{125,126}. MAIT cells are decreased in the colonic mucosa after TNBS treatment, and transfer of Ja33TCR⁺ MAIT cells reduces the severity of the disease, weight loss, intestinal bleeding and colonic inflammation. In experimental autoimmune encephalomyelitis (EAE)¹²⁶, the mouse model of MS, transgenic expression of the invariant Vα19TCR in T cells protects mice from EAE induction, and adoptive transfer of liver invariant Vα19TCR⁺ T cells into wild-type mice with EAE reduces the disease severity. Conversely, EAE in *Mr1*^{-/-} mice is exacerbated¹²⁶. The suppression of EAE by invariant Vα19TCR⁺ T cells is associated with reduced expression of pro-inflammatory molecules and increased production of IL-10 by B cells, through an inducible T cell co-stimulator-dependent interaction with invariant Vα19TCR⁺ T cells¹²⁶. IL-10 can inhibit excessive gut inflammation¹²⁷ and is implicated in IgA production¹²⁸. A regulatory role of IL-10-secreting MAIT cells is also suggested by their presence in the adipose tissue of lean individuals but their reduced frequency in obese individuals⁷³. In vitro experiments also support the immunoregulatory function of MAIT cells from both patients with MS and healthy adults⁹⁵, as they suppress IFNγ production by conventional T cells in a cell contact-dependent manner.

In a longitudinal study, the circulating MAIT cell frequency was shown to be lower in 1-year-old children who later develop asthma than in 1-year-olds who do not. A higher MAIT cell frequency is correlated with stronger IFNγ secretion by CD4⁺ T cells, suggesting a protective role of MAIT cells associated with a potent T_H1-type immune response¹²⁹. It would be interesting to determine whether there is a link between these observations in children and the hygiene hypothesis, implying that activation of the immune system at a young age can promote immunoregulatory mechanisms¹³⁰ and that MAIT cells could represent an ideal immune cell type to maintain regulatory function in the whole body.

Prevention of T1D by MAIT cells. Concerning T1D, an initial study suggested a protective role for MAIT cells, as diabetes is delayed in Vα19TCR-transgenic NOD mice¹³¹. More recently, analysis of *Mr1*^{-/-} NOD mice confirmed the protective role of MAIT cells, as these mice develop accelerated diabetes, despite the potential pathogenic role of MAIT cells in the pancreas⁷⁰. Indeed, this study suggests that MAIT cells could have a protective role in the gut mucosa through their production of IL-17 and IL-22, two cytokines required for the maintenance of gut barrier integrity. In *Mr1*^{-/-} NOD mice, exacerbation of diabetes development is associated with a loss of

gut barrier integrity, allowing the translocation of bacterial compounds and subsequent dendritic cell activation in pancreatic lymph nodes⁷⁰. Of note, a protective role of IL-17 has been established in several intestinal inflammation models¹¹⁶, and IL-22⁺ MAIT cells have been observed in the blood of patients with UC¹⁸ and AS⁸⁴.

Protective role of MAIT cells in the liver. In patients with nonalcoholic fatty liver disease (NAFLD), MAIT cells accumulate around fatty hepatocytes. Although mouse and human studies have demonstrated the profibrogenic properties of MAIT cells, *Mr1*^{-/-} mice fed a methionine–choline-deficient diet show enhanced steatosis, are more prone to liver injury and have more CD11c⁺ M1-like inflammatory macrophages but fewer CD206⁺ M2-like macrophages⁹³. Future studies will be needed to confirm this protective role of MAIT cells in a more relevant model of nonalcoholic steatohepatitis, and the contribution of IL-17 needs to be evaluated, as it is a major driver of NAFLD progression^{132,133}. In addition, it remains unclear whether gut-derived bacterial components impact MAIT cell activation during NAFLD.

Patients with alcoholic liver disease, either severe alcoholic hepatitis or end-stage cirrhosis, are at high risk of developing bacterial infections, because of a leaky gut, gut dysbiosis, increased bacterial translocation and dysregulation of immune cell functions. In in vitro experiments, stool extracts from patients with alcoholic liver disease induced MAIT cell depletion and suppressed their antibacterial cytokine (IL-17, TNF and IFNγ) responses and the expression of PLZF and RORγt, which are involved in MAIT cell maturation and effector functions⁸³. These data suggest that MAIT cell functional defects in patients with alcohol-related liver disease could contribute to this population's high susceptibility to bacterial infection, which is also described during autoimmune hepatitis⁸². Interestingly, prophylactic administration of antibiotics to patients with severe alcoholic cirrhosis, a treatment usually administered so as to avoid bacterial infections, partially prevents blood MAIT cell reduction and activation⁶⁹.

Together, MAIT cells can have multiple functions, and their pathogenic or protective role may rely on their activation or exhaustion status, tissue localization and cytokine profile, and on the chronicity of the disease. Even though the characterization of blood MAIT cells might not be as informative as tissue analysis, blood MAIT cells do seem to reflect the status of MAIT cells in inflamed tissues.

MAIT cells in human cancers

Recent studies have investigated the role of MAIT cells in cancer, and the observations reported are similar to those made in autoimmune and inflammatory diseases (TABLES 3,4). Circulating MAIT cell frequency is decreased in patients with colorectal cancer (CRC)^{134–136}, hepatocellular carcinoma (HCC)¹³⁷ and lung cancer¹³⁶. MAIT cells accumulate in tumour tissues from patients with colon adenocarcinomas¹³⁸ and CRC, particularly in those with advanced CRC^{134–136}, and in hepatic metastatic lesions in patients with CRC¹³⁹. In contrast, in

Table 4 | MAIT cells in cancer

Cancer	MAIT cell frequency		MAIT cell phenotype		MAIT cell effect on disease	Possible mechanisms of action	Refs
	Blood	Tissues	Blood	Tissues			
Kidney	↓	↑	ND	ND	ND	Increased MAIT cells in tumour tissue	160
Brain	↓	↑	ND	ND	ND	Increased MAIT cells in tumour tissue	160
Lung	↓	ND	ND	ND	ND	ND	136
Breast	↔	ND	ND	ND	ND	Increased IL-17 produced by breast duct MAIT cells	136,161
Colorectal	↓	↑	ND	ND	Protective?	<ul style="list-style-type: none"> Increased MAIT cells in tumour tissue Increased CD69 and PD-1 expression Decreased IFNγ production 	134–136,138
Hepatic metastasis	ND	↑	ND	ND	Protective and/or deleterious?	<ul style="list-style-type: none"> Increased MAIT cells in tumour tissue Decreased IFNγ production and increased IL-17 production 	139
Hepatocellular carcinoma	↓	↓	Exhausted	Exhausted	Deleterious?	<ul style="list-style-type: none"> Decreased MAIT cells in tumour tissue, more IL-8 produced (pro-tumour cytokine) High MAIT cell frequency associated with poor clinical outcome 	137
Graft-versus-host disease (in individuals treated for cancer with bone marrow transplant)	↓	ND	ND	ND	Protective	Low MAIT cell frequency associated with poor clinical outcome	140–142

Exhausted denotes cells with impaired cytokine production and function. Deleterious? denotes a deleterious effect is suggested or hypothetical and not yet demonstrated. Protective? denotes a protective effect is suggested or hypothetical and not yet demonstrated. ↑, increased frequency; ↓, decreased frequency; ↔, no change; IFN γ , interferon- γ ; MAIT, mucosal-associated invariant T; ND, not determined; PD-1, programmed cell death 1.

patients with HCC, the frequency of MAIT cells was decreased in the liver, and in the tumour, when compared with healthy controls. As we observed in immune-mediated diseases, this could reflect increased apoptosis of MAIT cells, possibly as a result of their chronic activation. Indeed, in patients with HCC, tumour-infiltrating MAIT cells had an exhausted phenotype, with high PD-1, CTLA4 and TIM3 expression, and produced less IFN γ and IL-17 than MAIT cells from healthy donors¹³⁷. Similarly, MAIT cells from colonic adenocarcinomas produced less IFN γ than MAIT cells from healthy colons¹³⁸, suggesting that tumour-infiltrating MAIT cells might be altered due to tumour microenvironmental factors. Furthermore, tumour-infiltrating MAIT cells produced less IFN γ than MAIT cells from healthy livers in response to recombinant IL-12 and IL-18 (REF.¹³⁹). As was observed in other diseases, circulating MAIT cells in patients with CRC produce more IL-17 and less IFN γ and TNF¹³⁴. Of note, chemotherapy does not alter the ability of MAIT cells to produce IFN γ , nor their frequency¹³⁹, in agreement with their expression of the multidrug resistance transporter ABCB1 (REF.¹¹). High frequencies of tumour-infiltrating MAIT cells are associated with poor clinical outcomes for patients with HCC. In contrast, in the context of haematological malignancies, a higher frequency of MAIT cells after allogeneic haematopoietic cell transplantation is associated with less graft-versus-host disease (GVHD)^{140,141}. Mouse studies have also shown a beneficial role of MAIT cells against GVHD and highlighted their role in maintaining gut integrity¹⁴². All these studies suggest that MAIT cells could represent attractive therapeutic targets; however, their precise role in cancer needs further investigation.

Conclusion and perspectives

MAIT cells provide efficient protection against acute bacterial and viral infections. However, in chronic pathological situations, including chronic viral infections and autoimmune, inflammatory and metabolic diseases, MAIT cells can play a pathogenic role by sustaining inflammation and cytotoxicity. This ambivalent function is reminiscent of NKT cells, which share many features with MAIT cells^{22,143,144}, although the ligands recognized by the two innate-like T cell populations are quite distinct. Several endogenous ligands of NKT cells have been described^{145,146}, however, the MAIT cell endogenous ligands remain to be identified. Although both MAIT and NKT cells are conserved in mammals, MAIT cells are much more abundant than NKT cells in humans¹⁴⁴. Moreover, MAIT cell localization in the gut mucosa and their recognition of bacterial ligands might play a key role in sensing any modification of the microbiota composition¹⁴⁷, either commensal or pathogenic, which might reflect the infectious, metabolic and general health status of the individual (BOX 2). Further studies should be performed to provide a better understanding of the functional plasticity of MAIT cells, the epigenetic regulation of MAIT cell function, the impact of their TCR repertoire, and their interactions with other immune and non-immune cells in physiological and pathological situations. Many of the data presented in this Review highlight MAIT cells as biomarkers in various diseases; however, new insights into their biological functions are still awaited, to inform the design of therapeutic approaches based on MAIT cell manipulation.

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