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From the diet to the nucleus: Vitamin A and TGF- β join efforts at the mucosal interface of the intestine

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Abstract

The Vitamin A metabolites, including retinoic acid (RA), form ligands for retinoic acid-related nuclear receptors and together they play pleiotropic roles in various biological processes. Recently, we described that RA also functions as a key modulator of transforming growth factor-beta (TGF- β)-driven immune deviation, capable of suppressing the differentiation of interleukin-17 secreting T helper cells (T_H17) and conversely promoting the generation of Foxp3⁺ T regulatory (Treg) cells. This review will focus on the role of RA in the reciprocal TGF- β driven differentiation of T_H17 and Treg and on the importance of such regulatory mechanism to control a functional immune system, in particular at the mucosal interface of the intestine.

Keywords

vitamin A; gut tropism; Foxp3; T_H17; nuclear receptors

Natural and induced regulatory T cells

Burnet's clonal selection theory introduced the idea that through self/non-self discrimination, auto-reactive cells would be eliminated in order to achieve tolerance [1]. In the past 30 years however, it has become clear that autoreactive T cells also exist in normal individuals, and that some of these are beneficial and required to avoid the occurrence of autoimmune diseases [2-6]. Beneficial autoreactive T cells are actively selected for in the thymus, where they adapt to their phenotype and acquire regulatory functions. Because of this, they have been characterized as natural regulatory T cells (nTreg). In contrast to self-reactive thymocytes that are deleted during conventional negative selection, precursors of nTregs require strong self-agonist selection signals to gain their regulatory capacity to maintain self-tolerance (Fig.1). Regulatory T cells were initially described by Sakaguchi and coworkers as CD4⁺CD25⁺ T cells [7]. Later, Sakaguchi's and Rudensky's groups further defined the forkhead-winged helix transcription factor family member, Foxp3, as the crucial driver of the naturally-occurring regulatory T cell-lineage [8,9]. The observation that patients with the X-linked fatal autoimmune disease, IPEX (immune dysregulation, polyendocrinopathy, enteropathy, X-linked) syndrome, lack functional Foxp3 underscores the importance of this transcription factor as the central key player in the functional regulation of nTreg [10,11]. In mice, a similar massive and fatal lymphoproliferative disease develops spontaneously in the mouse mutant strain,

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scurfy, which also results from impaired Foxp3 function. Introduction of a functional Foxp3 transgene can correct the autoimmune phenotype, exemplifying the importance of Foxp3 and Treg cells in systemic immune-regulation and tolerance [10].

In addition to self-tolerance a functional immune system also needs to be able to tolerize non-self antigens that do not impose a threat to the individual. Such harmless non-self antigens are abundant in the intestine where numerous, often beneficial commensal bacteria colonize the colon and where digested food is continuously absorbed via the small intestine epithelium. An effective immune-regulation is a condition *sine qua non* for the gut physiology [12-15] and the importance of Treg cells to control and prevent aberrant immune responses directed towards self or non-self antigens and to establish tolerance has been demonstrated at length [16].

Although self-specific thymus-derived nTregs can suppress effector T cells with other specificity, they need to be activated first by their cognate self-antigens. In addition, since nTregs functionally differentiate in the thymus during agonist selection with self-antigens, the differentiation of nTregs in the thymus with specificity towards non-self antigens is very unlikely. Instead, a substantial number of studies have clearly demonstrated that naïve Foxp3⁻, primarily non-self-reactive, CD4⁺ T cells can be “converted” by TGF-β into induced Foxp3⁺Treg (iTreg) in response to self or non-self antigens encountered in the periphery [17-22] (Fig.1). Such iTreg are especially important to maintain tolerance towards antigens introduced post thymically via the intestine (Fig.2) [21,22]. Consistent with this, our group and those of Belkaid and Powrie have clearly demonstrated that iTregs are preferentially induced at the mucosal induction sites of mesenteric lymph node (MLN) and *lamina propria* (LP), rather than in spleen or peripheral lymph nodes (LNs), reinforcing that the intestine is a privileged site for iTreg differentiation [23-25]. In this context, Lafaille's group has further shown, using mice that lack nTregs, that peripheral *neoconverted* Foxp3⁺ iTreg cells are effective and sufficient to mediate oral tolerance, which can be defined as diminished systemic immune responses for an antigen previously contacted via the oral route [21]. Recently these findings were extended by the demonstration that mice which cannot induce iTreg are unable to mediate oral tolerance, suggesting the iTreg are not only sufficient, but also crucial for oral tolerance induction in this model [22].

Through the expression of tight junctions (TJs), epithelial-associated DCs are able to establish rigid contacts with the neighboring epithelial cells, while extending their dendrites to sample luminal antigens, including food particles and whole bacteria [26]. Upon activation, DC mature and migrate to regional LNs, where they can present processed antigens to naïve T cells. This migration requires the expression of the chemokine receptor, CCR7, by the mucosal DCs (Fig. 2). Consistent with this, CCR7⁺ but not CCR7⁻ MLN DCs, have vacuoles containing intestinal epithelial-cell derived debris, suggesting their intestinal origin [27]. Presentation of gut luminal antigens by these mucosal DCs plays crucial roles in the development of oral tolerance and in the generation of iTregs [13,28,29]. Consistent with this, CCR7^{-/-} mice are inefficient in inducing oral tolerance, likely due to the fact that antigen-carrying CCR7-deficient DCs cannot migrate to the MLN and induce T cell-dependent oral tolerance [29].

A balance between protective and suppressive immunity in the intestine

In addition to the requirement for tolerance to maintain the integrity of the barrier, the intestine, which forms the largest entry surface for pathogens, also requires the most effective immune protection, indicating that a critical but flexible immune balance has to be in place. An important molecule in this context is TGF-β, abundantly produced in the gut and crucial for both systemic and mucosal immune-regulation. Among multiple roles, TGF-β has the capacity to block T_H1 and T_H2 differentiation and to convert naïve CD4 cells into Foxp3-expressing iTreg cells [30]. Paradoxically, TGF-β also displays pro-inflammatory roles and in the presence

of pro-inflammatory cytokines such as IL-6, IL-21, IL-1 β and TNF, TGF- β promotes the conversion of naïve T cells into T_H17 [31-33]. This contrasting deviation puts TGF- β as a principal controller of immune responses and underscores a central role of this cytokine in orchestrating the pro- and anti-inflammatory nature of adaptive immunity. Nowhere else is a critical regulation of this TGF- β dependent balance of more significance than at the mucosal interface of the gut, where efficient immune protection against pathogens has to coincide with maintenance of the mucosal barrier integrity.

The reciprocal developmental pathways for the generation of pro-inflammatory effector T_H17 and immune suppressive iTreg, imposes a dichotomy in their generation in addition to their functional antagonism. Therefore, TGF- β -mediated iTreg cells and T_H17 effectors arise in a mutually exclusive fashion, depending on whether they are activated in the presence of TGF- β alone or TGF- β together with pro-inflammatory cytokines [32]. The constant exposure to luminal antigens creates a so-called “physiological chronic inflammation” in the gut, which could easily favor the TGF- β driven effector differentiation at the expense of iTreg. Nevertheless, under normal steady state conditions, immune responses in the gut are under control and iTregs tend to preferentially accumulate at this site.

We recently identified a metabolite of the nutrient vitamin A, retinoic acid, as a key regulator of TGF- β -dependent immune responses, capable of inhibiting the TGF- β /IL-6-driven induction of pro-inflammatory T_H17 cells and promoting the TGF- β -dependent peripheral differentiation of anti-inflammatory Foxp3⁺ iTregs [23] (Fig.2). Since vitamin A cannot be synthesized by the human body and it must be absorbed by the intestine from the diet, this means that this intestinal induced regulatory mechanism allows for environmental factors to modulate aberrant endogenous immune responses from the outside in. The induced immune tolerance is not necessarily irreversible and in the presence of innate danger signals RA effects might diminish or synergize with innate responses to promote or enhance protective immunity.

Retinoids and their nuclear receptors: from the diet to the nucleus

Vitamin A is initially absorbed from the diet and via the function of various enzymes, including the ubiquitous alcohol dehydrogenases (ADH) and the more restricted retinal dehydrogenases (RALDH), it is further metabolized to RA. Two isoforms of RA have been identified *in vivo* (*all-trans-RA* (*at-RA*) and *13-cis-RA*) but the 9-cis-RA isoform has never been detected *in vivo*. All RA isoforms, can bind to homo- or hetero-dimers of nuclear receptors, such as retinoic acid receptors (RARs) and retinoid X receptors (RXRs). These nuclear receptors are able to function as ligand-activated transcription factors through binding to DNA response elements (RARE) [34,35]. Depending on the type of ligand interaction as well as on the availability of co-activators (CoAs) and co-repressors (CoRs), nuclear receptor-ligand complexes are able to perform either transcriptional repression or activation of target genes [36]. Due to its role as a required heterodimer partner, RXR also influences the transcriptional control of many other nuclear receptors such as vitamin D (VDR) and thyroid hormone receptors in addition to RARs. This may have important additional repercussions in T cell differentiation, since for example, Vitamin D3, is able to regulate transcription factors, such as NF-AT, NF- κ B, and, importantly, TGF- β signaling molecules, SMADs [37-39]. Specifically, Smad3 was reported to act as a CoA for ligand-induced transactivation of VDR [39]. Finally, VD3 is also described as a potent inducer of the anti-inflammatory cytokine IL-10 [40], reinforcing the network of immune-regulation associated with retinoic acid receptors. It is noteworthy that although it was reported that RA directly enhanced LPS-induced IL-10 expression in a cell line [41], our data suggest that RA efficiently suppresses TGF- β /IL6-induced IL-10 production (Mucida, D et al., *manuscript in preparation*), which might be correlated with this uncommon characteristic of RXR/VDR heterodimer action [42]. RXRs can also interact with members of the orphan nuclear peroxisome proliferative activated receptor family, PPAR, which have important

functions in immune-regulation that are closely linked to retinoic acid-related pathways. Recently PPAR- δ was reported to play a role in the suppression of IL-17 and IFN- γ production in colitis [43].

Immune-regulatory functions of RA and TGF- β

Similar to the vitamin A metabolites and their nuclear receptors, members of the TGF- β superfamily are pleiotropic regulators of numerous biological processes. In many of their functions, TGF- β and retinoids synergize or counteract each other reflecting an intense relationship between members of these two superfamilies of gene regulators.

In addition to their crucial roles in development, TGF- β and RA are involved at almost every level of immune differentiation and function, affecting passive immunity as well as innate and adaptive immunity. Both, TGF- β and RA are actively produced by the intestinal epithelium and both play important roles in mucosal epithelial cell differentiation and in maintaining the integrity of its barrier function (Fig 2).

TGF- β signaling and nuclear hormone receptor regulation also greatly influence multiple functions of the innate immune system. An intensive crosstalk between nuclear receptors and factors of the innate immune system, for example IFN regulatory factor 3, a key transcription factor involved in the induction of antiviral genes, may lead to repression of innate immune responses and inflammation and conversely to the repression of nuclear hormone receptor signaling [44,45]. This crosstalk has been implicated in the pathogenesis of atherosclerosis and Reye's syndrome and other pathogen-associated metabolic and developmental disorders [45]. The cytosolic pattern recognition receptor, retinoic acid-inducible gene I (RIG-I), also plays important role in the cytosolic recognition of invading pathogens and the induction of the retinoic acid early inducible gene 1 (RAE-1), which forms a potent ligand for the NK receptor NKG2D, makes target cells highly sensitive to NK cell killing. TGF- β negatively regulates the antigen presentation, phagocytosis and expression of costimulatory molecules by DCs and macrophages and, similarly to RA, TGF- β can negatively regulate the production of inflammatory cytokines such as IL-1, TNF- α and IFN- γ , by antigen presenting cells and NK cells [41,46-50].

In the adaptive immune system, TGF- β and RA positively regulate secretory IgA production, important in mucosal barrier function and antibacterial protection, by actively promoting IgA-class switching [51-54]. In addition to secretory/mucosal IgA-class switching, RA also induces migration and homing of B cells to the LP of the intestine [54].

On T cells, gut specific homing receptors, including $\alpha 4\beta 7$ and CCR9, are induced by RA released during priming by MLN, LP and Peyer's patch (PP) DCs [55-58]. On the other hand, TGF- β , which is produced by many cells in the intestine including epithelial cells, macrophages, DCs and subsets of intraepithelial lymphocytes (IEL)[24,59-62], induces the integrin $\alpha E\beta 7$ (CD103). The combination of TGF- β and RA results in further upregulation of both integrins, $\alpha 4\beta 7$ and $\alpha E\beta 7$, but downmodulates the expression of CCR9 [23].

TGF- β and RA play also major roles in functional differentiation of T cells. In addition to the RA mediated reciprocal differentiation of TGF- β dependent iTreg/T_H17 cells, TGF- β was shown to be crucial for early nTreg development as well, indicating that TGF- β is important for both nTreg and iTreg differentiation [63]. Similar to TGF- β , RA also strongly inhibits the T_H1 cytokine IFN- γ synthesis, but in contrast, RA has the capacity to stimulate T_H2 differentiation under certain conditions [64]. The effect of RA on T_H2 development is dependent on the stage of T helper differentiation and RA enhances IL-4 production only when added after the initial activation of naïve CD4 cells *in vitro* [65]. When added at the time of

the initial stimulation, RA rather suppresses both T_H1 and T_H2 differentiation, indicating that RA could have divergent effects depending on the context of exposure [66,67].

Reciprocal regulation of TGF- β mediated T_H17 /Treg differentiation by RA

iTreg cells and T_H17 cells arise in a mutually exclusive way, depending on whether they are generated in the presence of TGF- β or TGF- β and the pro-inflammatory cytokine IL-6. This remarkable pro- and anti-inflammatory immune deviation is of particular relevance at the mucosal surface of the intestine and the recent finding that RA plays a key role in controlling this critical TGF- β dependent immune balance, implies an important role for RA to mediate mucosal immune regulation (Fig. 2). Consistent with this, several groups, including ours, demonstrated that iTregs were preferentially induced by mucosal DCs rather than spleen or peripheral LN DCs, reinforcing that the intestine is a privileged site for Treg induction (Fig. 2) [24,25]. Among the mucosal DCs, CD103⁺ DCs seem to be especially capable of releasing RA during priming. Coombes et al. showed that RA-production by CD103⁺ but not CD103⁻ MLN DCs, efficiently converted naive CD4⁺ T cells into Foxp3⁺ T cells in a TGF- β -dependent fashion [24]. In contrast however, Sun et al. described that whereas LP CD103⁺ DCs could induce Foxp3⁺ T cells in the absence of exogenous TGF- β , both CD103⁺ and CD103⁻ LP DC-populations were highly efficient at inducing Treg cells in the presence of exogenous TGF- β [25]. More recently, Denning et al. extended and added complexity to these findings. They reported that LP macrophages are also potent regulatory antigen presenting cells, able to convert naive T cells into Foxp3⁺Treg cells in an IL-10-, RA- and TGF- β -dependent manner. On the other hand, LP CD11b⁺CD11c⁺ DCs are pro-inflammatory, inducing IL-17-producing T cells but few Tregs. Denning et al. also reported that LP CD11b⁻CD11c⁺ cells express high levels of CD103 however they did not detect any increased ability for these CD103⁺CD11b⁻ DCs to induce Tregs [60].

In contrast to the enhanced Foxp3 induction by MLN DCs, we showed that they had less capacity to promote T_H17 differentiation as compared to their spleen counterparts. In the presence of exogenous RA, however, spleen DCs readily induce iTregs but show much reduced capacity to promote T_H17 differentiation. Conversely, the addition of RAR inhibitors impaired Treg differentiation by MLN DCs but enhanced the T_H17 induction. Using an infectious mouse model, we further showed that exogenous RA inhibits the induction of T_H17 cells *in vivo* whereas injection of RAR antagonists resulted in a decrease of Foxp3⁺Treg cells in the LP [23]. These observations clearly demonstrated the capacity of RA to directly control the reciprocal differentiation of TGF- β driven Treg and T_H17 cells.

The abundant production of TGF- β and RA in the mucosa and the ability of RA to promote TGF- β -dependent iTregs could thus directly be related to the high frequency of Foxp3-expressing Treg cells in the intestine of normal mice. Furthermore, the synergistic effect of TGF- β and RA to promote anti-inflammatory but suppress pro-inflammatory immune responses might be central for oral tolerance induction and therefore, mucosal and systemic immune regulation [23,25]. It is also of importance to note that RA mediated the differentiation of a stable TGF- β induced iTreg lineage *in vitro*, which upon transfer efficiently prevented colitis induction *in vivo* indicating that in addition to functional iTregs, RA mediates epigenetic imprinting that is critical for establishment of a stable Treg lineage *in vivo*.

The RA mediated enhanced TGF- β induced Foxp3 expression was drastically reduced in the presence of anti-IL-2 antibodies or when IL-2-deficient naive CD4 T cells were used *in vitro*. In the later case, the addition of exogenous IL-2 restored RA-effects [23]. Conversely, the RA-mediated suppression of IL-17 production by either CD4 or CD8 T cells was inhibited in the presence of anti-IL-2 antibodies or when IL-2 deficient naive T cells were used [23]. Although it was previously shown that the addition of exogenous IL-2 also promotes TGF- β

induced iTreg differentiation *in vitro*, the direct effects of RA and exogenous IL-2 appear distinct. Whereas the combination of IL-2 and TGF- β induce mostly CD103⁻Foxp3⁺ Treg cells, RA and TGF- β induced preferentially CD103⁺Foxp3⁺ cells. In addition, since RA also induces the expression of the gut homing receptors, CCR9 and α 4 β 7, in combination, RA and TGF- β readily induce iTregs with a gut-specific tissue tropism.

The observation that in the absence of DCs, RA also mediates reciprocal TGF- β dependent differentiation of naïve T cells *in vitro*, indicates that RA directly affects the transcriptional control within the T cells. We showed that indeed RA directly inhibits retinoic acid orphan receptor gamma T (ROR γ t) which is induced by TGF- β and further enhanced by IL-6 signaling and which is required for efficient T_H17 differentiation [68]. It is not known, however, whether RA antagonistic effects on IL-6 signaling extend to the recently described IL-21 pathway of T_H17 differentiation [69-71]. In addition to the negative regulation of IL-17 transcription, RA also has direct positive regulatory effects for the transcription of Foxp3.

Recently, it was shown that CD11c^{hi}CD11b^{hi} LP DCs, when stimulated by TLR5-ligand flagellin, promoted a modest differentiation of antigen-specific T_H17 and T_H1 cells and addition of RAR-antagonist LE540 inhibited this IL-17 production [72]. Unexpectedly, LPS did not induce measurable IL-17 production, but addition of a low-dose (1nM) RA induced some IL-17 production in the presence of LPS whereas a higher dose of RA (10 μ M), inhibited IL-17 production. Based on this, it was concluded that low-dose RA induces T_H17-development, whereas a high dose of RA is suppressive for IL-17 expression [72]. It is important, however, to point out here that previous work, including ours, describing the suppressive effects of RA on T_H17 development in the absence of TLR stimulation, reported RA-mediated suppression of IL-17 in all doses examined, ranging from 1nM to 100nM [23] and 10 μ M [66]. A typical dose-response curve of T_H17-suppression by RA was also observed by Kattah and coworkers using a range from 1nM to 1 μ M in human cells [73]. In addition, the release of physiological amounts of RA by mucosal DCs in T cell/DC co-cultures also inhibited IL17-production and enhanced Foxp3 induction, whereas addition of RAR-antagonist in these cultures reversed this effect [23-25]. It does seem that “physiological amounts” of RA released by gut-derived DCs suppress rather than induce IL-17 production but enhance TGF- β mediated Foxp3 induction. It is possible, however, that innate stimuli, such as TLR-signals, may induce contrasting effects of RA on DCs and/or T cells [74].

Transcriptional regulation of Foxp3 and IL-17 by RA

The transcription factors STAT5 and STAT3 are important for the transcription of Foxp3 and IL-17, respectively [68,75,76]. The enhanced expression of Foxp3 in the presence of RA suggests a potential relationship between STAT5 and RARs in a similar fashion as the cooperation between STAT3 and ROR γ t (Fig.3). It is therefore perhaps not a coincidence that ROR γ t displays strong homology with the RARs and also appears to function in the context of transcriptional activators and repressors [77]. Consistent with STAT5 and RAR cooperation they can physically interact with each other *in vivo* to promote RAR-mediated transcription (Fig. 3) [78]. Furthermore, the STAT5 consensus-binding site directly overlaps with RAR-response elements, which may lead to coordinated transcription activity rather than competition for the same site [78]. The cooperation between STAT5 and RARs results in STAT5-enhanced responsiveness of the RARs to RA induced transcription of target genes [78]. RAR and STAT5 can also bind the same repressor of transcription, silencing mediator of thyroid and retinoid receptors, SMRT, which can be released upon RA engagement. The RA mediated effects on Treg and T_H17 differentiation may thus reflect an intense communication between the STAT and RAR families of transcription factors for the differentiation of T lymphocytes [79].

In addition to direct inhibition of ROR γ t by RA, it is also possible that RA indirectly suppresses T_H17 differentiation via the enhanced transcription of Foxp3. Indeed, it was recently shown that ROR γ t and Foxp3 can be co-expressed in naive CD4⁺ T cells upon TGF- β exposure and that TGF- β -induced Foxp3 inhibits IL-17 induction by direct suppression of ROR γ t [80-82]. Consistent with additional STAT-independent effects of RA, it was shown that with *in vitro* culture of STAT3 conditional knockout CD4 T cells (in this case, IL-17 production is abolished [76]), RA still enhanced Foxp3 expression in the presence of TGF- β , suggesting that RA can induce Foxp3 independently of IL-6 signaling. Conversely, using STAT5 conditional knockout CD4 T cells (Foxp3 expression is decreased), 1 μ M RA still suppressed IL-17 production in the presence of TGF- β and IL-6. It is important to note, however, that in this study, total, but not naïve, CD4 STAT5-deficient T cells were used and exogenous IL-2 was added, which could mask the effects of RA on naïve T cell differentiation [66]. Finally, and in contrast to our observations, it was shown in that study that also in the presence of anti-IL2 antibodies, RA could still inhibit IL-17 production (although less efficiently) and enhance TGF- β induced Foxp3 expression [66]. It is possible that the high dose of RA used (1 μ M) [66] could bypass the requirement for IL-2 observed in our previous study [23].

In addition to IL-6, T_H2 cytokine IL-4, also inhibits TGF- β -mediated Foxp3 induction via IL-4 receptor activated STAT6, which directly binds to the Foxp3 promoter (Fig.3) [67]. RA on the other hand can reverse this IL-4-mediated inhibition of Foxp3 expression by engaging RAR α /RXR α heterodimers that interfere with the silencing capacity of STAT6 on Foxp3 induction [67].

Finally, the synergy between RA and TGF- β is likely controlled by the Smad signaling molecules (Fig.3) that act downstream of the TGF- β receptor, and/or with the transcription factor Runx3, which is mediating the TGF- β induced CD103 expression and which physically interacts with Smads to cooperate in TGF- β mediated signaling [83].

Concluding remarks

A functional immune system is central for efficient protection of the organism against invasive pathogens as well as invasive transformed autologous cells. Aberrant immune responses directed towards harmless non-self agents or normal endogenous cells can lead to severe immune pathology and autoimmunity. Many regulatory mechanisms are in place and in particular Foxp3⁺ regulatory T cells are indispensable to prevent excessive and self-destructive immune responses. The intestine, which forms the largest entry surface for invading pathogens but also contains a vast load of harmless antigens derived from colonizing commensal bacteria and absorbed nutrients, imposes a unique challenge for the immune system. In particular, the ability to mount a highly efficient protective immune response needs to coincide with the steady state prerequisite of tolerance towards innocuous antigens. An improper balance between inflammatory and suppressive immunity can jeopardize mucosal homeostasis and destroy the integrity of the mucosal barrier. The intake of vitamin A as part of the food absorption by the intestine and the ability of the vitamin A metabolite, retinoic acid, to drive the differentiation of induced Foxp3⁺ regulatory T cells and to suppress the differentiation of pro-inflammatory effector cells, provides an expedient mechanism to secure tolerance when protective immune responses are not advantageous. The plasticity of the system and the ability of innate and pro-inflammatory signals to override the RA controlled immune suppression provides the mucosal immune system with a key control to allow for protective immunity in the face of steady state immune tolerance.

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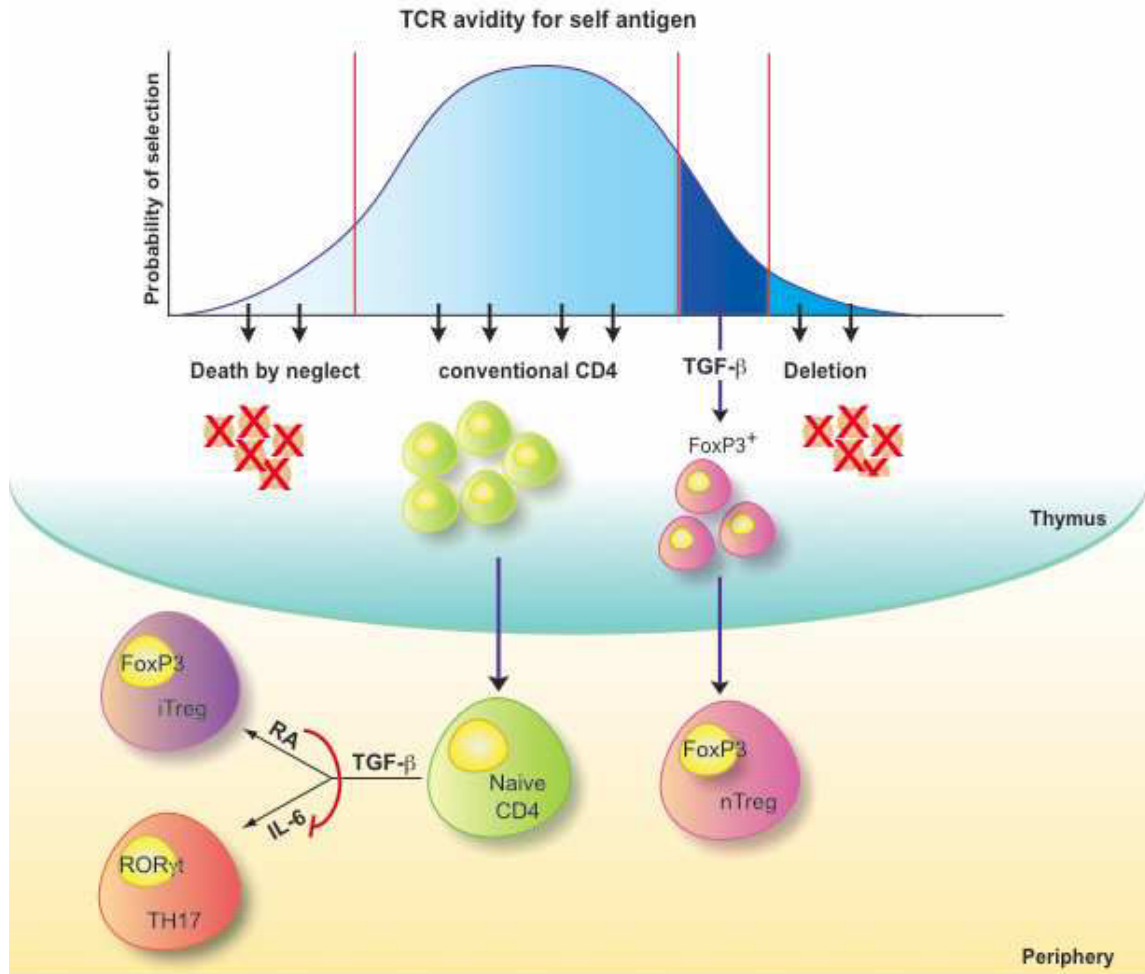


Figure1. Development of natural and induced regulatory T cells

Thymus-derived “natural” regulatory T cells (nTreg) are actively selected upon high avidity TCR/self peptide-loaded MHC interactions, express the forkhead-winged helix transcription factor family member, Foxp3, and are fundamental in the process of self-tolerance, so-called “dominant tolerance”. Conventionally selected naïve Foxp3⁻ CD4⁺ T cells can be “converted” into induced Foxp3⁺Treg (iTreg) in response to self or non-self antigens encountered post thymically in the periphery. TGF-β is required for both early nTreg development in the thymus and for peripheral induction of iTreg. TGF-β is also able to promote, in the presence of inflammatory cytokines such as IL-6, the development of pro-inflammatory T_H17 cells. The vitamin A metabolite, retinoic acid (RA), is capable of inhibit the TGF-β/IL-6–driven induction of pro-inflammatory T_H17 cells and simultaneously promote the TGF-β-dependent peripheral differentiation of anti-inflammatory Foxp3⁺ iTregs.

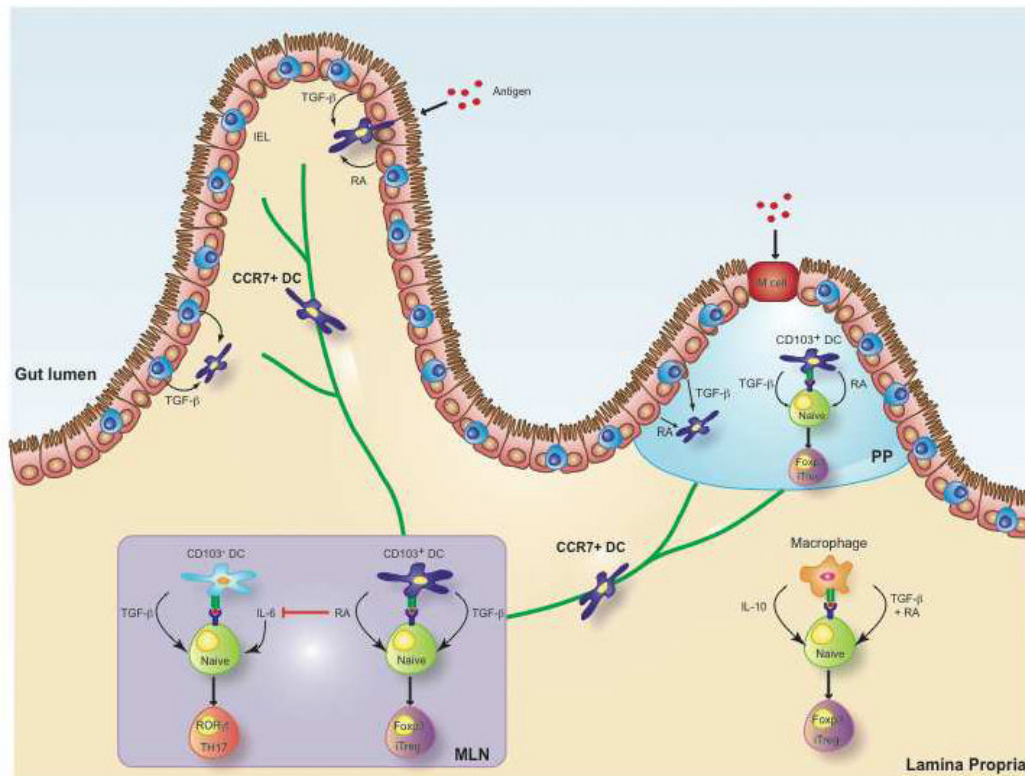


Figure 2. Role of retinoic acid in mucosal immune-regulation

Under the influence of intestinal epithelial cells and intra-epithelial lymphocytes, *lamina propria* and Peyer's patch CD103⁺ DCs acquire their ability to produce retinoic acid (RA). LP DCs that have acquired the expression of the chemokine receptor CCR7 are able to migrate through the afferent lymphatic vessels into the mesenteric lymph nodes and influence T-cell activation and differentiation during priming of naïve T cells. Gut-derived CD103⁺ DCs and LP macrophages efficiently promote the generation of Foxp3⁺iTreg in a RA-, TGF-β-, and also IL-10 (in the case of LP macrophages) dependent manner. In contrast, release of RA by these DCs suppresses the TGF-β- and IL-6- dependent differentiation of TH17 cells.

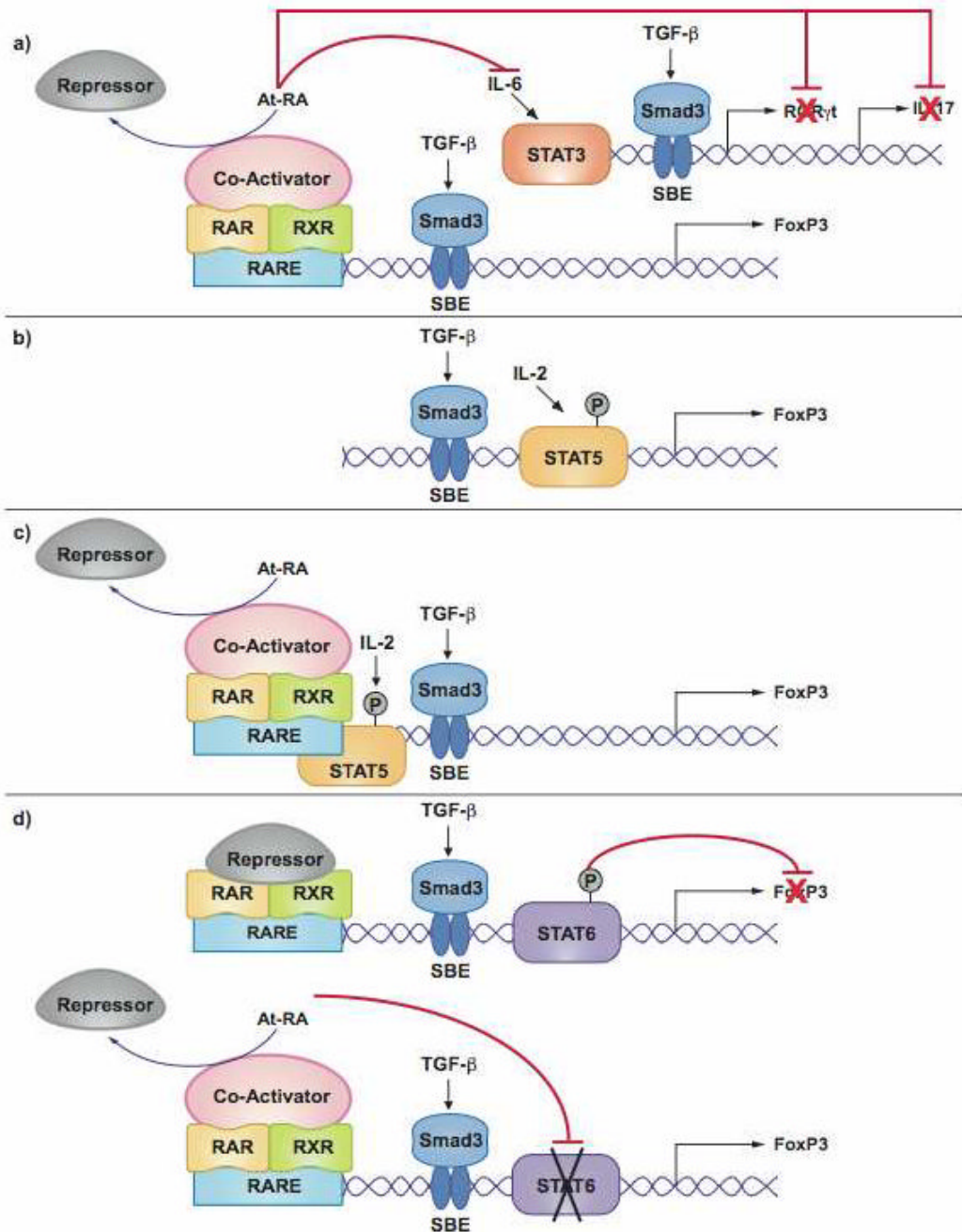


Figure 3. Transcriptional regulation of Foxp3 and ROR γ t/IL-17 mediated by RA
 Retinoic acid receptors (RARs) and retinoid X receptors (RXRs) are nuclear receptors able to function as ligand-activated transcription factors through binding to retinoic acid DNA response elements (RARE). Upon ligand (*at*-RA) binding, RAR/RXRs release co-repressors (CoRs) and recruit co-activators (CoAs). This figure shows possible mechanisms by which RA binding to its nuclear receptors can affect the induction of Foxp3 and IL-17. **A.** RA could enhance TGF- β signaling via the TGF- β signaling molecules, Smad2 and 3 and SBE (Smad-binding elements) and thus indirectly increase TGF- β -dependent Foxp3 induction. In contrast, RA may also repress different target genes in the T_H17-development pathway, including STAT3, ROR γ t and IL-17. Conversely, the suppression of these molecules could also result

in enhanced TGF- β induced Foxp3 expression. **B.** The induction of Foxp3 expression in naïve T cells requires both TGF- β signaling and IL-2-dependent STAT5 phosphorylation and binding in the Foxp3 promoter. **C.** STAT5 and RAR interaction may lead to coordinated transcription activity of target genes, therefore enhancing TGF- β -induced Foxp3 expression. **D.** IL-4 is able to inhibit TGF- β -mediated Foxp3 induction via IL-4 receptor activated STAT6, which directly binds to the Foxp3 promoter. RA binding to RAR α /RXR α heterodimers can interfere with the silencing capacity of STAT6 resulting in increased Foxp3 induction.