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DENDRITIC CELL SUBSETS AS VECTORS AND TARGETS FOR IMPROVED CANCER THERAPY

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SUMMARY

Current active immunotherapy trials have shown durable tumor regressions in a fraction of patients. However, clinical efficacy of current vaccines is limited, possibly because tumors skew the immune system by means of myeloid-derived suppressor cells, inflammatory type 2 T cells and regulatory T cells (Tregs), all of which prevent the generation of effector cells. To improve the clinical efficacy of cancer vaccines in patients with metastatic disease, we need to design novel and improved strategies that can boost adaptive immunity to cancer, help overcome Tregs and allow the breakdown of the immunosuppressive tumor microenvironment. This can be achieved by exploiting the fast increasing knowledge about the dendritic cell (DC) system, including the existence of distinct DC subsets. Critical to the design of better vaccines is the concept of distinct DC subsets and distinct DC activation pathways, all contributing to the generation of unique adaptive immune responses. Such novel DC vaccines will be used as monotherapy in patients with resected disease and in combination with antibodies and/or drugs targeting suppressor pathways and modulation of the tumor environment in patients with metastatic disease.

Keywords

dendritic cells; cancer; vaccines; priming

1. INTRODUCTION

Vaccines against infectious agents demonstrate the power of manipulating the immune system. Vaccines have spared countless numbers of people from polio, measles, tetanus etc (Nossal 1997), even though they have not been designed according to immunological principles (Doherty et al. 2006). Immunology has the potential to identify vaccines, i.e., antigen-specific, durable, non-noxious preventions and therapies for infections, cancer, allergy, autoimmunity, transplantation. This has formed a conceptual basis for the development of therapeutic vaccines in cancer. Molecular identification of human cancer antigens has ushered in a new era of antigen specific cancer immunotherapy specifically targeting these antigens. Initial attempts (e.g. peptides, DNA vaccines, viral vectors and first generation DC-based vaccines) have thus far met with a limited success in the clinic.

However, cancer vaccines are in a renaissance era due to recent clinical trials showing promising immunological data and some clinical benefit to the patients. For example, an active immunotherapy product, sipuleucel-T (APC8015) based on antigen-loaded and GM-CSF activated PBMCs, appears to contribute to prolonged median survival in phase III trials in patients with prostate cancer (Higano et al. 2009). Similarly, a randomized phase II trial of a poxviral-based vaccine targeting PSA (PROSTVAC) in men with metastatic castration-resistant prostate cancer showed improved overall survival in patients who received PROSTVAC compared to patients receiving control vectors (Kantoff et al. 2010). While these first generation positive randomized phase II/III clinical trials need further analysis and mechanistic studies, they underline the therapeutic potential of the immune system that can be tapped into. Vaccines act through DCs which induce, regulate and maintain T cell immunity. Here we summarize our recent studies aimed at a better understanding of the DC system to unravel the pathophysiology of cancer and to design novel cancer vaccines.

2. DENDRITIC CELLS

Generating the right type of immune response can be a matter of life and death. In leprosy, for instance, the tuberculoid form of the disease is characterized by a Type 1 response which keeps the disease in check, while the lepromatous form induces an often fatal Type 2 response (Yamamura et al. 1991). These responses are under the control of DCs (Banchereau and Steinman 1998; Steinman and Banchereau 2007). DCs reside in peripheral tissues and in lymph nodes where they are poised to capture antigens (Ags). DCs present processed protein and lipid Ags to T cells via both classical (MHC class I and class II) and non-classical (CD1 family) antigen presenting molecules (Heath and Carbone 2009) (Figure 1). In the steady state, non-activated (immature) DCs present self-antigens to T cells, which leads to tolerance (Hawiger et al. 2001; Steinman et al. 2003). DCs induce immune tolerance in a number of ways including i) T cell deletion (Fairchild and Austyn 1990; Zal et al. 1994; Volkman et al. 1997); ii) induction of T cell unresponsiveness (Hawiger et al. 2004); and iii) activation of regulatory T cells (Tregs) (Jonuleit et al. 2000; Akbari et al. 2001; Wing and Sakaguchi 2010; Zheng et al. 2010). Once activated (mature), antigen-loaded DCs are geared towards the launching of antigen-specific immunity (Finkelman et al. 1996; Brimnes et al. 2003) leading to the T cell proliferation and differentiation into helper and effector cells. DCs are also important in launching humoral immunity partly due to their capacity to directly interact with B cells (Jego et al. 2005; Qi et al. 2006) and to present unprocessed antigens (Zhong et al. 1997; Wykes et al. 1998; Bergtold et al. 2005; Batista and Harwood 2009).

2.1 Human Dendritic Cell Subsets

To allow resistance to infection and tolerance to self, DCs are endowed with two critical features: subsets and functional plasticity (Steinman and Banchereau 2007). The two major subsets are the myeloid DCs (mDCs) and the plasmacytoid DCs (pDCs). The best studied human mDC subsets are those from skin, where three subsets can be identified. The epidermis hosts only Langerhans Cells (LCs) while the dermis displays two mDC subsets, CD1a⁺ DCs and CD14⁺ DCs, as well as macrophages (Zaba et al. 2007; Klechevsky et al. 2008; Merad et al. 2008; Nestle et al. 2009).

2.1.1 Dermal DCs, antibody responses and IL-12—In the mid 90, we observed that CD14⁺ DCs derived from CD34⁺ hematopoietic progenitor cells (HPCs) induce CD40-activated naïve B cells to differentiate into IgM-producing plasma cells through the secretion of IL-6 and IL-12 (Caux et al. 1997). A decade later, we found that CD14⁺ DCs, but not LCs, induce naïve CD4⁺ T cells to differentiate into cells with properties of T follicular helper cells (Tfh) (Klechevsky et al. 2008), a CD4⁺ T cell subset specialized in B

cell help (King et al. 2008; Fazilleau et al. 2009). There, CD4⁺ T cells primed by CD14⁺ DCs help naïve B cells to produce large amounts of IgM, and switch isotypes towards IgG and IgA. Our recent studies in human indicate that acquisition of Tfh phenotype and function depends on IL-12p70 (Schmitt et al. 2009).

Thus, IL-12 appears to contribute to humoral immunity in humans through a direct path in DC-B interaction, and an indirect path in DC-T cell interaction and induction of Tfh cells. These findings might explain the modest clinical efficacy of systemic IL-12 administration in cancer patients (Motzer et al. 2001; Cheever 2008). Furthermore, the injection of IL-12 into tumor sites of head and neck cancer patients resulted in the activation of B cells in the draining lymph nodes, which was associated with their infiltration into tumor sites and tumor regression (van Herpen et al. 2008).

2.1.2 LCs and CD8⁺ T cell responses—LCs induce a robust proliferation of naïve allogeneic CD8⁺ T cells when compared to CD14⁺ DCs (Klechevsky et al. 2008). Furthermore, when pulsed with MHC class I peptides derived from tumor or viral antigens, LCs are far more efficient than CD14⁺ DCs in the priming of antigen-specific CD8⁺ T cells. LCs are also efficient in cross-presenting peptides from protein antigens to CD8⁺ T cells. CD8⁺ T cells primed by LCs show high avidity in tetramer binding assays and express higher levels of cytotoxic molecules, such as granzymes and perforin. Accordingly, they are remarkably more efficient in killing target cells; in particular tumor cells that express low level of peptide/HLA complexes (Klechevsky et al. 2008). IL-15 might explain the remarkable effects of LCs on the development of Cytotoxic T Lymphocyte (CTL) responses (Mohamadzadeh et al. 2001; Dubsky et al. 2007; Klechevsky et al. 2009). Thus, the two different arms of adaptive immunity, i.e., humoral and cellular arms, might be differentially regulated by the two skin mDC subsets (Figure 2). Such framework might be of capital importance for the understanding of the immune alteration in malignancy and for development of novel and improved vaccination strategies against cancer, as well as chronic infections.

2.1.3 Plasmacytoid DCs—Plasmacytoid DCs (pDCs) are considered as the front line in anti-viral immunity owing to their capacity to rapidly produce high amounts of type I interferon (Siegal et al. 1999; Liu 2005). Similar to mDCs, pDCs display a remarkable functional plasticity. Thus, pDCs exposed to viruses, such as live influenza virus, are able to launch memory responses by inducing the expansion and differentiation of antigen-specific memory B and T lymphocytes into plasma cells (Jego et al. 2003), and CTLs (Fonteneau et al. 2003; Di Pucchio et al. 2008), respectively. On the contrary, pDCs activated with CpG or IL-3/CD40L induce in vitro IL-10-secreting regulatory CD4⁺ T cells (Ito et al. 2007) as well as suppressor CD8⁺ T cells through the expression of ICOS ligand (Gilliet and Liu 2002).

Human pDCs, in fact, are composed of two subsets, distinguished by the expression of CD2 (Matsui et al. 2009). CD2^{high} pDCs are more potent than the CD2^{low} pDCs to induce allogeneic T cell proliferation. These different functional properties of CD2^{high} pDCs and CD2^{low} pDCs are associated to distinct transcription profiles, differential secretion of IL12 p40 and differential expression of co-stimulatory molecule CD80 on activation. Additional studies will be necessary to understand the biological role of these two pDC subsets.

2.2 DCs in tumor environment

Numerous studies in humans have concluded that DCs can infiltrate tumors. We found that breast cancer tumor beds are infiltrated with immature DCs. In contrast, mature DCs are found in the peri-tumoral areas in ~60% of cases (Bell et al. 1999). A number of studies have suggested that DCs can contribute to tumor development. Our studies in breast cancer

indicate that tumor cells polarize mDCs into a state that drives the differentiation of naïve CD4⁺ T cells into IL-13-secreting T cells (Aspord et al. 2007). These Type 2 T cells in turn facilitate breast tumor development in xenograft model as it can be partly inhibited by administration of IL-13 antagonists (Figure 3). The role of Th2 cells was further established in a spontaneous mouse breast cancer model, where Th2 cells facilitate the development of lung metastasis through macrophage activation (DeNardo et al. 2009). In several other mouse tumor models, IL-13 produced by NKT cells induces myeloid cells to make TGF- β that inhibits CTL functions (Berzofsky and Terabe 2008). Thus, type 2 cytokines are involved in tumorigenesis through various mechanisms. mDCs can also have direct interactions with tumor cells as shown in multiple myeloma where they directly promote the survival and clonogenicity of tumor cells (Kukreja et al. 2006; Bahlis et al. 2007).

pDCs have been found in approximately 10% of breast carcinomas and are associated with poor prognosis (Treilleux et al. 2004). The infiltrating pDCs produce little type I IFN upon TLR ligation (Hartmann et al. 2003). This inhibition appears to depend on the ligation of ILT7 on pDCs binding by BST2 expressed on tumor cells (Cao et al. 2009). Likewise, in ovarian carcinoma, tumor-infiltrating pDCs do not induce effector CD8⁺ T cell responses, but rather promote the differentiation of IL10⁺ CCR7⁺ CD8⁺ Tregs (Wei et al. 2005). Finally, pDCs may promote tumor angiogenesis by the secretion of proangiogenic cytokines (Curiel et al. 2004; Coukos et al. 2005).

DC can fight back tumors at least through two pathways: an indirect one with the induction of potent CTL responses, and a direct one through DC-dependent tumor cytotoxicity. For example, pDCs appear to directly contribute to the anti-tumor activity of in vivo-administered Imiquimod (TLR7 ligand), which is used for the treatment of basal cell carcinoma (Urošević et al. 2005; Panelli et al. 2007; Stary et al. 2007).

Clearly, understanding the functions of DCs in the tumor bed represents an important area of future investigations and exploitation for therapy. An interesting strategy would be to rewire their molecular pathways from “pro-tumor” DCs into “anti-tumor” DCs.

3. DENDRITIC CELLS IN VACCINATION AGAINST CANCER

3.1 Outcomes of current DC vaccination trials

Ex vivo-generated DCs have been used as therapeutic vaccines in patients with metastatic cancer for over a decade and early studies have been discussed in detail elsewhere (Palucka et al. 2007). While a fraction of patients can experience durable tumor regressions (Palucka et al. 2006), the most common outcome of the current DC vaccination protocols is a demonstration of expanded antigen-specific immunity, most often using IFN- γ ELISPOT, but no durable objective tumor regression.

Altogether, three outcomes emerge from our studies:

1) *No immune response.* Patients of this group usually progress quickly. These patients mount immune responses to control antigens such as KLH or viral peptides (Flu-M1 or CMV). In vitro experiments indicated that T cells of several patients can be primed to differentiate into CTLs with specificity for multiple melanoma antigens (Berard et al. 2000). Thus, tumor antigen-specific CD8⁺ T cells are kept anergic rather than deleted. This inability to mount immune responses to tumor antigens in vivo might be at least partly related to the presence of tumor antigen-specific Tregs (Vence et al. 2007; Andrews et al. 2008). Tregs limit the onset of protective immunity through several mechanisms, for example by eliminating DCs in lymph nodes (Boissonnas et al. 2010). As discussed later, the control of Tregs becomes a key target to address for the coming vaccination trials. 2)

Immune response without clinical response. The most common outcome of current DC vaccination protocols is the induction of immune responses in the absence of clinical responses. This might in part be explained by the quality of the elicited T cells including their capacity to migrate into tumors and penetrate tumor stroma (Gajewski 2007). Improved immunomonitoring is expected to provide insights into the mechanisms of immune efficacy as discussed hereunder (Butterfield et al. 2008; Tahara et al. 2009). 3) *Immune response and clinical response.* Vaccination with DCs can elicit therapeutic immunity. These patients represent a formidable opportunity for the development of cancer immunotherapy. The challenge is two-fold. First, to establish the immunological mechanism that allowed tumor eradication. Second, we need to find ways to increase the fraction of patients experiencing durable tumor regression and/or prolonged survival.

3.2 The quality of elicited antigen-specific immune responses

Establishing causative links in clinical studies is a difficult task which often requires large patient cohorts. The current data suggest an association between the tumor-specific CD8⁺ T cell responses and clinical outcomes. In our view, four critical components will determine whether the induced immune response will be therapeutic: 1) the quality of elicited CTLs; 2) the quality of induced CD4⁺ helper T cells; 3) the elimination and/or non-activation of Tregs; and 4) the breakdown of immunosuppressive tumor microenvironment.

Indeed, the immune responses elicited by the first generation DC vaccines might not be of the quality required to allow the rejection of bulky tumors. For example, the induced T cells might not migrate into the tumor lesions (Appay et al. 2008; Harlin et al. 2009). Furthermore, low avidity T cells might be unable to recognize peptide-MHC class I complexes on tumor cells and/or to kill them (Appay et al. 2008). Finally, the tumor microenvironment might inhibit effector T cell functions, for example by action of myeloid derived suppressor cells and Tregs as summarized in recent reviews, respectively (Gabrilovich and Nagaraj 2009; Menetrier-Caux et al. 2009).

The recent progresses in immunomonitoring of specific immune responses in the blood and at the tumor site should help us address these questions (Palucka et al. 2006; Vence et al. 2007; Butterfield et al. 2008; Janetzki et al. 2009; Tahara et al. 2009). Modern approaches including polychromatic flow cytometry rather than the analysis of a single cytokine (e.g., IFN- γ ELISPOT) and/or frequency of tetramer positive cells will contribute to a better assessment of the quality of the immune responses elicited in the patients (Kammula et al. 1999; Lee et al. 1999). Indeed, several studies, mostly performed in the context of HIV vaccines, have led to the conclusion that a mere measurement of the frequency of IFN- γ secreting CD8⁺ T cells is insufficient to evaluate the quality of vaccine-elicited immunity (Wille-Reece et al. 2006; Appay et al. 2008; Seder et al. 2008).

4. BUILDING ON DENDRITIC CELL SUBSETS TO IMPROVE CANCER VACCINES

4.1 Optimal DCs

The results summarized above prompted us to hypothesize that DCs with the properties of LCs might prove to be the best ones for the generation of strong cellular immunity (Figure 2). In line with this, the combination of cytokines used to differentiate monocytes into DCs play a critical role in determining the quality of the elicited T cell responses. For example, DCs generated with GM-CSF and IL-15 display the phenotype and characteristics of LCs. In particular, they are more efficient in priming melanoma-antigen specific CD8⁺ T cells in vitro than DCs derived with GM-CSF and IL-4 (Mohamadzadeh et al. 2001; Dubsy et al. 2007). Thus, vaccination with IL15-DCs might elicit stronger CD8⁺ T cell responses that

might lead to improved clinical responses. We are currently initiating such a clinical trial in patients with malignant melanoma. The selected method for activating DCs also represents a critical parameter is the DC activation pathway. First, immature (non-activated) DCs induce antigen specific IL-10 producing T cells (Dhodapkar et al. 2001; Dhodapkar and Steinman 2002). Second, IL-4 DCs activated with a cocktail of IFN- α , polyI:C, IL-1 β , TNF, and IFN- γ induce up to 40 times more melanoma-specific CTLs in vitro than DCs matured with the “standard” cocktail of IL-1 β /TNF/IL-6/prostaglandin E₂ (PGE₂) (Mailliard et al. 2004; Fujita et al. 2009; Giermasz et al. 2009). Additional studies will be necessary to establish the therapeutic value of these newer generation DC vaccines in patients. These studies are critical to the understanding of the human immune system because they permit us to assess in vivo the type of immune responses elicited by human DCs generated in different cytokine environments.

This in turn is essential for building a novel approach to vaccination that is based on the delivery of antigens directly to DCs in vivo using chimeric proteins that are made of an anti-DC receptor antibody molecularly fused to a selected antigen (DC targeting). Studies in mice demonstrate that the specific targeting of antigen to DCs in vivo results in considerable potentiation of antigen-specific CD4⁺ and CD8⁺ T cell immunity if the DC maturation signal is provided (Hawiger et al. 2001; Bonifaz et al. 2002; Bonifaz et al. 2004). Otherwise, tolerance ensues (Hawiger et al. 2001). Thus, selection of appropriate adjuvant is also a critical parameter for the induction of the immunity of the desired type. Although TLR-ligands are widely considered to promote protective immunity against infectious agents, selecting the appropriate ligand will be critical. For instance, TLR2 ligation, which promotes the induction of Tregs rather than Th1 or Th17 cells (Manicassamy et al. 2009), does not appear to be a preferred option for cancer vaccines.

These pioneering studies have been already extended to demonstrate the targeting of tumor antigens to DCs (Caminschi et al. 2009) and Langerhans cells (LCs) in animal models (Flacher et al. 2008; Flacher et al. 2009) and the generation of anti-tumor immunity (Wei et al. 2009). The therapeutic success of these vaccines will build on the recent knowledge and progress in our understanding of the biology of human DC subsets, cutaneous myeloid DCs (mDCs) in particular.

4. 2 “Ideal” antigens

Assuming that appropriate solutions are identified to reverse immunosuppression, there is a need for an “ideal” set of target antigens. An “ideal” antigen is one which is necessary for cancer cells to survive and/or for which strong immunity able to reject the tumor and prevent its growth can be elicited.

Candidate tumor antigens include: i) unique (mutated) antigens; and ii) shared self-antigens including cancer/testis antigens and tissue differentiation antigens (Gilboa 1999; Vlad et al. 2004; Boon et al. 2006; Parmiani et al. 2007). The choice between these types of antigens for vaccination could be viewed as a choice between inducing immunity (mutated antigens) or breaking tolerance and inducing autoimmunity (self antigens). The debate about which type of antigen will be more efficient is still open. Mutated antigens are postulated to present several advantages, for example their specific T cell repertoire should not be deleted as they are not recognized as “self” by immune cells (Parmiani et al. 2007). Shared antigens are attractive as they might allow us to establish “generic” vaccines, however the enthusiasm for these antigens might be dampened because of their: i) relatively weak immunogenicity due to the negative selection of high affinity auto-reactive cells; and ii) the existence of antigen specific T regs (Hoos et al. 2007).

Perhaps the most compelling evidence of active in vivo tumor antigen-specific immune responses arises from the study of paraneoplastic neurologic disorders (PNDs) that led to the discovery of onconeural antigens (Darnell 1996). PNDs develop as remote effects of systemic malignancies. The discovery of onconeural antibodies led to the proposal that paraneoplastic cerebellar degeneration (PCD), associated to breast and ovarian cancer, is an autoimmune disorder mediated by the humoral arm of the immune system. These antibodies permitted the cloning of the cdr2 antigen, a protein with a coil/leucine zipper domain. It has now been shown that the disease is due to the development of cdr-2 specific CD8⁺ CTL (Albert et al. 1998). The list of onconeural antigens is growing and, besides cdr2, two other antigens such as Nova and amphiphysin appear as potential targets of the immune system (Floyd et al. 1998; Rosin et al. 1998).

An important shift in the selection of antigen targets might be brought about by the identification of cancer stem cells (Jordan et al. 2006; Polyak and Hahn 2006; Rossi et al. 2008). While a majority of studies have focused on eliminating mature cancer cells with limited proliferation capacity, it seems more efficient to target the self-renewing cancer stem cells. The importance of stem cell associated antigens in malignancy can be best illustrated by the presence of SOX-2-specific immunity in patients with monoclonal gammopathy (Spisek et al. 2007). This immunity is lost in patients who developed multiple myeloma suggesting differential antigenic targets at pre-malignant and malignant stages. In fact, the major factor from the immunization point of view is the linkage between expression of genes associated with pluripotency and those expressed in cancer. Ideal target genes would be those shared between cancer cells and embryonal cells, which are necessary for cancer cell survival but not expressed in adult stem cells (Dhodapkar 2010).

Thus far, all antigenic targets are protein antigens whose peptides can be presented on the cell surface in the form of complexes with classical MHC molecules (Townsend et al. 1985). However, tumors express altered lipids and sugars that can be bound by CD1 molecules on APCs and presented to NKT cells as well as T cells (Beckman et al. 1994; Fujii et al. 2002; Hava et al. 2005). These lipid antigens might possibly be harnessed for improved vaccination.

4. 3 Combining DC vaccines with other therapies

In view of the remarkable diversity of regulatory/suppressive pathways present in patients with metastatic cancer, any durable clinical response elicited by vaccination with DCs is already a remarkable achievement. However, to improve the outcomes in metastatic disease, DC vaccines need to be combined with other therapies that offset the suppressive environment created by the tumor (Dougan and Dranoff 2009). Such combination regimens will involve several drugs that target different pathways (Figure 4).

In particular antibodies, or other soluble antagonists such as engineered receptors, can be exploited for the blockade of suppressive cytokines in the tumor microenvironment such as IL-10 (Moore et al. 2001), IL-13 (Terabe et al. 2000), TGF- β (Li et al. 2005; Terabe et al. 2009) and VEGF (Gabilovich 2004; Rabinovich et al. 2007). They can also be used to block inhibitory ligand:receptor interactions (Melerio et al. 2007) by acting on antigen presenting cells such as tumor or DCs (for example anti-PD-L1) or on lymphocytes as illustrated by anti-CTLA-4 (Peggs et al. 2006; Peggs et al. 2009) and/or anti-PD1 (Day et al. 2006; Curran et al. 2010; Pilon-Thomas et al. 2010). In contrast, agonistic antibodies (Gabilovich 2004; Rabinovich et al. 2007) might further promote co-stimulation of effector T cells as for example with anti-CD137 (Watts 2005), a ligand for 4-1BB (Maus et al. 2002). Just as different tumors are treated with different combinations of cytostatic drugs and targeted therapies, we foresee development of clinical protocols combining DC vaccines with individualized adjunct therapies.

CONCLUDING REMARKS

The considerable progresses made in the knowledge of DC biology as well as effector/regulatory T cell biology clearly open the avenues for development of vastly improved clinical protocols. These, optimized vaccines eliciting strong and long-lived antigens-specific CD8+ T cell immunity will be offered to patients with early stage disease. For patients with late stage disease strategies that combine novel highly immunogenic vaccines and immunomodulatory antibodies will have high impact on enhancing therapeutic immunity in cancer by simultaneously increasing the potency of beneficial immune arms and offsetting immunoregulatory pathways (Figure 4). These optimized therapeutic strategies will be tailored to the patient and to the specific suppressive pathways that the patient displays (Figure 5).

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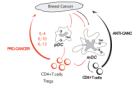
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**Figure 1. Dendritic cells**

DCs reside in the tissue where they are poised to capture antigens (Geissmann et al. 2010). During inflammation, circulating precursor DC enter tissues as immature DC (Geissmann et al. 2010). DCs can encounter pathogens (e.g.: viruses) directly, which induce secretion of cytokines (e.g.: IFN- α); or indirectly through the pathogen effect on stromal cells. Cytokines secreted by DCs in turn activate effector cells of innate immunity such as eosinophils, macrophages and NK cells. Microbe activation triggers DCs migration towards secondary lymphoid organs and simultaneous activation (maturation). These activated migratory DCs that enter lymphoid organs display antigens in the context of classical MHC class I and class II or non-classical CD1 molecules, which allow selection of rare circulating antigen-specific T lymphocytes. Activated T cells help drive DCs toward their terminal maturation, which allows lymphocyte expansion and differentiation. Activated T lymphocytes traverse inflamed epithelia and reach the injured tissue, where they eliminate microbes and/or microbe-infected cells. B cells, activated by DCs and T cells, migrate into various areas where they mature into plasma cells that produce antibodies that neutralize the initial pathogen. Antigen can also reach draining lymph nodes without involvement of peripheral tissue DCs and be captured and presented by lymph node resident DCs (Itano et al. 2003).

**Figure 2.**

DC s as tools for vaccination. We envision that targeting antigens and activation of distinct mDC subsets, with different specializations, will result in the generation of a broad and long lived immune protection. Thus, the most efficient cancer vaccines might be those that will target LCs thereby allowing the maximal stimulation of cellular immune responses and the generation of long-term memory protection.

**Figure 3.**

DCs as targets for therapy. Cancer cells attract immature DC possibly through chemokines such as MIP3 alpha and/or SDF-1. The DC can then be either blocked or skewed in their maturation, for example by VEGF, leading to induction of polarized CD4+T cells that promote the expansion of cancer cells (pro-cancer) at the expense of CD8+T cells that can cause tumor regression (anti-cancer). An interesting strategy would be to rewire their molecular pathways from “pro-cancer” DCs into “anti-cancer” DCs for example with antibodies or DC activators.

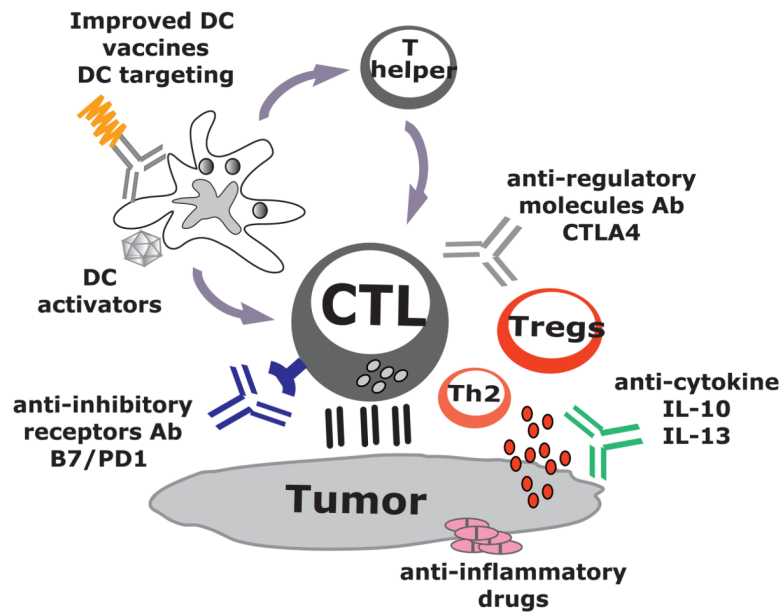


Figure 4.

DC vaccines in combination therapies. Current active immunotherapy trials have shown durable tumor regressions in a fraction of patients. However, clinical efficacy of current approaches is limited, possibly because tumors invade the immune system by means of myeloid-derived suppressor cells, inflammatory type 2 T cells and regulatory T cells (Tregs). To improve the clinical efficacy of immunotherapies, we need to design novel and improved strategies that can boost adaptive immunity to cancer, help overcome Tregs and allow the breakdown of an immunosuppressive tumor microenvironment. This can be achieved by developing combination therapies targeting these three major components.

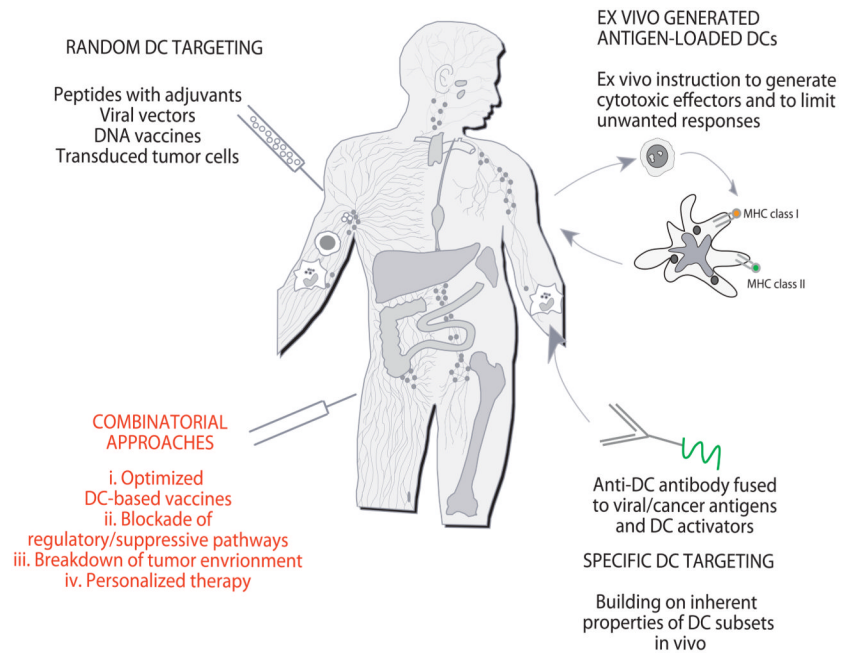


Figure 5. Approaches to DC-based immune intervention in cancer

1) Vaccines based on antigen with or without adjuvant that target DCs randomly. That might result in vaccine antigens being taken up by a “wrong” type of DCs in the periphery which might lead to “unwanted” type of immune response. Vaccine antigens could also flow to draining lymph nodes where they can be captured by resident DCs; 2) Vaccines based on ex-vivo generated tumor antigen-loaded DCs that are injected back into patients; and 3) specific in vivo DC targeting with anti-DC antibodies fused with antigens and with DC activators. 4) Next generation clinical trials will test optimized DC vaccines combined with patient-adjusted approaches to block Tregs and to breakdown the tumor environment. These therapies will be tested in pre-selected patients thereby leading to personalized therapy.