

Dendritic cell-based immunotherapy

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Immunotherapy using dendritic cell (DC)-based vaccination is an approved approach for harnessing the potential of a patient's own immune system to eliminate tumor cells in metastatic hormone-refractory cancer. Overall, although many DC vaccines have been tested in the clinic and proven to be immunogenic, and in some cases associated with clinical outcome, there remains no consensus on how to manufacture DC vaccines. In this review we will discuss what has been learned thus far about human DC biology from clinical studies, and how current approaches to apply DC vaccines in the clinic could be improved to enhance anti-tumor immunity.

Keywords: dendritic cell; immunotherapy

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Introduction

Originally described by Ralph Steinman and Zanvil Cohn in 1973, dendritic cells (DCs) have a critical role in mediating innate immune response and inducing adaptive immune response. Since then, DCs, often referred to as “Nature’s adjuvant” have been recognized as the most potent antigen-presenting cells (APCs), capable of activating both naive and memory immune responses. DCs have a superior capacity for acquiring and processing antigens for presentation to T cells and express high levels of costimulatory or coinhibitory molecules that determine immune activation or anergy [1]. Almost 40 years later, the importance of DCs was recognized when the Nobel Prize for Medicine or Physiology was awarded to Ralph Steinman in 2011 for his discovery of these critical innate immune cells.

DC immunology

DC ontogeny and characteristics

DCs are a sparsely distributed, heterogeneous group of specialized APCs that originate in the bone marrow from CD34⁺ stem cells. Data obtained from *in vitro* human and mouse studies and *in vivo* animal studies, and limited *in vivo* human studies, provide support for our understanding of DC differentiation. The most recently accepted

model proposes that monocytes and DCs originate from the same common progenitors called the monocyte and DC progenitors (MDPs). These two cell types diverge in the bone marrow when the MDPs give rise to monocytes and the committed DC progenitors (CDPs). The CDPs give rise to pre-DCs that migrate out of the bone marrow to produce the two major sub-populations of DCs [2]. Stromal cell culture systems comprising hematopoietic stem cells cultured with mouse bone marrow stromal cells and stem cell factor (SCF), granulocyte-macrophage colony-stimulating factor (GM-CSF) and Flt3L have identified a definitive DC precursor population that gives rise to DC subsets found in the blood [3-5]. Although this model has largely been studied in mouse, studies in humans have confirmed these findings as well. Human granulocyte monocyte DC precursors sequentially develop into monocyte DC precursors, which subsequently give rise to common DC progenitors that are restricted to produce the three major subsets of DCs: CD1c⁺ DCs and CD141⁺ DCs (which are together considered conventional DCs or cDCs), and plasmacytoid DCs (pDCs). A migratory phenotype (hpre-cDC) has also been identified in human cord blood, bone marrow, blood, and peripheral lymphoid organs, which sustains the cDC pools through differentiation. Furthermore, Flt3L given systemically to humans has been shown to increase the pre-cDC pool [4, 6].

Phenotypically, human DCs lack lineage (Lin) markers (CD3, CD19, CD14, CD20, CD56, and glycophorin A), but constitutively express major histocompatibility complex (MHC) class II [7, 8]. cDCs and pDCs repre-

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sent the two major types of DCs in the blood and lymphoid tissue. cDCs are MHC-II⁺CD11c⁺ and are further subdivided into CD1c⁺ and CD141⁺ subsets. Three DC populations can also be distinguished by molecular signatures: CD1c⁺ DCs express IRF4, Notch2, Rbpj, and Klf4; CD141⁺ DCs express IRF8, batf3, Bcl6, and Flt3; and pDCs express IRF8, Bcl11a, Spi-B, E2-2, Runx1, and IL-3RA [9, 10]. CD1c⁺ DCs are the predominant subset, whereas the CD141⁺ DCs are a minor population, at least in the blood. CD141⁺ DCs are believed to be the human equivalent of mouse CD8⁺ DCs, which have the ability to cross present cell-associated antigens to CD8⁺ T cells. CD1c⁺ DCs express toll-like receptor (TLR) 1-8 and when stimulated, can secrete interleukin-12 (IL-12), tumor necrosis factor- (TNF α), IL-8, and IL-10. CD141⁺ DCs express TLR3 and 8 and secrete high levels of type I interferon upon stimulation with synthetic dsRNA poly-ICLC [11]. This DC subset is also known for producing high levels of IL-29 or type III interferon in response to TLR3 activation [12]. CD141⁺ DCs exclusively express Clec9A (DC NK lectin group receptor-1), an endocytic receptor that renders cells more capable of taking up and presenting antigens derived from necrotic cells [13, 14], and also XCR1. Although both CD1c⁺ and CD141⁺ DCs can cross-present antigens to CD4⁺ and CD8⁺ T cells, CD141⁺ DCs may be more efficient, although this may depend on the type and form of antigens and how they are accessed [15].

pDCs are defined as Lin⁻MHC-II⁺CD303⁺CD304⁺ cells. They are major effector cells in immune responses due to their ability to produce up to 1000-fold more type I interferons (IFN- α/β) in response to viral infections than other cell types [16]. pDCs can also acquire antigens through, e.g., receptor-mediated endocytosis or uptake of dying cells, although not as efficiently as cDCs [17]. Moreover, pDCs can rapidly cross-present antigens, including components of influenza virus, to CD8⁺ T cells [18], after processing in endosomal-type vesicles. pDCs express high levels of TLR7 and TLR9, which enable them to recognize viral and self nucleic acids [19]. Although cDCs can be found in almost every peripheral tissue as well as in lymphoid organs, pDCs have a more restricted distribution. They are found mostly in the T cell area of lymphoid organs such as the lymph nodes, tonsils, spleen, thymus, BM, and Peyer's patches, the blood, and some peripheral tissues including the liver and nasal mucosa. pDCs can activate melanoma-specific CD8⁺ T cell responses [20], but they may also inhibit anti-tumor immune responses. pDCs with diminished capacity to produce IFN α have been found in many tumors [20, 21] and the accumulation of pDCs expressing indoleamine 2,3-dioxygenase (IDO), which is responsible for the deg-

radation of tryptophan (an amino acid essential for T cell proliferation) and implicated in the generation of regulatory T cells, has been observed in tumor-draining lymph nodes [22-25]. Therefore, modulating pDCs may be an important step in generating effective anti-tumor immunity.

There are five major types of DCs found in the skin: Langerhans cells (LCs), CD14⁺ DCs, CD1a⁺ DCs, CD1c⁺ DCs, and CD141⁺ DCs. LCs are the major APCs found in the epidermis of the skin. They are positive for CD45, MHC-II, epithelial cell adhesion molecule, Langerin, and contain large granules called Birbeck granules. LCs acquire antigens in the epidermis and transport them to the regional lymph nodes where they present the antigens to T cells to initiate immune responses. During inflammation, pDCs and inflammatory DCs are also recruited to the skin. Inflammatory DCs are CD11c⁺⁺MHC⁺⁺CD40⁺CD80⁺CD86⁺ and are differentiated from the circulating monocytes in the blood only during inflammation/infection to infiltrate the site of inflammation/infection [26]. Dermal DCs are thought to comprise a heterogeneous mixture of CD14⁺ cells, the less characterized CD1a⁺ DCs, CD1c⁺ DCs as well as CD141⁺ DCs, but most recently, dermal CD14⁺ cells have been shown to be tissue-resident monocyte-derived macrophages [27].

Lymphoid tissues have CD1c⁺ and CD141⁺ DCs; some of the secondary lymphoid tissues have specialized resident DC subsets. For example, CD103⁺ DCs have been found in the mesenteric lymph nodes (MLNs) [28] where they have a crucial role in the induction of tolerance to commensal bacteria and food antigens. In addition, these cells may be similar to the CD103⁺ DCs found in murine MLNs that have an enhanced ability to metabolize vitamin A to generate retinoic acid, which drives the differentiation of gut-homing regulatory T cells [29].

The concept of maturation

In their resting state, DCs are considered to be immature but primed to acquire antigens *in situ* through a variety of receptors and mechanisms. Upon exposure to "activating stimuli," DCs undergo a complicated series of phenotypic and functional changes referred here to as "activation" and "maturation", respectively [30]. The process of DC activation is an intricate and tightly controlled differentiation process that is closely associated with antigen acquisition. Activation is characterized by the upregulation of chemokine receptors (e.g., CCR7), adhesion molecules, costimulatory molecules (CD54, CD80, and CD86), immunoproteasomes and MHC class I and II molecules, all essential for migration of the cells to the lymphoid tissues and optimal activation of the immune responses. Cytokines produced during this process

influence the immune responses generated by subtypes of CD4⁺ T cells such as T helper 1 (Th1) cells (by IL-12), Th2 cells, and regulatory T cells. The role of DCs in priming Th9 or Th22 cells is still under investigation. In addition, DCs qualitatively and quantitatively orchestrate the type of immune responses that develop by activating naive and memory B cells [31], natural killer (NK) cells (via the action of IL-12, IL-15, and type I IFNs) [32], and NKT cells (through antigen presentation on the CD1 molecule) [33]. DC maturation is characterized by the reduction in phagocytic capacity, enhancement in antigen processing and presentation, improved migration to lymphoid tissues, and increase in the capacity to stimulate B and T cells. Maturation is induced by microbial products that trigger the activation of pattern recognition receptors such as TLRs [34], or activation of intracellular sensors such as RIG-I [35] or inflammasome [36], or by the action of inflammatory molecules (TNF α , IL-1, IL-6, and IFN α) produced by the cells of the immune system or by damaged tissues [37]. Dead cells can also release factors that activate DCs (e.g., heat-shock proteins, RNA, and DNA) [38].

Antigen acquisition and presentation

DCs take up antigens through phagocytosis, micro- or macro-pinocytosis, and endocytosis using Fc receptors (Fc γ receptor type I or CD64 and Fc γ receptor type II or CD32), integrins ($\alpha_v\beta_3$ or $\alpha_v\beta_5$), C-type lectin receptors (CLRs, including mannose receptor and DEC205), apoptotic cell receptors, and scavenger receptors. Antigens are either processed into peptides via the endogenous pathway for presentation on MHC class I molecules to CD8⁺ T cells, or processed via the exogenous pathway for presentation on MHC class II molecules to the CD4⁺ T cells. Alternatively, DCs can also process antigens via two cross presentation pathways [39]: the cytosolic pathway and the vacuolar pathway [40]. In the cytosolic pathway, antigens are transferred to the cytoplasm, where they are processed in the proteasome before being loaded onto the newly formed MHC class I molecules; and this process may involve the participation of the ER machinery. The vacuolar pathway is less defined but is thought to occur in the endocytic compartments because the pathway is resistant to proteasome inhibitors, but sensitive to inhibitors of lysosomal proteolysis, and depends on cathepsin S. Recently it has been found that TLR signaling influences the maturation of phagosomes and induces the accumulation of MHC class I molecules in the phagosomes for optimal cross presentation [41]. Although the underlying mechanisms of cross presentation are still being clarified, the ability of DCs to utilize this process to activate CD8⁺ T cells is in many cases well established [42-44]. DCs

can present antigens in various forms besides conventional peptides. For example, phosphopeptides and citrullinated antigens have been shown to be presented on HLA molecules and recognized by T cells [45, 46]. These antigens, which can be expressed on cancer cells, are novel targets of immunotherapy (i.e., NCT01846143). DCs can also process lipid antigens and present them on the CD1d molecule to activate NKT cells. Furthermore, they can recognize and acquire antigens containing carbohydrate structures via CLRs including MMR (macrophage mannose receptor), DEC205, and DC-SIGN (dendritic cell-specific intercellular adhesion molecule-3-grabbing non-integrin). CLRs can function as endocytic receptors to internalize antigens for antigen processing and presentation, but some of the CLRs, e.g., MICL (myeloid C-type lectin-like receptor) and DICR (dendritic cell immunoreceptor), may be inhibitory [47].

Induction of immunity

Upon maturation, DCs migrate to the secondary lymphoid tissues such as the lymph nodes (capturing antigen from the skin and solid organs), the spleen (capturing antigen from the blood), or the Peyer's patches (capturing antigen from the gut lumen) where they come in contact with T and B cells (reviewed in Benvenuti [48]). cDCs migrate through the afferent lymph from non-lymphoid tissues to the T cell-rich areas of lymph nodes. pDCs, which also migrate into T cell areas of secondary lymphoid tissues, do so through high endothelial venules of lymph nodes and the marginal zone of the spleen, likely using CCR7 and CD62-L [49]. Both activated blood cDCs and pDCs can migrate in response to lymph node homing chemokines (CCL19 and CCL21) through the expression of CCR7. CCR7 expression and CD141⁺ DC infiltration both have been shown to correlate with intratumoral T cell infiltration and better clinical outcome in melanoma [50]. Most recently, it was shown in the mouse that strategically localized DCs residing in the lymph nodes can induce T cell responses much sooner and independently of migratory DC [51]. These DCs reside in the lymphatic sinus endothelium, where they can "scan" lymph to capture soluble lymph-borne antigens to rapidly initiate immune responses without delay. It is conceivable certain antigen adjuvant platforms can be developed to access these DC subsets.

Through their TCRs, T cells specifically recognize antigens bound on MHC molecules on the surface of DCs. Peptides bound on MHC class I molecules are recognized by CD8⁺ T cells, whereas peptides bound on MHC class II molecules are recognized by CD4⁺ T cells. Activation of T cells is dependent on the intensity and duration of DC-T cell interactions, mediated through

the immunological synapse (IS). IS forms as the result of cytoskeletal reorganization within the T cell leading to the dynamic clustering of T cell surface receptors and signaling molecules into supramolecular activation clusters, which provide an optimal environment for signaling molecules downstream of the TCR. The upregulation of costimulatory molecules (CD40, CD86) and MHC molecules during DC maturation is critical for making stable and long-lasting contacts with T cells through the IS, which is required for T cell expansion and differentiation into memory and effector T cells.

DCs can also activate naïve and memory B cells [31], primarily through their ability to stimulate CD4⁺ T cells (such as T follicular helper cells), which induce B-cell growth and antibody production. Antibody class switching is influenced through the production of factors that activate and induce B-cell proliferation (B-Lys and APRIL) [52]. The follicular DCs, which are present in the germinal centers of lymph nodes, support maintenance of B-cell memory via formation of multiple antigen-antibody complexes and continuous stimulation of B cells. DCs also activate NK cells via IL-12, IL-15, and type I IFNs [32]. IL-12 produced by DCs potentiates the cytolytic activity of NK cells. Conversely, the interaction with NK cells can induce further DC maturation. The production of XCL1 and XCL2 by NK cells, as well as CD8⁺ T cells, can recruit and initiate specific responses in XCR1-expressing DCs [53, 54]. Finally, DCs activate NKT cells through the expression of invariant CD1 molecules and presentation of glycolipid molecules [33]. Therefore, DCs have important roles by mobilizing multiple arms of both the innate and adaptive immune responses.

In addition to inducing immunity, DCs mediate peripheral tolerance and prevent autoimmunity [55, 56]. Immature DCs do not activate T cells due to their low expression levels of MHC and costimulatory molecules. DCs can also express IDO in response to certain stimuli, consequently driving T cells to undergo cell cycle arrest or apoptosis possibly due to a counter-regulatory effect. Moreover, the metabolites produced by tryptophan catabolism can exert direct cytotoxic effects on T cells [57, 58]. Through production of IDO, DCs can induce the differentiation of regulatory T cells [59, 60], which can infiltrate several types of tumors [61, 62]; they can also exert their effects through TGF β , IL-10, and CTLA-4, among other mechanisms, to inhibit proliferating T cells [56, 63]. Migratory DCs have a specialized role in mediating peripheral tolerance by inducing the generation of regulatory T cells [64]. As part of immune surveillance, immature DCs acquire self-antigens and in the absence of maturation stimuli, these DCs induce tolerance to

these self-antigens. DCs can also mediate tolerance to self-antigens through the uptake of apoptotic cells via receptors including LOX-1, CD36, integrins ($\alpha_v\beta_3$ and $\alpha_v\beta_5$), and complement receptors (CR3 and CR4). Binding of apoptotic cells to CR3 on DCs leads to impaired maturation of the DCs in response to lipopolysaccharide (LPS) stimulation and an impaired priming and activation of the memory T cell responses [65]. Furthermore, binding of apoptotic microparticles to CD44 can inhibit human DCs [66]. Altogether, these properties of DCs make them ideal tools for inducing tolerance in the settings of transplantation and autoimmunity.

DC immunotherapy

The role of the immune system in eliminating tumors has been established in several studies. First, incidences of spontaneous regression of metastatic melanoma have been proposed to be caused by the body's immunologic responses [67]. Second, the existence of immunogenic tumor-associated or tumor-specific antigens [68] that are recognized by B and T cells has been reported. Third, knockout mice lacking components of the IFN signaling pathway, perforin, or recombination activating gene 1/2 have significantly higher risk of developing tumors [69]. Finally, adoptive transfer of tumor-infiltrating lymphocytes as well as engineered tumor antigen-specific T cells has led to the regression of tumors in melanoma patients [68]. Altogether, these studies prove a protective role of the immune system in eliminating tumors. The complex relationship between the immune system and tumors is now described as consisting of three phases of cancer immunoediting: elimination, equilibrium, and escape [70]. The first phase involves induction of the anti-tumor immune responses whereby both the innate and adaptive arms of the immune systems are activated to destroy the tumors. Equilibrium is the second phase when the immune system exerts a selection pressure on the tumor. Tumor cells that have become resistant to this selection pressure enter the third phase, escape. It is during this third phase that tumors grow and expand beyond control, and become malignant. Given the central role of DCs in initiating immune responses and surveillance, investigators have theorized that DCs would serve as an ideal tool for boosting endogenous anti-tumor responses that can lead to the effective eradication of tumors. As a proof of principle, we conducted a controlled study in healthy subjects and showed that, a single injection of DC vaccine comprising antigen-pulsed DCs was sufficient to induce antigen-specific immune responses *in vivo* [71], whereas soluble antigen alone failed to induce immunity. This result indicates that DCs can be successfully loaded

with antigens, and following injection, traffic to the lymphoid tissues to activate antigen-specific T cells. In support of this notion, early clinical trials using *ex vivo*-generated DCs pulsed with tumor antigens demonstrated that immune responses could indeed be induced thus setting the stage to further manipulate DCs to enhance immune responses *in vivo* (reviewed in Constantino *et al.* [72]).

Despite many studies, however, the approach involving DCs remains to be optimized. Though the most commonly adopted method in clinical trials uses *ex vivo*-generated monocyte-derived DCs, it is currently unknown which primary DC subset is the best subset to use as an adjuvant or a target for vaccination. This is important to point out because monocyte-derived DCs are not comparable to the steady-state DC subsets present in the body; instead, they resemble more the cells that appear during an inflammatory response *in vivo*. With the plethora of

maturation stimuli available, more studies are required to compare the performance of these cells. In addition, the number of DCs to inject (0.3×10^6 cells to 200×10^6 cells per injection), the vaccination schedule (once every 2 weeks vs 3-4 doses or even up to 10 injections given every 3-4 weeks) and the route of injection (subcutaneous, intradermal, intranodal, intravenous, or even intratumoral) to induce the most potent and long-lasting immune response have also not been determined. Below, we discuss current and future approaches to apply DC-based and targeted vaccines in the clinic (Figure 1).

Current approaches

Ex vivo approaches to DC vaccination

Current methods for generating DCs used in the clinical trials include differentiation from monocyte precur-

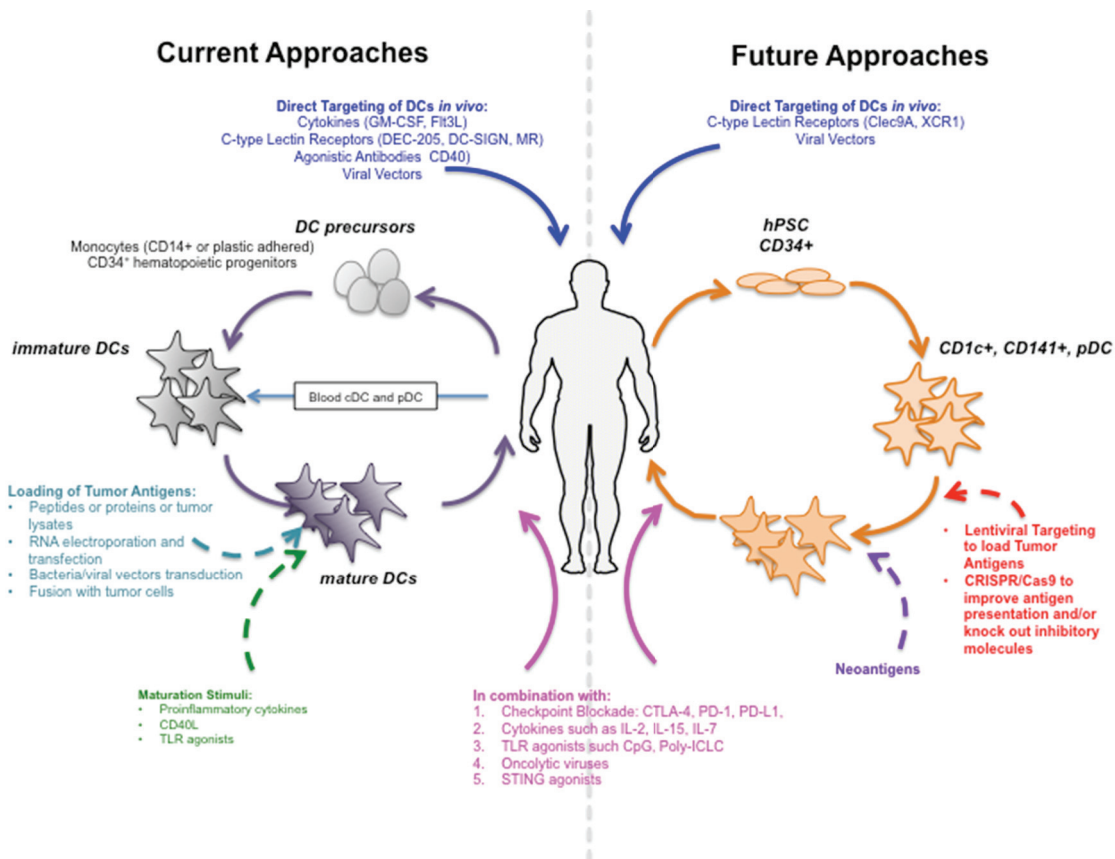


Figure 1 Current and future approaches for dendritic cell (DC)-based vaccination. The current approaches for DC-based vaccination are mainly based on antigen loading on *in vitro*-generated DCs from monocytes or CD34⁺ cells, activating them with different TLR ligands, cytokine combinations, and injecting them back to the patients. The *in vivo* targeting approaches comprise administering specific cytokines (e.g., Flt3L, GM-CSF) and targeting the DCs with antibodies to C-type lectin receptors or agonistic antibodies (e.g., anti-CD40) that are conjugated with antigen of interest. Future approach may target DC subsets based on their specifically expressed C-type lectin receptors or chemokine receptors. Another potential approach is the generation of genetically engineered DCs from induced pluripotent stem cells and use of neoantigen-loaded DCs for inducing better clinical outcome.

sors or CD34⁺ hematopoietic precursors, *in vivo* expansion of circulating DCs, and most recently, isolation and enrichment of circulating blood DC subsets. Although no direct comparison of all these methods of DC generation has been made in clinical trials, analyses of the transcriptional profiles of the DCs generated *ex vivo* using various methods have revealed fundamental differences from the *in vivo* DC subsets [73]. Monocyte-derived DCs cluster with CD34⁺-derived DCs [73] and are more closely related to macrophages [74]. Furthermore, immune system-related transcripts expressed by the *ex vivo*-generated DCs differ from those expressed by the *in vivo* DC subsets, suggestive of functional differences between them. Despite these differences, DCs derived using the most commonly used methods (monocyte-derived DCs and CD34-derived DCs) have been shown to stimulate antigen-specific T cell responses in both preclinical and clinical studies. The different methods of DC differentiation and their advantages are discussed below.

Monocyte-derived DCs Differentiation of DCs from monocytes in peripheral blood mononuclear cells (PBMCs) obtained from whole blood or leukapheresis is by far the most commonly used approach. DCs derived from this process are called monocyte-derived DCs (MDDCs). CD14⁺ monocytes are selected from PBMCs by plastic adherence or positive selection using immunomagnetic beads [75-79] and are induced to differentiate into immature CD14⁺CD83⁻ DCs in culture for several days in the presence of IL-4 and GM-CSF. The immature DCs are stimulated to mature by exposure to a maturation stimulus and simultaneously loaded with tumor antigen for an additional 1-2 days. The differentiated, matured, and antigen-loaded DCs are then harvested and cryopreserved in aliquots and are thawed at each scheduled vaccination date. This approach is both time-consuming and costly to undertake but is used in many treatment centers. The method has been adapted by some into a programmable closed culture system. Although autologous DC vaccination is the preferred approach, the use of allogeneic DCs has been explored for the treatment of renal cell carcinoma and demonstrated to be immunogenic [80]. This approach was also tested in acute myeloid leukemia (NCT01373515). Alternatively, umbilical cord blood-derived CD11c⁺ DCs can be a possible source of allogeneic DCs [81]. Allogeneic DCs may be an attractive source as they could be prepared from an unrelated healthy donor whose immune system has not been rendered dysfunctional as in cancer patients [82]. Finally, although DC production would still be expensive and time-consuming, the allogeneic approach would enable the off-the-shelf treatment that would be more viable and attractive for

pharmaceutical companies. Partially HLA-matched allogeneic DCs may yet be a useful source for vaccination as (a) HLA-mismatched DCs can induce a strong immune response and (b) the use of different DC donors will still boost the immune response to the antigen of interest [83, 84]. However, it is possible that this approach may not be feasible over a prolonged period due to antigen competition.

CD34⁺ precursors CD34⁺ precursors can be mobilized from the bone marrow prior to leukapheresis by the pre-treatment of granulocyte-colony stimulating factor. The harvested cells are further expanded *in vitro* for up to 12 days in the presence of GM-CSF, Flt3L, TNF α , TGF β , and SCF, producing a mixture of MDDCs that are phenotypically similar to epidermal LCs, and a large proportion of myeloid cells at different stages of differentiation [77]. Compared with MDDCs and dermal-interstitial DCs, mature LCs stimulate more effective CD8⁺ T cell responses, a process requiring IL-15 produced by the LCs [85]. In a clinical trial aimed to compare melanoma peptide-loaded LCs and MDDCs, LCs were more potent stimulators of antigen-specific T cell responses than MDDCs, but MDDCs supplemented with IL-15 stimulated significantly more antigen-specific effector memory T cells [86]. Perhaps retroviral transduction of LCs, which enables sustained antigen expression, may enhance their efficacy *in vivo* [87]; however, this approach has been less enthusiastically accepted. Most recently, it was shown that CD34⁺ cells (derived from cord blood, bone marrow, or peripheral blood) cultured with MS5 stromal cells in the presence of Flt3L, SCF, and GM-CSF can develop into all three major DC subsets [88]. This new method has allowed a clear delineation of how the three subsets develop from their progenitors. The authors showed that granulocyte-monocyte-DC progenitors develop into monocyte-DC progenitors, which give rise to monocytes and common DC progenitors. These common DC progenitors differentiate into the three DC subsets. In addition, the authors identified an immediate DC precursor population circulating in the blood and showed these cells differentiate from the common DC progenitor cells in response to Flt3L administration [4]. Adaptation of this method has allowed investigators to obtain higher numbers of each DC subset; e.g., yields of pDCs and CD141⁺ DCs can be increased by 1.5- and 9-fold, respectively. The CD1c⁺ DCs in this culture system are not very well characterized and they may be more close to the MDDCs than the CD1c⁺ DCs circulating in the blood. More importantly, these investigations have made it possible to adjust the methodology for good manufacturing practice (GMP) production, thus allowing the potential

comparison of all three DC subsets in the clinic for their relative immunogenicity. Significantly, this approach is also amenable to genetically manipulating DC subsets to improve their antigen presenting function.

Blood DCs The first FDA-approved cell-based therapy for the treatment of hormone-refractory prostate cancer, Provenge (Sipuleucel-T), is a CD54-enriched peripheral blood vaccine [89]. Provenge consists of a mixture of DCs, B cells, monocytes, and NK cells that have been cultured *ex vivo* with a recombinant fusion protein containing prostatic acid phosphatase (PAP) and GM-CSF, before the mixture is infused intravenously back into patients within 48 h of the leukapheresis collection [90]. Patients receive three freshly processed products; products two and three contain primed T cells in addition to activated APCs, which include DCs. phase I and II clinical trials have demonstrated that Provenge is safe and patients develop immune responses to the fusion protein [91]. Reactivity to the native protein also develops but it is less strong. The increase in CD54 expression upon culture with the fusion protein correlated with APC activation and was used as a surrogate marker for potency of the therapy [92]. The phase III IMPACT trial showed a 4.1-month improvement in median overall survival at 36 months and the survival rate was 31.7% in treated patients vs 23.0% in placebo patients [89]. In addition to inducing circulating humoral (with antigen spreading as evidenced by responses against PSA [93]) and cellular immune responses [94], Provenge has also been shown to induce T cell infiltration in the tumor microenvironment (TME) [95]. Provenge is currently being tested in combination with anti-PD1 (NCT01420965), anti-CTLA-4 (NCT01832870, NCT01804465), and IL-7 (NCT01881867) to see whether any of these combinations would further enhance its efficacy.

Administration of growth factor Flt3L induces the *in vivo* expansion of circulating DCs [96]. Daily administration of Flt3L for 10 days led to a 130-fold increase in CD1c⁺ DC number, 48-fold increase in CD141⁺ DC number, and 6- to 16-fold increase in pDC number from the mean baseline count. CD34⁺ cells also increased 23-fold from the mean baseline count in healthy human volunteers [97]. Flt3L administration may also increase the number of precursors to CD1c⁺ DCs and CD141⁺ DCs (the pre-cDCs) [4], and the number of precursors for pDCs, when combined with thrombopoietin, at least *in vitro* [98]. Clinical trials using Flt3L are currently ongoing in melanoma in combination with DEC205/NY-ESO-1 fusion protein and poly-ICLC (NCT02129075) with preliminary results confirming the immunogenicity and safety of the combined product. The treatment is also being given

intratumorally in low grade B-cell lymphoma in combination with poly-ICLC and radiation (NCT01976585), where the response rate is > 30% (J Brody, personal communication). Interestingly, intratumoral Flt3L not only mobilizes DCs into the TME but also achieves a systemic mobilization of DCs. Therefore, by increasing the DC numbers and localization to tumors and also providing a maturation stimulus, Flt3L may substantially improve anti-tumor T cell priming. The contribution of individual DC subsets in acquiring and presenting tumor-derived antigens in these settings remains to be determined. Nevertheless, such studies are enabling the analysis of DCs at the individual cell level by analyzing their phenotype and transcriptome through strategically obtained tumor biopsies. Flt3L has also been delivered as a single agent to patients with acute myeloid leukemia (NCT00006223) and colorectal cancer (NCT00003431); and has been given in combination with HLA-A2-restricted TAAs emulsified in Montanide ISA-51 adjuvant to patients with melanoma or renal cell cancer (NCT00019396).

CD1c⁺ DCs and pDCs Due to their paucity, blood DC subsets have not been rigorously compared and tested in clinical trials. In one study, however, autologous-activated pDCs loaded with tumor-associated peptides were evaluated in patients with melanoma. One group successfully isolated sufficient numbers of pDCs to test their immunogenicity *in vivo*. pDCs were isolated by leukapheresis using a CliniMACS isolation system with average purity of 75% and yields of 13×10^6 to 33×10^6 cells. Isolated pDCs were then cultured in the presence of IL-3 and loaded with HLA-A2-restricted gp100 and tyrosinase peptides. The study showed that despite the limited cell numbers, vaccination with autologous pDCs loaded with tumor-associated peptides was safe and induced antigen-specific CD4⁺ and CD8⁺ T cell responses in the patients [99]. This study was the first to show the feasibility of isolating primary DC subsets and using them for immunotherapy. A recent study by the same group demonstrated the successful isolation of autologous primary CD1c⁺ DCs from leukapheresis product using the CliniMACS isolation system from metastatic melanoma patients enrolled in the study [100]. Positively selected CD1c⁺ DCs had an average purity of 93% and yields of $27\text{--}96 \times 10^6$ cells. The isolated cells were cultured overnight in the presence of GM-CSF and loaded with HLA-A2-restricted gp100 and tyrosinase peptides. Patients received $3\text{--}10 \times 10^6$ DCs given intranodally every 2 weeks. The study demonstrated the feasibility and safety of isolation of primary CD1c⁺ DCs for vaccination. The vaccine induced peptide-specific CD8⁺ T cell responses in a small subset of patients. It remains to

be seen whether these cells will elicit a superior anti-tumor response than pDCs or even CD141⁺ DCs, which also produce type I and III IFNs but have the capacity to migrate from tumor tissues to draining nodes and to cross-present antigens to T cells.

Although most of clinical studies have cited the induction of immune responses by DC vaccine as a measure of their success, most T cell responses are evaluated after long *in vitro* stimulation cultures. And those who monitored the direct *ex vivo* response reported weak T cell reaction primarily involving the CD4⁺ cells, but not the CD8⁺ cells. Presumably the nature of antigen (tumor antigen to which patients may have already developed tolerance) and possibly the quality and amount of cells, as well as the lack of efficient migration to the lymph nodes may compromise the possibility of eliciting a potent response as was observed in healthy patients receiving DC vaccines when the antigens were technically neoantigens, e.g., keyhole limpet hemocyanin (KLH) or antigens derived from pathogens [71]. In an unpublished study from our group, our DC vaccine induced very weak *ex vivo* CD8⁺ T cell responses to HLA-A2-restricted melanoma peptides (Melan-A, gp100, tyrosinase, MAGE-A3, and NY-ESO-1) but very strong CD8⁺ T cell responses to KLH, consistent with this hypothesis. Indeed, DCs seem to be quite capable of eliciting CD8⁺ T cell responses to tumor-derived neoantigens, suggesting that the choice of antigen dictates the effectiveness in priming T cell responses [101]. Studies with Provenge using a fusion protein of PAP-GM-CSF also support this assertion.

Maturation of DCs

Mature DCs have enhanced expression of costimulatory molecules, produce cytokines and chemokines necessary for the efficient activation of T cell responses [42], and can migrate to the lymphoid tissues [102]. In contrast, immature DCs fail to induce antigen-specific responses [103] and may in fact induce the differentiation of regulatory T cells [56, 104]. In clinical studies, only the mature, peptide-loaded DCs had the capacity to induce antigen-specific T cell responses in healthy subjects [71] and patients with metastatic melanoma [105].

Maturation stimuli DCs can be matured in various ways. In the laboratory, TLR agonists such as LPS (TLR4), poly IC (TLR3), and resiquimod (TLR7) are commonly used as DC activators. In the clinic, a commonly used maturation cocktail of proinflammatory cytokines TNF α , IL-1 β , and IL-6 combined with prostaglandin E₂ (PGE₂) was initially established as the gold standard for maturation [106]. This cocktail induced the upregulation of MHC class I and II molecules, CD40,

CD80, CD86, and CCR7 but failed to effectively induce IL-12p70 [106]. However, when compared with other DC maturation stimuli (CD40L trimer, poly IC, and LPS), the cytokine cocktail induced the most uniform maturation in terms of upregulation of DC maturation markers, with the highest yield and recovery; it also stimulated the highest levels of allogeneic T cell proliferation and cytokine production, and it induced priming of Th1 responses [106]. Other studies have indicated that PGE₂ may induce differentiation of regulatory T cells and Th2 responses [107], IDO expression [108], and is responsible for the lack of IL-12p70 production [109]. However, PGE₂ has also been shown to have important roles in promoting the migration of DCs into lymphoid tissues through the upregulation of CCR7 on DCs [110], and in enhancing T cell proliferation through the induction of OX40L, CD70, and 4-1BBL on DCs [111].

Alternative methods of maturing DCs have also been explored. CD40 ligand (CD40L) is expressed primarily by activated T cells and B cells, and binds to its receptor CD40 on DCs. Engagement of CD40L with CD40 on DCs induces upregulation of costimulatory molecules and secretion of cytokines such as IL-12 [112]. An irradiated CD40L-expressing K562 cell line together with IFN γ has been used to mature DCs for vaccination in melanoma patients. The DC vaccine produced IL-12p70, and the amount of IL-12p70 produced positively correlated with the induction of CD8⁺ T cell responses to HLA-A2-restricted gp100 as measured by tetramer binding assays [113]. TLR agonists have also been used to activate DCs. Activation of the TLRs on DCs can induce DC maturation, upregulation of costimulatory molecules, and production of cytokines and chemokines [114]. Moreover, simultaneous triggering of different TLRs on DCs mediates synergistic effects resulting in the “superinduction” of IL-12 [115, 116]. Consequently, TLR agonists have the potential of inducing optimal DCs to stimulate an effective immune response while concurrently conditioning the environment *in vivo* to favor the development of immune response. For MDDCs, the TLR3 agonist poly-ICLC combined with TNF α , IL-1 β , IFN α , and IFN γ has been used in clinical trials to mature DCs. DCs matured with this cocktail are called α -type-1 polarized DCs (α DC1). They produce high levels of IL-12p70, migrate in response to CCR7 ligand, and induce cytotoxic T lymphocyte-mediated responses against tumor-associated antigen (TAA) [117, 118]. Administration of α DC1 in patients with high-grade glioma proved to be safe and immunogenic and resulted in progression-free status in 9/22 patients lasting at least 12 months [119]. A clinical grade TLR4 agonist LPS has also been used to mature DCs for vaccination in ovarian cancer patients

[120]. LPS-matured DCs produced IL-12p70 and induced tumor-specific T cell responses in melanoma patients. A modified, less toxic version of LPS called 3-*O*-deacylated monophosphoryl lipid A (MPLA), which is derived from *Salmonella minnesota* R595, has been approved for use in human and is used in combination with Cervarix. MPLA matures DCs and together with IFN γ was demonstrated to produce high levels of IL-12 and induce CD8⁺ T cell responses *in vitro* [121]. A new synthetic TLR4 agonist, glucopyranosyl lipid A, has been used in combination with anti-DEC205 that targets HIV gag p24 to DCs in mouse studies. It upregulates CD86 and CD40 and induces IL-12p70 production from DCs *in vivo* [122]. It is now being used in human studies as an immune modulator with and without anti-PD-1 (NCT02501473) to modify the TME in follicular non-Hodgkin's lymphoma.

Alternatively, a cocktail of commonly used preventative vaccines consisting of BCG (Bacille Calmette-Guerin)-SSI, Influvac, and Typhim, which contains agonists of TLRs, has been used to mature DCs *in vitro*. The DCs matured with this cocktail showed high expression of CD80, CD83, and CD86, was able to migrate, and produced IL-12p70 [123]. This cocktail of preventative vaccines was tested in a clinical trial to mature DCs for vaccination in melanoma patients. The DC vaccine induced T cell responses that coincided with longer overall survival; however, local and systemic grade 2 and 3 adverse events were observed in a number of patients and were attributed to the presence of BCG in the maturation cocktail, precluding any future use of the cocktail [124].

Another potential factor that can affect the quality of the DC vaccine is the timing of the maturation process. DC vaccines used in clinical trials are typically matured for 24-48 h and cytokine assessments are performed during that period [125]. However, most of the cytokines produced after DC maturation are produced within 24 h and they only regain the capacity to produce cytokines after the T cell encounter and CD40L ligation [126]. Thus, maturation protocols that are < 24 h may be more ideal so that the DC vaccines would still retain their ability to produce cytokines upon injection *in vivo*.

Therefore, the most ideal maturation method is one that induces the most effective APCs in order to induce the most successful immune response *in vivo*.

Antigen loading

DCs are typically loaded through incubation with peptides, proteins, RNA, or autologous/allogeneic tumor cells [79]. Peptides are loaded directly on the MHC molecules on the surface of the DCs, whereas the use of proteins and tumor cells requires processing into peptides

before loading on the MHC molecules.

Short synthetic peptides (8-15 aa) that correspond to defined CD8⁺ T cell epitopes within the defined TAAs have been used in clinical trials. However, use of these short peptides necessitates the knowledge of patient's HLA haplotypes and the defined epitopes that would bind to these specific haplotypes. Most recently, the use of synthetic long peptides (SLPs) that are 28-35 aa long has gained some favor. Due to their length, SLPs are preferentially taken up, processed, and presented by DCs through cross presentation [127] leading to the activation of CD8⁺ T cell responses as well as CD4⁺ T cell responses; however, the advantages of SLPs have been underappreciated until recently [128]. Clinical trials using SLPs covering the entire length of HPV-16 E6 and E7 sequences emulsified in Montanide ISA-51 adjuvant in patients with advanced HPV-16⁺ cervical cancer showed that the vaccine was safe and immunogenic [129, 130]. In another study, 79% of patients vaccinated with the HPV-16 SLPs with Montanide ISA-51 adjuvant had objective responses after 12 months with no evidence of virus in the original lesions in five patients [131]. On the basis of the success of these studies, vaccine formulations using SLPs have been further explored in cervical [132], ovarian [133, 134] and colorectal cancer [135, 136] and HIV infection [137] with evidence of induction of CD4⁺ and CD8⁺ T cell responses. An study of DCs with SLPs from NY-ESO-1 and Melan-A/MART-1 is currently being conducted in melanoma. The study is a randomized two arm study comparing the immunogenicity of DC vaccine to Montanide ISA-51 (NCT02334735), and should determine whether DCs can cross-present long peptides to T cells as has been shown to be the case when Montanide is used as the adjuvant.

DCs loaded with proteins [138] or lysates of autologous or allogeneic whole tumors or tumor cell lines [76, 139-142] have also been used in treating numerous cancers. The major advantages of this method are that (1) multiple epitopes can be presented on MHC molecules of different haplotypes, thus having the potential of inducing both CD4⁺ and CD8⁺ T cell responses to a wide spectrum of antigens, and (2) the necessary requirement for processing results in prolonged antigen presentation [143]. One way to enhance immunogenicity is to increase the exposure/release of damage-associated molecular patterns (DAMPs) derived from tumor cells [144] to enhance DC maturation [145]. Coincidentally, the most common procedure used for killing of autologous or allogeneic whole-tumor cells or tumor cell lines relies on multiple cycles of freeze-thaw, which have been demonstrated to release endogenous DAMPs [146]. Potent CD4⁺ and CD8⁺ T cell responses were detected in ovar-

ian cancer patients vaccinated with DCs activated with hypochlorous acid-oxidized autologous whole-tumor lysate generated via multiple cycles of freeze-thaw [120]. Most recently, a study in a mouse model of high-grade glioma showed that hypericin-based photodynamic therapy-activated DC vaccines increased survival, which was accompanied by an immunostimulatory shift in the brain contexture from regulatory T cells to Th1/cytotoxic T cells/Th17 cells [147]. Alternatively, DCs have been fused to autologous tumor cells with polyethylene glycol. The fusion allows for the tumor to be accessible to both the endogenous and exogenous pathways for activation of both CD4⁺ and CD8⁺ T cell responses. This approach was tested in patients with multiple myeloma and shown to be safe and effective, resulting in both CD4⁺ and CD8⁺ T cell responses [148]. These various approaches of antigen loading are currently being explored in larger studies. A phase II multicenter study in multiple myeloma using DCs fused to tumor cells led by Dr David Avigan is set to open. A phase III study using DCs with autologous tumor lysates (DCVAX-L) of glioblastoma is currently ongoing (NCT00045968).

Bacterial or viral vectors have been evaluated to target DCs with tumor antigens. For example, vectors derived from bacteria such as BCG, *Listeria monocytogenes*, *Salmonella*, and *Shigella*, and viruses including Canarypox virus, Newcastle disease virus, vaccinia virus, Sindbis virus, yellow fever virus, human papillomavirus, adenovirus, adeno-associated virus, and lentiviruses have all been explored in this capacity [75, 149-154]. These vectors have distinct advantages including the ability to insert genes encoding TAAs and to remove genes encoding virulence or replication factors for safety, and/or the ability to induce DC maturation. Human DCs infected with a killed but metabolically active (KBMA) *L. monocytogenes* strain encoding an epitope of Melan-A/Mart-1 induced the maturation of DCs, which were able to prime Mart-1-specific CD8⁺ T cells, resulting in the lysis of patient-derived melanoma cells *in vitro*. Furthermore, KBMA bacteria efficiently targeted APCs *in vivo* to induce protective antitumor responses in a mouse tumor model [155]. However, pre-existing immunity against a vector may ultimately limit its ability to induce immune responses *in vivo*. Lentiviral vectors have several properties that offer distinct advantages over current vectors used in clinical trials [156]. Although lentiviral vectors are typically less immunogenic, they can activate the immune system via endosomal or cytoplasmic sensors, i.e., TLRs, RIG-I, and PKR, etc. Use of lentiviral vectors can prime tumor-specific T cell responses *in vitro* [157], activate CD8⁺ T cells, and inhibit the growth of pre-existing tumors in mouse models [158, 159]. They

have the additional capacity of transducing non-dividing cells such as MDDCs, which are usually propagated from CD14⁺ or CD34⁺ progenitors [160-163]. Lentiviral vectors have been successfully used in adoptive T cell therapy to transduce T cells to express chimeric antigen receptors [164]. Most recently, a new technology was developed involving transduction of CD14⁺ monocytes with lentiviral vectors co-expressing GM-CSF/IL-4 and a melanoma antigen to generate so called "SmartDCs" (self-differentiated myeloid-derived antigen-presenting cells reactive against tumors-DCs). *In vitro* studies have shown that the SmartDCs expressed high levels of CD80, CD86, and those that were transduced with the tyrosinase antigen (SmartDC-TRP2) stimulated antigen-specific T cell responses [165]. It is possible that viral vectors may augment DC vaccination in the context of virus-induced inflammation and modulate the immunosuppressive TME [166] as has been observed with the use of oncolytic viruses.

DCs can also be transfected with mRNAs encoding TAAs. A number of studies have confirmed the ability of these DCs to induce potent tumor antigen-specific T cell responses *in vivo* [167-170]. Transfection of DCs with mRNA can be accomplished with the use of a cationic lipid, i.e., DOTAP, or electroporation. Electroporation has been demonstrated to be the most efficient method to introduce mRNA into DCs [171] and has been successfully used in preclinical [172-174] and clinical [175] studies. Autologous DCs electroporated with RNAs encoding CD40L and HIV antigens were demonstrated to be safe and capable of inducing HIV-specific immune responses [176]. Results from a phase II study using DCs co-electroporated with amplified tumor RNA and synthetic CD40L RNA (AGS-003) in combination with Sunitinib in metastatic clear cell renal cell carcinoma showed that the vaccine was well tolerated and that 62% of vaccinated patients experienced clinical benefit. Furthermore, the magnitude of increases in effector/memory T cells correlated with the overall survival [177]. The phase III trial (NCT01582672) is currently ongoing and a phase II trial was recently opened for non-small cell lung cancer (NCT02662634).

DC vaccines have typically targeted viral antigens, cancer-testis antigens (CT antigens), overexpressed antigens, and differentiation antigens [178]. The development of deep sequencing technologies has made it possible to identify unique mutations within the protein-encoding parts of the tumor genomes and to predict potential antigens. These antigens are not expressed in the normal human genome and therefore serve as effective stimulators of naive antigen-specific T cells [179]. In support of this, whole-exome sequencing and RNA

transcript analysis showed that tumors with high mutational burden had more CD8⁺ T cells [180] and that the cytolytic activity of NK and CD8⁺ T cells correlated with the mutational burden [181]. Moreover, three studies have suggested that mutational burden in melanoma and lung cancer correlates with improved survival in response to checkpoint inhibitors such as anti-CTLA-4 and anti-PD-1 antibodies [182-184]. Mutational burden alone does not necessarily correlate with improved survival however, and the response rate can be influenced by immune suppressive factors in the TME [185]. Altogether these studies suggest that mutated antigens deserve to be tested in the clinic where they could prove to be far more immunogenic than conventionally used antigens, e.g., CT antigens, overexpressed antigens, and differentiation antigens.

Indeed, DCs matured with GMP-grade CD40L-expressing K562 cells, IFN γ , poly IC, and R848, and loaded with neoantigen peptides derived from missense mutations in the tumor primed or boosted HLA-A2-restricted CD8⁺ T cell responses in patients with melanoma [186]. Additional factors such as pretreatment with cyclophosphamide and checkpoint blockade antibodies to eliminate or reduce regulatory T cells, and administration of DCs via intravenous infusion to allow better trafficking to secondary lymphoid tissues, may have contributed to the success of this study. These very encouraging results indicate that DC vaccination can be used to stimulate neoantigen-specific T cell responses in patients. Given that only a subset of predicted neoantigens were shown to be immunostimulatory there is a need to further refine prediction tools to identify high affinity immunogenic neoantigens that can prime both CD4⁺ and CD8⁺ T cell responses.

As noted earlier, DCs have the capacity to present non-conventional antigens. Newer classes of antigens include phosphopeptides, which are synthesized as a result of deregulated phosphorylation, citrullinated peptides, and antigens derived from noncoding DNAs that are expressed in cancer cells. In the latter case these include antigens expressed by endogenous retroviral remnants. It remains to be seen where such antigens will fit into the realm of effective anti-tumor antigens in DC-based cancer vaccines.

Administration and migration of DCs

Migration to the lymph nodes is critical for inducing immune responses. DC vaccines have been administered intradermally, subcutaneously, intravenously, intranodally, or intratumorally, but the optimal route of administration has yet to be established. Large amount of intradermally administered 111-indium-labeled DC vaccine

loaded with melanoma peptides remained at the injection site, lost viability, and was cleared within 48 h, with < 5% reaching the draining lymph nodes [187]. Intratumoral administration of DC vaccines showed retention at the injection site with little detected in the draining lymph nodes [188, 189], indicating failure of the vaccines to reach their targets.

Newer strategies include administration of DC vaccines via more than one route, i.e., intradermally plus intravenously to induce a systemic response, and administration directly into the lymph nodes (intranodally). Intranodally administered DC vaccine loaded with melanoma peptides redistributed to multiple lymph nodes within 30 min of injection; however, despite direct delivery of DCs into the lymph nodes, immunologic responses elicited were comparable to [187] or not better [189] than intradermally administered DC vaccines. Many factors likely contributed to this outcome. One possibility is the technical difficulty of properly injecting a vaccine to lymph nodes; improper injection could disrupt the lymph node architecture. Another possibility is that intranodal injection delivers all the vaccines into the lymph nodes; whereas with intradermal injection, only the viable, mature, and fully functional DCs migrate into the lymph nodes to stimulate T cell responses. It may not matter how many DCs reach the lymph nodes as one study has shown that reducing the number of cells injected improved migration of DCs [190]. Pre-conditioning the site of injection has been identified as a way to increase DC vaccine migration. First tested in mice by Martin-Fontecha *et al.*, preconditioning with the cytokine TNF or activated DCs improved lymph node homing and efficacy of DC vaccines [6]. Similarly, topical application of TLR7 agonist imiquimod, which recruits T cells and DCs into the skin, also enhanced DC vaccine migration [7]. In a recently published study, Mitchell *et al.* [191] showed that pre-conditioning the site of DC vaccine injection with tetanus/diphtheria toxoid (Td) vaccine improved DC migration in patients with glioblastoma multiforme by inducing CCL3 levels resulting in CCL21 upregulation and increased lymph node migration of the DC vaccine. The increased DC vaccine migration was associated with improved overall survival. However, the immunogenicity may partly be attributed to the viral antigen targeted by the DC vaccine. Therefore, generating a local inflammatory response at the site of the DC vaccine injection may serve to improve DC migration. Further studies are required to comprehend how these strategies might be incorporated into DC vaccines in order to enhance the arrival of DCs into the lymph nodes. Other unknown factors are whether these DCs directly elicit T cell responses as suggested in animal studies [192] or if a component of

the maturing DCs is in turn cross-presented by resident DCs either through uptake of dying cells, trogocytosis or through a process called “cross-dressing” [193]. In animal tumor models, it is clear that the CD103⁺ DCs acquire antigen from tumor cells and migrate to the draining lymph nodes to activate CD8⁺ T cells [50]. Intravenous injection of DCs has the advantage that injected cells can rapidly access secondary lymphoid tissue. Mouse studies of melanoma have shown that intra-tumorally injected mature DCs can modulate the TME, reversing an immune suppressive TME into one that supports T cell attraction and ultimately tumor control [194]. In this case a deficiency of CCL4 due to Wnt signaling prevented the recruitment of DCs into the tumor bed, which could be bypassed by restoring DC numbers in the tumor itself. The administration of Flt3L under such circumstances together with supplying maturation stimuli may alleviate these barriers *in vivo* [195].

In vivo approaches to DC vaccination

Direct targeting of antigens to *in vivo* DCs to induce tumor-specific immune responses [196, 197] bypasses the expensive, labor-intensive, and often difficult to standardize and scale up *ex vivo* DC generation process. *In vivo* targeting allows for vaccines to be produced on a larger scale and perhaps, more importantly, allows for direct stimulation of the activation of natural DC subsets at multiple sites *in vivo*. Early approaches of targeting *in vivo* DCs involved engineering irradiated tumor cells to secrete GM-CSF [198, 199] to stimulate the recruitment and enhance the function of APCs. Delivered in a neoadjuvant setting, GVAX comprising an allogeneic pancreatic cell line secreting GM-CSF resulted in the recruitment of T cells into pancreatic tumors [200]. In a phase II trial of GVAX combined with and without recombinant live attenuated *L. monocytogenes* engineered to secrete mesothelin (CRS-207) and with low dose cyclophosphamide, CD8⁺ T cell responses induced were associated with longer overall survival in patients with advanced pancreatic cancer [201]. However, a phase 2b study failed to show improved overall survival in patients treated with the combination or CRS-207 alone compared with chemotherapy. Prolonged GM-CSF production in the TME has been associated with disease progression in some experimental models and in a phase III trial in patients with hormone therapy-refractory prostate cancer [199], possibly due to recruitment of myeloid suppressor cells or differentiation of myeloid precursors into immature tolerogenic DCs [202, 203]. The effect of GM-CSF is therefore context-dependent. Systemically elevated GM-CSF is not effective whereas when locally delivered, as in the form of GVAX, can recruit DCs.

Newer approaches involve the *in vivo* targeting of DCs via ligation of CD40, and CLR. Agonist CD40 antibodies have been used to mimic the activation of DCs, which is required for the release of pro-inflammatory cytokines and enhancement of T cell activation. CD40 antibody was given after chemotherapy in mouse models and was shown to have a synergistic effect on inducing CD8⁺ T cell responses that eliminated the tumor [204]. A phase I study of the agonist CD40 antibody in combination with gemcitabine in patients with advanced pancreatic cancer demonstrated some clinical activity with associated cytokine release syndrome that is manageable in an outpatient setting [205]. Interestingly, anti-CD40 may modulate the TME by activating monocytes [206]. In a mouse model, CD40 agonistic antibody could overcome resistance to checkpoint blockade and improve survival [207]. The production of anti-CD40 antibodies that preferentially target DCs and convert M2 macrophages into M1 macrophages, and have reduced toxicity *in vivo*, remains challenging.

CLRs are also attractive targets as DC subsets are known to express different CLRs, which are involved in the recognition and capture of many glycosylated antigens [208]. Studies in animals targeting antigens to CLRs (DEC205, Langerin, and Clec9A) resulted in effective generation of T cell responses [209]. Antigen targeting using Clec9a in animal models showed the induction of cellular and humoral immune responses, which is essential for effective vaccination [210–212]. Antigen-specific T cell responses were reported in humanized mouse models of Clec9a-mediated antigen targeting indicating the potential of clinical application of this strategy [213].

Recent studies using DEC205-targeting antibodies have successfully induced immune responses to both cancer antigens [214] and HIV antigens [215, 209]. Although the majority of these studies were done in mice, human studies using antigens targeted to DC-SIGN [216], MR [217], and DEC205 [218] have demonstrated successful induction of tumor-specific T cell responses in subsets of patients, although in some the responses were weak. The correlation with clinical responses remains unclear and larger studies will be needed to further evaluate the efficacy. Clinical trials of anti-DEC205-NY-ESO-1 are currently ongoing in ovarian cancer (NCT02166905), acute myeloid leukemia (NCT01834248), various solid cancers (NCT01522820, NCT02661100), and melanoma (NCT02129075). Preliminary analysis of the clinical trial in melanoma showed that Flt3L with anti-DEC205-NY-ESO-1 treatment considerably increased the numbers of innate immune cells and induced NY-ESO-1-specific immune responses (N Bhardwaj, personal communication). Another potential approach is via targeting of antigen

with the chemokine for XCR1, a chemokine receptor exclusively expressed on CD141⁺ DCs [219–221].

Newer approaches of targeting *in vivo* DCs may involve delivery of tumor antigens via systemic administration of RNA lipoplexes. The lipoplexes protect the RNA from degradation, whereas the RNA triggers activation of pDCs and the subsequent release of type I IFNs [222]. A phase I dose escalation study evaluating the safety of intravenous administration of the RNA-LPX vaccine, which encodes four TAAs in melanoma patients (NCT02410733) is ongoing. Preliminary immune monitoring data from the study shows induction of IFN α and strong antigen-specific T cell responses in the first three patients evaluated [222].

Finally, given their safety profile, DC vaccines are being tested in immune prevention. Immune prevention is an intervention designed to halt cancer recurrence. It is currently being evaluated in patients with Lynch syndrome or colorectal cancer with microsatellite instability. These patients have germline mutations in DNA repair genes. In an ongoing clinical trial (NCT 01885702) mutations in various genes, e.g., genes encoding caspase, TGF β receptor, and CEA, have been identified in these patients. Some of these patients have T cells that recognize these mutated antigens in the circulation suggesting the development of spontaneous immunity and justifying an immunization approach in these high-risk cohorts [223]. Other studies have delivered DC vaccine into ductal carcinoma *in situ* lesions targeting HER-2/neu. A decline and/or eradication of HER-2/neu expression was demonstrated [224]. A long-term goal, therefore, in high risk of recurrence settings is to immunize patients prophylactically.

Future approaches

The safety of DC vaccines has been well established in clinical trials conducted thus far. Observed side effects are relatively mild and transient; they usually include fever, injection site reactions, adenopathy, and fatigue. The immunogenicity of DC vaccines has also been established in most clinical trials with evidence of induction of tumor-specific T cell responses in many patients. A comprehensive meta-analysis of DC vaccination for prostate cancer and renal cancer has shown that DC vaccines have induced tumor-specific immune responses in 77% and 61% of the vaccinated patients, respectively [225]. However, in our opinion, the clinical efficacy of DC vaccines remains unsubstantiated. Many reasons account for this but the most likely contributing factor is the quality of the DC vaccine. We still lack a consensus on the optimal method of DC preparation for majority of the DC

vaccines described thus far. The DC vaccines may be phenotypically mature as evidenced by upregulation of costimulatory molecules [113]. They may also produce IL-12, a key factor for inducing vaccine efficacy [113]. But it is unknown how much IL-12 is actually sufficient and whether the DC vaccine will still be producing the IL-12 upon injection or encounter with T cells. Given the correlation between IL-12 production and vaccine efficacy, it will be important to know that IL-12 is indeed still in production after injection [113]. Furthermore, the majority of the DC vaccine may not migrate to the lymph nodes to stimulate the T cells, and the DC vaccine itself may not directly function to prime immunity, which requires cross-presentation by lymph node resident DCs instead [4]. Those that do reach the lymph nodes, i.e., after exposure to PGE₂, may induce regulatory T cells and Th2 responses. Additional factors that contribute to the limited success of the DC vaccines are intrinsic to the tumors. The TME employs mechanisms that include loss of tumor antigen expression, alteration of MHC molecules, lack of costimulation, expression of inhibitory ligands, induction of regulatory T cells, expression of IDO, and/or production of immunosuppressive cytokines [58, 226–229]; these may inhibit the capacity of DC vaccines to induce immune responses against the tumor cells. Therefore, studies focused on optimizing the DC preparation in combination with approaches that overcome immune evasive mechanisms by the TME will likely improve efficacies of DC vaccines *in vivo*. Newer strategies addressing these issues are in development.

In combination

Combination strategies may be the key to the success of DC vaccines. The FDA approvals of checkpoint blockades made the most significant impact on immunotherapy but not just as a monotherapy option and also as part of combination therapy. A recent study showed that DCs electroporated with mRNAs encoding CD40 ligand, CD70, constitutively activated TLR4 and one of four melanoma-associated antigens (gp100, tyrosinase, MAGE-A3, or MAGE-C2) fused to an HLA class II-targeting signal (DC-LAMP) (also called TriMixDC-MEL) in combination with ipilimumab in patients with pretreated advanced melanoma achieved a very encouraging overall response rate of 38% [230]. A number of DC vaccine combination studies with other checkpoint blockade agents are currently ongoing (NCT02677155, NCT01067287, and NCT01441765). A systematic approach to testing the combination is needed to determine the optimal timing of vaccination or whether using one or two checkpoint inhibitors would be more effective.

Reducing tumor-elicited suppression first may be the

key to the success of DC vaccination. By first modulating the TME to make it more receptive, DC vaccination may have a better chance to induce a more effective immune response that can infiltrate the tumors. Talimogene laherparepvec (T-VEC or OncoVEX), derived from human herpes simplex virus 1 and encoding GM-CSF, became the first oncolytic virus that was FDA-approved for the treatment of metastatic melanoma [231]. Oncolytic viruses have had an impact on melanoma presumably through their effects not only directly on tumor cells but also by activating innate immunity and inducing tumor-specific immunity thus effectively reducing immune suppression in the TME [232]. Alternatively, intratumoral approaches that mimic viral infection such as intratumoral injections of poly-ICLC [233] or CpG [234], but without its associated side effects of inducing dominant antiviral immunity would be preferable [235]. Another possible method to modulate the TME is through direct activation of the STING pathway. Intratumoral administration of STING agonists stimulated IFN β production leading to regression of established and distant tumors in mouse models [236]. Finally as alluded to earlier, intratumoral administration of Flt3L may have multipronged effect of increasing DC numbers, mobilization of DCs into the TME, and providing a maturation stimulus, which altogether can substantially improve anti-tumor T cell priming.

New DCs

Another tactic being explored is the differentiation of DCs from human pluripotent stem cells (hPSCs) including induced pluripotent stem cells and embryonic stem cells. There are currently two methods for differentiating DCs from hPSCs: one involves the use of embryoid bodies, an aggregate-like structure mimicking embryonic development, and the other depends on co-culture with stromal cell lines. Both involve a multi-step process utilizing various growth factors such as BMP-4, VEGF, GM-CSF, SCF, Flt3L, and IL-4 at key intervals to induce the differentiation [237]. Most importantly, this novel source of DCs has potential for large-scale production using bioreactors [238], unlike the currently used methods, which are highly dependent on operational quality standards. Most recently, a lentiviral vector was used to transduce hPSCs to express the tumor antigen MART-1. The transduced hPSCs were differentiated into DCs that expressed typical DC markers and matured in response to TNF stimulation. Furthermore, the DCs were able to prime MART-specific CD8⁺ T cell responses [239]. Therefore, this new technology has the potential to improve DC vaccination by ensuring an unlimited source of DCs with the added feature of being able to transduce the antigen directly, ensure presentation on MHC class I

and stimulate CD8⁺ T cell responses. The development of stem cell-derived DCs for clinical purposes will also enable the isolation of DC subsets that are profoundly more immunogenic, e.g., the CD141⁺ DC subset.

The clustered regulatory interspaced short palindromic repeat-associated 9 (Cas9) system is a powerful and versatile genetic engineering technology that gives the most exceptional control over genome editing. Briefly, the guide RNA binds to the target sequence and subsequently, Cas9, an endonuclease, targets the specific DNA sequence that pairs with the guide RNA [240]. This technology can be used to manipulate DCs to prevent expression of inhibitory molecules (i.e., PD-L1) and cytokines (i.e., IL-10) to improve their effectiveness *in vivo*, or to preferentially drive CD8⁺ T cell differentiation. Our laboratory has successfully applied this approach to delete target genes in DCs.

Conclusion

The plethora of different methods used in clinical trials and lack of controlled comparative studies make it virtually impossible to determine the most effective method to use DC vaccines. However, these earlier studies have provided valuable information and contributed to expansion of our knowledge of DCs and tumor biology. Combinatorial approaches and development of new technologies should continue to improve efficacy of DC vaccines in the clinic. Phase III trials testing the efficacy of MDCCs-based vaccines pulsed with autologous tumor preparations in renal cancer (NCT01582672) and GBM (NCT00045968) are ongoing and should provide valuable information.

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