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Regulation of TGF β in the immune system: An emerging role for integrins and dendritic cells

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ABSTRACT

Regulation of an immune response requires complex crosstalk between cells of the innate and adaptive immune systems, *via* both cell-cell contact and secretion of cytokines. An important cytokine with a broad regulatory role in the immune system is transforming growth factor- β (TGF- β). TGF- β is produced by and has effects on many different cells of the immune system, and plays fundamental roles in the regulation of immune responses during homeostasis, infection and disease. Although many cells can produce TGF β , it is always produced as an inactive complex that must be activated to bind to the TGF β receptor complex and promote downstream signalling. Thus, regulation of TGF β activation is a crucial step in controlling TGF β function. This review will discuss how TGF β controls diverse immune responses and how TGF β function is regulated, with a focus on recent work highlighting a critical role for the integrin $\alpha\nu\beta$ 8 expressed by dendritic cells in activating TGF β .

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TGF β structure and signalling

TGF β is the prototypical member of the TGF β family of cytokines, which consists of 33 members and includes the activins, inhibins, bone morphogenetic proteins (BMPs) and growth and differentiation factors (GDFs) (Derynck and Miyazono 2008). There are three isoforms of TGF β (TGF β 1, 2 and 3), all of which have overlapping but non-redundant functions (Annes et al. 2003). Importantly, all three TGF β isoforms are produced as inactive complexes, which must be activated to bind to their receptors. Thus, TGF β genes encode a single product consisting of an N-terminal propeptide called latency associated peptide (LAP) and a C-terminal active cytokine moiety (herein referred to as active TGF β). After translation, the LAP-active TGF β product dimerises *via* disulphide bond formation. LAP and active TGF β are then cleaved from each other

in the Golgi by the enzyme furin, but remain non-covalently associated in a conformation that blocks active TGF β binding to its receptor (Fig. 1) (Annes et al. 2003). Recent structural data show that LAP forms the arms of a 'straightjacket' that wrap around active TGF β to mask the receptor binding sites and keep the complex in an inactive form (Shi et al. 2011). Thus, TGF β can only signal *via* the TGF β receptor and trigger biological effects once the complex is activated.

The TGF β receptor complex consists of two receptor subunits, TGF β receptor (TGF β R) I and II. Initial engagement of active TGF β with TGF β RII causes a conformational change in the receptor which facilitates dimerisation with TGF β RI (Fig. 1). A dimer of TGF β RII and TGF β RI forms a tetrameric complex bound to the active TGF β dimer, to initiate downstream signalling pathways (Kang et al. 2009).

Classically, TGF β receptor signalling occurs by activating the Smad-dependent intracellular signalling pathway. Both TGF β RI and II are serine-threonine kinases, and upon formation of the active TGF β -TGF β R complex, the cytoplasmic domain of TGF β RII phosphorylates the cytoplasmic domain of TGF β RI, which recruits either Smad2 or Smad3 to TGF β RI. Smad2/3 is then phosphorylated by TGF β RI, and a homodimer binds to Smad4 before shuttling to the nucleus where it acts as a transcription factor to regulate gene transcription (Fig. 1) (Shi and Massagué 2003). The interaction of Smad4 with Smad2/3 can be blocked by the inhibitory Smad, Smad7, which can act as an important negative regulator of TGF β signalling (Monteleone et al. 2008). However, TGF β can also



Abbreviations: TGF β , transforming growth factor β ; BMP, bone morphogenetic protein; GDF, growth and differentiation factor; TGF β R, TGF β receptor; LAP, latency-associated peptide; Treg, regulatory T-cell; nTreg, natural regulatory T-cell; iTreg, induced regulatory T-cell; EAE, experimental autoimmune encephalitis; DC, dendritic cell; LC, Langerhans cells; AHR, airway hyper-responsiveness.

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Fig. 1. Activated TGFβ signals *via* Smad-dependent and independent pathways. Once the latent complex of TGFβ is activated, active TGFβ binds to TGFβRII, resulting in recruitment of a TGFβRI. Phosphorylation of TGFβRI by TGFβRII allows either SMAD2 or SMAD3 to bind. TGFβRI phosphorylates SMAD2/3, which dissociates from the receptor complex and binds SMAD4. SMAD7 is able to disrupt SMAD2/3 attachment to the receptor complex. SMAD4 in complex with SMAD2/3 translocates to the nucleus where they decorate SMAD-binding elements of gene promoter regions. Constitutive dephosphorylation of SMADs results in their export from the nucleus, ensuring tight regulation of TGFβ signalling. The complex is able to recruit coactivators or corepressors to either up- or down-regulate expression of a wide range of genes, dependent on the cell type. TGFβ can also signal through poorly understood SMAD-independent pathways including MAP kinase, PI3 kinase, Wnt and other pathways.

propagate signals *via* a number of Smad-independent routes including the MAP kinase, PI3-Kinase and Wnt pathways (Guo and Wang 2009). Thus, as TGF β can regulate numerous intracellular signalling pathways and many cells express TGF β Rs, this pleiotropic cytokine needs to be carefully regulated. As will be discussed, a key regulatory step in the TGF β signalling pathway is at the level of cytokine activation.

Regulation of immunity by TGFβ

Although TGF β can have diverse effects on multiple cell types, the cytokine plays a non-redundant, crucial role in regulating immunity. TGF β is produced by, and has diverse functional effects on, many cells of the immune system. The most highly expressed isoform of TGF β in the immune system is TGF β 1, and mice lacking this isoform die early in life from multi-organ inflammation (Shull et al. 1992; Kulkarni et al. 1993). Thus, TGF β has classically been proposed to have important anti-inflammatory effects in the immune system. However, it is clear that TGF β can have both proand anti-inflammatory effects depending on the context in which it is acting. Although it should be noted that virtually all cells of the immune system can produce and respond to TGF β to some extent (see Li et al. 2006a for detailed review), below we highlight the important roles of TGF β in regulating T-cells and dendritic cells, the key players in driving the adaptive immune response.

TGFβ regulation of T-cells

CD4+ T-cells

A major focus has been placed on the role of TGF β in regulation of CD4+ T-cell responses. Multi-organ inflammation observed in TGF β 1–/– mice is completely rescued if mice are crossed to a MHCII knockout background, highlighting a crucial role for TGF β in regulating pathological CD4+ T-cell responses (Letterio et al. 1996). Additionally, early *in vitro* evidence suggested that TGF β can directly downregulate both Th1 and Th2 cell differentiation by suppressing T-bet and GATA-3 expression respectively (Li et al. 2006a). More recently, generation of mice lacking TGFβRII specifically on T-cells has re-inforced the importance of TGFβ in regulating T-cell responses *in vivo*, as mice develop multi-organ inflammation similar to that seen in TGFβ1–/– mice (Li et al. 2006b; Marie et al. 2006). Disease development correlated with an enhanced Th1 response; however, in the absence of the Th1 master transcription factor T-bet, mice lacking TGFβRII on T-cells developed multi-organ inflammation associated with an enhanced Th2 response (Li et al. 2006b). Together, the data support a key role for TGFβ in attenuating both Th1 and Th2 CD4+ T-cell activation to maintain immune homeostasis.

In addition to the direct effects on effector Th1 and Th2 responses, TGF β can promote immunosuppression *via* direct induction of regulatory T-cells (Tregs), a key CD4+ T-cell subset involved in the suppression of self-harmful immune responses. Tregs are either produced in the thymus during T-cell development (natural or nTregs) or induced from naive CD4+ T-cells in the periphery (so-called induced or iTregs) and are marked by expression of the transcription factor Foxp3 (Sakaguchi and Powrie 2007). Although earlier evidence suggested that TGF β played no role in the development of nTreg in the thymus (Marie et al. 2005, 2006; Li et al. 2006b), it now appears that TGF β is involved in the early differentiation of T-cells to nTregs in the thymus. Thus, neonatal mice lacking TGF β RI on T-cells show reduced levels of thymic Foxp3+ Tregs, but these are expanded by enhanced IL-2 production in the first week of life (Liu et al. 2008).

However, it is clear that TGF β plays a fundamental role in the induction of iTregs in the periphery. In combination with IL-2 *in vitro*, TGF β directly promotes expression of Foxp3 in CD4+ T-cells, converting them to a regulatory phenotype (Chen et al. 2003). Both TGF β 1–/– mice and mice lacking TGF β RII on T-cells show reduced Foxp3+ Treg numbers in the periphery (Marie et al. 2005, 2006; Li et al. 2006b), suggesting a key role for TGF β in induction/maintenance of Tregs *in vivo*. Induction of iTregs is particularly

prevalent in the intestine, where they are required to prevent reactions against commensal bacteria and innocuous food antigens. In addition to induction and maintenance of Foxp3 expression, TGF β has also been shown to be important in the functional ability of Tregs to suppress immune responses (Marie et al. 2005; Fahlén et al. 2005).

In contrast to the negative regulatory role in CD4+ T-cell responses, TGFB has been shown to be a key cytokine in the differentiation of pro-inflammatory Th17 cells in mice and humans (Littman and Rudensky 2010). In combination with the inflammatory cytokines IL-6, IL-1B IL-21 and/or IL-23, TGFB promotes expression of the master transcription factor of Th17 cells, RORyt (Korn et al. 2007). Although recent work has suggested that Th17 cells can differentiate into potent effector cells in the absence of TGFB (Ghoreschi et al. 2010), many studies have demonstrated an important role for TGF β in development of Th17 cells *in vivo*. For example, blockade of TGFβ signalling in T-cells (*via* expression of a dominant-negative TGFBRII) or local administration of anti-TGFβ blocking antibodies prevents the development of Th17 cells and protects mice from disease during experimental autoimmune encephalitis (EAE) (Veldhoen et al. 2006). Similarly, mice lacking TGF β production in T-cells show reduced differentiation of Th17 cells and protection from EAE (Li et al. 2007). Indeed, it appears that Th17 cells themselves are an important source of TGFβ that acts in an autocrine manner to maintain Th17 cells in vivo (Gutcher et al. 2011).

In addition to promoting Th17 cell responses recent data suggest that, in combination with IL-4, TGF β can drive differentiation of a novel CD4+ effector T-cell subset called Th9 cells, which are characterised by expression of IL-9 (Dardalhon et al. 2008; Veldhoen et al. 2008). In the presence of IL-4, TGF β induces expression of the transcription factor PU.1, which can directly drive expression of IL-9 (Chang et al. 2010). Th9 cells have effector properties and are involved in responses to helminth infection and induction of tissue inflammation (Dardalhon et al. 2008; Veldhoen et al. 2008). Thus, it is clear that TGF β can both inhibit and promote differentiation of effector CD4+ Th cell subsets.

Interestingly, the levels of active TGF β present during an immune response may be important in determining whether TGF β promotes or inhibits T-cell responses. At low concentrations, TGF β synergises with pro-inflammatory cytokines to promote Th17 differentiation by inducing ROR γ t expression (Zhou et al. 2008). However, at higher concentrations TGF β promotes Foxp3 induction and iTreg formation (Zhou et al. 2008). Thus, although the significance *in vivo* remains to be determined, these data suggest that alterations in local active TGF β concentrations may have profound effects on whether CD4+ T-cells are pushed towards a suppressive or effector phenotype.

Other T-cell subsets

In addition to regulating CD4+ T-cell subsets, TGF β plays important roles in controlling other T-cell compartments. Thus, TGF β signalling is involved in CD8+ T-cell development in the thymus, and restrains CD8+ T-cell activation in the periphery of mice (Li et al. 2006b; Marie et al. 2006). Also, TGF β limits the expansion of CD8+ T-cells during *Listeria monocytogenes* infection by inducing apoptosis during clonal expansion (Sanjabi et al. 2009). TGF β signalling is also important for NKT cell homeostasis and is required for the thymic development of canonical CD1d-restricted NKT cells, sometimes referred to as invariant NKT (iNKT) cells (Li et al. 2006b; Marie et al. 2006; Doisne et al. 2009; Havenar-Daughton et al. 2012). Conversely, TGF β signalling is required for the suppression of pathogenic effector function in an NK1.1⁺ T-cell subset (Marie et al. 2006). Finally, TGF β has recently been shown to be critical in the formation of CD8 $\alpha\alpha$ + intraepithelial lymphocytes in the intestine, which are proposed to be important in regulation of mucosal immune responses (Konkel et al. 2011).

Taken together, it is clear that TGF β plays essential roles in the regulation of T-cell development and function, acting to tune immune responses by promoting or suppressing different subsets during homeostasis and infection.

TGF β regulation of dendritic cells

The action of TGF β on dendritic cells (DCs) can both promote and inhibit DC-mediated immune responses. TGF β can downregulate the antigen-presenting function and expression of co-stimulatory molecules by DCs *in vitro* (Strobl and Knapp 1999). Mice expressing a dominant negative version of TGF β RII under the control of the CD11c promoter do not show any differences in DCs during homeostasis (Laouar et al. 2005) but are more prone to the development of EAE due to enhanced Th1 and Th17 responses (Laouar et al. 2008). Thus, TGF β appears to be important in regulating the ability of DCs to control self-reactive T-cells during models of autoimmunity. However, the exact mechanisms by which TGF β regulate DC-mediated responses *in vivo* require further exploration.

Although TGF β does not appear to be important in the development and homeostasis of conventional DCs, TGFB plays an important role in the homeostasis of Langerhans cells (LCs), a unique subset of DCs found in the epidermis of the skin (Kaplan 2010). TGF β 1–/– mice completely lack LCs (Borkowski et al. 1996), suggesting an important role for TGF β in the development/maintenance of LCs in the skin. Indeed, subsequent work has shown that TGFB1 production by LCs acting in an autocrine manner is important for LC homeostasis, as expression of a dominant negative TGFBRII in LCs (Kaplan et al. 2007) or deletion of the TGFBRI specifically in LCs (Zahner et al. 2011) results in very low LC numbers in mice. Ablation of TGF^β signalling in total skin DCs (by crossing mice expressing a conditional allele of the TGFBRI receptor to mice expressing CD11c-Cre) confirmed that TGFβ is required to maintain LCs in skin (Kel et al. 2010). The authors showed that TGFBRI deletion in skin DCs was incomplete in neonatal mice, but increased in the first week of life and correlated with a reduction in LC numbers (Kel et al. 2010). Lack of TGFB signalling in skin DCs also enhanced LC migration and maturation in steady state (Kel et al. 2010). Together, the data suggest that TGF β production and signalling are crucial for the development, maintenance and function of LCs in vivo.

TGF β production in the immune system

Many different immune cell types are capable of producing TGFB (Li et al. 2006a). Studies to investigate the important cellular sources of TGFβ *in vivo* have highlighted an important role for Tcell produced TGFB. Thus, conditional deletion of TGFB1 expression in CD4+ and CD8+ T-cells (by crossing mice carrying a conditional floxed allele of TGFβ1 with mice expressing CD4-Cre) resulted in age-related autoimmune disease, characterised by T-cell activation and colitis (Li et al. 2007). This phenotype is similar to that seen in mice expressing a dominant negative TGFBRII in T-cells (Gorelik and Flavell 2000) suggesting that TGFB acts upon T-cells in an autocrine fashion. However, as the autoimmune phenotype observed is not as severe as that in either TGF $\beta 1$ –/– mice or mice completely lacking TGFBRII expression in T-cells, this suggests that there are other important cellular sources of TGF β that act to control T-cells. Indeed, to date, TGFB knockout in any one single cell type has not recapitulated the severe phenotype observed in TGF β 1–/– mice and mice lacking TGF β RII in T-cells, strongly suggesting that there are multiple important cellular sources of TGF β production *in vivo*. The autocrine function of TGF β 1 in both T cells

(Li et al. 2007; Gutcher et al. 2011) and LC (Kaplan et al. 2007) suggests that TGF β often acts over short distances on local cells, which may be an important mechanism of spatially restricting the actions of this potent, multi-functional cytokine.

Activation of TGF $\!\beta$ in the immune system- a key role for integrins

As TGF β is secreted as an inactive complex, regulation of TGF β activation is crucial in controlling TGF β function. Early data showed that the latent complex of TGF β could be activated by high temperature, acidic pH and various proteases *in vitro* (Annes et al. 2003). However, the importance of these mechanisms *in vivo* remain to be determined. Recently, a fundamental role for members of the av integrin family in activating TGFb has been identified.

Integrins are a family of transmembrane receptors composed of an α and β subunit, with 24 different integrins expressed in mammals (Humphries et al. 2004). Integrins are important adhesion and signalling receptors, mediating both cell–cell and cell-extracellular matrix adhesion and conveying bi-directional signals across the plasma membrane (Hynes 2002). Several integrins are also capable of binding to a classical tri-amino acid binding motif, Arg-Gly-Asp (RGD) present in the LAP region of TGF β 1 and TGF β 3. *In vitro* binding studies have suggested that members of the integrin α v family (α v β 1, α v β 3, α v β 5, α v β 8) and integrin α 8 β 1 can interact with the RGD site of LAP in latent TGF β 1 and 3 (Worthington et al. 2011a). Although there is no evidence that either α v β 1 or α 8 β 1 binding to latent TGF β can result in activation, strong data now exist suggesting a critical role for other α v integrins in regulating TGF β activity and function during health and disease.

Seminal work by John Munger's group first proposed a fundamental role for integrins in activating TGFB1 to control immune homeostasis. Knock-in mice were generated that expressed latent TGFB1, but with a point mutation in the RGD integrin binding site to RGE. Thus, mice expressed latent TGFB1 but in a form that could not be bound to or activated by integrins (TGF $\beta 1^{RGD \rightarrow RGE}$ mice) (Yang et al. 2007). Despite expressing similar levels of latent TGFβ1 to control animals, TGF β 1^{RGD \rightarrow RGE mice showed a remarkably simi-} lar phenotype to mice completely lacking TGF^β1 production, dying from multi-organ inflammation early in life (Yang et al. 2007). Subsequent work has shown that integrins $\alpha v\beta 6$ and $\alpha v\beta 8$ are the key activators of TGF β 1 in the steady state immune system, as a combined lack of function of these integrins recapitulates the phenotype seen in TGF β 1-/- and TGF β 1^{RGD \rightarrow RGE mice (Aluwihare et al.} 2009). Thus, integrins $\alpha v\beta 6$ and $\alpha v\beta 8$ play a non-redundant role in the activation of TGF β 1 in the immune system which is required to prevent self-harmful immune responses.

Integrin $\alpha v\beta 8$ -mediated TGF β activation in the immune system: a crucial pathway in the regulation of immune homeostasis

After establishing a crucial role for integrins $\alpha\nu\beta6$ and $\alpha\nu\beta8$ in activating TGF β *in vivo* to maintain immune homeostasis, an important question became where are these integrins expressed and how does activation of TGF β regulate immunity to prevent inflammation? Expression of integrin $\alpha\nu\beta6$ is normally restricted to epithelial cells (Busk et al. 1992), suggesting an important role for epithelial cell-activated TGF β in regulating immune cells to maintain homeostasis. Expression of integrin $\alpha\nu\beta8$ is more wide-spread, with expression seen in locations such as cells of the central nervous system, vascular epithelial cells and astrocytes in the brain (Milner et al. 1997; Nishimura et al. 1998; Zhu et al. 2002; Cambier et al. 2005), mesangial cells in the kidney (Khan et al. 2011) and fibroblasts and epithelial cells in the airway (Araya et al. 2006). Interestingly, integrin $\alpha v\beta 8$ is expressed by cells of the immune system, most prominently in CD4+ T-cells and in DCs (Travis et al. 2007).

To reveal potential function of integrin-mediated TGF β activation by cells of the immune system, conditional KO mice were generated by crossing mice expressing a conditional floxed β 8 integrin allele (Proctor et al. 2005) with mice expressing Vav1-Cre to delete integrin $\alpha\nu\beta$ 8 in all leukocytes. As the β 8 subunit only forms an integrin heterodimer with $\alpha\nu$, this approach allowed specific deletion of integrin $\alpha\nu\beta$ 8. Such mice developed an agerelated wasting disorder associated with T-cell activation and aberrant T-cell associated antibody production and by ~6 months of age developed severe colitis (Travis et al. 2007). Similar results were observed in mice lacking all $\alpha\nu$ integrins in leukocytes (Lacy-Hulbert et al. 2007). Thus, expression of integrin $\alpha\nu\beta$ 8 by leukocytes is key to maintaining T-cell homeostasis and preventing inflammation of the intestine.

Additional work determined the cell type of the immune system that expressed functionally important integrin $\alpha\nu\beta8$. Lack of integrin $\alpha\nu\beta8$ expression in T-cells (using CD4-Cre-mediated deletion of a conditional $\beta8$ allele) caused no phenotype in mice (Travis et al. 2007). However, when integrin $\alpha\nu\beta8$ was deleted in CD11c+ cells, which are predominantly DCs (*via* expression of CD11c-Cre), mice developed an identical wasting and inflammatory disorder to mice lacking the integrin on all leukocytes (Travis et al. 2007). The phenotype was correlated with a decreased ability of DCs to activate TGF β , and also a reduced level of Foxp3+ Tregs in the large intestine (Travis et al. 2007). Similar results were obtained when all $\alpha\nu$ integrins were deleted from myeloid cells (using LysM-Cre) (Lacy-Hulbert et al. 2007). Thus, integrin $\alpha\nu\beta8$ -mediated TGF β activation by CD11c+ DCs is a crucial pathway in the maintenance of immune homeostasis in the intestine.

A key role for integrin $\alpha \nu \beta 8$ -mediated TGF β activation by specialised DCs of the intestine

The intestine is a site of high immune load, given the trillions of microorganisms that form the microflora, and the diverse array of dietary antigens. Specialised immune pathways therefore exist in the intestine to promote tolerance in the face of the heavy antigenic load, with an important role for TGFB (Maloy and Powrie 2011). Interestingly, a recently identified tolerogenic subset of DCs in the intestine, marked by expression of CD103, are specialised to induce Foxp3+ Tregs. This enhanced Treg induction is linked to their enhanced ability to produce retinoic acid, a metabolite of Vitamin A, and is TGFβ-dependent (Coombes et al. 2007; Sun et al. 2007). However, how TGF β was activated in the intestine to drive Treg induction was not understood. We have recently shown that CD103+ intestinal DCs, as well as producing more retinoic acid, are specialised to activate TGFB which correlates with high expression of integrin $\alpha v\beta 8$ (Worthington et al. 2011b). Indeed, enhanced TGFβ activation by CD103+ gut DCs promotes Foxp3+ Treg induction even when the actions of retinoic acid are blocked, and loss of integrin $\alpha v\beta 8$ expression by CD103+ intestinal DCs ablates their enhanced ability to activate TGFB and induce Foxp3+ Tregs (Worthington et al. 2011b; Païdassi et al. 2011). Thus, integrin $\alpha\nu\beta$ 8-mediated TGF β activation is upregulated by specialised DC subsets in the intestine, and this pathway is crucial in maintaining immune homeostasis in the gut.

Interestingly, although mice lacking the TGF β -activating integrin $\alpha\nu\beta$ 8 on immune cells develop autoimmunity, this is not as severe as mice that lack global function of integrins $\alpha\nu\beta$ 6 and $\alpha\nu\beta$ 8 (Aluwihare et al. 2009). Global integrin $\alpha\nu\beta$ 6–/– mice develop only mild inflammation of the lung and skin (Huang et al. 1996). Crossing these mice with mice lacking integrin $\alpha\nu\beta$ 8 on leukocytes, although speeding the onset of colitis compared to mice lacking



Fig. 2. Integrin $\alpha\nu\beta$ 8-mediated TGF β activation is mediated by, and acts upon DCs to regulate pro- and anti-inflammatory immune responses. Latent TGF β binds integrin $\alpha\nu\beta$ 8 integrin νia an RGD motif in the LAP region of the latent complex, resulting in the release of active TGF β which can orchestrate pro- or anti-inflammatory responses. Pro-inflammatory immunity is stimulated when (1) Th17 cells are induced from naive CD4+ T-cells DC-mediated TGF β activation νia integrin $\alpha\nu\beta$ 8, which can result in diseases such as EAE and AHR and (2) TGF β is activated by lung fibroblast-expressed integrin $\alpha\nu\beta$ 8, with TGF β acting in an autocrine manner to stimulate chemokine secretion and DC recruitment to the lung. Anti-inflammatory immunity is stimulated when TGF β activated by DC-expressed integrin $\alpha\nu\beta$ 8: (3) induces Foxp3+ Tregs and (4) directly inhibits T-cell activation.

integrin $\alpha\nu\beta$ 8 on leukocytes, does not result in rapid multi-organ inflammation observed in the total absence of integrins $\alpha\nu\beta$ 6 and $\alpha\nu\beta$ 8 (Dean Sheppard and Mark Travis, unpublished data). Thus, expression of integrin $\alpha\nu\beta$ 8 on non-leukocyte cells appears to be important in the maintenance of immune homeostasis. Data does suggest that integrin $\alpha\nu\beta$ 8 can activate TGF β in a number of nonimmune cell types including airway fibroblasts and epithelial cells (Mu et al. 2002; Araya et al. 2006, 2007; Kitamura et al. 2011), astrocytes (Cambier et al. 2005; Hirota et al. 2011), and Muller glial cells and neurons (Arnold et al. 2012). However, the exact non-immune cell types required for integrin $\alpha\nu\beta$ 8-mediated TGF β activation to maintain immune homeostasis remain to be determined.

Integrin $\alpha v \beta 8$ -mediated TGF β activation by dendritic cells in inflammatory disorders

In addition to playing an important role in dampening immune responses to maintain homeostasis, activation of TGFB by DCs is crucial in promoting inflammation during autoimmunity. As mentioned earlier, as well as having important suppressive effects on CD4+ T-cells, TGFβ can also promote inflammatory T-cell responses by promoting differentiation of Th17 cells. It is now apparent that integrin $\alpha v\beta 8$ -mediated TGF β activation by DCs is important in the development of Th17 cells. DCs that lack integrin $\alpha v\beta 8$ expression have a reduced ability to induce Th17 cells in vitro, and this defect can be rescued by the addition of active TGF β (Melton et al. 2010). Mice lacking integrin $\alpha v \beta 8$ expression on DCs also have reduced numbers of Th17 cells in vivo, and are protected from symptoms during EAE, a model of autoimmunity known to involve Th17 cells (Melton et al. 2010). Similar results are observed in mice lacking all α v integrins on myeloid cells, and when integrin-latent TGF β interactions were blocked by administration of an RGD peptide (Acharya et al. 2010). Thus, TGF β activation by DCs can drive Th17-mediated inflammation during autoimmune disease.

TGF β activation by DCs is also important in the development of allergic disease in the lung. Mice lacking integrin $\alpha\nu\beta$ 8 on DCs display reduced airway hyper-responsiveness (AHR), a hallmark of allergic asthma, during lung antigen challenge models (Kudo et al. 2012). Reduced AHR is associated with impaired Th17 cell induction, which results in attenuated IL-17-mediated smooth muscle contraction in the airways (Kudo et al. 2012). Interestingly, lung inflammation observed during challenge was similar between control mice and mice lacking integrin $\alpha v\beta 8$ on DCs. Thus, Th17 cells induced in the lung *via* DC-mediated TGF β activation appear to play a specific function in regulating AHR during pulmonary antigen challenge rather than driving a broader inflammatory response.

In addition to driving activation of TGF β via integrin $\alpha \nu \beta 8$, it has recently been shown that DCs themselves can be influenced by $\alpha\nu\beta$ 8-mediated TGF β activation in the lung. Thus, during airway remodelling (a hallmark of lung diseases such as asthma and chronic obstructive pulmonary disease) airway fibroblasts upregulate integrin $\alpha v\beta 8$ to activate TGF β (Kitamura et al. 2011). This increased activation of TGFB promotes production of the chemokines CCL2 and CCL20 by the fibroblasts, enhancing DC migration to the lung to boost adaptive immune responses (Kitamura et al. 2011). Given the known importance of integrin $\alpha v\beta 6$ expression by lung epithelial cells in regulation of lung pathology during various pulmonary diseases (Sheppard 2004), and recent data suggesting an important role for integrin $\alpha v\beta$ 5-mediated TGF β activation by airway smooth muscle cells in mediating pathology during asthma (Tatler et al. 2011), it appears that there are multiple integrin-mediated mechanisms to regulate TGFβ activity in the lung during disease. Thus, it will be interesting to determine potential cross-talk between these related mechanisms of TGFB activation in the lung, and how this impacts on lung pathology during pulmonary disease.

Conclusions

Given its crucial and multi-functional role in controlling immunity, it is unsurprising that the action of TGF β is carefully regulated, not only at the level of cytokine production but also by activation of the latent complex. Recent data highlighting a crucial role for integrins in activating TGF β has highlighted not only a novel function for integrin family members *in vivo*, but also a key immunoregulatory mechanism that controls both tolerogenic and inflammatory immune responses. The finding that DCs are especially important in activating TGF β via integrin $\alpha v\beta 8$ has uncovered a novel pathway that allows intricate regulation of T-cell responses during health and disease (Fig. 2). However, many important questions remain. How does integrin binding to the latent complex of TGFB result in TGFB activation? Mechanisms proposed include contraction induced stretching of the latent complex after integrin engagement (for integrins $\alpha v\beta 3$, $\alpha v\beta 5$ and $\alpha v\beta 6$) and protease recruitment and cleavage of the complex (for integrin $\alpha v\beta 8$) (Worthington et al. 2011a). However, whether these mechanisms are important in vivo, and whether they are conserved between different cell types in different tissues of the body remains to be seen. Additionally, are there subsets of cells throughout the body that are specialised to activate TGFB via different integrin heterodimers, akin to the upregulation of integrin $\alpha v\beta 8$ seen in CD103+ DCs in the intestine? Furthermore, how are these pathways altered during disease? Regardless, TGFB-activating integrins are becoming potential therapeutic targets in a range of immunological disorders, to either stimulate anti-inflammatory pathways or inhibit pro-inflammatory pathways during progression of chronic disease.

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