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REVIEW

Targeting myeloid-derived suppressor cells augments antitumor activity against lung cancer

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Correspondence: Sherven Sharma Molecular Gene Medicine Laboratory, Veterans Affairs Greater Los Angeles Healthcare System, Los Angeles, CA 90073, USA Tel +1 310 478 3711 ext 41863 Fax +1 310 268 4807 Email ssharma@mednet.ucla.edu Abstract: Lung cancer evades host immune surveillance by dysregulating inflammation. Tumors and their surrounding stromata produce growth factors, cytokines, and chemokines that recruit, expand, and/or activate myeloid-derived suppressor cells (MDSCs). MDSCs regulate immune responses and are frequently found in malignancy. In this review the authors discuss tumor-MDSC interactions that suppress host antitumor activities and the authors' recent findings regarding MDSC depletion that led to improved therapeutic vaccination responses against lung cancer. Despite the identification of a repertoire of tumor antigens, hurdles persist for immunebased anticancer therapies. It is likely that combined therapies that address the multiple immune deficits in cancer patients will be required for effective therapy. MDSCs play a major role in the suppression of T-cell activation and they sustain tumor growth, proliferation, and metastases. Regulation of MDSC recruitment, differentiation or expansion, and inhibition of the MDSC suppressive function with pharmacologic agents will be useful in the control of cancer growth and progression. Pharmacologic agents that regulate MDSCs may be more effective when combined with immunotherapies. Optimization of combined approaches that simultaneously downregulate MDSC suppressor pathways, restore APC immune-stimulating activity, and expand tumor-reactive T cells will be useful in improving therapy.

Keywords: MDSCs, antigen-presenting cells, natural killer cell activation, T-cell activation, immunotherapy

Introduction

Lung cancer is a challenging health problem, with more than 1.1 million deaths attributed to lung cancer worldwide each year.¹ With current therapy, the long-term survival rate for the majority of lung cancer patients is low; thus, new therapeutic strategies are needed. Immunotherapy is an option; however, activation of the immune system alone is not sufficient for antitumor activity. Combined therapies that boost immune activation and reverse mechanisms of immune suppression ultimately will prove beneficial in the fight against lung cancer.

Modifications in oncogenes and tumor suppressor genes and/or epigenetic changes lead to tumor progression and local tissue invasion that cause the persistence of inflammatory cellular tumor infiltrates. The tumor causes the cellular infiltrates to sustain dysregulated inflammation, which immunologically is unresponsive to the tumor. Characterization of the infiltrate-tumor interactions may aid in the design of specific personalized anticancer treatments by reprogramming the tumor microenvironment. A significant proportion of the inflammatory cellular tumor infiltrates are myeloid-derived suppressor cells (MDSCs), which maintain an

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immunosuppressive environment,²⁻⁴ supporting tumor expansion in the surrounding tissue and at distant sites by facilitating angiogenesis and metastases.

MDSCs and the tumor microenvironment

MDSCs are a heterogeneous population of bone marrowderived cells comprising myeloid precursors of macrophages, granulocytes, and dendritic cells (DCs). In mice, MDSCs are identified by the cell surface expression of Gr1 and CD11b. Recently, MDSCs have been subcategorized based on the expression of Ly6C and Ly6G within the Gr1 cell population. CD11b+Ly6G+Ly6C^{lo} cells with multilobed nuclei and granulocytic-like morphology are called granulocytic MDSCs, and CD11b+Ly6G-Ly6Chi cells are referred to as monocytic MDSCs.5 In humans, MDSCs are identified by surface expression of CD33, and the lack of expression of markers of mature myeloid and lymphoid cells. They are typically CD11b+CD33+CD34+CD14- and vary in CD15, CD124, CD66, and major histocompatibility complex (MHC) class II expression.⁶ Melanoma MDSCs express CD14+CD11b+HLA-DR^{10/neg},7 and CD11b+CD14-CD15+CD33+ MDSCs have been described in non-small cell lung cancer patients.8 Because MDSC gene expression varies in different tumor types, the identification of a unique set of markers for human MDSCs has been challenging. Thus, phenotypic characteristics and the functional ability to suppress T cells are used to define MDSCs.

MDSC accrual, expansion, and activation in the tumor are dependent on tumor- or stromal-derived growth factors, cytokines, and chemokines. These include prostaglandin E2 (PGE2), matrix metalloproteinases, transforming growth factor beta (TGF- β), interleukin (IL)-10, vascular endothelial growth factor (VEGF), IL-1 beta (IL-1 β), IL-4, granulocyte-macrophage colony-stimulating factor (GM-CSF), IL-6, IL-13, S100A8/A9, stem cell factor, CCL2, CXCL5, CXCL12, toll-like receptor agonists, and tumor-derived HSP72.9-11 GM-CSF supports the survival and expansion of MDSCs in the tumor microenvironment.¹² The sources of GM-CSF include tumors and activated immune effectors such as T cells, natural killer (NK) cells, and DCs. IL-1 β recruits MDSCs; mice inoculated with tumor cells secreting IL-1ß exhibit elevated levels of MDSCs. IL-1βinduced MDSCs have elevated levels of reactive oxygen species with enhanced T-cell suppression.¹³ Furthermore, IL-6 secreted by tumors increases the frequency of MDSCs in tumors.14 Proinflammatory S100 proteins also cause MDSC

accumulation and immune suppression.^{15,16} The S100A8/A9 proteins are members of a large family of inflammatory and noninflammatory proteins produced by neutrophils, activated monocytes, tumor cells, and MDSCs. MDSCs have receptors for S100A8/A9 complexes and exhibit autocrine enhancement of S100A8/A9 levels. Antibody receptor blockade reduces the MDSC frequency in tumorbearing mice. Mice that are genetically S100A9 deficient are resistant to tumor challenge but become susceptible if MDSCs are adoptively transferred from wild-type mice. S100A8/A9 proteins affect MDSCs through two known mechanisms: (1) they block the further differentiation of myeloid precursors to differentiated DCs and macrophages through a signal transducer and activator of transcription 3-dependent mechanism and (2) they increase tumor MDSCs through a nuclear factor kappa-light-chain-enhancer of activated B cells-dependent pathway.^{15,16} This suggests that S100 protein-mediated accumulation of tumor MDSCs may serve as a useful therapeutic target to reduce immune suppression. Inflammation-mediated S100A8/A9 induction, MDSC accumulation, and the resulting immune suppression may represent a common mechanism of cancer-related escape from immune surveillance.

MDSC mechanisms of immune modulation

MDSCs suppress T-cell activation through several mechanisms (Figure 1). MDSCs exhibit control of T-cell responses via two enzymes: (1) arginase-1 (ARG1), which depletes L-arginine, and (2) inducible nitric oxide synthase 2 (iNOS), which generates nitric oxide.¹⁷ TGF-β and IL-10 induce MDSC ARG1, which suppresses CD4 and CD8 T-cell activation. Interferon gamma (IFN-y) and tumor necrosis factor alpha induce MDSC iNOS, which releases nitric oxide and induces T-cell apoptosis.¹⁸

MDSCs also deplete cysteine, an essential amino acid for T-cell activation.¹⁹ T cells obtain their cysteine from extracellular sources (specifically, DCs and macrophages), which convert methionine to cysteine and reduce cystine to cysteine during normal antigen processing and presentation. MDSCs depend on the conversion of cystine to cysteine. In the tumor microenvironment large numbers of MDSCs deplete the available cysteine, which deprives DCs and macrophages of this important molecule and indirectly lowers DC and macrophage production of cysteine and T-cell activation.19

MDSC-mediated downregulation of T-cell L-selectin (CD62L) impairs T-cell migration and activity.²⁰ CD62L is a plasma membrane molecule necessary for the homing of naive

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Figure I Myeloid-derived suppressor cells (MDSCs) suppress antitumor activity.

Notes: MDSCs are recruited and expanded in the tumor through the induction/production of transforming growth factor beta (TGF- β), stem cell factor (SCF), cyclooxygenase 2 (COX-2), prostaglandin E2 (PGE2), interleukin (IL)-6, granulocyte-macrophage colony-stimulating factor (GM-CSF), IL-10, IL-1 beta (IL1- β), vascular endothelial growth factor (VEGF), S100 proteins, and Snail. T-cell activation is suppressed by MDSC-mediated deprivation of L-arginine and cysteine from the environment, production of reactive oxygen species (ROS) and peroxynitrite, downregulation of L-selectin (CD62L) and the T-cell receptor-associated zeta chain, and the induction of T regulatory (Treg) cells through MDSC IL-10 and TGF- β production. MDSCs suppress natural killer (NK) cell cytotoxicity and NK interferon gamma (IFN- γ) production and they inhibit M1 macrophages (M1 Macs). MDSC expansion and IL-10 production inhibit dendritic cell (DC) antigen presentation. MDSC depletion enhances therapeutic vaccination responses. **Abbreviations:** APC, antigen-presenting cell; ARG, arginase; EMT, epithelial-mesenchymal transition; NO, nitric oxide.

T cells to lymph nodes for activation; MDSC downregulation of CD62L on naive T cells reduces this process.²⁰

The MDSC release of reactive oxygen species and peroxynitrite inhibits CD8⁺ T cells by catalyzing the nitration of the T-cell receptor, thereby preventing T-cellpeptide-MHC interactions.²¹ MDSCs also downregulate the T-cell receptor–associated zeta chain²² required for T-cell activation. MDSCs directly induce T regulatory (Treg) cells through the production of IL-10, TGF- β , or ARG1.¹⁰ The Treg cells actively downregulate the activation and expansion of antitumor reactive T²³ and NK cells.²⁴ MDSCs polarize T cells toward a type 2 tumor-promoting phenotype by IL-10 production and downregulation of macrophage IL-12.²⁵

T-cell nonresponsiveness to tumor antigens has been shown to be an early event in tumor progression.²⁶ Although T-cell tolerance in cancer has been shown to be mediated by host antigen-presenting cells (APCs),²⁷ the nature of these APCs has not been clear. Recent evidence suggests that MDSCs may represent a population of APCs responsible for induction of antigen-specific CD8 T-cell tolerance in cancer.²¹ CD8 T-cell responses in vivo are induced by professional APCs that present exogenous antigens in a MHC class I-restricted manner.28 This has been referred to as cross-priming or representation. DCs are the host APCs responsible for cross-priming by presenting epitopes obtained from apoptotic cells. However, in tumor-bearing hosts, there is a state of T-cell unresponsiveness that is regulated by DCs. The tumor not only fails to provide the inflammatory signals needed for efficient DC activation but also inhibits DC differentiation and maturation through IL-10²⁹ and VEGF.³⁰ Myeloid precursor differentiation into granulocytes, macrophages, and DCs is frequently dysregulated in the tumor, leading to an accumulation of MDSCs. Immature DCs have little or no expression of costimulatory molecules such as CD80, CD86, and CD40 on their surface and they produce little or no IL-12, which is required to support T-cell proliferation and for leading to tolerance. MDSCs also impair innate NK cells by inhibiting their cytotoxic ability and IFN- γ production. $^{\rm 31}$

Lung cancer genetic signatures that drive immune suppression

The authors' laboratory has been evaluating tumor signatures that maintain tumor growth through the modulation of immune activity. The authors have reported that the constitutive overexpression of tumor cyclooxygenase 2 (COX-2) leads to an immunosuppressive network in lung cancer.32,33 Overexpression of tumor COX-2 is associated with decreased host immunity³²⁻³⁴ and metastasis.35 COX-2 has been shown to modulate MDSC activity in lung cancer through the production of high levels of ARG1, which blocks T-cell function.³⁶ Until recently, the mechanism that induces ARG1 in MDSCs was unknown. Utilizing mice bearing the Lewis lung carcinoma (3LL), Rodriguez et al³⁶ showed that maintenance of MDSC ARG1 expression was independent of T-cell cytokine but was dependent on tumor-derived PGE2. These results demonstrated a new pathway of prostaglandin-induced immune dysfunction and provided a novel mechanism to explain the antitumor benefits of COX-2 inhibitors. The present authors have demonstrated that the tumor microenvironment induced DC immunosuppressive effects that promoted tumor growth and that tumor COX-2 inhibition restored DC APC and antitumor activity.34

The complex interactions between MDSCs and Treg cells are yet to be fully defined; however, it is evident that MDSCs promote Treg cell development in vivo. Tumor-reactive T cells are found in lung cancer tissue but fail to respond to tumor because of CD4+CD25+ Treg cells.³⁷ CD4+CD25+ Treg cells play an important role in the maintenance of immunologic self-tolerance and in suppressing antitumor immune responses mediated via self-reactive lymphocytes.23 Reducing the number of Treg cells or abrogating their activity within the tumor environment could induce effective tumor immunity in otherwise nonresponding hosts by activating tumor-specific as well as nonspecific effector cells. Studies have demonstrated that tumor COX-2 expression contributes to decreased host antitumor immune responses by affecting the frequency and activity of CD4+CD25+FOXP3+ Treg cells.38,39 COX-2 inhibition reduced Treg cell frequency and activity, attenuated Foxp3 expression in tumor-infiltrating lymphocytes, and decreased tumor burden. Transfer of Treg cells or administration of PGE2 to mice receiving COX-2 inhibitors reversed these effects.38 These studies were the first documentation that COX-2 inhibition downregulated tumor-induced Treg cell activity, leading to the restoration of antitumor responses.

In addition to COX-2, the present authors are currently evaluating the effect of genetic programs that regulate epithelial-mesenchymal transition (EMT) on tumor growth and metastases in lung cancer. Cancer cells acquire the ability to progress, invade, and metastasize by undergoing the process of EMT, by activating transcription factors (eg, Snail, Twist, Zeb, Slug) that repress epithelial cadherin.^{40,41} Snail expression in primary non-small cell lung cancer has been associated with a shorter overall survival,⁴² and in melanoma, Snail expression has been demonstrated to be important in EMT-induced metastases.⁴³ The authors have demonstrated that tumor Snail expression alters tumor growth and metastasis by affecting MDSCs in the tumor microenvironment.44 Compared with controls, the tumors of mice implanted with 3LL Snail knockdown cells decreased MDSC frequency, activity, and intracellular expression of ARG1 but increased CD107a cytolytic marker expression on CD8 T cells. Snail knockdown tumor cells demonstrated reduced subcutaneous growth and lung metastases.

MDSC depletion and antitumor activity in lung cancer

Although immunotherapy for lung cancer has potential, to date there has been no randomized trial reporting documented survival benefits.45 Tumor-induced immune suppression through MDSCs could contribute to the limited efficacy of the approaches. In a recent study, the authors tested the hypothesis that activating immune cells through therapeutic vaccination and disruption of MDSC-mediated immune suppression would improve antitumor activity in a murine lung cancer model.⁴⁶ To initiate antigen-specific responses, a cellular vaccine, comprising bone marrow adherent cells pulsed with the model ovalbumin (OVA) antigen expressed by OVA-modified 3LL cells, was utilized. The bone marrow adherent cells were pulsed with the OVA antigen to allow for antigen processing and presentation and were then injected subcutaneously into mice on the contralateral flank of the established tumor to initiate antigen-specific antitumor immune responses. To circumvent MDSC-mediated immune suppression, MDSCs were depleted with monoclonal antibodies (anti-Gr1 or anti-Ly6G).

Broad depletion by antibodies that targeted Gr1 or Ly6G on the MDSC population led to a decrease in MDSCs in the tumor and systemically in the blood, spleen, and bone marrow. The APC activity in the tumor was reduced as the tumor progressed; however, MDSC depletion led to increased frequency and activities of APC, NK cell, and T-cell effectors. Along with the increased NK and T-cell

10

activities, there was an increase in the frequency of apoptotic tumor cells and a concomitant reduction in tumor burden and migration of tumor cells from the primary tumor site to the lung. The reduction in tumor growth and abrogation in tumor cell migration may be explained by an increase in the frequency of activated T and/or NK effector-mediated tumor apoptosis and/or T or NK IFN-y-mediated antiangiogenesis. Following MDSC depletion, the antiangiogenic chemokines CXCL9 and CXCL10 were increased and the pro-angiogenic cytokines VEGF-A, angiopoietin 1, angiopoietin 2, CXCL2, and CXCL5 were markedly reduced. Accompanying this profile was a reduction in the endothelial markers MECA-32 in the tumor but there was also an increase in CXCR3 expression. MDSC depletion led to an influx of IFN-y-producing activated T and NK cells that promoted angiostasis in the tumor by altering the balance of pro- and antiangiogenic chemokines. This suggests that MDSC depletion not only improves APC, NK cell, and T-cell immune activities but also promotes antiangiogenesis, leading to a more effective tumor control.

Although the depletion by antibodies affects other Gr1or Ly6G-expressing monocytes, the data suggest that the depletion of MDSCs is beneficial in eliciting anticancer effects and are consistent with studies by other groups.^{47,48} Therapeutic vaccination led to decreased tumor burden without complete eradication of the tumors. Depletion of MDSCs in combination with the vaccine led to a 20-fold reduction in tumor burden in 50% of the mice and to complete eradication of tumors in the remainder of animals. These findings suggest that an effective attack necessitates changes in multiple components of the immune system in parallel, leading to immune cell activation and to the disruption of the regulatory mechanisms that limit antitumor immune responses. MDSC depletion reprograms the tumor niche by altering the inflammatory responses, allowing tumor immune destruction and generation of immunological memory.

Conclusion

Despite the identification of a repertoire of tumor antigens, hurdles persist for immune-based anticancer therapies. Tumor-induced immune suppression contributes to the limited efficacy of the current approaches. More effective immunotherapeutic strategies will likely result from an improved understanding of the mechanisms that sustain tumor growth. It is likely that combined therapies that address the multiple immune deficits in cancer patients will be required for effective therapy. MDSCs play a major role in the suppression of T-cell activation and they sustain tumor growth, proliferation, and metastases. Regulation of MDSC recruitment, differentiation or expansion, and inhibition of the MDSC suppressive function with pharmacologic agents will be useful in the control of cancer growth and progression. Pharmacologic agents that regulate MDSCs may be more effective when combined with immunotherapies. Optimization of combined approaches that simultaneously downregulate MDSC suppressor pathways, restore APC immune-stimulating activity, and expand tumorreactive T cells will be useful in improving therapy.

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Disclosure

The authors report no conflicts of interest in this work.

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12