

## Original Research Article

## Recent Advances on T Regulatory Cells

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## ARTICLE INFO

*Article History***Received:** October 14, 2019**Accepted:** December 25, 2019**Volume:** 1**Issue:** 1

## KEYWORDS

T regulatory cells (Tregs), Myeloid derived suppressor cells (MDSCs), antigen presenting cells (APCs), dendritic cells (DCs), forkhead box P3 (FoxP3+), natural Tregs (nTregs) and induced Tregs (iTregs)

## ABSTRACT

Active suppression of tumour-reactive T cells can limit both immune surveillance and immunotherapy. Development of cancer results into the generation of immune suppressive network to inhibit anti-tumour activity and evade the host's immune response, which eventually facilitates tumour progression. Emerging evidence indicates that immunosuppressive cells; T regulatory cells (Tregs) may be largely responsible for inhibiting host T-cell activity against tumour-associated antigens and impair the effectiveness of anticancer immunotherapeutic approaches. Reducing the deleterious effects of Treg may increase the success of various immunotherapeutic modalities in cancer. However, an increase in our understanding of the local tumour microenvironment and the exact mechanisms of Treg induction and/or expansion in peripheral blood and tumour microenvironment of cancer patients should provide opportunities to test different treatments to target these immunosuppressive cells and alter the balance in favor of generating effective anti-tumour immune responses. The purpose of this study is to provide information about the phenotype, role and the function of Tregs and the mechanisms of their expansion in cancer.

## 1. Background

## 1.1. Introduction

Numerous approaches have been explored to harness the potency of the immune system to target cancer. To date, these have been essentially focused on enhancing the immunogenicity of the tumour or the induction and expansion of immune effectors to potentially target and eradicate the tumour. One plausible hypothesis is that the local tumour environment is strongly immune suppressive suggesting that there is a need to overcome immunosuppression mechanisms (Talmadge, 2007). It is hypothesised that as a tumour evolves during the early stages of neoplastic disease, there is an orchestrated generation of a pro-inflammatory microenvironment which provides mitogenic stimuli, supports neo-angiogenesis and circumvents T cell mediated antitumour responses (Dunn, Bruce, Ikeda, Old, & Schreiber, 2002). Evidence indicates that increase the level of immunosuppressive cells; Tregs and MDSCs may be responsible for the inhibition of host T cell activity against tumour-associated antigens (TAA). In addition, it is proposed that an increase in the numbers of these immune suppressive cells could negatively impair the effectiveness of anti-cancer immunotherapeutic approaches.

To clarify these roles and immunosuppression activity of Tregs, we have to understanding first of all; the interaction between immune cells (Tregs) and cancer cells. And secondly we need to know the functional activity of these immunosuppressive cells and reduce their effects, which may increase the success of design various and effective immunotherapeutic strategy in cancer patients. This review will focus on the immunosuppressive cells (Tregs) phenotype, functional activity and their roles in normal and cancer.

## 2. T regulatory cells (Tregs)

The importance of Tregs in both normal and pathological conditions (including tumour and microbial immunity, transplantation and allergy) is now better known (Yamaguchi, Wing, & Sakaguchi, 2011).

### 3. Definition of Tregs

Tregs cells were discovered by (Sakaguchi, Sakaguchi, Asano, Itoh, & Toda, 1995) group as double positive cells with CD4 and CD25 (CD4<sup>+</sup>CD25<sup>+</sup>) that main function was inhibiting the activity of T cells (Sakaguchi et al., 1995). Tregs cells are heterogeneous subset of cells composed of different phenotypes and functions (Mohr, Malhotra, Mayer, Gorochoy, & Miyara, 2018). Moreover, there is also CD8<sup>+</sup> Tregs but still not well described yet (Holderried, Lang, Kim, & Cantor, 2013; H.-J. Kim, Verbinen, Tang, Lu, & Cantor, 2010). In the presented study will be focused on CD4<sup>+</sup> Tregs and it is believed that there are two main subpopulations of CD4<sup>+</sup> Tregs with different origins; natural Tregs (nTregs) and induced Tregs (iTregs).

### 4. Phenotype of Tregs

Tregs express the canonical Treg markers such as, interleukin-2 (IL-2; CD25) receptor  $\alpha$ -chain, FoxP3, glucocorticoid-induced TNF receptor (GITR) and cytotoxic T-lymphocyte associated protein 4 (CTLA4; CD152), the TNF receptor superfamily member 25 (TNFRSF25; DR3), CD73 and CD39 (Deaglio et al., 2007; B.-S. Kim et al., 2015; Lin et al., 2013; Madireddi et al., 2017; Zhao, Liao, & Kang, 2017). In addition, nTregs display higher levels of programmed cell death-1 (PD-1; CD279), neuropilin 1 (Nrp1), Helios, glycoprotein A repetitions predominant (GARP; LRRC32), lymphocyte activation antigen-3 (LAG-3; CD223) and latency-associated peptide (LAP) compared with iTregs (M.-L. Chen, Yan, Bando, Kuchroo, & Weiner, 2008; Huard, Prigent, Tournier, Bruniquel, & Triebel, 1995; Y. C. Kim et al., 2012; Lin et al., 2013; Sun, Jin, & Li, 2016; Tran, Andersson, Wang, et al., 2009; Weiss et al., 2012; Yadav et al., 2012).

### 5. Classification of Tregs

Treg cells are mainly subdivided into two population nTregs and iTregs as discussed in the following sections.

- a. nTregs is also named as thymus-derived nTreg cells as it develops in the thymus and it is originally characterised as a subset of CD4<sup>+</sup> T cells constitutively expressing high levels of the interleukin-2 (IL-2) receptor  $\alpha$ -chain (CD25) (Lin et al., 2013). They are antigen-specific, play a key role in maintaining self-tolerance, and comprise approximately 5-10% of total peripheral CD4<sup>+</sup> T cells. The main characteristic phenotype of these cells that they express the forkhead box P3 (FoxP3<sup>+</sup>) proteins and also known as the winged helix transcription factor, this is crucial for the development and function of nTregs. and it was demonstrated that retroviral gene transfer of FoxP3 confers suppressor function on peripheral naïve CD4<sup>+</sup>CD25<sup>-</sup> T cells (Fontenot, Gavin, & Rudensky, 2003; Hori, Nomura, & Sakaguchi, 2003; Rudensky, 2011). In addition to FoxP3, approximately 70% of CD4<sup>+</sup> Tregs express high level of the Ikaros family member Helios (Thornton et al., 2010). The co-expression of Helios<sup>+</sup> makes CD4<sup>+</sup> Tregs more effectively suppress cytokine production by T cells (Himmel, MacDonald, Garcia, Steiner, & Levings, 2013; Sugita et al., 2015) as well as stabilize Helios<sup>+</sup> CD4<sup>+</sup> Tregs under inflammatory conditions than Helios<sup>-</sup> CD4<sup>+</sup> Tregs (Dhuban et al., 2015; Dijke et al., 2016; Sharma et al., 2013). Studies demonstrated that deficient mice with Helios cells develop autoimmune with low suppressive activity of CD4<sup>+</sup> Tregs (H.-J. Kim et al., 2015; Sebastian et al., 2016).

On the other hand, coexpression of Helios and FoxP3 in murine CD4<sup>+</sup> Tregs increases the suppression activity of Tregs compared with Foxp3 alone (Fu et al., 2012). Additionally, it has been shown that Helios is also critical for CD8<sup>+</sup> Treg to mediate its function (H.-J. Kim et al., 2015).

- b. iTregs, also known as adaptive or periphery Tregs (pTregs) which are generated in peripheral lymphoid tissue by the conversion of conventional naïve CD4<sup>+</sup>FoxP3<sup>-</sup> T cells to FoxP3<sup>+</sup> Tregs (Li & Zheng, 2015). iTregs develop outside the thymus upon the activation of conventional T cells by polyclonal or antigen-specific activation in the presence of specific immune suppressive cytokines and exerting their function *in vivo* in a cytokine-dependent manner such as transforming growth factor beta (TGF- $\beta$ ) (Bluestone & Abbas, 2003; de Lafaille & Lafaille, 2009; Gallimore & Simon, 2008; Schallenberg, Tsai, Riewaldt, & Kretschmer, 2010). TGF- $\beta$  is a multifunctional cytokine that has effect on wide range of cell growth and differentiation, it has been shown to induce the conversion of peripheral CD4<sup>+</sup>CD25<sup>-</sup> naïve T cells to CD4<sup>+</sup>CD25<sup>+</sup>FoxP3<sup>+</sup> Tregs (W. Chen et al., 2003; Filippi et al., 2008). Three subsets of CD4<sup>+</sup> Tregs (Tr1, Th3 and iTr35) and CD8<sup>+</sup> Tregs (CD8<sup>+</sup>CD28<sup>-</sup>) as discussed in brief subsequently:

- i. Type 1 regulatory T cells (Tr1), it lacks the expression of FoxP3<sup>+</sup> but it secretes high levels of interleukin (IL-10), this is induced by antigen stimulation via an IL-10-dependent process both *in vitro* and *in vivo* (Grazia Roncarolo et al., 2006; Hawrylowicz & O'garra, 2005; Povolieri et al., 2013). Moreover, Tr1 can be induced and mediated by molecules such as cyclooxygenase 2 (COX-2) that expressed by tumour cells (Akasaki et al., 2004; Bergmann, Strauss, Zeidler, Lang, & Whiteside, 2007).

- II. T helper cells (Th3), like Tr1, it also lacks FoxP3<sup>+</sup>, but it secretes a high level of TGF- $\beta$ , Th3 cells are induced as a result of antigen administration, in presence of TGF- $\beta$  and IL-4 (Inobe et al., 1998; Povoleri et al., 2013; Weiner, 2001).
- III. iTr35 cells do not express FoxP3 and they are induced and mediate their suppression activity via the inhibitory cytokine IL-35 (Collison et al., 2010; Povoleri et al., 2013). The exact role and function of iTr35 cells in immune response is still elusive and need further investigations.
- IV. CD8<sup>+</sup> Tregs are recently described T cells that function to control and regulate various T cell responses, in particular, play critical role in cancer and autoimmunity (Chaput et al., 2009; Gupta & Agrawal, 2020). CD8<sup>+</sup> Tregs are induced following repeated activation of CD8<sup>+</sup> T cells with antigen and this activation leads to generation of suppressive T cells with phenotype CD8<sup>+</sup>CD28<sup>-</sup> (Jiang et al., 1998).

Both nTregs and iTregs suppress a variety of physiological and pathological immune responses in a contact- and/or cytokine-dependent manners, with a wide potential immune suppressive molecules that processes FoxP3<sup>+</sup> Tregs in particular, reviewed largely (Shevach, 2009; Q. Tang & Bluestone, 2008).

## 6. Tregs expansion in cancer

In terms to the roles of Tregs cells in cancer, both nTregs and iTregs are thought to potentially contribute to its development by promoting tumour-specific T cell tolerance, inhibiting anti-tumour immune responses (Shevach, 2009). In addition to Treg cells, both CD4<sup>+</sup> and CD8<sup>+</sup> T suppressive cells have been described in a number of cancers (Andersen et al., 2009; Bailur, Gueckel, Derhovanessian, & Pawelec, 2015; Chaput et al., 2009; Preston et al., 2013; Tatsumi et al., 2003; Woo et al., 2001; Yamamoto et al., 2012; Zhou et al., 2020). Importantly, a large number of FoxP3<sup>+</sup> Treg infiltrate into tumours, and systemic removal of these cells enhances natural as well as vaccine-induced anti-tumour T cell responses, more details can be found in this review (Nishikawa & Sakaguchi, 2010). Studies demonstrated that Tregs are expanded in peripheral blood and tumour microenvironment in cancer patients and this usually correlates with poor prognosis and reduced survival (Chaudhary & Elkord, 2016; Filaci et al., 2007; Jacobs et al., 2010; Whiteside, 2015). It seems that tumours have diverse immunosuppressive mechanisms and networks to escape the antitumour immunity (Demaria, 2013). However, Dannull and colleagues have reported that vaccine efficacy (mediated antitumor immunity) was boosted in cancer patients following the depletion of Tregs (Dannull et al., 2005). The available evidence suggest that Tregs are implicated in the immunopathology of cancer and their depletion may enhance the immunotherapy efficacy.

Given the background on Tregs in cancers, however, it is still unknown whether systemic or local (tumour microenvironment) sources of Tregs are the key players. At least four potential sources for Tregs in the tumour microenvironment have been identified including 1) trafficking, 2) differentiation, 3) expansion by dysfunctional antigen-presenting cells (APCs) and 4) conversion of T cells into Tregs by TGF- $\beta$  (Zou, 2006). The fourth mechanism, conversion of conventional T cells into Tregs has been supposed as a potential mechanism for Treg expansion in cancers and this was supported by *in vitro* and mouse model studies with no data from *in vivo* or human studies to support its role in enriching peripheral and tumor-infiltrating Tregs. Moreover, studies reported that some soluble mediators such as prostaglandin (E<sub>2</sub>), vascular endothelia growth factor (VEGF) and TGF- $\beta$  were detected at high levels in certain tumours and their roles may have negative effects on the APC and dendritic cells (DCs) maturation and functions (Gomella, Sargent, Linehan, & Kasid, 1989; Smyth et al., 2003; Wada et al., 2009) and more details can be found on the following review (Zhao et al., 2017).

Studies have been shown that reported that chemokine gradients in the tumour microenvironment can attract circulating Tregs. Expression of different chemokine receptors, such as CXCR3, CCR4, CCR6 and CCR9 were shown to promote nTreg subsets accumulation in non-lymphoid tissues, in order to exert tissue-specific immunoregulation (Hirahara et al., 2006; Kleinewietfeld et al., 2005; Santodomingo-Garzon, Han, Le, Yang, & Swain, 2009; Yamazaki et al., 2008). In addition to CCR6, it has been demonstrated to be specific for effector/memory nTregs, which associated with increased levels of IL-10 secretions (Guo et al., 2008; Kleinewietfeld et al., 2005).

Indeed, chemokine ligand (CCL22), which can be produced by tumour cells and tumour-associated macrophages, and mediates the specific recruitment chemokine receptor CCR4 Tregs into the tumour (Gobert et al., 2009; Klarquist et al., 2016). Moreover, it suggested that the reason for the relative increase in Treg levels in cancer patients with renal cell carcinoma (RCC) could be associated with the upregulation of antiapoptotic genes and downregulation of pro-apoptotic genes (Jeron et al., 2009). However, it is worth to mention that there is a better understanding of antigen recognition and interactions by CTL

or Th cells, of note, the requirement for antigen specificity for Tregs and their suppressive actions and mechanisms remains largely elusive. In addition, tumour-associated antigens (TAA) that expressed by tumour cells may have a critical role in the recruitment, maintenance and expansion of Tregs at the sites of tumour and this may indicate that Tregs to function as anti-tumour immune response (Curiel, 2008; Wang & Wang, 2007). Therefore, it is important to determine the TAA that recognised by Tregs, it has been shown in some mouse model studies that a superior suppressive activity of antigen-specific Tregs over non-antigen-specific Tregs (Samy, Parker, Sharp, & Tung, 2005). Previous studies showed that GARP, LAP and CD137 (4-1BB) were defined as Treg-cell activation markers following polyspecific Treg-cell activation (Schoenbrunn et al., 2012; Tran, Andersson, Hardwick, et al., 2009; Tran, Andersson, Wang, et al., 2009). In this regard, it has been demonstrated that GARP and LAP can be used as pure alloantigen-specific Treg cells. Same study demonstrated that naïve and activated Treg cells do not express CD40 ligand (CD154) (Noyan et al., 2014). Moreover, there are some ligands that expressed tumour infiltrating lymphocytes and recognized by CD4<sup>+</sup> Tregs of cancer patients. Such these ligands LAGE1 (a homolog of NY-ESO-1) and ARTC1 were previously identified as ligands for tumour-specific CD4<sup>+</sup> T cells that share the common characteristics of nTregs (Wang et al., 2004; Wang, Peng, Guo, Shevach, & Wang, 2005). It may indicate that generation of CD4<sup>+</sup> T-cell clones that have the same antigen specificity but with helper or suppressor functions.

Despite the growing understanding of how Tregs are increased in the cancer setting, the exact mechanism involved in elevating their levels in cancer patients remains unclear. Investigation of whether the increased numbers of Tregs are nTregs or iTregs would provide greater insights into potential mechanisms of cancer evasion. Additionally, exploring which of these mechanisms may have a role in a specific cancer setting is of particular interest as some of the processes involved could be amenable to therapeutic intervention.

## 7. Suppression mechanisms of Tregs

Tregs can inhibit various type of immune cells including B cells, natural killer cells (NK) cells, natural killer T (NKT) cells, monocytes, dendritic cells (DCs) as well as CD4<sup>+</sup>, and CD8<sup>+</sup> T cells (A. Schmidt, Oberle, & Krammer, 2012). The exact suppression mechanism of Tregs is still incompletely understood, however, there are different mechanisms of Treg cells mediated its suppression activities that have been described; 1) suppression via co-inhibitory receptors (this review will only discuss this suppression mechanism in subsequent sections), 2) suppression via cytokines secretion including; TGF- $\beta$ , IL-10 and IL-35, and there are other mechanisms that have been described in more details elsewhere (A. Schmidt et al., 2012; Zhao et al., 2017).

### 1. Treg-mediated suppression via co-inhibitory receptors

#### a. CTLA-4

Tregs cells have the ability to regulate APCs via CTLA-4. CTLA-4 or CD152 is constitutively expressed on activated Tregs cells and function to bind with high affinity with B7 (CD80 and CD86) rather than CD28 resulting in reduction for co-stimulatory capacity of APCs (Wing, Yamaguchi, & Sakaguchi, 2011). Moreover, expression of CTLA-4 on Tregs can control their functions. A study demonstrated that the presence of CTLA-4 expression on Tregs lead to prevention of inflammatory tissue attack in arthritis (Klocke, Holmdahl, & Wing, 2017). It reported that deletion of CTLA-4 led to Tregs growth but with low suppressive activity (Peggs, Quezada, Chambers, Korman, & Allison, 2009; E. M. Schmidt et al., 2009). A study on mice model demonstrated that inhibition of CTLA-4 by using CTLA-4 specific blocking antibody rapidly induce Tregs proliferation (A. L. Tang et al., 2008). On the other hand, blocking B7 molecules on target cells using CTLA-4-Ig resulted reduction in level of memory Tregs (with robust suppressive function) (Holt, Punkosdy, Glass, & Shevach, 2017; Mohr et al., 2018). Taken together, deletion of CTLA-4 increases Tregs level whereas number of Tregs decrease as a result of loss of CD28. Moreover, monoclonal antibodies targeting the expression CTLA-4 have been shown a potent immunotherapy for some tumours (Eggermont et al., 2016; Robert et al., 2019; Robert et al., 2015).

#### b. PD-1

Apart from CTLA-4, an inhibitory receptor PD-1 (CD279) is also expressed on activated B cells, CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the periphery (Nishimura & Honjo, 2001). It shares similarity with CTLA-4 a member of CD28 family which provides a negative signal when it binds with its cognate ligands (PD-L1 and PD-L2) (Keir, Butte, Freeman, & Sharpe, 2008). A study demonstrated that positive Tregs for PD-1 displays strong suppression activity during viral infections (Ha et al., 2016). In addition, it has been shown that the inhibition of PD-1 contributed to enhancement of Tregs and production IL-10 (Woods, Ramakrishnan, Sodr , Berglund, & Weber, 2017). Combination of PD-1 with cytokine TGF- $\beta$  led to the downregulation of TGF- $\beta$ -mediated signals and thus induced the differentiation of naïve T cells to iTregs (Francisco et al., 2009). Nowadays, extensive research is going on on targeting PD-1 as a new immunotherapy besides CTLA-4 (Alsaab et al., 2017; Hodi et al., 2016; Postow et al.,

2015; Routy et al., 2018), however, a number of patients failed to respond to anti-PD-1 immunotherapy (Bu, Mahoney, & Freeman, 2016).

c. CD39 and CD73

CD39 and CD73 (also named as ectonucleotidases) have been suggested as functional markers and highly expressed on Tregs cells. Both molecules play critical roles in conversion of ATP-induced proinflammatory to adenosine-induced anti-inflammatory state (Cekic & Linden, 2016). In addition, Tregs can suppress the proliferation of effector T cells via two mechanisms a) exosomes (containing both CD39 and CD73) and b) production of adenosine by CD39 and CD73 that expressed on Tregs cells in which bind with the adenosine receptors (A2A) on effector T cells, as a result, induce the intracellular cyclic AMP (cAMP) levels that inhibit effector T cells function (Gruenbacher et al., 2016). For more details, readers are advised to look at these references (Leonard et al., 2017; Zhao et al., 2017).

d. T cell Ig and ITIM domain (TIGIT)

TIGIT is defined as a new immune checkpoint, and expressed on natural killer cells (NK), nTregs and Tr1 (Anderson, Joller, & Kuchroo, 2016; Lozano, Dominguez-Villar, Kuchroo, & Hafler, 2012). TIGIT binds with cognate ligands CD155 and CD112 on the APCs and tumour cells (Joller et al., 2014; Kurtulus et al., 2015). Studies have demonstrated that TIGIT/CD226 has a superior immunosuppressive activity in vitro function to inhibit T cells (Joller et al., 2014; Manieri, Chiang, & Grogan, 2017).

## 8. Conclusion

Tregs are heterogenous cells in phenotype, classification, function and their suppression pathways in cancers. Large evidence indicates that Treg cells may be responsible for inhibiting host T-cell activity against tumour-associated antigens and decrease the efficacy of immunotherapy approaches. The current review attempt to enhance our knowledge of Tregs cells, definition, phenotype, classification and suppression functions based on recent studies. Understanding the exact mechanisms of Treg cells function in tumour microenvironment of cancer patients may support the future studies in developing different immunotherapeutic strategy for cancer patients.

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