

# Dendritic Cells: Arbiters of Immunity and Immunological Tolerance

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Dendritic cells (DCs) link innate immune sensing of the environment to the initiation of adaptive immune responses. Given their supreme capacity to interact with and present antigen to T cells, DCs have been proposed as key mediators of immunological tolerance in the steady state. However, recent evidence suggests that the role of DCs in central and peripheral T-cell tolerance is neither obligate nor dominant. Instead, DCs appear to regulate multiple aspects of T-cell physiology including tonic antigen receptor signaling, priming of effector T-cell response, and the maintenance of regulatory T cells. These diverse contributions of DCs may reflect the significant heterogeneity and “division of labor” observed between and within distinct DC subsets. The emerging complex role of different DC subsets should form the conceptual basis of DC-based therapeutic approaches toward induction of tolerance or immunization.

In 2011, the late Ralph Steinman was awarded the Nobel Prize in Physiology or Medicine for his role in the discovery of dendritic cells (DCs) and of their importance in initiating the adaptive immune response. Although DCs were first observed in the skin by Paul Langerhans more than 100 years before his work, Steinman and Zanvil Cohn were the first to show the unique function of DCs. DCs are key sentinel cells that possess distinct “stellate” morphology and unparalleled ability to stimulate naïve T cells (Steinman 2007). Subsequent work has established that DCs primarily serve as a bridge between the innate and adaptive immune systems without engaging directly in effector functions. Such specialization in sentinel activity at the expense of effector function sets DCs apart

from other immune cell types, and it should guide our understanding of DC biology and potential applications.

From the early *in vitro* studies (Knight et al. 1982) to more recent intravital microscopy (Shakhar et al. 2005), DCs have been observed to continuously interact with T cells even in the absence of infection. Indeed, DCs bearing self-antigen have been shown to interact with T cells in the steady state (Scheinecker et al. 2002). Thus, DCs represent obvious candidates to enforce peripheral T-cell tolerance by continuously presenting self- or innocuous antigens (Ag) to T cells in the absence of costimulation and/or activating cytokines. This “tolerogenic” role of DCs could therefore be used as a therapeutic tool to induce or restore tolerance as necessary

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in autoimmune diseases, allergy, etc. (Steinman et al. 2003).

As discussed below, the important contribution of DCs to T-cell tolerance has been confirmed by independent approaches such as genetic or antibody-mediated Ag targeting to DCs. However, the role of DCs in both immunity and tolerance appears complex and highly dependent on genetically and functionally distinct DC subsets. Although extensive “division of labor” exists between and within these subsets, the existence of a unique tolerogenic DC subset or state remains in question. Furthermore, genetic ablation studies revealed that DCs play an essential role in T-cell physiology yet appear largely dispensable for central or peripheral tolerance. This may create significant hurdles to the tolerogenic applications of DCs; on the other hand, they represent supreme candidates in “tolerance-breaking” applications such as antitumor vaccination.

## DC LINEAGE AND SUBSETS

### DCs—a Common Cell Lineage

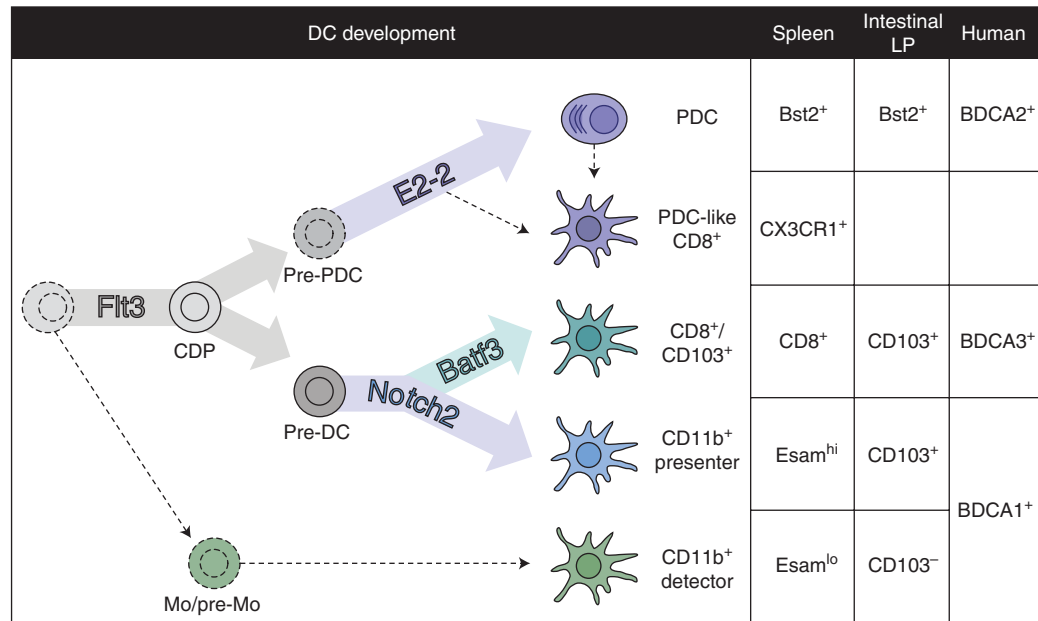
DCs are present throughout the body, including environmental interfaces such as the intestine, filtering organs, and lymphoid organs (Merad and Manz 2009). These cells can be broadly categorized into two classes: classical or conventional DCs (cDCs) and plasmacytoid DCs (pDCs). The cDCs are highly effective at Ag presentation and T-cell stimulation, even in the absence of intentional activation (Steinman 2012). The pDCs are specially equipped for the secretion of type I interferon (interferon  $\alpha/\beta$ , IFN-I) and other cytokines (Liu 2005); they present Ag inefficiently in the steady state but are fully capable of Ag presentation after pathogen-induced activation (Villadangos and Young 2008). Thus, both DC classes share essential DC functions such as highly efficient pathogen recognition, lack of obvious effector function, and the capacity to mobilize and activate multiple innate and adaptive immune cell types.

Recent evidence supports the definition of DCs as a distinct immune cell lineage that includes pDCs, cDCs, and subsets thereof (Geiss-

mann et al. 2010; Liu and Nussenzweig 2010). Progenitor cell populations giving rise to all DC subsets have been identified in the bone marrow, such as the common dendritic cell progenitor (CDP) (Naik et al. 2007; Onai et al. 2007). The development of CDP and its DC progeny is regulated by cytokine Flt3 ligand (Flt3L) and its receptor Flt3, and several transcription factors such as PU.1 and IRF8 are required in multiple DC subsets and/or developmental stages (Fig. 1) (Belz and Nutt 2012). The affiliation of pDCs with the DC lineage has been controversial, given that pDCs lack several essential DC features such as dendritic morphology and high MHC class II expression (Reizis et al. 2011). Moreover, unlike cDCs that undergo terminal differentiation in the periphery, pDCs complete their development in the bone marrow. However, this has been recently attributed to the role of a specific transcription factor, E2-2, in pDC development. The induction of E2-2 in DC progenitors diverts pDCs from the “default” DC pathway and specifies lymphocyte-like morphology and other distinct pDC features (Cisse et al. 2008). Indeed, the loss of E2-2 from mature pDCs causes their full phenotypic and functional conversion into cDC-like cells, further supporting the close genetic relationship between pDCs and cDCs (Ghosh et al. 2010).

## DC SUBSETS AND HETEROGENEITY

Murine cDCs have been traditionally categorized into two distinct subsets, the CD8<sup>+</sup> (CD103<sup>+</sup> in tissues) DCs and CD11b<sup>+</sup> “myeloid” DCs. The CD8<sup>+</sup>/CD103<sup>+</sup> subset appears highly efficient at Ag cross-presentation to cytotoxic CD8<sup>+</sup> T lymphocytes, which may be particularly important during intracellular infections and tumor surveillance (den Haan et al. 2000). The identification of transcription factor Batf3 as a “master regulator” of the CD8<sup>+</sup>/CD103<sup>+</sup> subset strongly supports its unique genetic identity and functionality as a major Ag cross-presenting cell type (Hildner et al. 2008). The precise function of CD11b<sup>+</sup> DCs remains less well understood, although they are generally believed to prime CD4<sup>+</sup> T-cell responses (Dudziak et al. 2007).

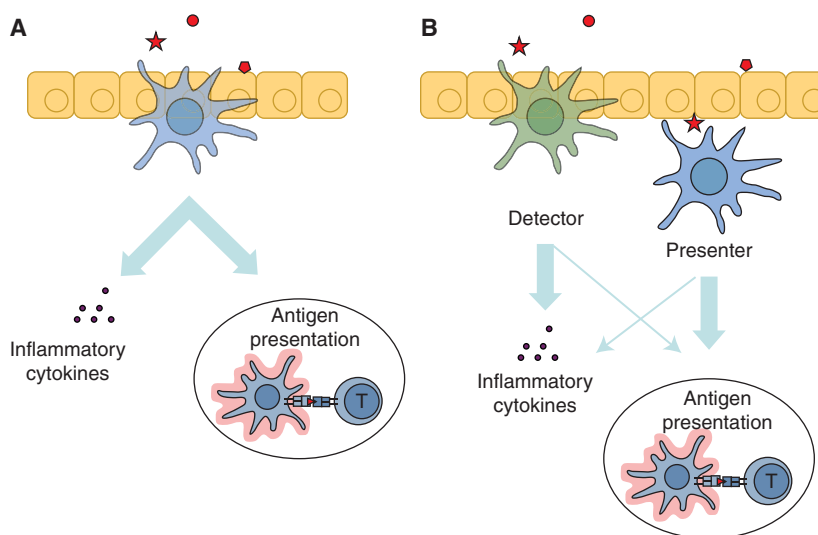


**Figure 1.** Dendritic cell development and subsets. Shown are functionally and genetically distinct DC subsets identified in the mouse spleen and intestinal lamina propria (LP), their major developmental regulators, and key surface markers. The known counterparts of these subsets in the human peripheral blood are also indicated.

Importantly, the traditional categorization appears to mask the significant heterogeneity that exists within each cDC subset. This is particularly evident in the murine spleen, where cDCs were thought to comprise only two cDC subsets of common origin. However, a significant proportion of splenic CD8<sup>+</sup> DCs were found to develop independently of Batf3. The Batf3-independent CD8<sup>+</sup> DCs resemble pDCs in their genetic makeup and dependence on E2-2, and may represent “by-products” of pDC development (Bar-On et al. 2010). These cells are unable to cross-present Ag, and their precise function in immunity, if any, remains unclear. Furthermore, splenic CD11b<sup>+</sup> DCs are also comprised of two distinct cell types whose origin and function differ significantly (Lewis et al. 2011; Kasahara and Clark 2012). Some CD11b<sup>+</sup> DCs develop through a Notch2- and lymphotoxin-β receptor-dependent pathway, show typical DC morphology and expression profile, and are required for efficient CD4<sup>+</sup> T-cell priming. Conversely, the Notch2-independent CD11b<sup>+</sup> DCs appear more related to monocytes in their

origin and expression profile, and show more robust secretion of inflammatory cytokines (Lewis et al. 2011). These results reveal a “division of labor” not only between DC subsets, but also within each “canonical” DC subset (Fig. 2).

A similar heterogeneity within DC subsets appears to exist in the periphery and is best documented in the intestine. The intestinal lamina propria (LP) includes three cDC types: the Batf3-dependent CD103<sup>+</sup>CD11b<sup>-</sup> DCs that are functionally similar to CD8<sup>+</sup> DCs in lymphoid organs (Edelson et al. 2010); the Notch2-dependent CD11b<sup>+</sup>CD103<sup>+</sup> DCs that migrate to mesenteric lymph nodes (Bogunovic et al. 2009) and maintain optimal CD4<sup>+</sup> effector T-cell numbers in the intestine (Denning et al. 2011; Lewis et al. 2011); and monocyte-derived CD11b<sup>+</sup>CD103<sup>-</sup> DCs that may be closely related to macrophages (Bogunovic et al. 2009; Schulz et al. 2009; Varol et al. 2009). These latter DCs appear to reside continuously in the LP and secrete the inflammatory cytokines that recruit and activate local immune cells. Whereas monocyte-derived DCs may sense the intestinal



**Figure 2.** The proposed “division of labor” within dendritic cell subsets. In a traditional view of DC function (A), a single DC detects pathogen, secretes inflammatory cytokines, and migrates to present Ag to naïve T cells. In a revised view (B), physically different DCs within the same subset detect pathogen directly and secrete inflammatory cytokines (“detector” DCs), whereas “presenter” DCs may receive pathogen-derived Ag indirectly through other cells, migrate, and present it (“presenter” DCs).

lumen contents through transepithelial processes (Niess et al. 2005), the CD103<sup>+</sup> DC subsets were recently shown to receive Ag through goblet cell-mediated transport and thereby initiate T-cell responses (McDole et al. 2012). Thus, DCs in tissues such as the intestine are genetically and functionally diverse, and show the same “division of labor” between Ag presentation and cytokine secretion.

### DCs IN HUMANS

Recent work confirmed that major DC subsets are genetically and functionally conserved between mice and humans. This is well illustrated by global gene-expression analysis, which reveals distinct and evolutionarily conserved expression profiles of DC subsets (Crozat et al. 2010b). Indeed, both murine and human pDCs are dependent on E2-2 and express a common set of E2-2-regulated genes (Cisse et al. 2008; Ghosh et al. 2010). Furthermore, several groups have established human BDCA-3<sup>+</sup> DCs as the cross-presenting equivalent to murine Batf3-dependent CD8<sup>+</sup>/CD103<sup>+</sup> DCs (Bachem et al.

2010; Crozat et al. 2010a; Jongbloed et al. 2010; Poulin et al. 2010, 2012). Finally, a major insight into DC biology has been provided by the identification of human patients with DC deficiency (Collin et al. 2011). In particular, patients with mutations in IRF8 lack pDCs, cDCs, and monocytes in circulation (Hambleton et al. 2011). These studies confirm that all DCs and monocytes comprise a unique branch of hematopoiesis, and highlight its evolutionarily conserved transcriptional regulation.

On the other hand, relatively little is known about the heterogeneity and functionality of human DC subsets, particularly in tissues. Several studies in human blood, spleen, and lymph nodes suggest a significant heterogeneity within the BDCA-1<sup>+</sup> DC subset corresponding to CD11b<sup>+</sup> DCs (MacDonald et al. 2002; Mittag et al. 2011; Segura et al. 2012). The function of these different DC populations and their relationship to the murine counterparts remain unclear. DCs capable of T-cell priming have been identified in the human intestinal LP (Bell et al. 2001), and heterogeneous expression of CD103 has been observed on human MLN

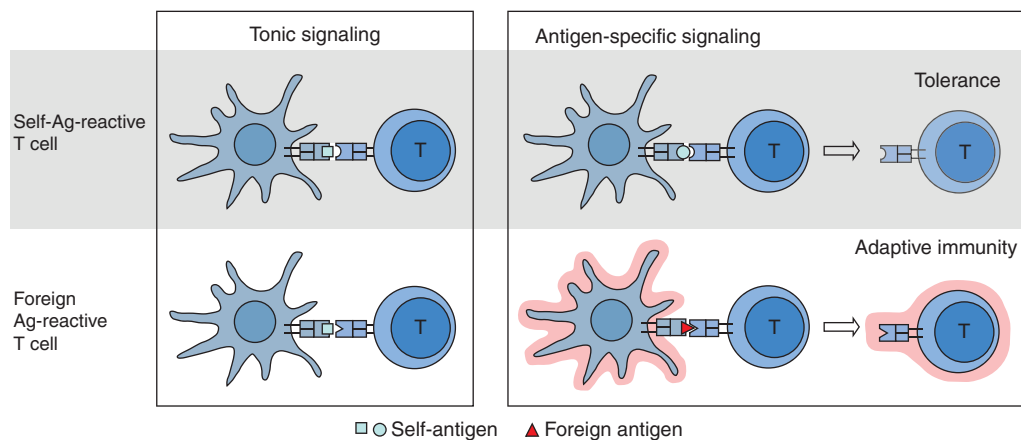
dendritic cells (Jaensson et al. 2008). An unknown fraction of these may correspond to the Batf3-dependent BDCA-3<sup>+</sup> subset (Poulin et al. 2012), although it is unclear whether it is uniformly CD103<sup>+</sup> in humans. Moreover, the human counterpart of T-cell-priming CD103<sup>+</sup> CD11b<sup>+</sup> LP DCs remains to be identified and characterized.

### DC SUBSET HETEROGENEITY: IMPLICATIONS FOR IMMUNITY AND TOLERANCE

It has been commonly assumed that the same physical cell of a given DC subset detects pathogens through pattern recognition receptors, secretes cytokines, migrates into the T-cell area of lymphoid organs, presents Ag, and directs T-cell priming. A similar scenario of steady-state migration and Ag presentation in the absence of pathogen detection would lead to T-cell tolerance (Fig. 3). However, the newly described heterogeneity of DCs in lymphoid organs and tissues suggests a more nuanced view of DC function. Thus, even within a given DC subset some DCs are more adept at pathogen detection and cytokine secretion in situ, such as the intes-

tinal CD11b<sup>+</sup>CD103<sup>-</sup> DCs and Notch2-independent splenic CD11b<sup>+</sup> DCs described above. Such DCs are related to monocytes in the origin and/or expression profile (Varol et al. 2010; Lewis et al. 2011), and we here propose to designate them as “detector” DCs. Conversely, a different DC population appears to capture Ag and present it to T cells after migration to lymphoid organs. The examples include CD11b<sup>+</sup>CD103<sup>+</sup> intestinal DCs (Bogunovic et al. 2009) and splenic CD11b<sup>+</sup> Esam<sup>hi</sup> DCs, both of which are Notch2-dependent (Lewis et al. 2011). These distinct DC populations, which are designated here as “presenters,” may benefit from the cytokine milieu created by the “detectors” at the recognition site.

Such “division of labor” between different DCs within each subset has major implications for our understanding and therapeutic use of DC functions. First, the proposed “detector” and “presenter” populations in each lymphoid organ and tissue must be identified and characterized. In animal models, this could be aided by genetic analysis using common regulators such as Notch2, which controls CD11b<sup>+</sup> “presenter” DCs in both spleen and intestine. In humans, this task is further complicated by limited access



**Figure 3.** Dendritic cells induce peripheral tolerance or immunity by directing the fate of antigen-specific T cells. Presentation by steady-state DCs of weakly agonistic self-peptides maintains tonic Ag receptor signaling and responsiveness of both normal and self-reactive T cells. Strongly agonistic self-peptides may tolerize self-reactive T cells when presented by steady-state DCs, whereas foreign peptides presented by activated DCs induce T-cell priming. The loss of DCs would lead to an unresponsive state in both normal and self-reactive T cells, and prevent self- as well as foreign Ag-specific T-cell responses.

to normal lymphoid organs and tissues, and by extensive variability reflecting individual age, health status, and genetic constitution (Mittag et al. 2011). Nevertheless, knowing the cell that one targets or injects appears as a prerequisite of any rational immunotherapy.

Finally, despite the extensive functional heterogeneity of DCs described above, a dedicated tolerogenic DC subset has not been clearly defined in the steady state. Several proposed tolerogenic or regulatory DC subsets (Zhang et al. 2004; Hadeiba et al. 2008) have not been supported by genetic analysis. For instance, the predicted role of intestinal CD103<sup>+</sup> DCs in regulatory T-cell (T<sub>reg</sub>) induction has been called into question by detailed analysis of specific CD103<sup>+</sup> DC subsets (Edelson et al. 2010; Denning et al. 2011; Lewis et al. 2011). Nevertheless, DCs with distinctly tolerogenic properties may arise in artificial genetic systems (Kriegel et al. 2012) or through specific manipulation in vitro (Morelli and Thomson 2007), and thus may be applied as a tolerance-inducing therapeutic tool.

## FUNCTION OF DCs IN IMMUNITY AND TOLERANCE

### DCs as Initiators of T-Cell Responses

The original in vitro observations on the supreme T-cell priming capacity of DCs (Steinman and Witmer 1978) have been strongly supported by genetic models in vivo (Sapozhnikov and Jung 2008). For instance, constitutive ablation of cDCs essentially abolishes the priming of allogeneic or Ag-specific T cells in the spleen (Birnberg et al. 2008). Furthermore, certain DC subsets have been shown to be essential in the response to specific pathogens. Thus, the cross-presenting Batf3-dependent DCs contribute to the cytotoxic T-cell response to West Nile and Sendai viruses (Hildner et al. 2008; Edelson et al. 2010). More recently, the same group showed that Batf3-dependent DCs are required for the T-cell-mediated control of *Toxoplasma* infection, owing to their unique capacity for IL-12 production (Mashayekhi et al. 2011). By comparison, much less is known about the role of CD11b<sup>+</sup> DCs in antimicrobial responses,

although the Notch2-dependent “presenter” subset appears important for optimal T-cell responses in the spleen and intestine (Lewis et al. 2011). Notably, human patients with IRF8 mutation T80A have a specific reduction of the CD11b<sup>+</sup> DC population and increased susceptibility to mycobacterial infections (Hambleton et al. 2011). These data suggest a major role of this subset in the immunity to intracellular bacteria, although the extent and mechanism of CD11b<sup>+</sup> DC reduction remain to be elucidated.

The pDCs secrete IFN-I in response to multiple viruses (Swiecki and Colonna 2010) and are particularly important for IFN-I-mediated innate control of acute cytopathic coronaviruses (Cervantes-Barragan et al. 2007, 2012). Although modest decreases in T-cell responses have been observed after transient ablation of pDCs (Swiecki et al. 2010; Takagi et al. 2011), their overall role in antimicrobial adaptive immunity remained moot. Recently, constitutive pDC ablation through E2-2 targeting revealed the key role of pDCs in T-cell response to persistent (but not acute) viral infection (Cervantes-Barragan et al. 2012). The pDCs were found to be essential for the priming of virus-specific CD4<sup>+</sup> T cells, even though MHC class II expression on pDCs was dispensable. Collectively, these results emphasize the role of DCs (including cDCs and pDCs) as a key link between innate immune recognition and adaptive immune response to infections. Furthermore, they show the importance of DC-derived cytokines such as IL-12 and possibly IFN-I in T-cell-mediated immunity, underscoring the likely importance of both “detector” and “presenter” DCs.

### The Subversion of DCs by Pathogens

As almost every functional part of the immune system, DCs are subverted by multiple pathogens. The mobility of DCs appears to be a particular advantage to several pathogens, which “hijack” migrating DCs to facilitate their spread. For example, by infecting pDCs, *Toxoplasma* may at once evade the IFN-I response and gain access to lymphoid and peripheral tissues (Bierly et al. 2008). Similarly, murine herpesvirus was shown to infect DCs and exploit their motility



to infect its ultimate target, B cells (Gaspar et al. 2011). Furthermore, cytokine-induced or genetic expansion of DCs increases pathogen burden in infections with intracellular bacteria such as *Listeria*, underscoring the role of DCs as “sentinels without armament” (Alaniz et al. 2004; Sathaliyawala et al. 2010). Conversely, Batf3-dependent CD8<sup>+</sup> DCs are required to transport *Listeria* into the splenic white pulp and initiate productive infection (Edelson et al. 2011). Thus, more DCs are not necessarily better, and this caveat must be taken into account in any therapeutic application of DCs.

### DCs in Tolerance: Central

In contrast to their emerging key role in antimicrobial immunity, the role of DCs in the steady-state immune tolerance is still poorly understood. One proposed mechanism whereby DCs might influence central tolerance is self-Ag presentation for negative selection of thymocytes. It has been proposed that thymic DCs either directly present or cross-present self-Ags acquired from medullary thymic epithelial cells (mTECs), which express many tissue-specific proteins in an Aire-dependent manner (Galegos and Bevan 2004). This model is supported by substantial evidence, suggesting that mTECs present self-Ag both directly and through thymic DCs (Hubert et al. 2011; Klein et al. 2011). A variant of this scenario suggests that mTECs recruit thymic DCs in an Aire-dependent manner and thereby facilitate the generation of T<sub>regs</sub> (Lei et al. 2011).

Another proposed mechanism of DC-mediated central tolerance is the recirculation of peripheral DCs into the thymus, which present peripheral self-Ag to induce clonal deletion or T<sub>reg</sub> generation (Proietto et al. 2009). The original demonstration relied heavily on the transfer of large numbers of cytokine-expanded DCs (Bonasio et al. 2006); nevertheless, this and subsequent studies (Proietto et al. 2008) showed that endogenous peripheral DCs migrate into the thymus, and may induce clonal deletion and/or T<sub>regs</sub> for certain model Ag. However, it is unclear whether this mechanism is relevant or operative at all except for special artificial con-

ditions. Indeed, constitutive depletion of cDCs did not induce an overt breakdown of central tolerance, and negative selection of model self-Ag was found to be normal (Birnberg et al. 2008). The migration-based tolerance induction is even more questionable for pDCs, which were studied in patently artificial conditions (Martin-Gayo et al. 2010; Hadeiba et al. 2012). Furthermore, the proposed Ccr9-mediated migration of murine pDCs into the thymus (Hadeiba et al. 2012) cannot operate in humans, because human pDCs do not express Ccr9. Overall, the role of endogenous thymic DCs in central tolerance is plausible but may be relatively subtle or restricted to only certain self-Ag.

### DCs in Tolerance: Peripheral

#### *The Role of Self-Antigen Presentation by DCs in Tolerance*

The first important evidence for the induction of self-tolerance by DCs came from the studies based on Ag targeting in vivo by DC-specific antibodies (Hawiger et al. 2001). This and subsequent studies (Hawiger et al. 2004; Dudziak et al. 2007) documented a profound T-cell tolerization to DC-targeted model Ag in the steady state. Although this approach is elegant and has potential therapeutic implications, some caveats should be kept in mind. First, the specificity of Ag targeting and the identity of targeted DC population have to be precisely defined, and may be critical for the outcome. For example, the DEC205 (Ly75) receptor used to target CD8<sup>+</sup> cDCs in the original studies is expressed on a variety of non-DC cell types including some macrophages and granulocytes. Another receptor used for targeting, Dcir2/33D1, is highly DC specific but marks a functionally distinct subset of CD11b<sup>+</sup> cDCs (Lewis et al. 2011; Kasahara and Clark 2012). Second, the identity of the targeted receptor and its potential signaling function may fundamentally influence the outcome of Ag targeting. Indeed, Ag targeting to pDCs using pDC-specific surface molecules SiglecH or Bst2 resulted in T-cell hyporesponsiveness or activation, respectively (Loschko et al. 2011a,b).



A compelling genetic way to test Ag presentation by steady-state DCs involved the Cre recombinase-induced expression of model Ag in DCs in vivo (Probst et al. 2003). Using this approach, it was shown that steady-state presentation of immunodominant virus-derived epitopes by DCs induces a profound CD8<sup>+</sup> T-cell unresponsiveness that could not be reversed by subsequent challenge with the virus. Further studies elucidated the mechanism of DC-induced T-cell unresponsiveness, including the expression of inhibitory molecules PD-1 and CTLA-4 on CD8<sup>+</sup> T cells and the induction of T<sub>regs</sub> (Probst et al. 2005; Schildknecht et al. 2010). Collectively, antibody-mediated and genetic Ag targeting suggest that DCs can induce peripheral T-cell tolerance to immunodominant epitopes.

### *The Impact of DC Loss on Immunity and Tolerance*

Given the evidence described above, it might be expected that the loss of DCs would cause a major breakdown of peripheral tolerance. Surprisingly, animals with Cre-mediated constitutive ablation of cDCs (but not of pDCs) had a relatively normal T-cell compartment without overt hyperactivation (Birnberg et al. 2008). Another study has claimed that constitutive ablation of DCs using a similar system causes autoimmune manifestations (Ohnmacht et al. 2009). However, this study neither documented the full course of the disease, nor provided any evidence for T-cell autoreactivity. It appears likely that the purported “autoimmune” disease was in fact a myeloproliferative syndrome caused by increased serum concentration of Flt3L in the absence of DCs (Birnberg et al. 2008; Hochweller et al. 2009; Bar-On et al. 2011). Similarly, human patients with monocyte and DC deficiency show increased Flt3L levels and the associated myeloproliferation, but no major autoimmune disease (Collin et al. 2011). Finally, constitutive DC ablation on the autoimmunity-prone Fas receptor-deficient background ameliorated rather than exacerbated the lupuslike disease (Teichmann et al. 2010). Thus, steady-state DCs have the ability to toler-

ize T cells, yet their actual role in peripheral tolerance appears neither essential nor dominant.

These findings can be reconciled if one takes into account another major consequence of DC ablation, i.e., the rapid loss of T-cell responsiveness. It has long been recognized that T cells require “tonic” signaling through the T-cell receptors for their survival and optimal functionality. It was recently shown that DCs provide a major source of such signals, so that DC ablation rapidly causes T cells to become unresponsive (Hochweller et al. 2010). Thus, autoreactive T-cell clones may receive two kinds of signals from DCs: a tolerizing signal from the self-Ag and a tonic signal from weakly agonistic MHC-peptide complexes. In the absence of DCs, these T cells would be relieved of the negative signal but also deprived of the positive signal, resulting in the net absence of self-reactivity. Another important aspect is the nature of self-Ag presented by DCs in the periphery. By analogy to T-cell selection in the thymus, strongly agonistic self-Ag would induce tolerization, whereas weakly agonistic self-peptides would provide a positive tonic signal (Garbi et al. 2010). Not surprisingly, model studies use unusually strong immunodominant epitopes and thus may predominantly reveal the negative signal.

### *DC-Intrinsic Breach of Immune Tolerance*

Contrary to the loss of DCs, it has been suggested that DC accumulation owing to defective apoptosis causes autoimmunity (Chen et al. 2006; Stranges et al. 2007). However, the DC-specific nature of apoptosis blockade has not been established in either model, and the mechanism of the proposed loss of tolerance in the steady state remains moot. On the other hand, the changes of DC functionality may breach T-cell tolerance and induce inflammation and/or autoimmune manifestations. This was first shown by DC-specific deletion of  $\alpha_V\beta_8$  integrin, which is required for the activity of immunosuppressive cytokine TGF $\beta$  on T cells (Travis et al. 2007). Similarly, the loss in DCs of A20, a negative regulator of the NF- $\kappa$ B pathway, causes widespread immune activation and variable manifestations of autoimmunity (Hammer





et al. 2011; Kool et al. 2011). In the intestine, DC-specific loss of Stat3 or  $\beta$ -catenin makes DCs refractory to IL-10 or Wnt signaling, respectively, causing or predisposing to inflammation (Manicassamy et al. 2010; Melillo et al. 2010). These studies used broad gene deletion in most DCs including pDCs and all cDC subsets, warranting further investigation into the DC subset(s) responsible for the phenotype.

It should be noted that most of these molecules are general negative regulators of immune activation, and their function is by no means restricted to DCs. Indeed, broad deletion of  $\alpha_V$  integrins, Stat3 or A20 from the myeloid lineage may cause even more pronounced inflammation and/or autoimmunity (Takeda et al. 1999; Lacy-Hulbert et al. 2007; Matmati et al. 2011). In that respect, a very interesting case is presented by Blimp-1 (Prdm1), a transcriptional repressor required for B- and T-cell differentiation. The loss of Blimp-1 enhanced IL-6 secretion by DCs and resulted in autoantibody production and other lupus-like manifestations in female (but not male) mice (Kim et al. 2011). This phenotype recapitulates the striking prevalence of lupus in females, and suggests that DCs may be ultimately responsible for this mysterious feature of the disease. Collectively, these results reveal elaborate DC-intrinsic molecular mechanisms that are essential to prevent aberrant DC activation and the ensuing breach of immunological tolerance.

### DCs and $T_{\text{regs}}$

As part of their tolerogenic function, DCs were proposed to mediate the homeostasis of regulatory T cells in the periphery. Steinman and colleagues showed that DCs can induce  $T_{\text{regs}}$  in vitro, especially when combined with strong  $T_{\text{reg}}$ -inducing stimuli such as TGF $\beta$  and retinoic acid (Yamazaki et al. 2003, 2008; Tarbell et al. 2004; Sela et al. 2011). However, the exact role of DCs in  $T_{\text{reg}}$  induction in vivo remains to be fully elucidated. Indeed, the absence of DCs leads to only a modest reduction in  $T_{\text{reg}}$  numbers (Birnberg et al. 2008; Darrasse-Jeze et al. 2009), although DCs were necessary for homeostatic proliferation of  $T_{\text{regs}}$  after their depletion (Suff-

ner et al. 2010). The contribution of DCs to  $T_{\text{reg}}$  maintenance is mediated through the costimulatory molecules CD80/CD86 expressed on DCs (Bar-On et al. 2011). However, when separated from myeloproliferation caused by DC loss, the reduction of  $T_{\text{regs}}$  does not cause spontaneous autoimmunity or lymphocyte hyperactivation (Bar-On et al. 2011). Similar observations have been made in humans with DC and monocyte deficiency (Collin et al. 2011).

Conversely, it was shown that administration of Flt3L leads to an expansion of both DCs and  $T_{\text{regs}}$  (Darrasse-Jeze et al. 2009; Swee et al. 2009; Collins et al. 2011). In particular, Flt3L treatment led to the expansion of CD103<sup>+</sup>CD11b<sup>-</sup> and CD103<sup>+</sup>CD11b<sup>+</sup> DCs in the intestinal LP and increased  $T_{\text{reg}}$  numbers, correlating with reduced severity of ileitis in a Crohn's disease-prone mouse (Collins et al. 2011). Furthermore, Flt3L administration also enhanced survival from graft-versus-host disease, presumably through the induction of  $T_{\text{regs}}$  (Swee et al. 2009). These studies convincingly documented  $T_{\text{reg}}$  expansion and overall tolerogenic environment following Flt3L administration in vivo, which has been interpreted as a simple consequence of increased DC numbers. However, this explanation is subject to major caveats. For instance, Flt3L causes skewing of DC populations toward the CD8<sup>+</sup> cDC lineage (O'Keeffe et al. 2002; Vollstedt et al. 2004); in addition, it may expand non-DCs including various myeloid cell types. Most importantly, Flt3L-expanded DCs may not be functionally equivalent to the steady-state DCs, e.g., owing to Flt3L-induced mTOR signaling (Sathaliyawala et al. 2010). Thus, the increase in DC numbers as a necessary and sufficient cause of  $T_{\text{reg}}$  expansion remains to be formally proven.

Interestingly, DC cell numbers increase after depletion of  $T_{\text{regs}}$ , suggesting that  $T_{\text{regs}}$  regulate DC expansion (Kim et al. 2007). It has been shown that DC expansion in the absence of  $T_{\text{regs}}$  occurs through a Flt3-dependent pathway (Liu et al. 2009). Thus,  $T_{\text{regs}}$  may participate in feedback control of DC activity through a yet unknown Flt3-dependent mechanism. The pDCs have been implicated into  $T_{\text{reg}}$  induction in several specialized models such as organ



transplantation (Ochando et al. 2006) and neural inflammation (Irla et al. 2010). On the other hand, pDC-deficient animals have normal  $T_{reg}$  compartments and no apparent T-cell activation or autoimmunity (Cervantes-Barragan et al. 2012; K Lewis, unpubl.). Altogether, current evidence suggests that DCs contribute to the induction and/or maintenance of peripheral  $T_{regs}$ , which in turn provide a negative-feedback signal to limit DC generation. However, it appears unlikely that DCs are absolutely required for  $T_{reg}$  homeostasis, or that DCs regulate  $T_{regs}$  preferentially compared to effector T cells.

### CONCLUDING REMARKS: TOWARD DC-BASED IMMUNOMODULATION

The evidence reviewed above suggests essential but complex contributions of the DC lineage to almost every aspect of T lymphocyte homeostasis and responses (leaving aside the cross talk of DCs with many other cell types). These contributions depend on the activation state as well as on DC class (e.g., classical vs. plasmacytoid), subset (e.g., Batf3-dependent cross-presenting DCs vs.  $CD11b^+$  DCs), and heterogeneity within the subset (e.g., Notch-dependent vs. independent  $CD11b^+$  DCs). In particular, DCs show extensive specialization and “division of labor” between the migratory “presenter” cells and the sessile “detector” cells generating local cytokine milieu. These complexities have been well appreciated in the development of clinical DC applications, in which different DC subsets may lead to different outcomes (Palucka et al. 2011).

As described above, no distinct tolerogenic subset or state of DCs has been clearly defined at the genetic level. Thus, therapeutic applications of DCs to induce tolerance may be relatively limited, and the advantage of DCs versus other (more abundant but potentially less immunogenic) cell types may have to be considered in every case. Nevertheless, certain clinical settings such as transplantation may provide fertile grounds for tolerogenic DC applications (Morelli and Thomson 2007).

On the other hand, the utility of DCs as cellular immunization vehicles and efficient “breakers” of tolerance remains unrivaled.

Among many potential uses of DC-based immunization, anticancer vaccines present the most promising venue (Palucka et al. 2010). Recent data suggest that endogenous DCs play an important role in antitumor immunity (Diamond et al. 2011; Fuertes et al. 2011), and that they may be functionally impaired by immunoevasive tumors (Engelhardt et al. 2012). Unleashing the capacity of DCs to present Ag and prime effector T cells to reverse the pathological tolerance to tumors may bring closer the ultimate success of cancer immunotherapy.

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