Inhibition of Tumor-Derived Prostaglandin-E2 Blocks the Induction of Myeloid-Derived Suppressor Cells and Recovers Natural Killer Cell Activity

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Abstract

Purpose: Increased frequencies of myeloid-derived suppressor cells (MDSC) correlate with poor prognosis in patients with cancers. Tumor-derived prostaglandin-E2 (PGE2) plays an important role in inducing MDSCs. However, the detailed mechanisms of this induction remain unknown. To develop targeted therapies for MDSCs, we sought to investigate the molecular basis of PGE2-regulated accumulation of MDSCs and their functional consequence on natural killer (NK) cell activity.

Experimental Design: The effects of PGE2 in inducing phenotypic, signaling, and functional alterations on monocytes were analyzed in vitro. Suppression of NK-cell activity by PGE2-treated monocytes was compared with that of freshly isolated CD14+HLA-DRlow CD11b+Ly6G-/Ly6Chigh monocytic MDSCs (moMDSC) from patients with melanoma. In addition, to explore the in vivo relevance of targeting PGE2 to reduce MDSC-mediated suppression of NK cells, we established a murine model, where tumor cells were disabled from cyclooxygenase-2 (COX-2) production.

Results: Patient-derived moMDSCs inhibited NK-cell activity through the production of TGFβ. In vitro, binding of PGE2 to EP2 and EP4 receptors on monocytes activated the p38MAPK/ERK pathway and resulted in elevated secretion of TGFβ. Similar to moMDSCs, PGE2-treated monocytes potently suppressed NK-cell activity through production of TGFβ. Furthermore, silencing COX-2 in murine 4T1 tumor cells reduced the accumulation of CD11b+Gr1+ MDSCs in the spleen, resulting in concomitant improved in vivo clearance of NK-cell sensitive YAC-1 cells.

Conclusions: Our results reveal an indispensable role of tumor-derived PGE2 in inducing MDSCs and suggest a favorable outcome of combining COX-2–targeted therapy and adoptive NK-cell transfer in patients with cancer.

Introduction

Tumor progression is frequently associated with a severe impairment of the host innate and adaptive immune response. In addition to immune-regulatory mechanisms directly mediated by tumor cells, a variety of suppressive immune cells, such as regulatory T cells, M2-like tumor-associated macrophages (TAM), and tolerogenic dendritic cells, have been reported to be expanded in tumor-bearing animals and patients with cancer (1).

Recently, myeloid-derived suppressor cells (MDSC) have been proposed as key immune regulators in various solid and hematologic malignancies (2). Although murine MDSCs are characterized as monocytic (moMDSCs; CD11b+Ly6G−/Ly6Chigh) and granulocytic (grMDSCs; CD11b+Ly6Ghigh/Ly6C+low) (3), the phenotypes of MDSCs in patients with cancer remains controversial. Distinct phenotypes of MDSCs are associated with different types of human cancers (2, 4). In patients with advanced stage melanoma, an accumulation of circulating CD14+HLA-DRlow moMDSCs were shown in peripheral blood (5–8), which demonstrated a potent suppression of autologous lymphocytes through production of TGFβ (5). In 2 separate studies, we showed that MDSC-derived reactive oxygen species (ROS), prostaglandin E2 (PGE2), and STAT-3 signaling contribute to the suppression of autologous T cells (6, 9). Other studies have highlighted a variety of suppressive mechanisms mediated by MDSCs, including indoleamine 2,3-dioxygenase (IDO), arginase-1, IL10, and iNOS (10, 11).

Increased numbers of MDSCs have been correlated with the disease stage in patients with breast cancer (8) and...
Translational Relevance
Myeloid-derived suppressor cells (MDSC) accumulate in high frequencies in peripheral blood of patients with cancer and mediate potent inhibition of adaptive antitumor immune responses. These cells are associated with the stage of cancers and have recently been shown to predict survival of patients with melanoma. However, it is less clear how MDSCs are induced in patients with cancers and how they affect the activity of human natural killer (NK) cells. In this study, we evaluated the importance and molecular property of tumor-derived PGE2 in enhancing the suppressive functions by MDSCs of NK cells. Collectively, our data suggest that a favorable clinical outcome may be achieved when COX-2 inhibition and NK adoptive transfer a combined in the treatment of cancers.

Survival of patients with ovarian cancer (12). In a recent study, Weide and colleagues demonstrated MDSCs, but not Tregs, predicted the survival of patients with advanced melanoma and were strongly correlated with the absence of antigen-specific T cells (13). In contrast, decreased frequencies of MDSCs were shown when patients with melanoma were treated with BRAF inhibitors (14), or anti-CTLA4 blocking antibodies (15).

Although the detailed molecular mechanisms remain to be elucidated, a number of inflammatory modulators, including GM-CSF (5, 16–19), IL6 (17, 19), and TNFα (19, 20) have been suggested to contribute to the induction and expansion of MDSCs. Recently, we and others proposed that PGE2 is indispensable for the induction and maintenance of MDSCs in patients with cancer. Coculture with tumor cells secreting PGE2 or treatment with PGE2 was shown to enhance the suppressive functions of monocytes on T cells. Furthermore, inhibition of cyclooxygenase-2 (COX-2), the enzyme catalyzing the synthesis of PGE2, potently rescued suppression of T cells mediated by MDSCs (9, 21).

Despite the recent advances in our understanding of how MDSCs suppress T-cell responses, much less is known how NK-cell responses are influenced by MDSCs. In tumor-bearing mice, MDSCs were shown to suppress NK-cell activation and cytolytic capacity via membrane-bound TGFβ (22, 23). In contrast, it has been proposed that MDSCs may also exert a stimulatory function on NK cells through ligation of the NKG2D receptor (24). In human cancers, it is less documented how MDSCs interact with NK cells. A recent study showed that suppression of NK cells by moMDSCs was mediated through the ligand of NKp30 in CD14HLA-DRneg fraction (Miltenyi Biotech) and moMDSCs were thereafter purified from the HLA-DR negative fraction using a Human Monocyte Isolation Kit (CD56<1%; Miltenyi Biotech) and moMDSCs were thereafter purified from the HLA-DR negative fraction using a Human Monocyte Isolation Kit (CD56<1%; Miltenyi Biotech). NK cells were enriched from the CD3 depurination reagents (Rosette-Sep, StemCell) during Ficoll gradient centrifugation (CD3<1%; Ficoll-Paque Plus, GE Healthcare). Next, HLA-DR<cells were sorted with anti-human HLA-DR microbeads (Miltenyi Biotech) and moMDSCs were thereafter purified from the HLA-DR negative fraction using a Human Monocyte Isolation Kit (CD56<1%; Miltenyi Biotech). NK cells were enriched from the CD3<1% CD14<1%HHLA-DR<1% fraction (Miltenyi Biotech). Cells were thereafter seeded for functional assays or frozen at −80°C for Western blot analysis. Peripheral blood mononuclear cells (PBMC) were isolated from healthy individuals by Ficoll gradient centrifugation as stated above. NK cells and monocytes were purified with a human NK cell or a CD14 Monocyte Isolation Kit, respectively (purity>95%; Miltenyi Biotech).

Mice
All animal experiments were conducted under ethical approved study (N603/12; Karolinska Institutet). BALB/c mice (8–10 weeks old) were purchased from Charles River Laboratory.

Mouse tumor cell line
Murine breast cancer cell line 4T1 stably expressing scrambled or shRNAs targeting COX-2 enzyme was provided by Dr. W. Schiemann (Case Comprehensive Cancer Center, Cleveland, OH) and maintained in DMEM supplemented with 10% FBS. YAC-1 cells were cultured in RPMI-1640 supplemented with 10% FBS.

Advanced melanoma is mediated via TGFβ. Furthermore, treatment with PGE2 conferred monocytes with an MDSC-like phenotype and intracellular signaling pattern enabling them to suppress NK-cell responses in a TGFβ-dependent manner. Finally, we demonstrate that shRNA-targeting of COX-2 in murine tumor cells result in a significant reduction of MDSC accumulation in spleen, with concomitant enhanced lysis of YAC-1 cells in vivo. To our knowledge, these observations for the first time reveal the direct involvement and molecular mechanism of PGE2 in stimulating the production of TGFβ from MDSCs and its importance in suppressing human NK cells. Given the prevalence of COX-2/PGE2 overexpression in tumors, these findings could provide fundamental knowledge on how MDSC-mediated immune suppression is initiated and provide targets for limiting the induction of MDSCs in various human cancers.
Proliferation assays

Purified monocytes were seeded in triplicates in 96-U-bottom well plates at 1:1 ratio with autologous NK cells in IMDM medium (Gibco) supplemented with 10% heat-inactivated human AB serum (referred as “medium” below), in the absence or presence of 50 nmol/L PGE2 and 200 IU/ml IL2. Next, reagents targeting different suppressive mechanisms were added, including 10 μg/ml neutralizing antibodies for PGE2 (Cayman Chemical) or TGFβ (R&D systems), 20 μmol/L selective antagonists for PGE2 receptors AH6809 (EP2), AH23848 (EP4), and L798106 (EP3) (all from Sigma-Aldrich). Following 2 days of coculture, 3H-thymidine (25 μCi per well) was added to the cells and harvested after additional 12 to 14 hours. Incorporation of 3H-thymidine in the harvested cells was analyzed by a MicroBeta scintillation counter (TRILUX 1450, PerkinElmer).

Chromium release assays

Following 48 hours of coculture with monocytes in the presence or absence of PGE2, NK-cell cytotoxicity was analyzed by chromium (51Cr) release assays (7 or 18 hours) against K562 (STR fingerprint in Supplementary Table S1) cells at a 5-to-1 effector–target ratio, as previously described (26). For the ex vivo studies, patient-derived moMDSCs and resting or IL2-activated NK cells were cocultured at 1:2 ratio, with the presence of 10 μg/ml neutralizing antibodies for TGFβ or PGE2, 200 IU/ml of the ROS scavenger catalase (Sigma-Aldrich) or superoxide dismutase (Sigma-Aldrich), 500 μmol/L arginine inhibitor N(o)-hydroxy-orn-arginine (nor-NOHA; Calbiochem) or iNOS inhibitor NG-monomethyl-l-arginine (L-NMMA; Sigma-Aldrich). Following overnight incubation, chromium release assays were performed as mentioned above.

Cytokine production assays

Levels of IFNγ released from patient-derived NK cells were quantified by ELISA assay after 20 hours (MabTech) after addition of K562, in the presence of moMDSCs or inhibitors indicated above. Alternatively, patient-derived NK cells were activated with 200 IU/ml IL2, and cocultured with autologous moMDSCs at 1:1 ratio and in the presence of TGFβ or PGE2 neutralizing antibodies and IFNγ release was analyzed after 48 hours.

Cocultures of NK cells and control or PGE2-treated monocytes were harvested and TGFβ concentration was measured using ELISA (R&D systems). For the patient-derived moMDSCs, cells were seeded in medium only, or in the presence of 10 μmol/L SB203580 (p38MAPK inhibitor; Sigma-Aldrich), 10 μmol/L H89 (protein kinase A inhibitor; Cayman Chemical), 10 μmol/L PD98059 (MEK1/2 inhibitor; Sigma-Aldrich), or DMSO (Sigma-Aldrich). Following 48 hours of coculture, supernatants were collected and evaluated for concentrations of TGFβ by ELISA. Moreover, IFNγ production during IL2 activation in the NK-moMDSC cocultures was quantified by ELISA (MabTech).

Western blot analysis

Monocytes treated with PGE2 (50 nmol/L) for 30 minutes, or freshly isolated moMDSCs were washed in PBS, and frozen at –80°C for protein analysis by Western blot analysis. Protein lysates were prepared from pellets and the blots were developed as previously described (27). In brief, equal amount of protein (50 μg) were loaded and detection of each protein were performed using primary monoclonal antibodies (mAb), including anti-p38MAPK, anti-phospho(Thr180/182)-p38MAPK, anti-ERK, anti-phospho(Thr202/Tyr204)-ERK, anti-AKT, anti-phospho(Ser473)-AKT (Cell Signaling). β-Actin was used as loading control (Sigma-Aldrich). The membrane was then stained with horse-radish-peroxidase-conjugated secondary antibodies (Cell Signaling), incubated in LumiLight Western blotting substrate (Roche Diagnostics), and recorded in a LAS3000 system (Fujifilm).

Flow cytometry and fluorochrome-conjugated antibodies

Detailed information of antibodies utilized in this study is summarized in Supplementary Table S2. Phenotypic changes of healthy donor-derived monocytes were compared with PGE2-treated monocytes (1 μmol/L). Moreover, surface molecules associated with NK-cell functions were also analyzed when cultured with control or PGE2-treated monocytes. In brief, cells were harvested, washed, and stained in FACS buffer (PBS containing 1% human serum albumin (HSA); Octapharma) for 20 minutes at 4°C. For measuring the intracellular expression of COX-2 enzyme, 4T1 tumor cells were fixed and permeabilized by BD CytoFix/CytoPerm buffer, followed by washing and staining steps using BD Perm/Wash buffer (BD Biosciences).

In experiments where murine Tregs were evaluated, extracellular molecules were first stained, and intracellular expression of Foxp3 was measured with BD transcription factor buffer set (BD Biosciences), according to the manufacturer’s protocol. Cells were acquired on a BD LSRII flow cytometer (BD Biosciences) and analyzed with FlowJo software (Tree Star).

NK-cell in vivo cytotoxicity

Because COX-2 silencing in 4T1 cells results to a delay of tumor growth in vivo (28), inoculated cell numbers were adjusted and all experiments were performed in mice bearing comparable tumor burdens. In brief, control, COX2–silenced 4T1 cells, or PBS was injected subcutaneously into the right flank of the mice. When average tumor size reached 300 mm3 (calculated by length × width × height × 0.5; ref. 29), murine NK-cell sensitive target YAC-1 cells were labeled with DiR near infra-red dye (1,1′-dioctadecyl-3,3,3′,3′-tetra-methylindotri-carbocyanine iodide; Invitrogen) for 10 minutes, followed by washing and intravenous injection into all mice, to evaluate the cytolytic activity of NK cells in vivo. The fluorescence intensity of YAC-1 cells was examined by in vivo live imaging using an IVIS camera platform (Caliper Life Sciences Benelux & Nordic) 1 and...
values. Histograms or images were chosen based on the average groups were presented as means and SD. Representative multiple inputs from separate experimental and control indicated in the figure legends). Unless otherwise stated, Student t-tion, thereafter differences among groups were analyzed by (GraphPad). All data were first tested for normal distribu-

5 hours after the injection. Lungs of mice bearing size-matched tumors were resected and intensity of YAC-1 cells was evaluated after 5 hours. Splenocytes were isolated from mice that did not receive YAC-1 injection, and measured for percentages of CD11b+ Gr1+ MDSCs, CD3- CD49b+ NK cells, and CD4+ CD25+ FoxP3+ regulatory T cells by flow cytometry as described above.

Statistical analysis
All data were first analyzed in the software mentioned above and summarized by Prism Version 6 software (GraphPad). All data were first tested for normal distribution, thereafter differences among groups were analyzed by Student t test or nonparametric, Mann–Whitney U tests (as indicated in the figure legends). Unless otherwise stated, multiple inputs from separate experimental and control groups were presented as means and SD. Representative histograms or images were chosen based on the average values.

Results
Melanoma-derived CD14+ HLA-DRlow+ moMDSCs potently inhibit NK-cell functions
To analyze the influence of MDSC on NK cells, patient-derived CD14+ HLA-DRlow+ moMDSCs and HLA-DR+ mononuclear cells were isolated and cocultured with autologous nonactivated NK cells. In the presence of moMDSCs, cytolytic capacity and production of IFNγ by NK cells were significantly inhibited (P < 0.01; Fig. 1A and B). Similar to nonactivated NK cells, the presence of moMDSCs during IL2-activation of NK cells resulted in significantly impaired cytotoxicity and IFNγ production by NK cells (P < 0.01 and P < 0.05, respectively; Fig. 1C and D). In contrast, HLA-DR+ cells did not show suppressive functions against NK cells (Supplementary Fig. S1A).

TGFβ is the major suppressive mechanism used by patient-derived moMDSCs
We next investigated the molecular basis of suppression mediated by patient-derived moMDSCs. As shown in Fig. 2A and B, blocking moMDSC-derived TGFβ significantly rescued the cytotoxicity (P < 0.05) and production of IFNγ by nonactivated NK cells (P < 0.01). Moreover, addition of the hydrogen peroxide scavenger catalase led to significant improvement of NK-cell-mediated cytotoxicity (Fig. 2A). Similarly, when TGFβ-blocking antibody was present in cocultures of IL2-activated NK cells and moMDSCs, NK-cell cytotoxicity (P < 0.01; Fig. 2C) and IFNγ released by NK cells (P < 0.05; Fig. 2D) were significantly improved. However, catalase did not show functional restoration of IL2-activated NK cells (data not shown). Of note, addition of neutralizing antibodies did not modulate cytolytic functions of IL2-activated NK cells in the presence of HLA-DR+ cells (Supplementary Fig. S1A).

PGE2-treated monocytes resemble patient-derived moMDSCs
We and others have previously shown that PGE2 contributes to the conversion of monocytes into MDSC-like cells (9, 21). To validate the similarities of PGE2-induced MDSCs and patient-derived moMDSCs, we evaluated the phenotypic alternations and PGE2-induced intracellular signaling pathways. The results demonstrated that PGE2 increased the expression of CD14 on monocytes whereas the expression of HLA-DR was downregulated (Fig. 3A and B). Expression of other myeloid cell markers, including
CD16, CD80, DC-Sign, CD33, and CD137 (4-1BB), was not altered between control-cultured and PGE2-treated monocytes (Supplementary Fig. S2A).

Intracellularly, p38MAPK and ERK pathways were phosphorylated in monocytes, as a result of PGE2 treatment (Fig. 3C). This was also the case in freshly isolated moMDSCs, in comparison to control HLA-DR+ cells (Fig. 3C). However, neither of the samples demonstrated phosphorylation of AKT proteins, even though total expression of AKT was upregulated in monocytes subsequent to PGE2 treatment (Fig. 3C).
in vitro culture (Fig. 3C). Because of these similarities with moMDSCs, PGE2-treated monocytes were regarded as "MDSC-like" cells.

**MDSC-like cells suppress NK cells through TGFβ**

To validate the suppressive functions of MDSC-like cells, we measured the proliferation, phenotype as well as cytolytic capacity of NK cells, when cocultured with MDSC-like cells or control monocytes. Because PGE2 directly suppressed proliferation of IL2-activated NK cells in a dose-dependent manner (Supplementary Fig. S2B), a noninhibitory level of PGE2 (50 nmol/L) was chosen for all subsequent assays. At this concentration, MDSC-like cells significantly (P < 0.05) reduced NK-cell killing capacity when present at a 1-to-1 ratio during NK-cell activation (Fig. 4A and Supplementary Fig. S2C). Moreover, MDSC-like cells significantly inhibited the proliferation of NK cells (P < 0.01; Fig. 4B). Notably, the expression of NKG2D, NKp46, and NKp44 were significantly reduced (P < 0.05) on IL2-activated NK cells, when MDSC-like cells were present. Although to a lower extent, coculture with monocytes downregulated the expression of these molecules (Fig. 4C).

Next, we asked what mechanisms were responsible for the suppressive actions mediated by MDSC-like cells. Similar to patient-derived moMDSCs, neutralization of TGFβ significantly restored NK-cell–mediated cytotoxicity (P < 0.01) and proliferation (P < 0.05) to similar levels to that of NK cells cultured alone (Fig. 4D and E). Importantly, neutralization of TGFβ or PGE2 had no direct impact on the cytotoxicity of NK cells (Supplementary Fig. S1B).

**PGE2 enhances the production of TGFβ by monocytes**

Because TGFβ plays a major role in suppression of NK-cell responses by MDSC-like cells and moMDSCs,
the accumulation of CD11b<sup>+</sup>Gr1<sup>−</sup> MDSCs (Fig. 5C), confirming the functional contribution of these pathways to MDSC-mediated suppression of NK cells in humans.

### Overexpression of COX-2 in tumor cells regulates MDSC accumulation and NK-cell cytotoxicity in vivo

To assess the functional consequences of COX-2 expression in inducing MDSCs in vivo, we used a murine model, where control or COX-2-silenced 4T1 mammary carcino-

we next evaluated if monocytes differ in their ability to produce TGFβ in the presence or absence of PGE2. Compared with untreated monocytes, MDSC-like cells produced significantly higher levels of soluble TGFβ (P < 0.05; Fig. 5A), which was abolished when cultured in the presence of neutralizing antibodies to PGE2 (P < 0.05; Fig. 5A). These observations were confirmed in the NK-MDSC-like cocultures, excluding the possibility of TGFβ-driven auto-regulation by PGE2-treated NK cells (Fig. 5B). In contrast, monocytes expressed low levels of membrane-bound TGFβ, which remained unchanged after PGE2 treatment (data not shown).

Protein kinase A is known to be one of the major intracellular regulators of EP2/EP4 signaling (30) and contribute to the activation of the p38MAPK pathway in myeloid cells (31). Therefore, chemical inhibitors of p38MAPK (SB203580), MAP or ERK kinase (MEK1/2, PD98085), protein kinase A (H-89), or DMSO were utilized. Indeed, interruption of these pathways resulted in significant attenuation of TGFβ release from patient-derived moMDSCs (Fig. 5C), confirming the functional contribution of these pathways to MDSC-mediated suppression of NK cells in humans.

Figure 5. PGE2 enhances the production of TGFβ in monocytes. A, monocytes isolated from healthy donors were cultured alone or in the presence of PGE2 (50 nmol/L, 48 hours) or together with a neutralizing antibody against PGE2 and analyzed for production of TGFβ (n = 3). B, production of TGFβ in cultures of NK cells alone or when cocultured with monocytes, in the presence or absence of PGE2 (50 nmol/L, n = 4). C, patient-derived moMDSCs (CD14<sup>+</sup>HLA-DR<sup>−</sup>/C3/C6/C3) were cultured in the presence of DMSO, SB203580 (p38MAPK), PD98085 (MEK1/2), or H-89 (protein kinase A) (all used at 10 μmol/L and analyzed for production of TGFβ (n = 5) relative to nontreated moMDSCs. Data, mean ± SD; Student t test; *, P < 0.05; **, P < 0.01.

Inoculation of control 4T1 cells resulted in a significant accumulation of CD11b<sup>+</sup>Gr1<sup>−</sup> MDSCs (P < 0.001; Fig. 6A), but a decrease in the percentage of NK cells in the spleen of BALB/c mice (P < 0.05; Fig. 6B). Notably, significantly fewer CD11b<sup>+</sup>Gr1<sup>−</sup> MDSCs (P < 0.01; Fig. 6A) and restored numbers of NK cells (P < 0.01; Fig. 6B) were found in spleens when COX-2 was not expressed by tumor cells. Although significantly higher percentages of regulatory T cells (CD4<sup>+</sup>CD25<sup>+</sup>FoxP3<sup>+</sup>, P < 0.01; Fig. 6C and Supplementary Fig. S3B) and lower percentages of CD3<sup>+</sup> T cells (P < 0.001; Supplementary Fig. S3C) were observed in mice inoculated with control 4T1 tumor cells, these cell populations were not regulated by tumor-derived COX-2 production (Fig. 6C and Supplementary Fig. S3C). Moreover, the percentage of CD4<sup>+</sup> T cells was comparable in all groups (Supplementary Fig. S3D).

To evaluate the cytolytic capacity of NK cells in vivo, naïve or mice bearing similar tumor burdens were challenged with DiR-labeled YAC-1 cells and the signal was evaluated by live imaging. Along with the decrease in MDSCs and increased NK-cell percentages in mice bearing COX-2-silenced 4T1 tumors, a significantly improved ability to reject NK-cell sensitive YAC-1 cells were observed (P < 0.05; Fig. 6D). Resection of lungs from YAC-1–infused mice bearing COX-2–silenced 4T1 tumor cells showed a pronounced reduction of YAC-1 cells after 5 hours, compared with the control group (Fig. 6E). No difference in the
fluorescence intensity of inoculated YAC-1 cells was observed within 1 hour after inoculation, showing that the fluorescence labeling intensity was similar among the groups (Supplementary Fig. S3E).

Discussion

In the recent years, there is an increasing interest in understanding the immune-regulatory role of MDSCs in cancers. NK-cell responses are negatively regulated by several immunosuppressive mechanisms used by tumor cells (32), regulatory T cells (33, 34), and MDSCs (23, 25). Although the mechanisms of how MDSCs suppress T cells have been extensively studied, much less is known how MDSCs regulate the activity of human NK cells. Here we provide evidence that support a direct involvement of PGE2 during the initiation and maintenance of NK-cell suppression by MDSCs.

Coculture of moMDSCs isolated from patients with melanoma with NK cells resulted in reduced cytolytic function and IFNg release of resting and IL2-activated NK cells. Because neutralization of moMDSC-derived TGFβ restored NK-cell activity, it demonstrated the importance of this cytokine in MDSC-mediated inhibition of NK cells. Moreover, ROS scavenger catalase significantly rescued the cytolytic activity of resting NK cells in the presence of moMDSCs. In contrast, production of IFNg from resting NK cells was not improved in the presence of catalase. This implies that although the cytotoxic function of resting NK cells is sensitive to ROS-mediated suppression, it is not the case for IFNg production. It is in line with earlier observations, where cytotoxic CD56dim NK cells are more susceptible to ROS-induced apoptosis, compared with cytokine-producing CD56bright NK cells (35, 36). The distinction in sensitivity to ROS between the 2 NK-cell subsets does not apply to IL-2–activated NK cells, where TGFβ, rather than ROS or other mechanisms, dominated the suppression of both cytotoxicity and cytokine production of NK cells.

Although strongly suggested by previous studies (5, 7), blockade of TGFβ secretion of CD14+ HLA-DRdim/moMDSCs did not result to rescued proliferation of purified human T cells (6, 9, 37). In the earlier studies, thymidine

Figure 6. Overexpression of COX-2 in tumor cells regulates MDSC accumulation and NK-cell cytotoxicity in vivo. Splenocytes were isolated from naïve BALB/c or mice bearing similar tumor burden (control or COX-2 silenced 4T1 cells). Percentages of MDSCs (CD11b+ Gr1+; A), NK cells (CD3+ CD49b+; B), or regulatory T cells (CD3+ CD4+ CD25+ FoxP3+; C) were determined by flow cytometry. D, quantitative analysis of YAC-1 cells in mice bearing control 4T1 or COX-2–silenced 4T1 tumors were examined by in vivo live imaging using the IVIS camera platform. E, lungs were resected and remaining YAC-1 cells were measured after 5 hours. Numbers of the dots in each group correspond to numbers of mice utilized in the experiments. Data, mean ± SD; Mann–Whitney nonparametric U test; *, P < 0.05; **, P < 0.01; ***, P < 0.001; n.s., not significant.
incorporation was evaluated in PMA-activated lymphocytes, when moMDSCs were present at different ratios. Therefore, TGFβ–mediated suppression of NK cells or their production of T-cell–stimulating cytokines in these experimental settings, as demonstrated in our study, may contribute to explain the existing discrepancies in the literature.

Production of arginase I and iNOS are 2 well-documented mechanisms that MDSCs use to suppress tumor-reactive T-cell responses. We here find moMDSC-mediated suppression of NK cells to be independent of these factors, concurrent with previous observations in patients with hepatocellular carcinoma (25). This can be explained by the heterogeneity of human MDSC population, in which granulocytic MDSCs are potent producers of arginase I, whereas suppression mediated by the monocytic subset is largely dependent on other factors, that is TGFβ (5) or ROS (38). Thus, because of the multifaceted functions of MDSCs, it is reasonable to propose that multiple suppressive factors need to be targeted to abolish MDSC-induced immune suppression.

In agreement with Lechner and colleagues, we have previously shown that coculture with a variety of tumor cell lines converted monocytes into MDSC-like cells, which mediated suppression of T cells through COX-2 production (9, 19). To elucidate the molecular basis of this induction, recombinant PGE2 was used as a surrogate for monocytic tumor cell coculture. Treatment of PGE2 maintained an immature status of monocytes, including high expression levels of CD14, but reduced expression of HLA-DR. Culture of monocytes in the presence of PGE2þ cells reduced this suppression. Notably, PGE2 treatment of PGE2 in inducing suppressive functions of MDSCs on NK cells, when moMDSCs were present at different ratios.

Phosphorylation of p38MAPK and ERK proteins has previously been shown to be associated with MDSCs in tumor-bearing animals (39, 40) and patients with cancer (41). However, the functional implications and molecular basis of these pathways remain unknown. Here, we provide a novel perspective, where elevated PGE2 levels in patients with cancer could activate these pathways in myeloid cells through the EP2 or EP4 receptor. Consequently, these cells are conferred with an immature MDSC phenotype as well as the ability to produce TGFβ. In addition, this may lead to a functional resistance to apoptosis of MDSCs, for example by Fas–FasL interactions (40), or a prolonged survival in tumor microenvironment, in comparison to other HLA-DR+ cells. Thus, targeting similar pathways in MDSCs with BRAF or MEK1/2 inhibitors may contribute to the clinical responses in patients with cancer (14).

Further studies of the role of PGE2 in inducing NK-suppressive MDSCs was addressed by an in vivo 4T1 mammary carcinoma model, which induced extensive accumula-

Disclosure of Potential Conflicts of Interest
No potential conflicts of interest were disclosed.

Authors’ Contributions
Conception and design: Y. Mao, D. Sarhan, B. Seliger, R. Kessling, A. Lundqvist
Acquisition of data (provided animals, acquired and managed patients, provided facilities, etc.): Y. Mao, D. Sarhan, A. Steven, R. Kessling, A. Lundqvist
Analysis and interpretation of data (e.g., statistical analysis, biostatistics, computational analysis): Y. Mao, D. Sarhan, A. Steven, B. Seliger, R. Kessling, A. Lundqvist
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Administrative, technical, or material support (i.e., reporting or organizing data, constructing databases): D. Sarhan
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