

Assessing T-cell responses in anticancer immunotherapy

Dendritic cells or myeloid-derived suppressor cells?

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Abbreviations: APC, antigen presenting cell; CTL, cytotoxic T lymphocyte; CTLA4, cytotoxic T lymphocyte-associated protein 4; DAMP, damage-associated molecular pattern; DC, dendritic cell; ESC, embryonic stem cell; G-CSF, granulocyte colony-stimulating factor; GM-CSF, granulocyte macrophage colony-stimulating factor; IDO, indoleamine 2,3 deoxygenase; IL, interleukin; iNOS, inducible nitric oxide synthase; LPS, lipopolysaccharide; Ly6C1, lymphocyte antigen 6 complex, locus C1; Ly6G, lymphocyte antigen 6 complex, locus G; MAMP, microbe-associated molecular pattern; M-CSF, macrophage colony-stimulating factor; MDSC, myeloid-derived suppressor cell; OVA, ovalbumin; PBMC, peripheral blood mononuclear cell; PD-1, programmed cell death 1; PD-L1, PD-1 ligand 1; PGE₂, prostaglandin E₂; PRR, pattern recognition receptor; TAA, tumor-associated antigen; TCR, T-cell receptor; TGFβ, transforming growth factor β; Treg, regulatory T cell

Since dendritic cells operate as professional antigen-presenting cells (APCs) and hence are capable of jumpstarting the immune system, they have been exploited to develop a variety of immunotherapeutic regimens against cancer. In the few past years, myeloid-derived suppressor cells (MDSCs) have been shown to mediate robust immunosuppressive functions, thereby inhibiting tumor-targeting immune responses. Thus, we propose that the immunomodulatory activity of MDSCs should be carefully considered for the development of efficient anticancer immunotherapies.

Introduction

Cancer is caused by the uncontrolled growth of transformed cells, initially forming a localized (primary) lesion and then colonizing distant organs (a process that is known as metastatic dissemination). Both these manifestations of cancer can significantly interfere with the physiological functions of the organism. Malignant cells exhibit consistent alterations in protein expression, survival and proliferation, mostly originating from the accumulation of mutations (reflecting a high degree of genetic instability) and epigenetic changes. Thus, neoplastic cells usually express an array of mutated proteins that provides

them with some degree of immunogenicity (quasi antigens). At least theoretically, such an acquired immunogenicity allows the immune system to identify and destroy cancer cells.

However, the immune system must overcome 2 major obstacles to effectively fight cancer. The first of such barriers is represented by the standard immunological tolerance toward self antigens, which impedes the activation of quasi-antigen-specific T cells. The second barrier stems from various immunosuppressive mechanisms set in place by neoplastic cells, which are largely responsible for the failure of conventional immunotherapeutic anticancer regimens. Such a systemic state of immunosuppression is caused by the expansion of potent immunomodulatory cells, including (but not limited to) myeloid-derived suppressor cells (MDSCs). Anticancer immunotherapy thus attempts at overcoming these obstacles by stimulating the immune system through a variety of procedures and interventions. In this regard, dendritic cell (DC)-based approaches deserve special attention. The refinement of ex vivo DC expansion protocols has boosted the development of several immunotherapeutic strategies against cancer, as DCs operate as central controllers of innate and adaptive responses. However, the administration of DCs loaded with tumor-associated antigens (TAAs) or previously treated with potent immunostimulants to cancer patients results in limited therapeutic responses, especially as compared with the expectations raised by preclinical data. Here, we review the systems that are currently available for the ex vivo evaluation of immunotherapeutic anticancer regimens, as well as the reasons for the limited predictive value of preclinical results obtained with this approach. We propose that testing immunomodulatory strategies ex vivo should focus not only on DCs but also on the immunosuppressive cells that are found in the tumor microenvironment, hence faithfully mimicking physiological

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conditions. This might allow for the identification (and further development) of immunotherapeutic regimens that are truly capable of overcoming the immunosuppressive effects of the tumor microenvironment.

Recognition of Tumor-Associated Antigens by the Immune System

Cancer immunotherapy relies on activating or boosting pre-existing tumor-specific cytotoxic T lymphocyte (CTL) responses. Generally, antigenic peptides are recognized by specific T-cell receptors (TCRs) once complexed with MHC molecules and exposed on the surface of antigen-presenting cells (APCs) (Fig. 1). The specific recognition of MHC-peptide complexes by TCRs results in the delivery of a first activating signal to T lymphocytes. However, such a signal is insufficient for the activation of T cells, implying that APCs must trigger additional signaling events to elicit antigen-specific immune responses.¹ These signals are delivered by the binding of a wide range of ligands to specific receptors, which can be stimulatory or inhibitory, expressed on the surface of T cells. The integration of these signal transduction cascades regulate the degree of T-cell activation.² A classical co-stimulatory interaction is represented by the binding of CD80, which is expressed on DCs, to CD28, which is found at the surface of T cells. Conversely, a strong inhibitory signal is delivered upon the interaction of CD80 with cytotoxic T lymphocyte-associated protein 4 (CTLA4) as well as upon the binding of programmed cell death 1 (PDCD1) ligand 1 (PD-L1, official name CD274) with PDCD1 (best known as PD-1) (reviewed in refs. 1 and 3). A variety of additional ligand-receptor interactions can regulate T-cell co-stimulation (Fig. 1). Finally, cytokines that are present in the microenvironment where antigen presentation occurs confer T cells with specific effector functions, mainly through the differentiation of specific CD4⁺ helper T-cell subtypes (reviewed in ref. 1).

The immune responses against infectious agents rely on the recognition of microbial molecules via pattern recognition receptors (PRRs) expressed by professional APCs, including DCs. The binding of such microbe-associated molecular patterns (MAMPs) to PRRs favors the maturation of DCs, resulting in the upregulation of co-stimulatory molecules, enhanced cytokine secretion and increased expression of MHC molecules (reviewed in ref. 4). As they mature, DCs migrate to secondary lymphoid tissues where they present antigenic peptides to antigen-specific T cells. However, most TAAs are either aberrantly overexpressed self proteins or quasi-antigens. Thus, the frequency of circulating TAA-specific T cells is low, mostly because of their efficient removal by clonal deletion in the thymus (central tolerance). The autoreactive T cells that survive thymic selections either bear TCRs that display a low affinity for cognate antigens or differentiate into regulatory T cells (Tregs).⁵ Nonetheless, the immune system is capable of recognizing, controlling, and eliminating cancer cells.^{6,7} Thus, the efficient activation of autoreactive TAA-specific T cells as well as the inhibition of the immunosuppressive activity of Tregs stand out as key goals for anticancer immunotherapy. Another significant challenge in this context is to neutralize the

profound state of systemic immunosuppression that characterizes cancer patients with a large tumor burden. Tumors actively secrete a variety of molecules that act in the bone marrow to divert myeloid differentiation, resulting in the accumulation of MDSCs.⁸ MDSCs are able to exit the bone marrow, distribute to peripheral organs, and actively infiltrate neoplastic lesions, hence inhibiting antitumor immune responses via both antigen-specific and non-specific mechanisms.⁹⁻¹¹ The expansion of MDSCs in cancer patients is responsible (at least for a large part) for the inefficacy of standard immunotherapeutic regimens. Thus, MDSCs have attracted great interest from the fields of experimental and clinical tumor immunology.¹² In summary, an ideal immunotherapeutic intervention against cancer would have to (1) stimulate the presentation of TAAs to T cells, while (2) counteracting the immunosuppressive activity of Tregs and MDSCs.

The Discovery of Dendritic Cells and their Impact in Biomedical Research

After the recent award of the Nobel Prize to Ralph Steinman (for the first time in history, posthumously), it is worthy to briefly comment on the impact that his work had on immunology. Indeed, although Langerhans cells (a particular type of DCs) had previously been described by Langerhans, the discovery of conventional DCs can be attributed to Steinman,¹³ who in 1973 demonstrated the capacity of these cells to either strongly activate¹⁴ or inhibit T cell-mediated immune responses.^{15,16} The characteristics of each DC lineage described since have extensively been described elsewhere.¹⁷

The line of investigation focusing on DC biology has significantly been stimulated in the mid-1990s, when systems for the differentiation of myeloid DCs *ex vivo* were first developed. These protocols relied on the murine bone marrow or purified monocytes as a starting material, and on granulocyte macrophage colony-stimulating factor (GM-CSF) as a central differentiation stimulus.^{18,19} Within 7 y from the publication of *ex vivo* DC differentiation protocols, the number of papers on DCs had increased from about 80 to approximately 500 per y, and this figure climbed further to 1000 per y 10 ys afterwards (Fig. 2). Because of its simplicity and reproducibility, the production of DCs *ex vivo* has *de facto* revolutionized the study of DC biology.

The differentiation of myeloid DCs *ex vivo* presents many advantages for biomedical research. First, cell numbers are not a limiting factor in this context. Indeed, close to 50×10^6 DCs can be easily obtained from the material collected from a single mouse within 1 wk. Although the human system is comparatively less efficient, mostly owing to a limited initial supply of peripheral blood mononuclear cells (PBMCs), it is still an amenable protocol of key clinical relevance. During differentiation, DCs tend to mature, implying that experimental assessments can easily be performed at different DC maturation stages. The large numbers of DCs that can be obtained *ex vivo* are compatible with their use for the systematic monitoring of immunomodulatory agents. DC cultures also provided a means to thoroughly study DC responses to large collections of maturation stimuli, and in

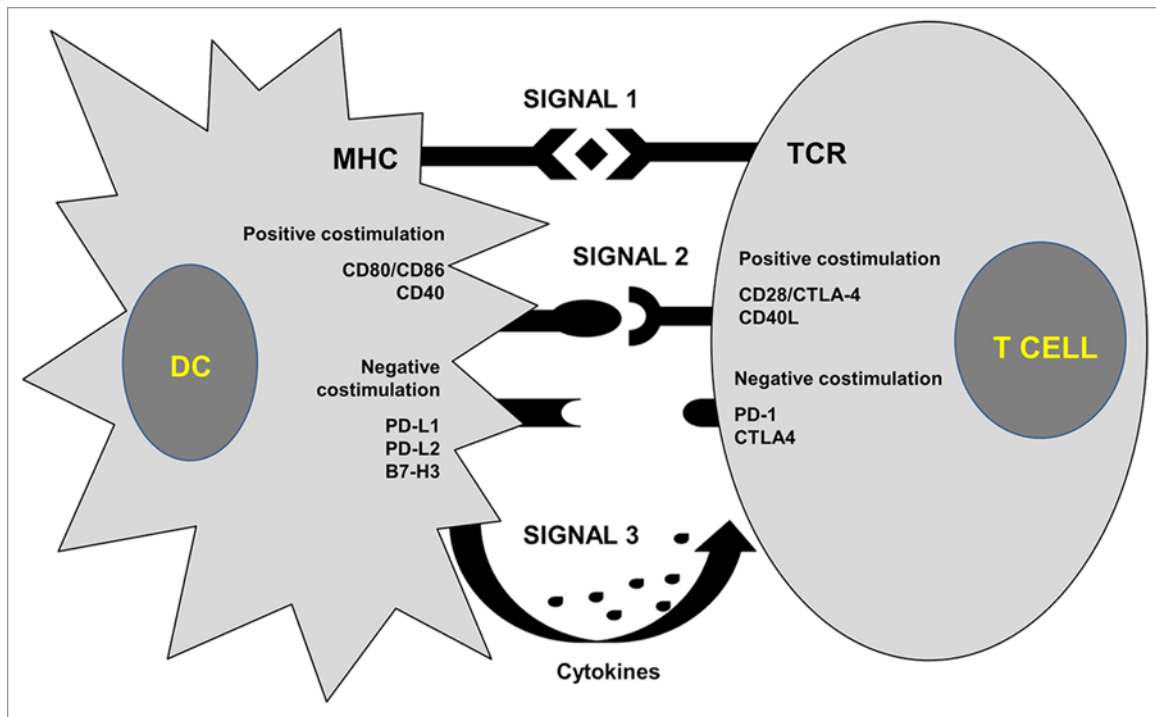


Figure 1. Activation of T cells by antigen-presenting cells. Antigenic peptides (rhomboids) complexed with MHC molecules on the surface of dendritic cells (DCs) are recognized by cognate T-cell receptors (TCRs), delivering a first activatory signal (signal 1) to T cells (top). A second signal (signal 2) is delivered to T cells upon the integration of positive (activatory) and negative (inhibitory) co-stimulation, originating from the interaction of specific receptors expressed on the T-cell surface and their ligands. A third signal (signal 3) is delivered by cytokines found in the microenvironment where antigen presentation occurs, which are often secreted by antigen-presenting DCs. Such a cytokine priming generally directs the polarization of T-cell responses. CD40L, CD40 ligand; CTLA4, cytotoxic T lymphocyte-associated protein 4; PD-1, programmed cell death 1; PD-L1, PD-1 ligand 1; PD-L2, PD-1 ligand 2.

particular how these agents influence intracellular signal transduction and antigen presentation by DCs (reviewed in ref. 4). These studies have generated a significant amount of experimental evidence in support of the so-called “danger signal” model, originally put forward by Charles Janeway and Polly Matzinger to explain the general regulation of immune responses.^{20,21} This model de facto establishes a link between innate and adaptive immunity. In brief, it postulates that APCs possess receptors for the recognition of a wide range of molecules that can be found in microbial products (MAMPs) as well as in cellular components that are released in the course of inflammation or trauma (the so-called damage-associated molecular patterns, DAMPs). The recognition of MAMPs and DAMPs triggers the phenotypic and functional maturation of DCs, resulting in the upregulation of both MHC and co-stimulatory molecules. Thus, only when professional APCs encounter danger signals of this type, their antigen-presenting capability is strongly improved. Although the danger model has weak points, it is both elegant and simple. The development of protocols for the differentiation of DCs ex vivo has allowed for the systematic study of the effects of MAMPs and DAMPs on DCs, in terms of functional responses and intracellular signaling (reviewed in ref. 22). Along similar lines, the refinement of procedures for the co-culture of DCs and T cells has allowed for the dissection of the cellular/molecular mechanisms of antigen-presentation and T-cell polarization, as recently reviewed by Liechtenstein and colleagues.¹

Dendritic Cell-Based Vaccines in Anticancer Immunotherapy

The possibility to differentiate DCs ex vivo in large-scale had an immediate therapeutic application: the development of anticancer vaccines based on these professional APCs. DCs are key regulators of immune responses and are possibly the immune cells with the most prominent adjuvant effects. Immature DCs exhibit an intense phagocytic activity, which provides them with a consistent amount of antigenic peptides, including TAAs, for loading on MHC molecules.¹⁷ Furthermore, DCs are highly susceptible to genetic engineering via viral vectors, including adenoviral, retroviral, lentiviral, and poxviral particles, as well as non-viral systems.^{17,23} Finally, the maturation state of DCs can be manipulated to boost or suppress immune responses.²² These APCs are therefore ideal vaccines for anticancer immunotherapy. DCs generated ex vivo have indeed been extensively employed to assess T-cell responses to infectious agents in vitro²⁴⁻²⁷ and harnessed for the development of anticancer immunotherapeutic regimens.²⁸⁻³⁴ Moreover, DCs have been successfully used to suppress immune responses, in both animal models and humans.^{15,16,35-40}

Undoubtedly, the refinement of DC production systems has promoted the use of DCs in various clinical settings, including anticancer immunotherapy.⁴¹ Since these methods have become part of the research routine, scientists generally follow the same

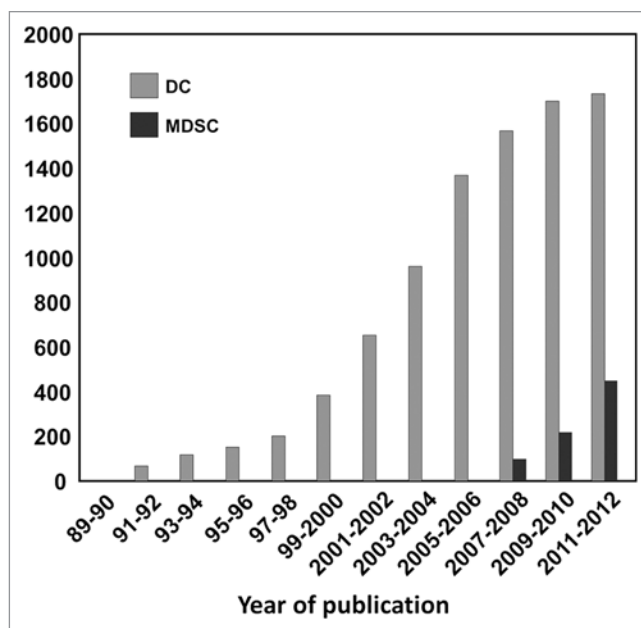


Figure 2. Number of publications dealing with conventional dendritic cells and myeloid-derived suppressor cells. The approximate number of publications dealing with conventional dendritic (DCs) or myeloid-derived suppressor cells (MDSCs), as retrieved by searching PubMed (<http://www.ncbi.nlm.nih.gov/pubmed>) for entries whose title contains the term “dendritic cell” or “myeloid-derived suppressor cells,” is represented as a function of publication biennium.

pipeline for developing new antineoplastic interventions, which involves the following steps: (1) treatment of DC cultures with a specific agent, to assess its ability to stimulate DC maturation or alter cytokine secretion; (2) assessment of how this agent influence DC-mediated antigen presentation to T cells, *ex vivo*; (3) testing of DCs *in vivo*, in experimental models that allow for the assessment of T-cell responses, and (4) testing of DCs activated according to optimal protocols in therapeutic/preventive experimental tumor models, *in vivo*.

Undoubtedly, a sizeable amount of preclinical research based on DCs amplified *ex vivo* has generated interesting insights into the immunobiology of DCs, often highlighting promising therapeutic activity.^{17,42-45} Surprisingly, however, these encouraging preclinical results have not always been translated into a clinical success in cancer patients.^{46,47}

Why are there such significant discrepancies between preclinical and clinical data on the immunotherapeutic anticancer profile of DCs? At least in part, this reflects the conditions employed to assess the antigen-presenting capacities of DCs *in vitro*, which are highly controlled. While this is certainly advantageous from an experimental point of view, this experimental setting may not appropriately reproduce the conditions in which tumor-specific immune responses occur. For instance, these assays often rely on well-defined, immunogenic TAAs or even model xenoantigens, such as ovalbumin (OVA).^{33,48,49} The use of tumor lysates to load DCs for clinical applications may also be disadvantageous, as tumor lysates (1) contain both defined and undefined TAAs, and (2) are generally tolerogenic, hence inhibiting the therapeutic activity of DCs unless additional immunostimulatory agents are also employed.⁵⁰⁻⁵⁶ Finally, upon infusion, DCs must counteract

the strongly immunosuppressive state of cancer patients. This is perhaps the most prominent factor accounting for the discrepancy between the preclinical and clinical immunotherapeutic activity of DCs against cancer.

From an experimental point of view, much effort and resources would be saved if candidate treatments were appropriately assessed early in the course of preclinical development. To this aim, it would be ideal to either establish DC-T cell co-culture assays in (immunosuppressive) conditions that closely resemble the tumor microenvironment, or utilize the immunomodulatory myeloid counterpart of DCs that robustly infiltrate neoplastic lesions: MDSCs.

MDSCs: Discovery and Definition

The dissemination of malignant cells from the primary lesion is in itself a key pathological feature of cancer. In addition, a major complication for cancer patients, especially individuals in the late stages of disease (when tumor load is generally consistent), is their state of systemic immunosuppression, which prevents the immune system from eliminating transformed cells. Growing neoplasms produce indeed a wide range of cytokines and metabolites that alter the differentiation of myeloid cells, facilitating the accumulation of cell populations exerting strong immunosuppressive effects.⁵⁷ Myeloid cells including tolerogenic DCs, tumor-infiltrating macrophages and granulocytes have been known for a long time to play an important role in various aspects of tumor progression, including neoangiogenesis. However, only recently specific subsets of myeloid cells have been identified as specialized immunosuppressive cells that accumulate in cancer patients.^{58,59} The recognition of MDSCs

as a specific cell lineage remains rather controversial. Still, recent experimental evidence indicates that MDSCs differ from other myeloid cells in many aspects. MDSCs comprise indeed a heterogeneous population of cells exhibiting cancer-specific phenotypic and functional characteristics. In mice, MDSCs were originally described as myeloid CD11b⁺ cells that express high levels of the granulocyte-specific epitope GR1. GR1 is shared by 2 surface markers, namely, lymphocyte antigen 6 complex, locus C1 (Ly6C1) and lymphocyte antigen 6 complex, locus G (Ly6G). Thus, based on the relative expression levels of these markers, murine MDSCs can be classified into a monocytic (Ly6C1^{high}Ly6G^{neg/low}) and granulocytic (Ly6C1^{high}Ly6G^{high}) subsets. As such, monocytic and granulocytic MDSCs can also be identified in cancer patients.⁶⁰ These cells, when isolated from tumor-bearing hosts, exhibit a pronounced capacity to functionally inhibit T cells, via both antigen-specific and non-specific mechanisms.⁶¹ The immunosuppressive functions of MDSCs mainly originate from the expression of enzymes that deplete the extracellular microenvironment of essential amino acids, such as inducible nitric oxide synthase (iNOS), indoleamine 2,3-deoxygenase (IDO) and arginase 1, as well as from the secretion of immunosuppressive cytokines such as interleukin (IL)-10 and transforming growth factor β (TGF β).⁶¹ The growing importance of MDSCs in biomedical research is well represented by the fact that, following their “official” definition, the number of papers dealing with these cells is steadily increasing (Fig. 2).

Differentiation of MDSCs Ex Vivo

As mentioned above, the establishment of a protocol for the amplification of conventional DCs ex vivo provided a significant boost to the corresponding area of biomedical research. The development of a similar system for the differentiation and amplification of MDSCs could hence entail another significant step forward. However, the current systems for the ex vivo differentiation of MDSCs are inefficient and fail to achieve significant levels of amplification. In addition, MDSCs isolated from tumor-bearing mice generally do not proliferate ex vivo in the presence of GM-CSF, and their survival is thus severely compromised.⁶² Considering the reports published so far, GM-CSF stands out as a key factor for MDSC differentiation, in vitro and in vivo. This is particularly interesting in view of the fact that GM-CSF-based therapies are currently used in patients affected by some types of neoplasms as an immunostimulatory regimen.^{63–66} Indeed, taking into consideration the key role of GM-CSF in the tumor-associated expansion of MDSCs, the administration of this cytokine as a standalone intervention might not be an ideal choice for anticancer immunotherapy, at least in some circumstances.^{67–69}

One of the first methods to obtain murine MDSCs from the bone marrow relied on high concentrations of GM-CSF coupled to lipopolysaccharide (LPS), a protocol allowing for the expansion of highly immature myeloid cells with robust immunosuppressive activities.⁵⁷ Possibly, these cells represented activated MDSCs, although nomenclature guidelines had not yet

been formulated at that time. Bone marrow-derived “immature” myeloid cells obtained with GM-CSF and LPS exerted strong immunosuppressive effects upon cell-to-cell contact (antigen-specific immunosuppression) and by expressing high levels of iNOS (non-specific immunosuppression). In line with these observations, the production of GM-CSF by breast carcinoma cells has soon been identified as one of the key drivers of MDSC expansion.⁷⁰ In this context, MDSCs could be generated upon the addition of GM-CSF to the bone marrow, and similar results could be obtained by using culture medium conditioned by breast carcinoma cells. Still, the authors of this paper only obtained 28% of differentiated cells exhibiting a CD11b⁺GR1⁺ phenotype, of which Ly6G^{high} cells did not exert immunosuppressive effects.⁷⁰ Moreover, the addition of GM-CSF-neutralizing antibodies did not completely abrogate the differentiation of MDSCs, suggesting that additional factors are required for the accumulation of these cells in cancer patients. Nevertheless, GM-CSF stands out as an important MDSC-polarizing cytokine produced by many cancer cells.^{57,70,71}

The addition of recombinant GM-CSF and IL-4 to culture media conditioned by various cancer cell lines (including EL4 lymphoma, LLC lung carcinoma, B16-F10 melanoma and C3 cervical carcinoma cells) reportedly induces the differentiation of MDSCs, correlating with the immunosuppressive effects exerted by tumor cells in vivo. However, also in this case the differentiation and proliferation of MDSCs ex vivo were rather inefficient. Indeed, relative amounts of MDSCs higher than 25–30% were hardly reached after 5 d of culture.⁹

A recent study has identified GM-CSF, granulocyte colony-stimulating factor (G-CSF) and IL-6 as factors that drive the differentiation of MDSCs from the bone marrow. In this context, the co-administration of GM-CSF and IL-6 combination generated MDSCs exerting robust immunosuppressive functions. Nonetheless, also the authors of this study pointed out that the recovery rate of MDSCs was comparable with the number of bone marrow cells initially plated.⁷² Thus, in this setting myeloid cell precursors had lost their proliferative capacity, differentiated and acquired potent immunosuppressive functions. Along similar lines, the differentiation of human MDSCs has been achieved upon 1 wk of incubation of CD33⁺ mononuclear cells with GM-CSF and IL-6.⁷³ Other cytokines found within neoplastic lesions and in the supernatants of cultured cancer cells have been shown to contribute to MDSC differentiation, although to a minor extent.⁷³ The authors of this study did not comment on the efficiency of differentiation, but pointed out that their cytokine cocktails also promoted the expansion of other cell lineages. Moreover, the expression of the transcription factor CCAAT/enhancer binding protein β (C/EBP β) was shown not to correlate with the immunosuppressive activity of MDSCs, in sheer contrast with previously published results.⁷²

The administration of recombinant GM-CSF coupled to macrophage colony-stimulating factor (M-CSF) drives the differentiation of murine MDSCs from the bone marrow in the absence of cancer cell-conditioned culture medium.⁷⁴ Interestingly, the addition of IL-13 appears to increase the

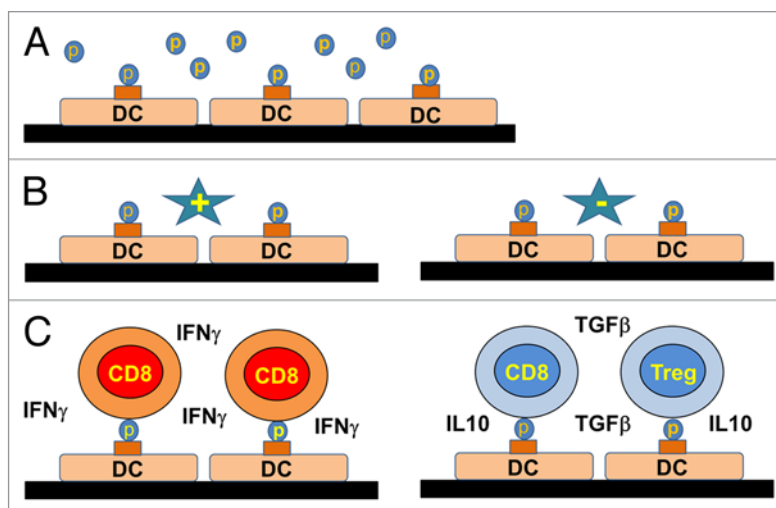


Figure 3. Dendritic cell-T cell antigen presentation assays for the preclinical evaluation of immunotherapeutic regimens. (A) Myeloid dendritic cells (DCs) differentiated ex vivo are loaded with the antigen of interest by overnight incubation. (B) Antigen-loaded DCs are then treated with a potential immunostimulatory (+) or immunosuppressive (-) agent. (C) Antigen-loaded DCs exposed to the agent of interest are incubated with transgenic antigen-specific CD8⁺ T cells. The elicitation of antigen-specific T-cell responses is then monitored by quantifying the production of interferon γ (IFN γ) by T cells and/or their proliferation. If necessary, the conversion of antigen-specific CD4⁺ T cells into regulatory T cells (Tregs) can also be monitored.

immunosuppressive activity of these cells, which have been shown to efficiently inhibit manifestations of the graft-vs-host disease in an arginase 1-dependent fashion. Nonetheless, also in this case the proportion of MDSCs obtained from bone marrow cell cultures did not exceed 40–50%.⁷⁴ Such a yield approaches those that would be required for clinical applications.

Another recent study developed a method to obtain large amounts of MDSCs from murine embryonic stem cells (ESCs) for therapeutic purposes.⁷⁵ Such ESC-derived MDSCs successfully inhibited graft-vs-host disease and were more immunosuppressive than tumor-derived MDSCs. However, the authors of this study used a mouse ESC line that overexpressed homeobox B4 (HoxB4), resulting in a net enhancement of myeloid differentiation. By this means, highly immunosuppressive MDSCs were obtained, including a Ly6C^{neg} subset that had never been described before in vivo.⁷⁵ This method to drive the differentiation of MDSCs is not straightforward and depends on a 3-stage process that involves complex cytokine cocktails. Still, this work demonstrated for the first time modified ESCs might constitute a source for high numbers of MDSCs.

Other protocols for the expansion of MDSCs ex vivo do not rely exclusively on recombinant GM-CSF, but also involve prostaglandin E₂ (PGE₂) and TGF β , both of which are abundant in tumor-derived exosomes.^{76,77} Indeed, antibody blocking either of these 2 molecules has been shown to abrogate the capacity of tumor-derived exosomes to drive the differentiation of MDSCs in vivo. Still, the production of MDSCs ex vivo by means of tumor-derived exosomes fail to yield murine MDSCs in percentages higher than 29% of cultured bone marrow cells.⁷⁶ PGE₂ in combination with GM-CSF, IL-4 and LPS has also been shown to promote the differentiation of human MDSCs ex vivo, presumably via a feedforward amplification loop resulting in the robust activation of cyclooxygenase 2 (COX2).⁷⁸

Putative Advantages of Using Large-Scale Ex Vivo MDSC-T Cell Assays

Many research groups, including us, routinely use bone marrow-derived or monocyte-derived DCs to assess the immunostimulatory potential of a range of interventions. To this aim, DCs are mainly employed in ex vivo presentation assays involving antigen-specific T cells (Fig. 3). These assays heavily rely on the presentation of model antigens such as OVA, as both CD4⁺ and CD8⁺ T-cell epitopes of OVA are well characterized. Other antigens including hemagglutinin and the nucleoprotein from influenza virus, as well as human molecules such as melan-A (MLANA, also known as MART1), are sometimes used in these assays.^{30,79} Frequently, we choose to engineer bone marrow-derived DCs with lentiviral vectors that drive the co-expression of OVA and DC maturation stimuli, allowing for the use OVA-specific transgenic CD4⁺ and CD8⁺ T cells in presentation assays.^{29,30,35} We have used this system to evaluate immunostimulatory as well as immunosuppressive treatments.⁴ Although the results of ex vivo antigen presentation assays correlate with therapeutic activity in some tumor models,²⁹ this does not always hold true. These inconsistencies are probably due to the strong immunosuppressive microenvironment generated by some neoplasms, which completely inactivate tumor-specific T cells. This might also explain discrepancies between the results of ex vivo assays and in vivo therapeutic profile of DC-based vaccines, as discussed above.

Based on these premises, we propose that the use MDSCs in antigen presentation assays would make up a more faithful model to assess the efficacy of immunotherapeutic anticancer regimens than that of DCs. Successful immunotherapeutic and chemotherapeutic approaches are known to inhibit the expansion and immunosuppressive activity of MDSCs.^{12,59,80} A range of treatments could therefore be systematically assessed

in MDSC-T cell antigen presentation assays to evaluate their ability to overcome MDSC-dependent immunosuppression. An experimental setup comparable to that generally employed for DC-T cell presentation assays (involving similar model antigens) could be applied to MDSC-T cell tests. However, as discussed above, the generation of cancer-specific MDSCs *ex vivo* is by far more cumbersome than that of DCs. In addition, the MDSCs obtained in this manner might lose their proliferative potential and/or plasticity, rendering their preservation in culture for the entire duration of the assay a challenge.

At least in part, these problems could be overcome with immortalized MDSC lines, as recently described by Apolloni and colleagues.⁸¹ One could argue that the genetic alterations required to immortalize these cells could significantly modify their phenotypic and functional profile. In particular, these authors employed *v-myc* and *v-raf*, which are indeed expected to affect several signal transduction cascades involved in proliferation and survival. As a matter of fact, immortalized MDSCs exhibited a modified phenotype, expressing increased levels of MHC class II molecules and no GR1.

Thus, what should we be looking for? Basically, a system that allows for the large-scale production of cancer-specific MDSCs *ex vivo*, similar to the protocols that have already been established for DCs. These MDSCs, which ideally should conserve their proliferative potential and differentiation plasticity, would constitute the basis for the development of an *ex vivo* test system that would faithfully resemble the tumor microenvironment. Moreover, a large-scale MDSC production system would allow for the high-throughput assessment of the effects of chemotherapeutic agents on MDSCs as well as for their proteomic/genomic profiling. All these strategies would accelerate the preclinical development of antineoplastic drugs, by favoring the early identification of agents that specifically target cancer-derived MDSCs.

In conclusion, the development of a protocol for the differentiation of DCs *ex vivo* has revolutionized

immunotherapy, setting up the basis for their application to neoplastic, infectious, and autoimmune disorders. We propose that a similar production system for MDSCs would be invaluable for the preclinical assessment of novel immunotherapeutic and chemotherapeutic agents. Such a system could be complementary to DC-T cell antigen presentation assays, in particular by mimicking robust immunosuppressive conditions. A significant effort is currently being devoted to the development of protocols for the expansion of cancer-specific MDSCs. Nonetheless, the methods published so far are quite heterogeneous, complicated, and generally characterized by low MDSC yields. To overcome this problem, some research groups have used modified ESCs that are particularly prone to myeloid proliferation or immortalized MDSC lines. However, the cells obtained with these systems may not resemble closely enough those that accumulate in the course of tumor progression and infiltrate neoplastic lesions *in vivo*. On a positive note, the area of tumor immunology specifically dealing with MDSCs is now speeding up and new isolation/production techniques may lead to the development of routinely applicable high-throughput MDSC-based T-cell assays.

Disclosure of Potential Conflicts of Interest

The authors declare no conflicts of interests.

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