**TIM-3, a Possible Target for Immunotherapy in Cancer and Chronic Viral Infections**

Roger Tieu¹, Praveen Kumar Amancha¹, François Villinger²* and Siddappa N. Byrareddy²*

¹Division of Microbiology and Immunology, Yerkes National Primate Research Center, USA
²Department of Pathology & Laboratory Medicine, Emory University School of Medicine, USA
*Corresponding author: Siddappa N. Byrareddy, Department of Pathology & Laboratory Medicine, Emory University School of Medicine, 101 Woodruff Circle, WMBR, Room 237A, Atlanta

**Introduction**

Upon exposure to non-self entities such as pathogens, a number of receptors including Pattern Recognition Receptors (PRRs) are triggered leading to a rapid response of the innate immune system [1]. This initial response paves the way for the elaboration of pathogen specific adaptive humoral and cell mediated responses via a complex network of co-stimulatory and inhibitory signals. The initial effector response by and large favors co-stimulatory signaling for the activation and proliferation of effector cells with the intent of eliminating the invading pathogen. After controlling or eliminating the threat, the absence of continuing antigen stimulation leads to elimination of most effector cells via apoptosis while a fraction of these cells become long term memory cells and homeostasis is reestablished. However, during the course of cancer and chronic persisting viral infections, this return to normacy is perturbed leading to immune dysfunction and dysregulation resulting in the continued presence of effector cells with poor function. The mechanisms leading to such muted phenotype are increasingly being characterized and involve the expression of receptors, which are thought to play a central role in this inhibition of function.

These inhibitory mechanisms are an integral and critical contributor to immune homeostasis, in minimizing collateral damage during extended inflammatory response as well as preventing autoimmunity by maintaining peripheral self-tolerance. Such pathways serve to restrain effector cell activity, mainly of T cells. Inappropriately activated lymphocytes recognizing self-antigen subsequently lose effector function and undergo apoptosis or energy. Such responses require constricted regulation and transient expression of inhibitory signaling. Prolonged and/or over expression of multiple inhibitory receptors have been associated with impaired immune function such as T cell exhaustion, a state characterized by poor effector function, a mechanism that is exploited by many persistent pathogens.

During chronic active infections, a progressive loss of function in pathogen-specific CD8+ T cells was documented following a hierarchy with an initial impaired production of IL-2 and cytotoxicity, followed by loss of TNF-α, and eventually IFN-γ at late stages [2], resulting in a condition known as exhaustion [2,3] (Figure 1). T cell exhaustion has been found to be associated with the up-regulation of inhibitory receptors in concert with their ligands also up-regulated on Antigen Presenting Cells (APCs) [3]. The initial marker associated with inhibition of function has been Cytotoxic T Lymphocyte Associated Antigen-4 (CTLA-4), a marker upregulated on the surface of activated T cells, with markedly higher affinity for CD80/CD86, the co-stimulator ligands to CD28 during the elaboration of an antigen specific response [4]. More recently, PD-1 (programmed death 1) was found to be selectively up-regulated by exhausted T cells during murine chronic LCMV (Lymphocytic Choriomeningitis Virus) infection, and blocking of this receptor with antibodies enhanced virus-specific CD8+ T cell response and decreased viral load [5]. This finding defined a novel role for inhibitory pathway involvement in controlling T cell response in chronic viral infection as well as the potential for targeting these same pathways. However, blockade of the PD-1 pathway only partially restored T cell functions, suggesting the involvement of other inhibitory pathways. Additional inhibitory receptors have been since identified which include T cell immunoglobulin and mucin domain 3 (TIM-3), CD244 (2B4), killer Cell Lectin like Receptor G1 (KLRG1), and Lymphocyte Activation Gene 3 (LAG3) [6-12].

**TIM-3, a receptor involved in the regulation of immune response**

TIM-3 is a type I transmembrane protein consisting of an N-terminal Immunoglobulin Variable (IgV)-like domain, a mucin domain with potential sites of O- and N-linked glycosylation, followed by a transmembrane domain, and a cytoplasmic tail with tyrosine phosphorylation motifs. The murine TIM-family genes were cloned from a segment of chromosome 11 homologous to human 5q23-35 associated with asthma susceptibility [13]. Polymorphisms in TIM-3 have been associated with development of airway hyper-

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**Abstract**

Effector T-cell responses are controlled by complex mechanisms involving various soluble factors and co-stimulatory and co-inhibitory molecules. These inhibitory receptors act as immune checkpoints and are extensively investigated as possible therapeutic targets, such as PD-1 and CTLA-4. Recently TIM-3 is also gaining prominence in tumor and chronic viral infection models as a candidate for immunotherapy in conjunction with other inhibitory receptors. This review discusses the recent findings on the expression of TIM-3 and its ligand in tumor and chronic viral infection.

**Keywords:** TIM-3; CTLA-4; Immunotherapy; Galectin-9
Galectin-9, a member of endogenous lectin acts as a ligand for TIM-3

Galectin-9 was identified as a ligand binding to carbohydrate motifs on IgV domain of murine TIM-3 [21]. Galectin-9 is an S-type lectin with two distinct carbohydrate recognition domains joined by a linker peptide [34], and can bind to glycans featuring galactose and other sugars. Target carbohydrates recognized by galectin-9 remain to be elucidated but structural analysis has revealed that galectin-9 can complex with poly-N-acetyllactosamine-containing structures [35]. Various tissues including the thymus express galectin-9 [34]. Given the nature of TIM-3/galectin-9 interactions, much effort has focused on identifying a specific galectin-9 expressing population. Initial reports have found Tregs and naïve CD4+ T cells to express galectin-9 though additional studies will be needed to elucidate in what context galectin-9 assumes an active role [36,37]. Interestingly, one study showed a lack of interaction between galectin-9 and TIM-3 from both murine and human origin [38]. However, it must be noted that this study utilized membrane-bound galectin-9 rather than the soluble form of galectin-9 used in the other studies, and such a difference in methodology may impact signaling and partly explain the discrepancies. Furthermore, structural analysis of TIM-3 revealed galectin-9 independent binding sites for the IgV region [39]. Tetramers of bacterially expressed TIM-3, which lack any glycosylation, bound to unknown ligand(s) on T cells, B cells, DCs, and macrophages. These studies warrant the investigation of additional ligands that may interact with the full-length TIM-3 protein, or its IgV or mucin domains, which have yet to be explored.

Stimulation of TIM-3 expressing T cells with galectin-9 resulted in decreased susceptibility to HIV infection, via downregulation of HIV-1 co-receptors, CCR5, CXCR4 and α4β7 on the surface of activated CD4+ T cells [40]. In HIV-infected T cells, the presence of galectin-9 limits viral replication. Conversely, resting T cells lacking TIM-3 expression exhibited increased HIV infection in the presence of galectin-9 in a PD1-dependent manner. A different study demonstrated the expression of TIM family proteins, including TIM-3, enhances HIV infection of CD4+ T cells but limits viral replication and release of virus from infected cells in an apoptotic-independent manner [41]. However, another study showed galectin-9 not as a pro-apoptotic agent, but rather as an inducer of pro-inflammatory cytokines from T helper cells in a TIM-3 independent manner [42]. In a controversial study, TIM-3 was shown not to interact with galectin-9 and in fact, blockade of TIM-3 signaling in human T cells did not restore effector function [38]. These findings suggest that the interaction of TIM-3 and galectin-9 is likely far more complex than a simple interaction and may heavily depend upon the context of the interaction.
Role of TIM-3 in cancer/tumor models and in chronic viral infections

Attempts to identify new Th1-specific cell surface proteins resulted in the identification of TIM-3 [18]. Treatment with antibody specific to TIM-3 exacerbated Experimental Autoimmune Encephalomyelitis (EAE) in mice [18], and increased the proliferation and activation of macrophages. Although the underlying mechanism was not elucidated, these studies suggested two possible mechanisms; 1) cross-linking of TIM-3 receptors on Th1 cells promotes an increased pro-inflammatory state, which may activate macrophages directly or indirectly; 2) Disrupting interactions between TIM-3 and its inhibitory ligand on macrophages allows for their activation in the presence of (or independent of) Th1 cells. TIM-3 expressing DCs can be either stimulatory or inhibitory depending on the microenvironment. A recent study illustrated that the expression of TIM-3 was markedly upregulated on tumor infiltrating DCs and that such expression correlated with inhibition of PRR mediated innate responses, independent from the effect of galectin-9. Instead, TIM-3 was shown to bind HMGB1 (high mobility group box protein 1) inhibiting the recruitment of nucleic acids from dying tumor cells as HMGB1/nucleic acid complexes to the endosome and thereby blocking the activation of the DCs [43]. Without proper stimulation of DCs, TIM-3 expressing tumor-specific CD8+ T cells will fail to escape their exhausted state [44]. In mice bearing solid tumors, a large fraction of tumor infiltrating lymphocytes were shown to be coexpressing PD-1 and TIM-3 [45]. This population exhibited a more severe exhausted state and a combined targeting of PD-1 and TIM-3 showed better control of the tumor [45]. Similar findings were noted for the Acute Myelogenous Leukemia (AML) model in mice [46].

Increased expression of TIM-3 has been observed on exhausted CD8+ T cells isolated from individuals infected with Human Immunodeficiency Virus (HIV) [16,47,48], Hepatitis B Virus (HBV) [49,50] and Hepatitis C Virus (HCV) [8,17,47] and this level of TIM-3 expression correlated with ineffective antiviral responses. Murine models of chronic viral infections have been used to observe altered patterns of effector function of TIM-3 expressing T cells in HBV [51], Herpes Simplex Virus (HSV) [52] and LCMV infection. Of note, one study demonstrated that HIV infection of CD4+ T cells does not induce expression of TIM-3 on T cells [53]. Instead, TIM-3 expression may be the result of immune activation through TCR/CD28 co-simulation and the presence of common γ-chain (γc) cytokines IL-2, IL-7, IL-15, and IL-2, but not IL-4 [53].

Recently, it has been suggested that TIM-3 and PD-1 expressing cells marked distinct populations, with a minority expressing both in HIV-infected individuals [16]. TIM-3 expression levels directly correlated with clinical prognosis and inversely with CD4+ T cell counts. Exhausted CD8+ T cells were characterized by increased expression of TIM-3, but blockade of the TIM-3 pathway restored partial function with increased production of IFN-γ in vitro. TIM-3 presents itself as a viable therapeutic target as demonstrated by enhanced HIV-specific T cell response. However, it remains unclear whether prolonged blockade of TIM-3 regulation of T cell functions may result in increased inflammation and potentially breaking of tolerance to self.

Resting T cells show minimal expression of TIM-3, which is upregulated only after prolonged stimulation. Unlike resting T cells, resting NK cells constitutively express TIM-3 [25,26]. TIM-3 upregulation appeared to mark the transition from CD56brightCD16- NK cells to mature CD56dimCD16+ NK cells [25]. Stimulation of TIM-3 expressing NK cells resulted in cytokine production and cytotoxicity. In a different study, the presence of galectin-9 significantly enhanced the ability of TIM-3 expressing NK cells to produce IFN-γ [26]. However, recent publications have reported conflicting findings regarding the role of TIM-3 on NK cells [27,54]. NK cells isolated from HBV patients had high expression levels of TIM-3, and blockade of TIM-3 signaling with TIM-3 blocking antibodies or soluble TIM-3 (sTIM-3) protein led to increased cytotoxicity and IFN-γ production in vitro, suggesting TIM-3 signaling plays a suppressive role for NK cell effector functions [54]. NK cells from advanced melanoma patients demonstrated an increased upregulation of TIM-3 and displayed an exhausted phenotype characterized by impaired cytotoxicity, IFN-γ production and cell proliferation [27]. Blockade of TIM-3 with various antibodies promotes internalization of the TIM-3 receptor and upregulation of the IL-2 Receptor (IL-2R). Increased density of IL-2R on the cell-surface renders NK cells more responsive to IL-2 stimulation, thus improving NK cell-mediated cytotoxicity, suggesting a potential mechanism linked to alternate signaling via the internalized TIM-3. However, similarities to prior studies are also apparent, with decreased NK cell cytotoxicity upon cross linking of TIM-3 by antibody or binding to soluble or membrane-bound galectin-9. Interestingly, the addition of galectin-9 blocking antibody did not affect cytotoxicity, suggesting TIM-3 negatively regulates N K cell function in a galectin-9-independent fashion, thus implicating additional TIM-3 ligands. It is important to note that the studies portraying TIM-3 as an NK cell activation marker only evaluated healthy donors, whereas the studies implicating TIM-3 as an NK cell suppression marker evaluated patients with chronic diseases. Such a distinction may play an important role in understanding the modular response of TIM-3 depending on the disease context.

TIM-3: target for immunotherapy

During chronic viral infections with sustained antigen stimulation, exhausted CD8+ and CD4+ T cells can co-express multiple inhibitory receptors, which can work synergistically to modulate the functional quality of these virus-specific T cells. The severity of the dysfunction has a direct correlation with the combined number of expressed and putatively triggered inhibitory receptors. Co-expression of TIM-3 and PD-1 on CD8+ T cells in a murine model of LCMV was associated with severe CD8 T cell exhaustion characterized by markedly diminished proliferation and cytokine production upon antigen stimulation [55]. TIM-3 and PD-1 function cooperatively to limit T cell function, since blockade of both pathways was markedly more effective than TIM-3 blockade or PD-1 blockade alone, which only partially restore T cell effector function. Upon dual blockade, T cell function restoration was significantly greater than blockade either single pathway suggesting additive or even synergy.

However, as alluded to above, TIM-3 blockade like any blockade of natural immune inhibitory mechanism will have to be carefully evaluated in vivo for potential exacerbation of inflammatory responses and the potential for breaking tolerance to self, if used for prolonged periods of time. In that regard, testing in murine models may or may not provide relevant findings for a translation to the clinic.
Future Directions

Many questions pertaining to TIM-3 remain unanswered. At the molecular level, the glycosylation pattern of the TIM-3 receptor remains to be elucidated. Carbohydrate moieties on TIM-3 play an important role in ligand binding as evident by galectin-9 binding the TIM-3 IgV domain in a carbohydrate-dependent manner [21]. Given that the TIM-3 mucin domain also carries carbohydrate moieties for ligand interactions, it would be of considerable interest to determine the role of the mucin domain as well as its potential ligands. The combined finding of a galectin-9 independent binding site on TIM-3 and that non-glycosylated soluble TIM-3 binds to numerous primary immune cells suggests that there are additional ligands involved in the immune regulation of various cell lineages [39]. The identification of an endogenous soluble TIM-3 is another intriguing finding that begs the question of whether this isoform may have a different glycosylation pattern [19].

Future investigations must keep in mind the context in which TIM-3 and soluble TIM-3 is expressed since their respective function appears to markedly vary depending on the disease, environment, and temporal context. Currently, it remains an open question as to whether the TIM-3 expressing T cells from cancer patients represent an equivalent functional population to TIM-3 expressing T cell from HIV infected people. In the light of unexpected and often times conflicting findings, it is important to keep in mind that species specific differences may exist and murine, rhesus macaque, and human biological pathways may not match up [56-58] as described previously (70-72).

Also the longevity of exhausted TIM-3+CD8+ T cells needs to be addressed: Thus, are these the same cells that are indeed being rescued or conversely, does blockade of TIM-3 (and other inhibitory markers) prevent newly activated cell from becoming exhausted? These are questions that need to be addressed in the context of immunotherapy aiming to reinvigorate of these populations in vivo. It may be possible to stimulate an effective response from T cells by inducing severe variations in levels of antigen followed by antigen specific stimulation, which is currently being explored with discontinuation of ART and administration of Fc-fusion proteins during SIV-infection [59,60] (Figure 2). Given that cases of spontaneous control of HCV have been reported, it is possible that exhausted T cells may be activated in response to an environmental cue [61]. Additional studies are needed to evaluate these concepts. Furthermore, it is unclear whether the dysfunctional states of CD4+ and CD8+ T cells are comparable. And other primary immune cell lineages including B cells [62], NK cells [27,54], and DCs [63] have been reported to exhibit exhaustion as well, the role of TIM-3 in these conditions remain to be elucidated.

Conclusion

Finally, it has become clear that TIM-3 plays an intricate role along with other receptors in balancing immune activation and suppression. Given the great complexity of the crosstalk between the various signaling pathways, computational analysis and systems biology will take upon more of a role in understanding immunological mechanisms and disease pathogenesis [64-67], especially those involving TIM-3. The integration of knowledge from all the “omics” fields, including genomic, transcriptomic, proteomic, metabolomic, and lipidomic, will be essential in dissecting the role of TIM-3.

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References

Siddappa N. Byrareddy


Siddappa N. Byrareddy


