

# Anticancer Chemotherapy-Induced Intratumoral Recruitment and Differentiation of Antigen-Presenting Cells

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## SUMMARY

The therapeutic efficacy of anthracyclines relies on antitumor immune responses elicited by dying cancer cells. How chemotherapy-induced cell death leads to efficient antigen presentation to T cells, however, remains a conundrum. We found that intratumoral CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells, which displayed some characteristics of inflammatory dendritic cells and included granulomonocytic precursors, were crucial for anthracycline-induced anticancer immune responses. ATP released by dying cancer cells recruited myeloid cells into tumors and stimulated the local differentiation of CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells. Such cells efficiently engulfed tumor antigens in situ and presented them to T lymphocytes, thus vaccinating mice, upon adoptive transfer, against a challenge with cancer cells. Manipulations preventing tumor infiltration by CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells, such as the local overexpression of ectonucleotidases, the blockade of purinergic receptors, or the neutralization of CD11b, abolished the

immune system-dependent antitumor activity of anthracyclines. Our results identify a subset of tumor-infiltrating leukocytes as therapy-relevant antigen-presenting cells.

## INTRODUCTION

Although it has been generally believed that cytotoxic antineoplastic agents mediate their therapeutic effects in a cancer cell-autonomous fashion, at least some chemotherapeutics inhibit tumor growth also indirectly, via the immune system (Mellman et al., 2011; Schreiber et al., 2011). Thus, a variety of transplantable or chemically induced primary mouse cancers respond more efficiently to anthracyclines when they develop in hosts carrying an intact immune system. In many cases, tumors evolving in immunodeficient mice fail to respond to anthracyclines in conditions in which the same tumors growing in immunocompetent mice do so (Apetoh et al., 2007; Casares et al., 2005; Ghiringhelli et al., 2009; Michaud et al., 2011). Accordingly, clinical studies indicate that anthracycline-killed tumor cells are particularly efficient in stimulating a therapeutic immune response in cancer patients (Zappasodi et al., 2010). Moreover, anthracycline-based neoadjuvant therapy of breast

cancer patients is more effective when the tumor is infiltrated by T cells before the beginning of chemotherapy (Denkert et al., 2010) as well as if chemotherapy causes a significant influx of CD8<sup>+</sup> T cells into the tumor bed and/or reduces the accumulation of immunosuppressive regulatory T (Treg) cells (Ladoire et al., 2011).

Several immune effectors that are essential for the therapeutic effect of anthracyclines have been characterized in mouse models. Indeed, the deletion of genes coding for several cytokines (*Il1b*, *Il17a*, *Ifng*) or their receptors (*Il1r*, *Il17r*, *Ifngr*) abolishes the anticancer effects of anthracyclines, and so does the injection of antibodies that neutralize IL-1 $\beta$ , IL-17, and IFN- $\gamma$  or that deplete CD4<sup>+</sup>, CD8<sup>+</sup>, and  $\gamma\delta$  T cells (Ghiringhelli et al., 2009; Ma et al., 2011; Michaud et al., 2011; Obeid et al., 2007). Functional experiments suggest a hierarchy of events whereby IL-1 $\beta$  produced by dendritic cells (DCs) allows for the activation of IL-17A-producing  $\gamma\delta$  T cells, which eventually sustain the accumulation of IFN- $\gamma$ -producing CD8<sup>+</sup> T cells into the tumor bed, 1 week after the chemotherapy has started (Ma et al., 2011).

In contrast to most other cytotoxic chemotherapeutics, anthracyclines stimulate immunogenic cell death. This type of cell death is characterized by a compendium of subtle biochemical changes in the properties of the plasma membrane as well as in the microenvironment of dying cancer cells (Galluzzi et al., 2012; Green et al., 2009). These changes include the preapoptotic exposure of calreticulin (CRT) on the cell surface (which facilitates the engulfment of portions of the dying cells by antigen-presenting cells, APCs) (Gardai et al., 2005) and the postapoptotic release of high mobility group box 1 (HMGB1) from the nucleus (which binds TLR4 receptors and hence stimulates antigen presentation) (Apetoh et al., 2007). We have found that the release of ATP by dying tumor cells is also essential for cell death to be perceived as immunogenic. Indeed, ATP stimulates the recruitment of inflammatory cells into the tumor bed and binds P2RX7 receptors on DCs, thereby facilitating the activation of the NLRP3 inflammasome and the consequent secretion of IL-1 $\beta$  (Ghiringhelli et al., 2009; Martins et al., 2009). Neoplasms in which the pathways leading to CRT exposure or ATP release have been interrupted are refractory to anthracyclines in vivo (Michaud et al., 2011; Panaretakis et al., 2009). Along similar lines, the elimination of TLR4 or P2RX7—as well as that of their intracellular signal transducers MYD88, NLRP3, and caspase-1—from the host immune system is detrimental for the efficacy of anthracycline-based chemotherapy (Ghiringhelli et al., 2009; Obeid et al., 2007).

An abundant literature has described different DC subsets that would be endowed with the capacity to capture dead cell-associated antigens and to stimulate or subvert anticancer immune responses in vivo (Mellman et al., 2011; Palucka and Banchereau, 2012). Beyond plasmacytoid DCs (pDCs), which are mostly tolerogenic (Matta et al., 2010), conventional DCs (cDCs) have been subclassified into inflammatory and BATF3-dependent DCs. The former exhibit a CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>+</sup> surface phenotype and play an important role in CD4<sup>+</sup> T cell and antimicrobial responses (Kool et al., 2008). The latter express CD8 $\alpha$  and/or CD103 and are particularly efficient at engulfing and cross-presenting antigens from dead cells to CD8<sup>+</sup> T lymphocytes (Albert et al., 1998; Hildner et al., 2008). The differentiation requirements and functional properties of these DC

subsets are highly heterogeneous, suggesting their specific involvement in peculiar sets of immune responses (Mellman et al., 2011; Palucka and Banchereau, 2012).

Here, we addressed the question as to how dead tumor cell antigens are presented to T cells upon anthracycline treatment in vivo. We observed that anthracycline-based chemotherapy stimulates the ATP-driven and purinergic receptor-dependent accumulation of CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> tumor-infiltrating leukocytes (TILs). This subset of TILs appeared particularly efficient at presenting tumor antigens to CD8<sup>+</sup> T cells in vitro and in vivo. Accordingly, experimental manipulations that suppress the recruitment of CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells into the tumor bed, such as the enzymatic degradation of local ATP, the pharmacological inhibition of purinergic receptors, and the injection of CD11b-neutralizing antibodies, all abolished the therapeutic effects of anthracyclines. Our results support the notion that CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> TILs are essential for the efficacy of anthracycline-based chemotherapy.

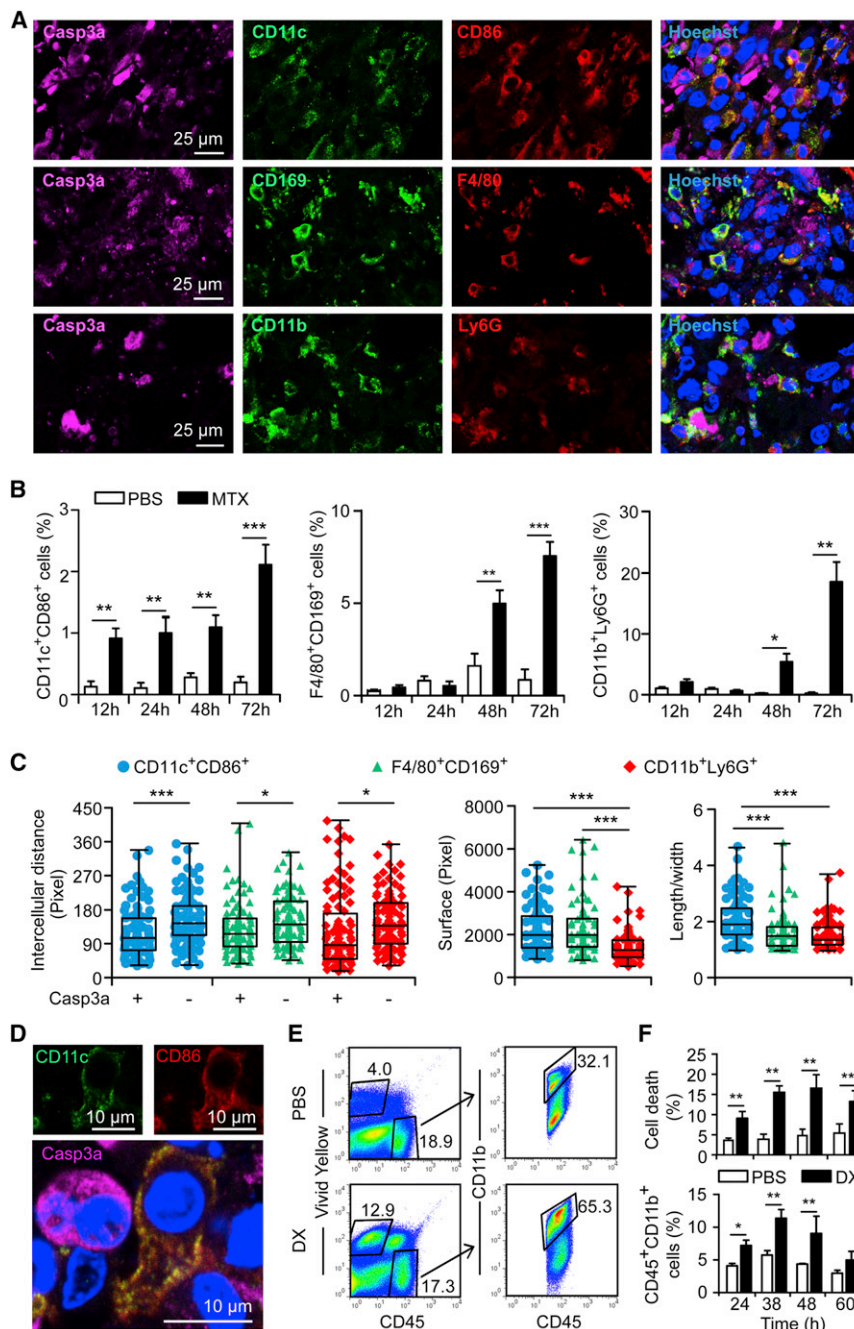
## RESULTS

### Chemotherapy-Induced Recruitment of Myeloid Cells into the Tumor Bed

When murine MCA205 fibrosarcomas growing in syngenic C57Bl/6 mice reached a diameter of 5–10 mm, they contained <1% cells with the surface phenotype of mature DCs (CD11c<sup>+</sup>CD86<sup>+</sup>), macrophages (F4/80<sup>+</sup>CD169<sup>+</sup>), and granulocytes (CD11b<sup>+</sup>Ly6G<sup>+</sup>) and <10% of cells bearing the general myeloid marker CD11b, as detectable by in situ immunofluorescence and confocal microscopy. Systemic chemotherapy with the prototypic anthracycline mitoxantrone (MTX) led to a rapid (12 hr) increase in the prevalence of CD11c<sup>+</sup>CD86<sup>+</sup> cells, followed by a slightly delayed (48 hr) accumulation of F4/80<sup>+</sup>CD169<sup>+</sup> and CD11b<sup>+</sup>Ly6G<sup>+</sup> TILs (Figures 1A and 1B). These three cell types were localized in the immediate proximity of dying tumor cells, as shown by the distance to the closest cancer cell positive for cleaved caspase-3 (a marker of apoptosis), which was significantly shorter than that to the closest living cancer cell (Figures 1C and 1D). As previously reported (Chow et al., 2011), CD11c<sup>+</sup>CD86<sup>+</sup> cells were significantly more elongated than F4/80<sup>+</sup>CD169<sup>+</sup> and CD11b<sup>+</sup>Ly6G<sup>+</sup> TILs, whereas both CD11c<sup>+</sup>CD86<sup>+</sup> and F4/80<sup>+</sup>CD169<sup>+</sup> cells were larger than CD11b<sup>+</sup>Ly6G<sup>+</sup> cells (Figure 1C). Mechanic and enzymatic dissociation of tumors followed by immunofluorescence and cytofluorometric analyses confirmed the recruitment of myeloid (CD45<sup>+</sup>CD11b<sup>+</sup>) cells, and more specifically DCs (CD11c<sup>+</sup>MHC-II<sup>+</sup>), bearing monocytic markers (Ly6C<sup>+</sup>7/4<sup>+</sup>), but not of other cell types such as Treg cells (CD3<sup>+</sup>CD4<sup>+</sup>FOXP3<sup>+</sup>) or Th17 cells (CD3<sup>+</sup>CD4<sup>+</sup>IL-17<sup>+</sup>), into the tumor bed 2 days after chemotherapy with doxorubicin (DX, another widely employed anthracycline), correlating with the induction of cancer cell death (Figures 1E and 1F; Figure S1 available online; data not shown). Altogether, these results indicate that distinct classes of myeloid cells accumulate in the tumor bed after chemotherapy.

### Critical Involvement of ATP and Purinergic Receptors in the Recruitment of Myeloid Cells

To elucidate the mechanisms accounting for the chemotherapy-induced recruitment of myeloid cells into the tumor bed, we



**Figure 1. Chemotherapy Induces the Accumulation of Myeloid Cells in Close Proximity to Dying Cells**

Wild-type C57BL/6 mice bearing MCA205 fibrosarcomas were treated with mitoxantrone (MTX) i.p. (A–D), doxorubicin (DX) i.t. (E, F), or with an equivalent volume of PBS, as a single injection. At the indicated time points (48 hr where not indicated), tumors were harvested and processed for the detection of the indicated proteins by immunofluorescence microscopy (A–D) or for cytofluorometric determinations (E, F). Hoechst 33342 was employed as nuclear counterstaining.

(A) Representative images (scale bars represent 25  $\mu$ m).

(B) Quantitative data on the frequency of the indicated cell populations (means  $\pm$  SEM, in 10 viewfields for at least  $n = 10$  distinct tumors).

(C) The indicated myeloid cell populations were quantitatively characterized for their intercellular distance from tumor cells displaying (+) or not (–) activated caspase 3 (Casp3a) (left), cell surface (middle), and the length/width ratio (right), as determined by morphometric methods. Boxplots report the lower and upper quartile plus the median value. Whiskers indicate min and max values.

(D) A CD11c<sup>+</sup>CD86<sup>+</sup> dendritic cell (DC) in the close proximity of a dying tumor cell (scale bars represent 10  $\mu$ m).

(E) Representative dot plots of cells from PBS-treated and DX-treated tumors are depicted, including dead cells that incorporate the exclusion dye vivid yellow and living CD11b<sup>+</sup>CD45<sup>+</sup> tumor-infiltrating leucocytes. Numbers indicate the percentage of cells found within the corresponding gate.

(F) The corresponding quantitative data (means  $\pm$  SEM).

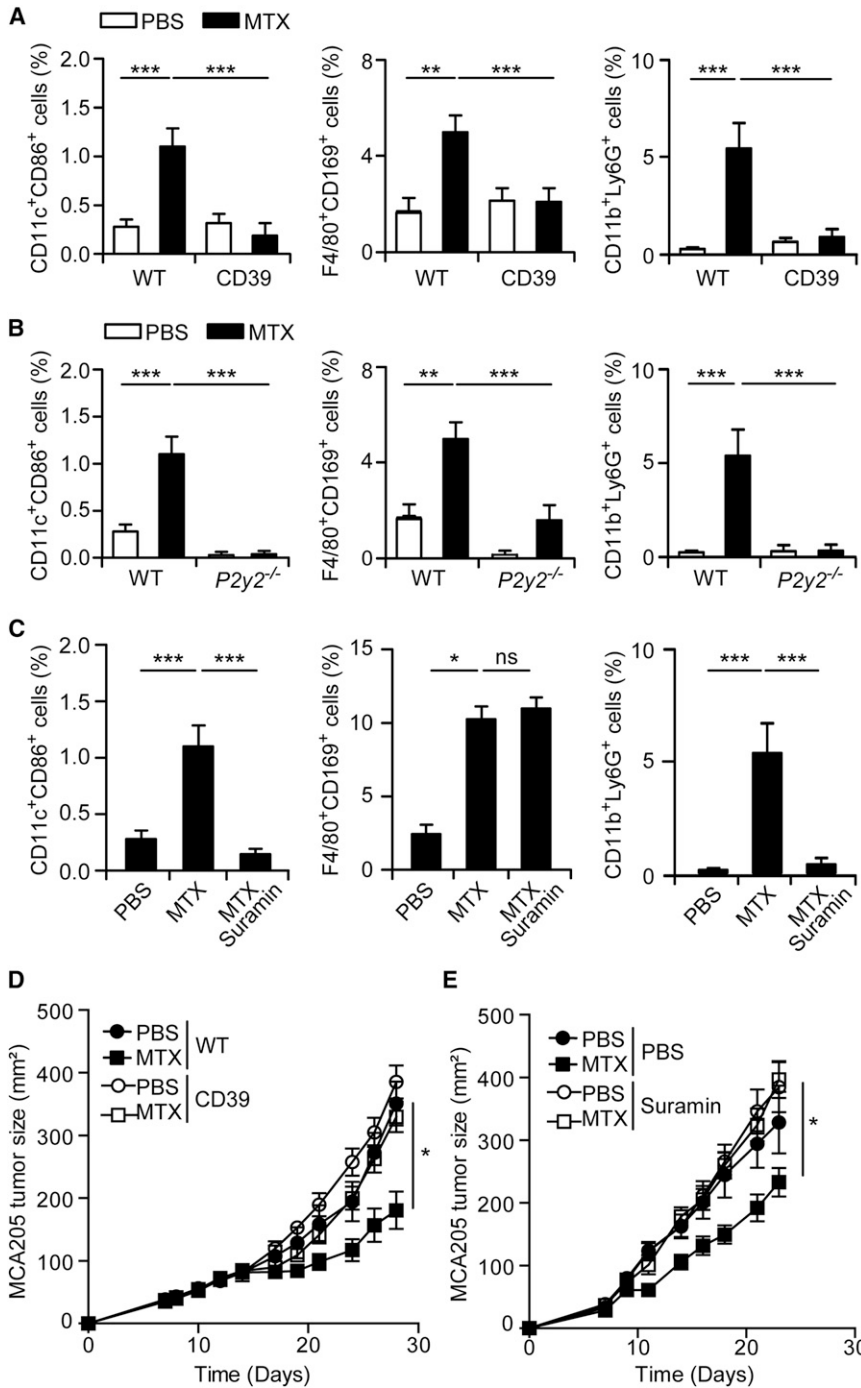
\* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$  (unpaired, two-tailed Student's  $t$  test). See also Figure S1.

took advantage of MCA205 cells that had been engineered to express the ecto-ATPase CD39 on the cell surface (Michaud et al., 2011; Stagg and Smyth, 2010). In response to MTX-based chemotherapy, CD39<sup>+</sup> MCA205 tumors failed to attract CD11c<sup>+</sup>CD86<sup>+</sup>, F4/80<sup>+</sup>CD169<sup>+</sup>, or CD11b<sup>+</sup>Ly6G<sup>+</sup> cells, whereas control tumors (derived from MCA205 cells transduced with an empty vector) did so (Figure 2A), indicating that ATP is essential for the recruitment of these myeloid cells in the tumor bed. Tumors developing in mice that lack the metabotropic purinergic receptor P2Y2 (which mediates the chemotaxis of monocytes toward apoptotic cells) (Chekeni et al., 2010; Elliott et al., 2009) also failed to accumulate CD11c<sup>+</sup>CD86<sup>+</sup>, F4/80<sup>+</sup>CD169<sup>+</sup>, and

local overexpression of CD39 (Figure 2D) and the administration of suramin (Figure 2E) abolished the growth-inhibitory effects of systemic MTX-based chemotherapy in a similar fashion, underscoring the functional importance of ATP and purinergic receptors for the therapeutic efficacy of anthracyclines.

**Identification of APCs among Tumor-Infiltrating Myeloid Cells**

The ablation of tumor-draining lymph nodes shortly after chemotherapy or even before tumor implantation failed to reduce the efficacy of anthracycline-based chemotherapy (Figure S3A). Similar results were obtained upon the injection of a lymphotoxin



**Figure 2. Obligate Contribution of ATP and Purinergic Receptors to Anthracycline-Induced Accumulation of Myeloid Cells and Chemotherapeutic Outcome**

(A–C) Wild-type (WT) (A–C) or *P2y2*<sup>-/-</sup> (B) C57BL/6 mice bearing WT (A–C) or CD39-expressing (CD39) (B) MCA205 fibrosarcomas received mitoxantrone (MTX) i.p. or an equivalent volume of PBS, either alone (A–C) or combined with suramin i.v. (C), and tumor-infiltrating leucocytes of the indicated phenotypes were quantified 48 hr later by immunofluorescence microscopy. Quantitative data are represented as means ± SEM; \**p* < 0.05, \*\**p* < 0.01, \*\*\**p* < 0.001; ns, nonsignificant (unpaired, two-tailed Student's *t* test).

(D and E) WT C57BL/6 mice were inoculated s.c. with WT (D, E) or CD39 (D) MCA205 cells and tumor size was routinely assessed. When tumor became palpable (surface area = 25 mm<sup>2</sup>), mice received MTX i.p. or an equivalent volume of PBS, either alone (D, E) or combined with suramin i.v. (E). Quantitative data are represented as means ± SEM; \**p* < 0.05 (Mann-Whitney U test). See also Figure S2.

against OVA-expressing MCA205 tumors from within the tumor bed (Figure S3E). In this model, the proliferation of intratumoral OT-I cells induced by chemotherapy was not reduced by the surgical removal of draining lymph nodes. Moreover, chemotherapy did not increase the frequency of CD11c<sup>+</sup>MHC-II<sup>+</sup> cells detectable in tumor-draining lymph nodes (data not shown). Altogether, these results suggest that the presentation of tumor antigens relevant to the success of anticancer chemotherapies may take place in the tumor bed itself, even in the absence of draining lymph nodes and tertiary lymphoid structures.

To determine the origin of intratumoral DCs (CD11c<sup>+</sup>MHC-II<sup>+</sup> cells), we established MCA205 cancers in CD45.1 mice, treated them by vehicle or chemotherapy, transplanted them 1 day later to isogenic CD45.2 C57BL/6 recipients, and measured the frequency of tumor-sectile (CD45.1<sup>+</sup>) versus freshly recruited (CD45.2<sup>+</sup>) DCs. The ratio of recipient-

β receptor-Fc fusion protein (LTβR-Fc), serving as a decoy for LTβ (Figure S3B), as well as after the simultaneous neutralization of IL-7 and IL-7R (Figure S3C), two measures that abolish the formation of tertiary lymphoid structures (Gräbner et al., 2009; Meier et al., 2007). Moreover, anthracycline-based chemotherapy failed to enhance (and actually it reduced) the number of TILs exhibiting the surface phenotype of lymphoid tissue inducers (CD45<sup>+</sup>B220<sup>-</sup>CD11c<sup>-</sup>CD3<sup>-</sup>CD4<sup>+</sup>CD127<sup>+</sup>) (Figure S3D), yet led to the activation of tumor-infiltrating CD8<sup>+</sup> T cells and adoptively transferred ovalbumin (OVA)-specific OT-I cells

derived (CD45.2<sup>+</sup>) versus donor-derived (CD45.1<sup>+</sup>) CD11c<sup>+</sup>MHC-II<sup>+</sup> DCs was increased by chemotherapy, supporting the notion that chemotherapy triggers the recruitment of DC precursors rather than a mere in situ differentiation of pre-existing myeloid cells (data not shown). To characterize the intratumoral APCs responsible for the chemotherapy-relevant presentation of tumor antigens, we engineered mouse CT26 colon cancers cells to express EGFP on the inner leaflet of the plasma membrane owing to the addition of a myristoylation motif (Figure 3A). Upon the administration of DX, a significant fraction of CD45<sup>+</sup>

TILs, especially among the CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> subset, became positive for EGFP (Figure 3B). In this context, CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells exhibited a stronger EGFP positivity and higher expression of CD11c than did CD11b<sup>+</sup>Ly6C<sup>lo</sup>Ly6G<sup>-</sup> cells and CD11b<sup>+</sup>Ly6G<sup>+</sup> granulocytes. Moreover, CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells clearly manifested the presence of EGFP in the cytoplasm, an indication of the uptake of antigens from tumor cells. Of note, the frequency of CD11b<sup>+</sup>Ly6C<sup>lo</sup>Ly6G<sup>-</sup> cells, which also engulfed tumor antigens, did not increase in response to chemotherapy (Figures 3C and 3D).

CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells accumulated in the tumor bed after chemotherapy in an ATP-dependent fashion (data not shown), mostly exhibited DC markers (CD11c<sup>+</sup>MHC-II<sup>int</sup>F4/80<sup>lo</sup>), and stained positive for 7/4, CD103, CD172a, CCR2, and CX3CR1, but lacked CD205 expression (Figure S4). When we treated palpable OVA-expressing MCA205 fibrosarcomas with anthracyclines, the frequency of Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells (as determined upon the cytofluorometric purification of the Ly6C<sup>hi</sup>Ly6G<sup>-</sup>, Ly6C<sup>lo</sup>Ly6G<sup>-</sup>, and Ly6G<sup>+</sup> subsets of CD11b<sup>+</sup>TILs) increased, and these cells turned out to be particularly efficient in stimulating the production of IFN- $\gamma$  by MHC class I-restricted OVA-specific OT-I cells *in vitro*. Moreover, Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells recovered from DX-treated OVA-expressing tumors were much more efficient than their Ly6C<sup>lo</sup>Ly6G<sup>-</sup> counterparts as well as than Ly6G<sup>+</sup> cells in stimulating the secretion of IL-2 by MHC class I-restricted OVA-specific B3Z hybridomas (Figure 3E). This effect was antigen specific (as shown by the fact that Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells recovered from OVA-negative tumors failed to stimulate B3Z cells) and required MHC molecules. Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs recovered from homozygous H2K<sup>bD</sup>-deficient hosts (Pérarnau et al., 1999) failed indeed to elicit IL-2 production by B3Z cells. Noteworthy, tumor or stromal cells (namely, the CD45<sup>-</sup> cell subset harvested from anthracycline-treated OVA-expressing MCA205 tumors) failed to directly activate OT-I or B3Z *ex vivo* (Figure 3E).

Next, we purified Ly6C<sup>hi</sup>Ly6G<sup>-</sup>, Ly6C<sup>lo</sup>Ly6G<sup>-</sup>, and Ly6G<sup>+</sup> TILs (all sharing the surface phenotype CD45<sup>+</sup>B220<sup>-</sup>CD3<sup>-</sup>CD11b<sup>+</sup>) from vehicle-treated or DX-treated OVA-expressing MCA205 tumors and adoptively transferred them into naive mice that were challenged 10 days later with living OVA-expressing MCA205 cells. Importantly, only Ly6C<sup>hi</sup>Ly6G<sup>-</sup> (but neither Ly6C<sup>lo</sup>Ly6G<sup>-</sup> nor Ly6G<sup>+</sup>) TILs harvested from anthracycline-treated tumors were capable of reducing the incidence and growth of tumors (Figure 3F), indicating that this population indeed contains functionally competent APCs. Accordingly, purified Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs exhibited a significant increase in the expression of genes relevant to MHC class I- or II-restricted antigen presentation after DX-based chemotherapy, *in vivo* (data not shown). Altogether, these results identify CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs as the cells that accumulate in the tumor bed to exert APC functions in response to anthracycline treatment.

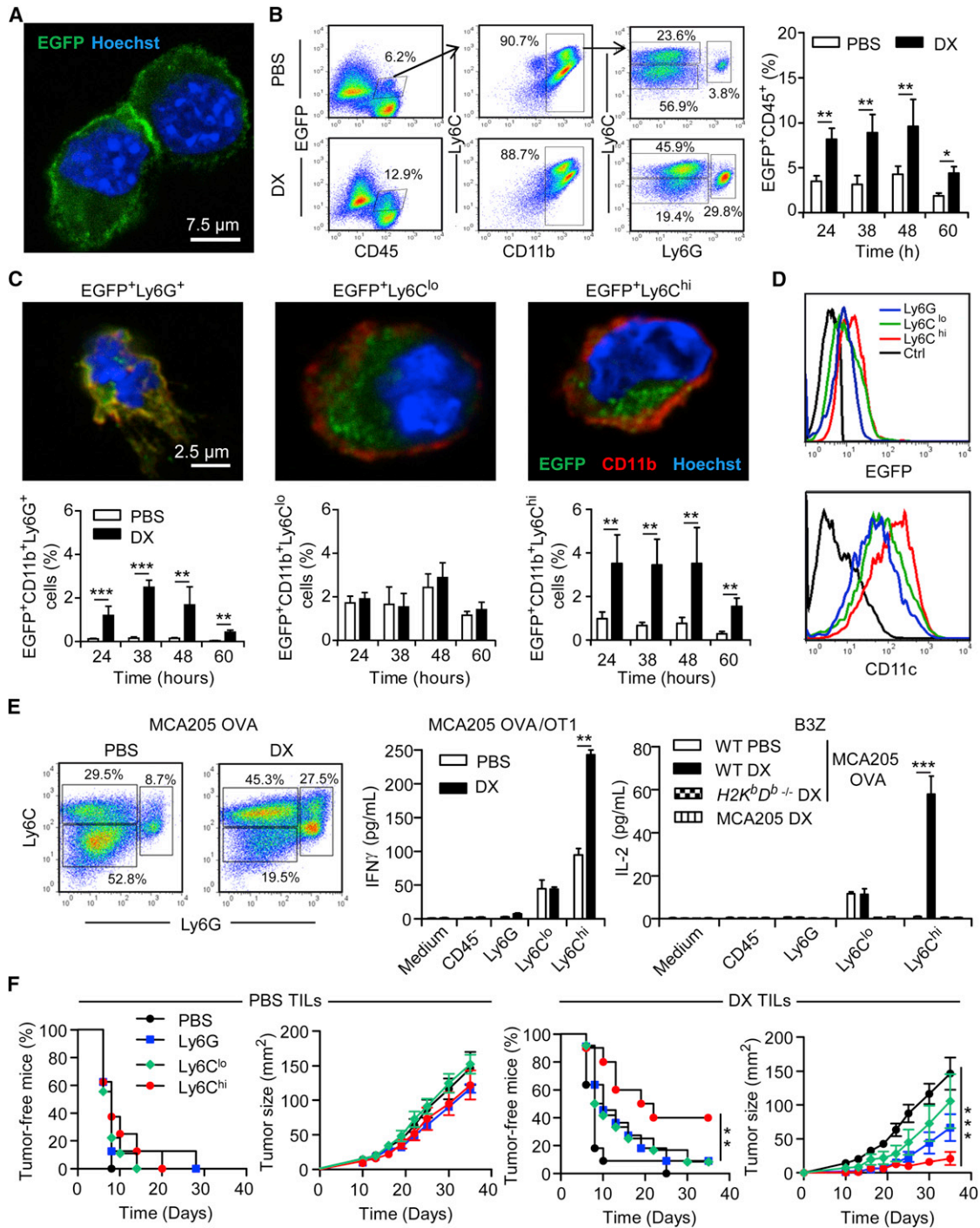
#### ATP-Dependent Differentiation of Myeloid Precursors into Dendritic Cells

Chemotherapy with anthracyclines induced the accumulation of CD34<sup>+</sup>c-Kit<sup>+</sup>Sca-1<sup>+</sup>CD11b<sup>int</sup>CD16/32<sup>+/-</sup> multipotent progenitors (MPPs) and CD34<sup>+</sup>c-Kit<sup>+</sup>Sca-1<sup>-</sup>CD11b<sup>int</sup>CD16/32<sup>hi/lo</sup> myeloid precursors (MPs) in tumors (data not shown). A more

detailed characterization of myeloid TILs (CD45<sup>+</sup>B220<sup>-</sup>CD3<sup>-</sup>CD11b<sup>+</sup>) revealed that several cell subsets, including Ly6C<sup>hi</sup>Ly6G<sup>-</sup>, Ly6C<sup>lo</sup>Ly6G<sup>-</sup>, and Ly6G<sup>+</sup> cells, contained a significant fraction of cells exhibiting a common myeloid progenitor (CMP) or granulocyte-monocyte progenitor (GMP) phenotype (CD34<sup>+</sup>c-Kit<sup>+</sup>Sca-1<sup>-</sup>CD16/32<sup>+</sup>), and more so after chemotherapy. Of note, the frequency of such CMP or GMP-like cells among Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs was reduced by the presence of the ecto-ATPase CD39 on the surface of tumor cells (Figure 4A). Quantitative RT-PCR analyses confirmed that transcription factors and regulators associated with the monocytic lineage (i.e., PU.1, Nab2) (Auffray et al., 2009; Laslo et al., 2006) were upregulated, in particular among Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs, after DX-based chemotherapy, and that this increase was abolished by the expression of CD39 (Figure 4B). The expression of transcription factors relevant for the differentiation of granulocytes (i.e., Gfi1) (de la Luz Sierra et al., 2010; Laslo et al., 2006) or the megakaryocytic-erythrocytic lineage (i.e., Gfi1b and Gata1) (Yu et al., 2009) was increased in response to DX, yet predominantly among Ly6G<sup>+</sup> cells and independently of CD39 (Figure 4B). In line with these observations, Ly6C<sup>hi</sup>Ly6G<sup>-</sup> (but not Ly6G<sup>+</sup>) TILs recovered from tumors and propagated in dedicated methylcellulose cloning conditions could form colonies that were highly proliferative (Figures 4C–4E). These proliferating cells retained GMP markers and acquired the morphology and phenotype of monocytes (Figure 4F and data not shown). Anthracycline-based chemotherapy dramatically increased the frequency of colony-forming units among Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs, an effect that was precluded by the expression of CD39 by tumor cells (Figure 4C). Thus, chemotherapy appears to induce the recruitment of GMPs with monocytic differentiation potential at the tumor site, in an ATP-dependent manner.

#### Chemotherapy Induced the ATP-Dependent Differentiation of CD11b<sup>+</sup>Ly6C<sup>hi</sup> Tumor-Infiltrating Leucocytes into Inflammatory DC-like Cells

We characterized the impact of the intratumoral milieu on the differentiation of Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs into bona fide DCs, placing special emphasis on ATP. Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs were recovered from tumors established in CD45.2 C57BL/6 mice and treated with anthracyclines and then adoptively transferred into MCA205 tumors growing in congenic CD45.1 mice that were treated with either vehicle (PBS) or DX. Upon transfer into tumors that did not receive chemotherapy, the initial CD45.2<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cell population mostly differentiated into CD45.2<sup>+</sup>Ly6G<sup>+</sup> granulocytes. In contrast, upon transfer into anthracycline-treated tumors, a fraction of Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs maintained the Ly6C<sup>hi</sup>Ly6G<sup>-</sup> phenotype and acquired surface markers of DCs (namely CD11c and MHC class II molecules) (Figures 5A–5C). Similar results were obtained in a syngeneic (CD45.2) system in which Ly6C<sup>hi</sup>Ly6G<sup>-</sup> or Ly6G<sup>+</sup> TILs recovered from anthracycline-treated tumors were labeled with CFSE and then transferred into anthracycline-treated MCA205 tumors. A substantial fraction of such Ly6C<sup>hi</sup>Ly6G<sup>-</sup>CFSE<sup>+</sup> cells acquired the expression of CD11c and CD86 (Figures 5D and 5E). Failure to provide chemotherapy as well as transfer into CD39-expressing tumors reduced the frequency of the adoptively transferred cells that could be recovered from the tumor bed (Figures 5E and 5F). As a control, the Ly6G<sup>+</sup> population, which did not contain significant



**Figure 3. Chemotherapy Promotes the Uptake and Presentation of Tumor Antigens by CD11c<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> Tumor-Infiltrating Leucocytes**

(A–D) BALB/c mice bearing MyrPalm-EGFP-expressing CT26 tumors were treated with doxorubicin (DX) i.t. or with an equivalent volume of PBS. At the indicated time points (38 hr, where not indicated), tumors were harvested and processed for immunofluorescence studies (A and C) or cytofluorometric determinations (B and D).

(A) MyrPalm-mEGFP-expressing mouse colon carcinoma CT26 cells, exhibiting a plasma membrane-restricted distribution of EGFP (scale bar represents 7.5  $\mu$ m).

(B) Representative dot plots of EGFP<sup>+</sup>CD45<sup>+</sup> tumor-infiltrating leucocytes (TILs) stained for CD11b, Ly6C, and Ly6G are illustrated on the left, and quantitative data are reported on the right.

(C) Uptake of tumor antigens (as exemplified by EGFP) by myeloid cells of the indicated phenotype (top, scale bar represents 2.5  $\mu$ m) and quantitative kinetic data (bottom).

(D) Representative cytofluorometric distribution of EGFP and CD11c mean fluorescence intensity for the indicated myeloid cell populations.

(A and C) Hoechst 333342 was employed as nuclear counterstaining.

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numbers of precursor cells (Figure 4C), failed to generate CD11c<sup>+</sup>CD86<sup>+</sup> cells upon adoptive transfer into anthracycline-treated tumors (Figure 5E). Altogether, these results suggest that chemotherapy-elicited ATP release within the tumor bed can skew the differentiation of CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> TILs toward a CD11c<sup>+</sup> DC-like phenotype, hence subverting their default pathway leading to the generation of Ly6G<sup>+</sup> granulocytes, which we observed within PBS-treated tumors (Figure 5C).

To get further insights into this issue, we cocultured control or CD39<sup>+</sup> MCA205 cells that were optionally treated with anthracyclines in vitro (CD39 expression abolished the anthracycline-driven accumulation of extracellular ATP; Figure 5G) with bone marrow-derived CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> cells, a substantial part (around 30%–40%) of which showed a GMP phenotype. Only anthracycline-treated CD39<sup>-</sup> (but not CD39<sup>+</sup>) tumor cells stimulated such a subset to express the monocyte-relevant transcription factor PU.1 (Figure S5). Along similar lines, only CD39<sup>-</sup> cells treated with anthracyclines (but neither anthracycline-treated CD39<sup>+</sup> cells nor untreated CD39<sup>-</sup> cells) enhanced the differentiation of CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup> bone marrow cells into CD11b<sup>+</sup>Ly6C<sup>+</sup> monocytes (de facto limiting granulocytic differentiation) and allowed for the generation of a significant number of CD11c<sup>+</sup>MHC-II<sup>+</sup> DC-like cells (Figures 5H and 5I). This skew in differentiation from Ly6G<sup>+</sup> granulocytes to cells with a CD11b<sup>+</sup>CD11c<sup>+</sup> DC phenotype was reduced by the pannexin 1 inhibitor DIDS (which limits ATP release from dying cells) and by several purinergic receptor antagonists including oxidized ATP and suramin, but neither by recombinant IL-1 receptor antagonist (IL-1RA) nor by the TNF- $\alpha$  antagonist etanercept (Figures 5J and 5K). These results underscore the importance of extracellular ATP and purinergic receptors for the generation of DCs in response to anthracycline-based chemotherapy.

#### Identification of the DC Subset Underlying Chemotherapy-Induced Anticancer Immune Responses

Multiple distinct DC subsets might contribute to the presentation of dead cell antigens to T cells. To identify the specific DC population that is relevant to anticancer chemotherapy, we first injected a PDCA-1-specific antibody into tumor-bearing mice that had received anthracycline-based chemotherapy. Although the PDCA-1-targeting antibody effectively depleted plasmacytoid DCs (data not shown), this treatment neither suppressed the chemotherapy-induced accumulation of CD11b<sup>+</sup>CD11c<sup>-</sup>GFP<sup>+</sup> TILs (in mice that express GFP under the control of the CD11c promoter) (Figure 6A) nor reduced the efficacy of anthracycline- and oxaliplatin-based chemotherapy (Figures 6B and 6C). Along similar lines, the deletion of the transcription factor *Batf3*, which is essential for the generation of CD8 $\alpha$ <sup>+</sup>CD103<sup>+</sup>Langerin<sup>+</sup> DCs (Hildner et al., 2008), failed to reduce MTX-

induced tumor infiltration by CD11b<sup>+</sup>CD11c<sup>+</sup> TILs (Figure 6D), and tumors implanted in wild-type and *Batf3*<sup>-/-</sup> mice responded to chemotherapy in an indistinguishable fashion (Figure 6E). In contrast, the injection of a CD11b-blocking antibody strongly reduced the frequency of CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells in the peripheral blood (Figure 6F), as well as of CD11b<sup>+</sup>CD11c<sup>-</sup>GFP<sup>+</sup> cells infiltrating MTX-treated tumors (Figure 6G). In accord with the finding that CD11b<sup>+</sup>CD11c<sup>+</sup> TILs are responsible for chemotherapy-related antigen presentation and antitumor protection in vivo (Figure 3), neutralization of CD11b abolished the anticancer effect of anthracyclines in four different tumor models, namely MCA205 fibrosarcomas and AT3 mammary cancers, both of which grow in C57BL/6 mice (Figures 6H and 6I), F244 sarcoma cells, which form tumors in 129/Sv mice (Figure 6J), and H2N100 mammary cancers, developing in BALB/c mice (Figure 6K). In contrast, the depletion of macrophages (with clodronate-containing liposomes) or neutrophils (with the 1A8 antibody) had no or only minor effects, respectively, on the efficacy of chemotherapy (Figure S6 and data not shown). Altogether, these results underscore the fundamental importance of inflammatory DC-like CD11b<sup>+</sup>CD11c<sup>+</sup> cells for the efficacy of anticancer chemotherapy.

#### DISCUSSION

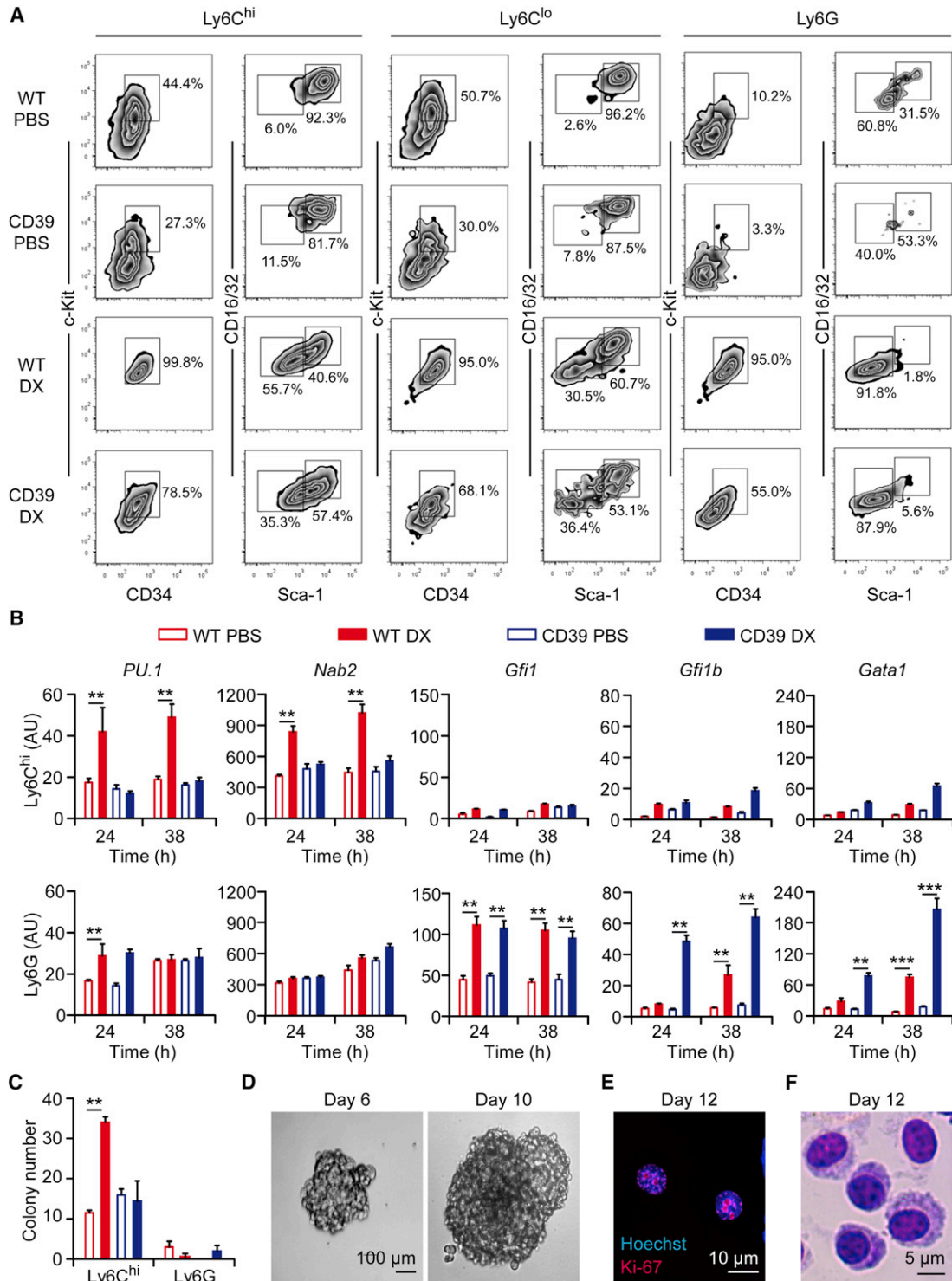
Our findings provide at least three lines of evidence indicating that a peculiar subtype of TILs, namely CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells, which share some of the characteristics of inflammatory DCs (Cheong et al., 2010), is particularly important for the induction of anticancer immune responses by anthracyclines. First, CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells were recruited into the tumor bed as early as 12 hr after chemotherapy. Second, manipulations aimed at preventing the intratumoral accumulation of such cells, including (1) the local overexpression of the ATP-degrading enzyme CD39, (2) the pharmacological blockade of purinergic receptors, and (3) the neutralization of CD11b with a specific antibody, abolished the immune-dependent anticancer effects of anthracyclines. Third, functional experiments revealed that CD11c<sup>+</sup>CD11b<sup>+</sup>Ly6C<sup>hi</sup> TILs are particularly efficient in capturing and presenting tumor cell antigens and that such cells—once adoptively transferred into naive mice—can confer protection against a challenge with living tumor cells.

ATP appears to be particularly important for the infiltration of myeloid TILs into the tumor bed upon anthracycline-based chemotherapy. Indeed, palpable tumors generated from cancer cells manipulated to overexpress the ecto-ATPase CD39 failed to recruit TILs of various types, including cells with a DC-like, macrophage-like, or granulocyte surface phenotype in response to anthracyclines. In line with recent results (Chen et al., 2006), P2Y2 turned out to be essential for the chemotherapy-induced

(B and C) Quantitative data are represented as means  $\pm$  SEM; \* $p$  < 0.05, \*\* $p$  < 0.01, \*\*\* $p$  < 0.001 (unpaired, two-tailed Student's  $t$  test).

(E) WT or *H2k<sup>b</sup>D<sup>b</sup>-/-* C57BL/6 mice bearing OVA-expressing (OVA) MCA205 fibrosarcomas were treated as in (A)–(D) and CD11b<sup>+</sup> TILs were purified by cytofluorometry 38 hr after chemotherapy upon staining with Ly6C- and Ly6G-specific antibodies (gating exemplified on the left). Sorted cells were cocultured with OT-I cells for 72 hr or with OVA-specific T cell hybridoma B3Z cells for 36 hr and supernatants were assessed for interferon- $\gamma$  (IFN- $\gamma$ , middle) and interleukin-2 (IL-2, right) levels. Numbers indicate the percentage of cells found within the corresponding gate. Quantitative data are represented as means  $\pm$  SEM; \*\* $p$  < 0.01, \*\*\* $p$  < 0.001 (unpaired, two-tailed Student's  $t$  test).

(F) Alternatively, PBS (negative control) or myeloid cells sorted as in (E) from PBS-treated (left) or DX-treated (right) tumors were inoculated in the footpad of naive WT C75BL/6 mice, which were rechallenged 10 days later with living OVA MCA205 tumor cells. Tumor incidence and growth were then routinely assessed. Results from one representative experiment out of three independent ones involving ten mice/group is depicted. \*\* $p$  < 0.01, \*\*\* $p$  < 0.001 (log-rank test). See also Figure S3.



**Figure 4. Chemotherapy-Induced and ATP-Dependent Enrichment of Granulocyte Monocyte Progenitors in Tumor-Infiltrating Leucocytes** C57BL/6 mice bearing wild-type (WT) or CD39-expressing (CD39) MCA205 fibrosarcomas were treated with doxorubicin (DX) i.t. or with an equivalent volume of PBS. (A) 24 hr later, tumors were harvested and the expression of hematopoietic progenitor markers (CD34, c-Kit, Sca-1, and CD16/32) on CD45<sup>+</sup>CD3<sup>-</sup>B220<sup>-</sup>CD11b<sup>+</sup> (Ly6C<sup>hi</sup>, Ly6C<sup>lo</sup>, or Ly6G<sup>+</sup>) tumor-infiltrating cells was analyzed by flow cytometry. Numbers indicate the percentage of cells found within the corresponding gate. Please note that the CD34<sup>+</sup>c-Kit<sup>+</sup>Sca-1<sup>-</sup>CD16/32<sup>+</sup> phenotype marks common myeloid progenitors (CMPs) as well as granulocyte-monocyte progenitor (GMPs). (B–F) Alternatively, tumors were harvested at the indicated time points (24 hr, where not indicated), and CD11b<sup>+</sup>Ly6C<sup>hi</sup> and CD11b<sup>+</sup>Ly6G<sup>+</sup> tumor-infiltrating cells were characterized by RT-PCR for the expression of transcriptional regulators (the monocytic factors PU.1 and Nab-2, the granulopoietic factor Gfi-1, and the erythroid and megakaryocytic factors Gata-1 and Gfi1b) (B) or used in clonogenic assays in methylcellulose-based medium (C–F).

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recruitment of TILs to the tumor bed. In contrast, pharmacological inhibition of purinergic receptors with suramin was more efficient in blocking the chemotherapy-elicited infiltration of tumors by DCs and granulocytes than that by macrophages.

Lymphadenectomy failed to affect the therapeutic response of mice to anthracyclines, nor did the neutralization or blockade of the  $LT\beta$ - $LT\beta R$  and  $IL-7$ - $IL-7R$  systems. These findings suggest that TILs accumulating in the tumor upon chemotherapy can mediate anticancer immune responses without the need to migrate into lymph nodes nor to form tertiary lymphoid structures. It appears therefore plausible that  $CD11c^+CD11b^+Ly6C^{hi}$  TILs may efficiently process and present tumor cell antigens (which they readily take up *in vivo*, as indicated by the transfer of cancer cell-derived EGFP into their cytoplasm) locally, within the tumor bed.

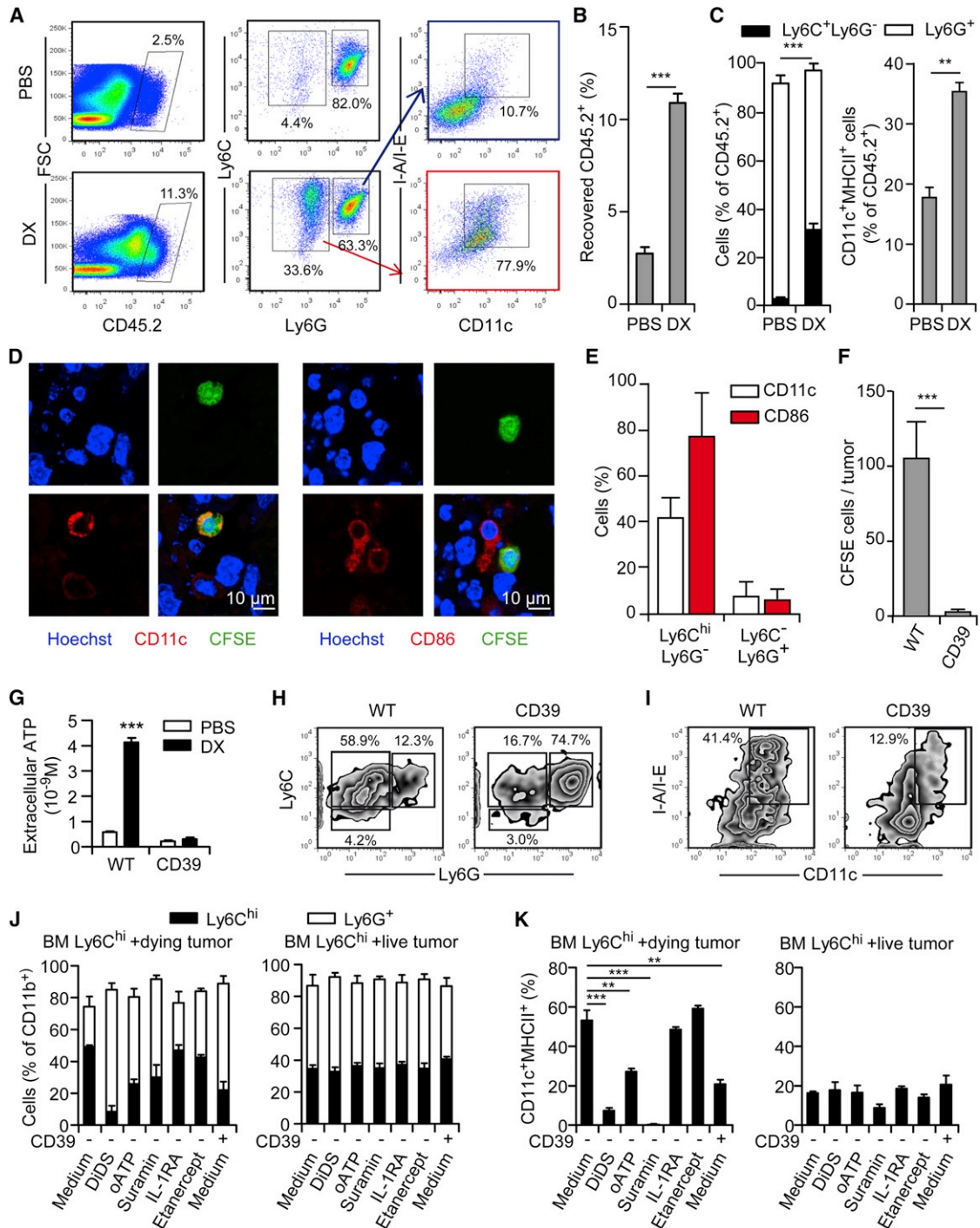
ATP can be released from dying cells in response to physiological cell death inducers (such as glucocorticoids acting on thymocytes) (Chekeni et al., 2010; Elliott et al., 2009) or therapeutic interventions including radiotherapy and chemotherapy (Martins et al., 2009). Within tissues, the concentration of extracellular ATP ( $[ATP]_e$ ) in the proximity of stressed and dying cells is believed to decrease along a steep gradient, owing to the presence of ecto-ATPases that are expressed by a variety of cell types (including  $FOXP3^+$  Treg and Th17 cells) (Borsellino et al., 2007; Chalmin et al., 2012; Schreiber et al., 2011) as well as of soluble, plasmatic ATPases (Yegutkin et al., 2003). Such a pattern of  $[ATP]_e$  may facilitate the migration of myeloid cells toward dying cells. Our results are compatible with the notion that ATP acts as a specific chemotactic (and perhaps chemotropic) signal that emanates from dying cells. Indeed,  $CD11c^+CD11b^+Ly6C^{hi}$  TILs, as well as other myeloid populations, were found within nests of dying cells, and—more precisely—in close proximity of cells that manifested apoptotic caspase-3 activation (Kroemer et al., 2007). Moreover, ATP may affect the differentiation of pluripotent myeloid cell populations recruited into the tumor bed, in line with its previously reported capacity to promote the differentiation of hematopoietic stem cells (Barbosa et al., 2011) and the cardinal importance of purinergic receptors for DC differentiation *in vitro* (Wilkin et al., 2001).  $CD11b^+Ly6C^{hi}Ly6G^-$  TILs were found to contain a subset of GMPs that became particularly enriched after chemotherapy, provided that CD39 was not expressed by cancer cells. In the absence of dying tumor cell-derived ATP,  $CD11b^+Ly6C^{hi}Ly6G^-$  TILs (as well as their bone marrow-derived counterparts) preferentially differentiated into granulocytes. The capacity of anthracycline-treated cancer cells to skew the differentiation of  $CD11b^+Ly6C^{hi}Ly6G^-$  TILs toward an inflammatory DC-like phenotype depended on ATP and purinergic receptors, both *in vitro* and *in vivo*. Altogether, these observations suggest a dual role for ATP in the tumor microenvironment. First, ATP would be essential for the recruitment of myeloid cells into the proximity of dying cells and, second, ATP would facilitate the local differentiation and maintenance of cells with the phenotype of inflammatory DCs.

Multiple distinct DC subsets have been involved in anticancer immunosurveillance (Palucka and Banchereau, 2012), and advanced tumors can actually convert tumor-infiltrating DCs into immunosuppressive cells (Gabrilovich et al., 2012). In the specific context of anthracycline-based chemotherapy, however, only inflammatory DC-like  $CD11c^+CD11b^+Ly6C^{hi}$  cells were found to be essential for optimal immune system-mediated anticancer responses. Indeed, several distinct manipulations suppressing the accumulation of such inflammatory DC-like cells in the tumor bed also abolished the antitumor activity of anthracycline-based chemotherapy. In contrast, the depletion of pDCs, neutrophils, or macrophages with antibodies targeting PDCA-1 and Ly6G or clodronate-containing liposomes, as well as the elimination of  $Batf3$ -dependent DCs, had no major impact on chemotherapeutic responses. Intriguingly, inflammatory DCs, which are particularly important for chemotherapy-elicited immunosurveillance, can differentiate in response to HMGB1 (Han et al., 2008), suggesting that HMGB1 released from dying cancer cells may further contribute to the maturation and/or differentiation of TILs toward such phenotype. Irrespective of these possibilities, our results unravel the essential contribution of one particular DC-like subset, namely the  $CD11c^+CD11b^+Ly6C^{hi}$  population, to the therapeutic anticancer immune response elicited by anthracyclines.

The question as to whether chemotherapy elicits *de novo* T cell priming and/or reactivates local effector and/or memory T cells remains elusive. As outlined by De Visser and colleagues (Ciampricotti et al., 2012), oncogene-addicted nonimmunogenic breast tumors fail to prime a significant T cell response when they are treated with anthracyclines. However, we speculate that chemotherapy-induced immunogenic cell death may at least reboot antigen-specific T cell responses that have been initiated prior to chemotherapy. Our findings indicate that (1) draining lymph nodes are dispensable for the immune system-mediated antitumor effects of anthracyclines, (2) inflammatory DCs accumulate within neoplastic lesions in response to anthracyclines, and these cells function as efficient APCs when extracted from tumor beds, and (3) no significant  $CD8^+$  T cell (OT-I) priming against the immunodominant epitopes of OVA is detectable in the spleen and lungs by day 7 after chemotherapy (data not shown), although such response occurs within the tumor. In line with these notions, pegylated  $IL-10$ -based therapy has been shown to reactivate intratumoral cancer-specific cytotoxic  $CD8^+$  TILs, to restore the *in situ* expression of Th1 cytokines, granzymes, and antigen-presentation molecules, and to culminate in the control of tumor growth, even in the absence of draining lymph nodes (Mumm et al., 2011). However, our data do not formally exclude the possibility that antigen presentation may take place at extratumoral sites, such as the bone marrow, to which tumor antigens may be transported by neutrophils (Duffy et al., 2012). Irrespective of this partially unresolved issue, we show here that anthracyclines promote a crucial ATP-dependent pathway whereby the intratumoral accumulation of GMPs and

(B and C) Quantitative data (means  $\pm$  SEM); \*\* $p < 0.01$ , \*\*\* $p < 0.001$  (unpaired, two-tailed Student's *t* test).

(D–F) Morphology of colonies generated from  $CD11b^+Ly6C^{hi}$  TILs (scale bar represents 50  $\mu m$ ) (D), Ki-67<sup>+</sup> dots in the nucleus of cells, as assessed by immunofluorescence microscopy upon Hoechst 33342 counterstaining (scale bar represents 10  $\mu m$ ) (E), and May-Grünwald-Giemsa staining of a cytospin prepared from these cells (scale bar represents 5  $\mu m$ , original magnification 400 $\times$ ) (F). See also Figure S4.



**Figure 5. Chemotherapy-Induced and ATP-Dependent Differentiation of CD11b<sup>+</sup>Ly6C<sup>hi</sup> Tumor-Infiltrating Leucocytes into Inflammatory DC-like Cells**

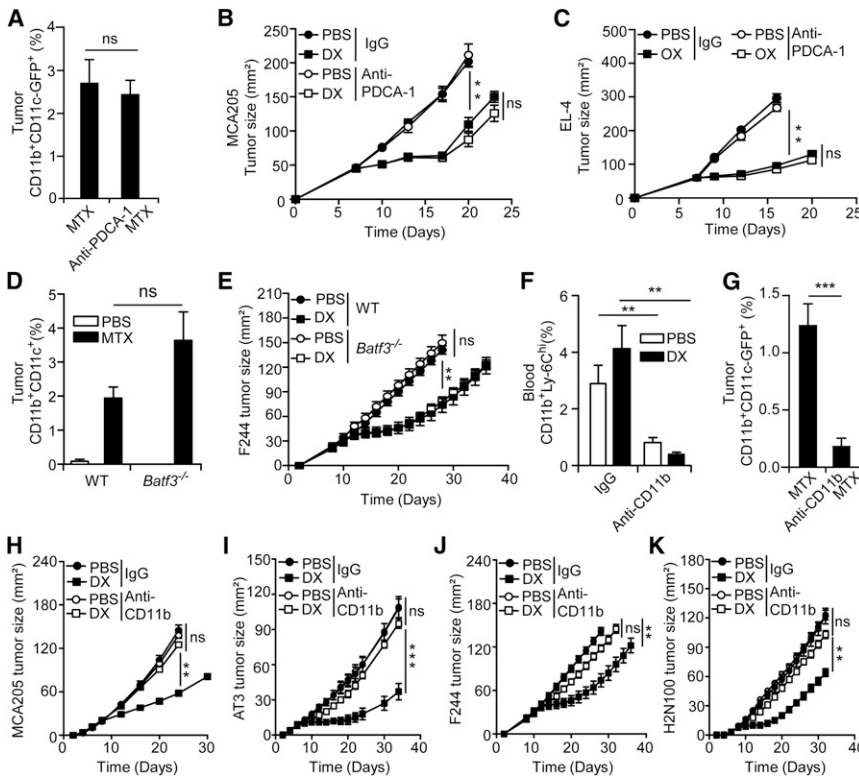
(A–C) CD11b<sup>+</sup>Ly6C<sup>hi</sup> cells purified from doxorubicin (DX)-treated tumors from C57BL/6 CD45.2 mice were injected into MCA205 tumors implanted in C57BL/6 CD45.1 mice that had been pretreated with DX i.t. or with an equivalent volume of PBS. After 48 hr, tumors were collected and the phenotype of transferred CD45.2<sup>+</sup> cells was analyzed by flow cytometry.

(A) Representative dot plots (numbers indicate the percentage of cells found within the corresponding gate).

(B and C) Quantitative data on the percentage of recovered CD45.2<sup>+</sup> cells (B) and on their differentiation into CD11b<sup>+</sup>Ly6C<sup>+</sup>Ly6G<sup>-</sup>, CD11b<sup>+</sup>Ly6C<sup>+</sup>Ly6G<sup>+</sup>, or CD11c<sup>+</sup>MHC-II<sup>+</sup> cells (C) (means ± SEM).

(D–F) CD11b<sup>+</sup>Ly6C<sup>hi</sup> or Ly6G<sup>+</sup> cells sorted from MCA205 fibrosarcomas treated with DX were labeled with CFSE and inoculated into wild-type (WT) or CD39-expressing (CD39) MCA205 fibrosarcomas that had been pretreated 48 hr earlier with mitoxantrone (MTX) i.p. or with an equivalent volume of PBS. 48 hr later, tumors from recipient mice were harvested and processed for immunofluorescence microscopy-assisted determinations. Representative images are depicted in

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**Figure 6. The Therapeutic Efficacy of Anthracyclines Relies on CD11b<sup>+</sup> Myeloid DCs but Neither on Plasmacytoid DCs nor on *Batf3*-Dependent DCs**

(A) CD11c-DTR/GFP mice bearing wild-type (WT) MCA205 fibrosarcomas were treated with mitoxantrone (MTX) i.p., combined with either PDCA-1-specific or isotype control antibodies (IgGs). Nine days later, the percentage of CD11c<sup>+</sup>CD11b<sup>+</sup>GFP<sup>+</sup> tumor-infiltrating cells was determined by immunofluorescence microscopy. Quantitative data are reported as means ± SEM; ns, nonsignificant (unpaired, two-tailed Student's t test). (B and C) C57BL/6 mice were inoculated with WT MCA205 fibrosarcoma (B) or EL-4 lymphoma (C) cells (day -7) and, on day 0, they were treated with doxorubicin (DX) i.t. (B) or oxaliplatin (OX) i.p. (C), combined with either PDCA-1-targeting antibodies or isotype control IgGs on days -1, 1, and 2. Tumor growth was routinely assessed throughout the experiment (n = 5–8 mice/group). Quantitative data are reported as means ± SEM; ns, nonsignificant, \*\*p < 0.01 (Mann-Whitney U test). (D) WT or *Batf3*<sup>-/-</sup> mice bearing F244 sarcomas were treated with MTX i.p. or an equivalent volume of PBS. Two days later, tumors were harvested and processed for the immunofluorescence microscopy-assisted detection of CD11b<sup>+</sup>CD11c<sup>+</sup> DCs within the tumor bed (n = 4–5 mice/group). Quantitative data are reported as means ± SEM; ns, nonsignificant (unpaired, two-tailed Student's t test).

(E) WT or *Batf3*<sup>-/-</sup> 129/Sv mice were inoculated with WT F244 sarcoma cells (day -7) and, on day 0, they were treated with DX i.t. or an equivalent volume of PBS. Tumor growth was routinely assessed throughout the experiment (n = 5–8 mice/group). Quantitative data are reported as means ± SEM; ns, nonsignificant, \*\*p < 0.01 (Mann-Whitney U test).

(F and G) CD11c-DTR/GFP mice bearing WT MCA205 fibrosarcomas were treated with MTX i.p., DX i.t., or with an equivalent volume of PBS, combined with either CD11b-specific antibodies or isotype control IgGs, which were administered 1 day before and on the same day as chemotherapy. 20 and 38 hr later, blood (F) and tumor (G) samples were collected and analyzed by flow cytometry and immunofluorescence microscopy, respectively, for the abundance of CD11b<sup>+</sup>Ly6C<sup>hi</sup> and CD11c<sup>+</sup>CD11b<sup>+</sup>GFP<sup>+</sup> cells, respectively. Quantitative data are reported as means ± SEM; \*\*p < 0.01, \*\*\*p < 0.001 (unpaired, two-tailed Student's t test). (H–K) C57BL/6 (H and I), 129/Sv (J), or BALB/c (K) mice were inoculated with WT MCA205 (H), AT3 (I), F244 (J), and H2N100 (K) cells (day = -7) and, on day 0, they were treated DX i.t. or an equivalent volume of PBS, combined with either CD11b-specific antibodies or isotype control IgGs (n > 5 mice/group). Quantitative data are reported as means ± SEM; ns, nonsignificant, \*\*p < 0.01, \*\*\*p < 0.001 (Log-rank Mann-Whitney U test).

See also Figure S6.

inflammatory monocytes in fine facilitate the local differentiation of inflammatory DCs and the activation of T cells against cancer.

**EXPERIMENTAL PROCEDURES**

**Tumor Chemotherapy Models**

For the establishment of syngenic solid tumors, CB57BL/6 mice received EL-4, AT3, WT MCA205, or OVA-expressing MCA205 cells, BALB/c mice were inoculated with H2N100, WT CT26, or MyrPalm-mEGFP-expressing CT26 cells, and WT or *Batf3*<sup>-/-</sup> 129/Sv mice were engrafted with F244 cells. Tumor growth was then monitored routinely by means of a caliper, and when the tumor surface

reached 25–45 mm<sup>2</sup> (day 0), mice received either DX, MTX, oxaliplatin, or equivalent volumes of solvent. When appropriate, mice received intravenous suramin or an equivalent volume of solvent, on the same day of chemotherapy. Alternatively, mice received on days -1, 0, and weekly thereafter until the end of the experiment CD11b-, PDCA-1-, or Ly6G-targeting antibodies. In some protocols, tumor-bearing mice received on day -1, 1, and twice weekly thereafter until the end of the experiment IL-7- or IL-7R-specific antibodies or LTβR-Fc fusion proteins (kindly provided by Y.-X. Fu, University of Chicago). All experiments were performed in accordance with the Federation of European Laboratory Animal Science Association (FELASA) guidelines and were approved either by the IGR Ethics Committee (CEEA IRCIV/IGR no. 26, registered with the French Ministry of Research) or by the Peter MacCallum Animal Experimentation Ethics Committee.

(D) (scale bars represent 10 μm), and quantitative data (means ± SEM, n = 2 independent experiments with 3–5 mice/group) are reported in (E) (percentage of CD11c<sup>+</sup> and CD86<sup>+</sup> cells) and (F) (number of CFSE<sup>+</sup> cells/tumor).

(G–K) WT and CD39 MCA205 tumor cells were treated with PBS or 25 μM DX for 20 hr, followed by the quantification of extracellular ATP by a luciferase-based assay (G) or the coculture with bone marrow-derived (BM) Ly6C<sup>hi</sup>Ly6G<sup>+</sup> cells for 5 days in the presence of 5 ng/ml GM-CSF and 5 ng/ml IL-4, alone or combined with 10 μM 4,4'-diisothiocyano-2,2'-stilbenedisulfonic acid (DIDS), 500 μM oxidized ATP (oATP), 2 mM suramin, 20 μg/ml interleukin-1 receptor antagonist (IL-1RA), or 20 μg/ml etanercept (H–K). (H) and (I) depict dot plots representative of the commitment toward the monocytic (Ly6C<sup>+</sup>Ly6G<sup>-</sup>) or granulocytic (Ly6G<sup>+</sup>) lineage, and toward a DC-like phenotype (CD11c<sup>+</sup>MHCII<sup>+</sup>), respectively. Numbers indicate the percentage of cells found within the corresponding gate. (J) and (K) summarize quantitative data (means ± SEM) on the percentage of monocytic (CD11b<sup>+</sup>Ly6C<sup>hi</sup>Ly6G<sup>-</sup>) (J), granulocytic (CD11b<sup>+</sup>Ly6G<sup>+</sup>) (J), and DC-like (CD11c<sup>+</sup>MHCII<sup>+</sup>) (K) cells obtained.

\*\*p < 0.01, \*\*\*p < 0.001 (unpaired, two-tailed Student's t test). See also Figure S5.

**Lymphadenectomy**

For lymph node removal, mice were anesthetized under continuous isoflurane flow and subjected to skin decontamination with 10% povidone-iodine (commercial Betadine solution). Thereafter, a narrow incision (3 mm) was made, and the inguinal lymph node was removed with dissecting forceps. Finally, the incision was closed with sterile skin closure clips. Sham-operated mice (without lymph node resection) were treated similarly, and the duration of surgery was standardized between the two groups. Placing mice on a heater plate set at 37°C until awakening facilitated recovery.

**Colony-Forming Assays**

Myeloid subpopulations sorted from tumors were resuspended in MethoCult GF M3434 medium (StemCell Technologies) and 1 ml cell suspension was dispensed into 35 mm vented Petri dishes (Greiner Bio-One) when bubbles dissipated. Cultures were then incubated for 7–14 days in standard conditions in the presence of sterile water to preserve humidity.

**ATP Quantification**

ATP in culture supernatants was quantified by means of the luciferin-based ENLITEN ATP Assay Kit (Promega), as per manufacturer's instructions, on a Fluostar multiwell plate reader (BMG Labtech).

**Statistical Analyses**

Unless otherwise indicated, results are expressed as means  $\pm$  SEM of  $n = 3$  parallel assessments. All experiments were repeated at least twice, yielding similar results. Normal distributions were compared by unpaired, two-tailed Student's *t* tests, and tumor growth curves were compared by the Mann-Whitney U test. Kaplan-Meier survival curves were analyzed with the log-rank test. Statistical analyses were performed by means of the software Prism 5 (GraphPad) or Excel 2007 (Microsoft). *p* values  $< 0.05$  were considered as statistically significant.

**SUPPLEMENTAL INFORMATION**

Supplemental Information includes Supplemental Experimental Procedures and six figures and can be found with this article online at <http://dx.doi.org/10.1016/j.immuni.2013.03.003>.

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