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# **Patrolling Monocytes Control Tumor Metastasis to the Lung**

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## **Abstract**

The immune system plays an important role in regulating tumor growth and metastasis. For example, classical monocytes promote tumorigenesis and cancer metastasis; however, how nonclassical "patrolling" monocytes interact with tumors is unknown. Here we show that patrolling monocytes are enriched in the microvasculature of the lung and reduce tumor metastasis to lung in multiple mouse metastatic tumor models. Nr4a1-deficient mice, which specifically lack patrolling monocytes, showed increased lung metastasis in vivo. Transfer of Nr4a1-proficient patrolling monocytes into Nr4a1-deficient mice prevented tumor invasion in lung. Patrolling monocytes established early interactions with metastasizing tumor cells, scavenged tumor material from the lung vasculature and promoted natural killer cell recruitment and activation. Thus, patrolling monocytes contribute to cancer immunosurveillance and may be targets for cancer immunotherapy.

> Monocytes and monocyte-derived macrophages play key roles in tumor progression (1-4). Classical "inflammatory" monocytes (CCR2<sup>high</sup>Ly6C<sup>+</sup> in mouse; CCR2<sup>high</sup>CD14<sup>+</sup>CD16<sup>-</sup> in humans) are recruited to tumor sites where they contribute to tumor macrophage content and promote tumor growth and metastasis (5, 6). In contrast, very little is known about the role

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of nonclassical "patrolling" monocytes (CX3CR1<sup>high</sup>Ly6C<sup>-</sup> in mouse;  $CX3CR1<sup>high</sup>CD14<sup>dim</sup>CD16<sup>+</sup>$  in humans) in the early growth and metastasis of tumors. Patrolling monocytes (PMo) are involved in the resolution of inflammation; they actively survey the endothelium of the vasculature where they scavenge microparticles and remove cellular debris (7-9). The orphan nuclear receptor Nr4a1 (also known as Nur77/TR3/Ngfi-b) is highly expressed in PMo compared to other immune cells and functions as a master regulator for the development of PMo in mice (10).

To investigate the actions of PMo during early tumor metastasis, we utilized mice expressing green fluorescent protein (GFP) under control of the Nr4a1 promoter (Nr4a1-GFP mice). In these mice, PMo but not  $Ly6C^+$  classical monocytes, are GFPhigh (10, 11). We focused our studies on the lung, which is a common site of tumor metastasis and is an important site of PMo activity (12-14). We confirmed that Nr4a1-GFPhigh cells in the lung were PMo by flow cytometry (fig. S1A-C). Tracking of Nr4a1-GFPhigh cells by confocal imaging in the lungs identified a large number of Nr4a1-GFPhigh PMo patrolling the microvasculature (Movie 1, Fig. 1A). Consistent with an important role for PMo in the lung vasculature, we found a 3-4 fold enrichment of Nr4a1-GFPhigh PMo in the lung compared to other tissues (fig. S1D).

To examine the interactions of PMo with tumor in vivo, we imaged Nr4a1-GFPhigh PMo in the lung after intravenous (IV) injection of Lewis Lung Carcinoma cells expressing red fluorescent protein (LLC-RFP). The number of Nr4a1-GFPhigh monocytes in the lung increased significantly at 24 hrs following injection of LLC-RFP cells, implying that PMo are actively recruited to the lung tumor environment (Fig. 1A). Within 4 hrs after tumor injection, most Nr4a1-GFPhigh monocytes decreased patrolling speed in the vasculature, and by 24 hrs they had arrested near lung tumor sites (Fig. 1B). The majority of Nr4a1-GFPhigh monocytes isolated from lung after LLC tumor transfer maintained their PMo phenotype, which we further confirmed in vitro (Fig. 1C, fig. S1).

Nr4a1-GFPhigh PMo were recruited to tumor cell clusters within 30 minutes after IV tumor injection, and recruitment continued for at least 7 days (Fig. 1D-E, Movie 2-5). Nr4a1- GFPhigh cells that were recruited to lung tumor sites were not positive for Ly6C/G (GR-1) in vivo, further confirming that these Nr4a1-GFPhigh cells are not Ly6G<sup>+</sup> granulocytes or Ly6C+ classical monocytes (Movie 4). The kinetics of PMo recruitment to the lung differed from that of  $Ly6C<sup>+</sup>$  monocytes (fig. S2A). At 7 days, there were significantly higher numbers of Nr4a1-GFP<sup>high</sup> PMo ( $\sim$ 24/100  $\mu$ m<sup>3</sup>) associated with tumor areas compared to tumor-free areas, confirming active recruitment of PMo to the tumor (Fig. 1D-E, Movie 5, fig. S1B).

Nr4a1-GFPhigh monocytes patrolling the vasculature at 4 hrs after tumor injection appeared to move towards and inhibit the attachment of tumor cells to the lung microvasculature (Movie 3). We examined if Nr4a1-GFPhigh PMo could extravasate outside the vasculature. We found that by 4 hrs post-tumor injection 10-20% of Nr4a1-GFPhigh PMo had extravasated at tumor sites (fig. S2B-C), and this increased to 40-50% PMo extravasation by 7 days. Together, these findings confirm that PMo establish early immune interactions with tumor cells and can extravasate and accumulate at tumor sites.

In order to determine if PMo have a major role in regulating tumor invasion, metastasis and growth in the lungs in vivo, we used Nr4a1-knockout ( $Nr4a1^{-/-}$ ) mice, which have a selective loss of PMo (10) (fig. S3).  $Nr4a1^{-/-}$  mice were IV injected with either syngeneic B16F10 melanoma cells expressing a luciferase reporter or LLC-RFP cells (Fig. 2A-B, fig. S4-5). As early as 24 hrs and up to 21 days after IV injection of B16F10 melanoma, we observed increased tumor invasion in the lungs of  $Nr4aI^{-/-}$  mice compared to control mice (Fig. 2A-B, fig. S4A-B). We observed no differences in either  $Ly6C^+$  monocyte or  $Ly6G^+$ granulocyte populations in the lungs of these mice at 7 days after tumor injection (fig. S4C). B16F10 tumor invasion appeared specific for the lung, as increased tumor metastasis was not observed in the liver (fig S4B). Additionally, increased spontaneous metastases to the lung were observed in  $Nr4a1^{-/-}$  mice after subcutaneous injection of B16F10 melanoma, suggesting that Nr4a1 expression is important for suppressing primary tumor metastasis to the lung (Fig. 2C). A similar early and sustained increase in lung metastasis in  $Nr4a1^{-/-}$ mice was also observed after intravenous LLC tumor transfer (fig. S5). We did not detect differences in vascular permeability in the lung between  $Nr4a1^{-/-}$  and control mice (fig. S6).

We next utilized the MMTV-PyMT model, in which female mice spontaneously develop mammary tumors that metastasize to the lung (15). To focus on Nr4a1 function exclusively in hematopoietic cells, we performed bone marrow transplants using either WT or  $Nr4a1^{-/-}$ bone marrow transferred into female recipient MMTV-PyMT mice. MMTV-PyMT mice receiving  $Nr4a1^{-/-}$  bone marrow developed significantly higher numbers of spontaneous metastases to lung, but no differences in primary mammary tumor growth compared to mice receiving WT bone marrow (Fig. 2D-E).

We further tested hematopoietic Nr4a1 function using B16F10 melanoma. Only mice receiving  $Nr4a1^{-/-}$  bone marrow had increased B16F10 tumor metastases, confirming that Nr4a1 expression in hematopoietic cells regulated tumor cell metastasis to the lung (fig. S7A-B). Analysis of immune cells isolated from lung tumors verified a selective loss of PMo in  $Nr4aT^{-/-}$  bone marrow transplanted mice (fig. S7C). In the 1:1 chimera mice, equal reconstitution of immune cells from each donor was observed (fig. S7D). However, PMo were derived almost exclusively from WT bone marrow, suggesting that the restoration of the Ly6C− monocyte population prevented tumor metastasis.

To confirm if Nr4a1 expressed in myeloid cells was regulating tumor metastasis to the lung, we examined two different myeloid-specific Nr4a1 conditional knockout models (CSF1R-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> and LysM-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup>). Deletion of Nr4a1 using CSF1R-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> and LysM-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> mice significantly reduced PMo in circulation (fig. S8), and increased tumor lung metastasis (Fig. 3A-B, fig. S9). Nr4a1 deletion using CSF1R-Cre or LysM-Cre also targets Nr4a1 in macrophages and Ly6C+ monocytes, so we cannot completely rule out effects of Nr4a1 in these cells. However, Nr4a1 expression in macrophages and  $Ly6C^+$ monocytes is relatively low suggesting limited Nr4a1 function (10, 16, 17). No differences in tumor metastasis were observed with T lymphocyte-specific Nr4a1 deletion (fig. S10). Collectively, our studies illustrate increased lung metastasis burden in the absence of Nr4a1 in myeloid cells in multiple cancer models.

To confirm a direct role for PMo in regulating tumor metastasis, wild-type Ly6C<sup>+</sup> or Ly6C<sup>−</sup> monocytes were adoptively transferred into recipient  $Nr4a1^{-/-}$  mice prior to tumor injection. A significant number of the transferred monocytes could be found in lung (fig. S11A-B). Reconstitution of PMo into  $Nr4a1^{-/-}$  mice prevented lung tumor metastasis (Fig. 3C-D). In contrast, transfer of Ly6C<sup>+</sup> monocytes into  $Nr4a1^{-/-}$  mice actually promoted tumor metastasis, consistent with known protumoral properties of this subset of monocytes (5, 18). The majority (80-90%) of transferred Ly6C<sup>+</sup> monocytes in circulation did not lose Ly6C expression (fig. S11C). Interestingly, transfer of PMo 24 hrs post-tumor injection into  $Nr4a1^{-/-}$  mice did not suppress tumor metastasis (fig. S11D), suggesting that PMo need to already be present and active in the vasculature in order to prevent early tumor metastasis. These data directly show that nonclassical PMo inhibit tumor metastasis to the lung.

PMo can act as "intravascular housekeepers" that scavenge microparticles and remove cellular debris from the microvasculature (7). Extracellular vesicles from tumors are important mediators of tumor metastasis, progression and immune suppression, and targeting their removal is an emerging target for cancer therapy (19, 20). We used highresolution confocal imaging to determine if PMo could engulf and remove tumor material from the lung vasculature. A sizable number of Nr4a1-GFPhigh PMo containing large amounts of LLC-RFP tumor material were observed at tumor sites in the lung 24 hrs after IV tumor transfer (Fig. 3E). Co-culture assays of mouse PMo with fluorescent-labeled tumor cells confirmed engulfment of large amounts of tumor material (Fig. 3F). Analysis of monocyte populations isolated from the lung at 24 hrs after IV LLC-RFP injection indicated that PMo preferentially took up tumor material, taking up  $\sim$  5-fold more tumor than did  $Ly6C<sup>+</sup>$  classical monocytes (Fig. 3G). PMo also preferentially took up significantly more B16F10-YFP tumor material with an average size of 1.39  $\mu$ m<sup>2</sup>, and an average total amount of tumor material per monocyte of  $1.92 \mu m^2$  (fig. S12A-B). The homologous human CD14 $\text{dimCD16}^+$  population of PMo, which similarly has high Nr4a1 expression (8, 21), also engulfed a large quantity of tumor material in vitro, suggesting analogous tumor engulfment function (fig. S12C). Moreover, PMo actively engulfed tumor material within classic endocytic compartments (22)(Fig. 4A). Collectively, these results demonstrate that Nr4a1 dependent PMo rapidly and preferentially endocytose tumor material.

We then asked how PMo recognized tumor cells to prevent metastasis in the lung. The chemokine receptor CX3CR1 is highly expressed on PMo and is important for their arrest at inflammatory sites (fig. S13A) (23-25). CX3CR1-deficient ( $Cx3cr1^{-/-}$ ) mice, which also have a significant reduction in PMo, exhibit a similar phenotype to  $Nr4a1^{-/-}$  mice—of increased tumor burden and metastasis to the lung (26, 27). Though PMo were reduced in numbers in the lung vasculature of  $Cx3cr1^{-/-}$  mice (about 30-50% reduction, fig. S13B; confirming previous reports) (27), a major proportion of the CX3CR1-deficient PMo remaining were observed patrolling the vasculature as has previously been observed (7). Unlike  $Cx3cr1^{+/-}$  or WT PMo,  $Cx3cr1^{-/-}$  PMo did not arrest near LLC tumor cells, and instead remained patrolling within the lung vasculature (Fig. 4B-C, Movie 6). Interestingly,  $Cx3cr1<sup>-/-</sup>$  PMo were not recruited to the lung 24 hrs after LLC tumor challenge, while Ly6C<sup>+</sup> recruitment was unaffected by loss of CX3CR1 expression (fig. S13B). Importantly,  $Cx3cr1^{-/-}$  PMo present in the lung showed defective engulfment of tumor material,

indicating that CX3CR1 expression on PMo is critical for mediating the sensing and uptake of tumor material (Fig. 4D).

CX3CL1, a ligand for CX3CR1, has been reported to be high in human and mouse lung (28). Using a CX3CL1-mCherry reporter mouse (29), we found that CX3CL1 was specifically expressed on  $CD31<sup>+</sup>$  endothelial cells (EC) at low levels in the lung microvasculature (Fig. 4E-F, fig. S13C). CX3CL1 expression was most prevalent in lung EC compared to EC in other tissues (fig. S13D), which may partially explain the enrichment and preferential function of PMo in the lung. CX3CL1 expression on lung EC increased in response to tumor challenge (Fig. 4E), and at sites of lung tumor metastasis (Fig 4F), consistent with reports of increased CX3CL1 during lung inflammation (30).

TLR7 has been linked to recruitment of PMo in response to kidney damage in mice (7). However, we found that TLR7 did not play a significant role in either recruitment of PMo to the lung after tumor injection (fig. S13B), or in the uptake of tumor material by PMo (Fig. 4D). We conclude that both CX3CR1 expression on monocytes and CX3CL1 expression by EC are critical for recruitment of PMo to sites of tumor extravasation to mediate the removal of tumor material from the lung. CX3CL1 expression by tumor cells (31) may also drive monocyte recruitment. In agreement with our findings, many studies have reported that CX3CL1 expression by either tumor cells or by tumor-associated cells is anti-tumoral and correlated with good prognosis (32-34). However, the function of the CX3CL1/CX3CR1 axis, particularly during later stages of tumor growth, is complex (35, 36).

Finally, we examined whether PMo can directly kill tumor cells. After multiple attempts using various experimental conditions, direct killing of tumor cells by PMo was not observed (fig. S14). However, PMo may be important for antibody-dependent cell-mediated cytotoxicity of either tumor cells or suppressive immune cells within the tumor environment (37, 38). In response to IV injected B16F10 tumor, PMo isolated from lung produced significantly higher levels of natural killer (NK) cell activation and recruitment-related chemokines CCL3, CCL4 and CCL5 compared to classical Ly6C+ monocytes (Fig. 4G)(39, 40). Correlating with this finding, myeloid-specific Nr4a1 knockout mice (CSF1R- $Cre^+Nr4a1<sup>f1/f1</sup>$ ) showed reduced NK cell recruitment to the lung in response to tumor (Fig. 4H), suggesting that PMo controlled the recruitment of NK cells to tumor sites. A similar reduction in NK cell recruitment, and CD44 activation (fig. S15A-B) was also found in lungs of PyMT mice that received Nr4a1-deficient bone marrow. However, Nr4a1 does not regulate NK cell development (fig. S15C). Uptake of tumor material by PMo does not require the presence of NK cells (fig. S16A-B). Importantly, NK cell depletion reduced the differences in metastasis between WT and CSF1R-Cre+Nr4a1 $f<sup>1/f1</sup>$  mice (fig. S16C). Thus, PMo inhibit metastasis, at least in part, through the regulation of NK cell recruitment and activity.

In summary, we demonstrate a cancer surveillance role of PMo wherein they prevent tumor metastasis to lung. PMo are actively recruited to lung metastasis sites in a CX3CR1 dependent manner, where they function to scavenge tumor material as well as recruit and activate NK cells that leads to the prevention of tumor cell metastasis (fig. S17). Selective

targeting by increasing PMo activity and/or their regulation by Nr4a1 may represent a novel therapy for preventing cancer metastasis to the lung.

#### **Supplementary Material**

Refer to Web version on PubMed Central for supplementary material.

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Please see detailed methods for this experiment in the Materials and Methods section of Supplementary Materials.

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### **One Sentence Summary**

Nonclassical patrolling monocytes aid in cancer surveillance by preventing tumor metastasis to the lung.

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**Fig. 1. Nr4a1-GFPhigh monocytes patrol the vasculature, and interact with tumor in the lung** (A) Quantification of Nr4a1-GFPhigh PMo per μl of blood volume in lung (Untreated), 4 hrs or 24 hrs after IV LLC-RFP transfer (n=5 mice per group). (B) Quantification of Nr4a1- GFPhigh monocyte movement in lung before (Untreated), 4 hrs or 24 hrs after LLC-RFP tumor injection. Monocyte tracks transposed to a common origin from a representative 20 min movie (left, scale bar=100 μm), and quantification of median speed of monocytes (right, combined speed data from analysis of 3 separate mice; \*=p<0.001 lower than untreated; \*\*=p<0.001 than 4hr tumor). (C) Representative gating of Nr4a1-GFPhighCD11b<sup>+</sup> cells from all Live  $CD45^{\circ}CD11c^{\text{low}}$  cells 24 hrs after IV LLC-RFP transfer. (D) Representative confocal image of Nr4a1-GFPhigh monocytes (Green) interacting with LLC-RFP cells (Red) in the lung 7 days after IV LLC-RFP transfer. Immune cells in the vasculature were labeled with IV injected anti-CD45 antibody (Blue). (E) Quantification of free (>100 μm from tumor site) and tumor-associated  $\langle$  <50 µm from tumor site) Nr4a1-GFPhigh monocytes in the lung at various time points after tumor injection (combined analysis of 5 mice per group; p<0.01 for each tumor-associated area compared to free tumor area for each time point).

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#### **Fig. 2. Increased lung metastasis of tumors in** *Nr4a1***−/− mice**

(A) In vivo luciferase detection in wild-type control (WT) and  $Nr4a1^{-/-}$  mice 24 hrs after IV injection of  $5\times10^5$  B16F10 melanoma cells expressing luciferase (Left) and quantification (Right) (\*=p<0.03, representative experiment with 5 mice per group). (B) In vivo luciferase detection (Left), and quantification (Right), in WT and  $Nr4a1^{-/-}$  mice 7 days after IV injection with  $3\times10^5$  B16F10-luciferase cells (\*=p<0.001, n=18 mice per group combined from 3 separate experiments). (C) Number of spontaneous tumor metastases per  $5000 \mu m^2$ of lung surface 28 days after SubQ injection of  $1 \times 10^5$  B16F10-YFP cells (\*=p<0.01, n=7 mice per group). (D-E) Lung tumor metastasis in MMTV-PyMT mice reconstituted with WT (WT:PyMT) or  $Nr4a1^{-/-}$  (Nr4a1<sup>-/-</sup> :PyMT) bone marrow. (D) Representative MMTV-PyMT mouse lung histology stained with hematoxylin and eosin (Left), and quantification of number of spontaneous lung metastases per 5000  $\mu$ m<sup>2</sup> of lung surface (Right). \*=p<0.05 n=12 for WT and n=15 for  $Nr4a1^{-/-}$ . (E) Quantification of primary breast tumor growth in MMTV-PyMT mice. N=12 for WT and n=15 for  $Nr4a1^{-/-}$ .

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#### **Fig. 3. Nr4a1-expressing patrolling monocytes reduce tumor metastasis and engulf tumor material in the lung**

(A) In vivo imaging (Left), and quantification (Right) of tumor in lungs of CSF1R-Cre<sup>−</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>−</sup>) or CSF1R-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>+</sup>) mice 7 days after IV injection of  $3\times10^5$  B16F10-luciferase tumor cells (n=6 mice per group, \*=p<0.01; experiment replicated 2 times). (B) Quantification of the number of tumor metastases per lungs of CSF1R-Cre<sup>−</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>−</sup>) and CSF1R-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>+</sup>) mice 7 days after IV injection of  $3\times10^5$  B16F10-YFP tumor cells (n=8 mice per group, \*=p<0.01). (C-D)  $Nr4a1^{-/-}$  mice were injected IV with 5×10<sup>5</sup> wild-type Ly6C<sup>-</sup> PMo, Ly6C<sup>+</sup> inflammatory monocytes, or PBS at day 0. On day  $1, 3 \times 10^5$  B16F10-luciferase tumor cells were injected IV and tumor metastasis and growth were measured by *in vivo* imaging at day 8. Representative in vivo imaging (C), and quantification (D) of B16F10-luciferase metastasis 8 days after monocyte transfer and 7 days after tumor transfer in wild-type (WT) or  $Nr4a1^{-/-}$  mice. (Combined data from 5 separate experiments with n=2 mice per group; \*=p<0.01 statistically different than WT; \*\*=p<0.05 statistically different than  $Nr4aT^{-/-}$ ). (E) Imaging of tumor material uptake in lung by Nr4a1-GFPhigh monocytes 24 hrs after IV injection of LLC-RFP tumor cells. Representative higher magnification images to right. (F) Uptake of LLC-RFP tumor material by CX3CR1-GFPhighLy6C− PMo after 24 hrs of coculture. Note that Nr4a1-GFP expression is primarily nuclear so monocyte cell membranes are not visible in images (G) Representative flow plot (Left) and quantification (Right) of tumor material uptake by all monocytes in the lung 24 hrs after IV tumor injection of  $3\times10^5$ LLC-RFP cells ( $n=4$  mice per goup;  $*=p<0.01$ ; experiment replicated 3 times).

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#### **Fig. 4. Patrolling monocytes detect tumor material in a CX3CR1-dependent manner and recruit NK cells to the lung tumor environment**

(A) Ratio of fluorescent intensity of tumor material engulfed by PMo (black) to fluorescent intensity of whole tumor (black and grey) 3 hrs after IV LLC tumor injection. LLC tumors were labeled with either CellTrace Violet control dye (Control) or a pH-sensitive pHrodo Red dye (pHrodo) and then IV injected in a 1:1 ratio into a wild-type mouse  $(n=3$  mice per group of representative experiment replicated 3 times). Representative tracking (B) and median speed (C) of  $Cx3cr1^{-/-}$  or  $Cx3cr1^{-/+}$  monocyte movement 24 hrs after IV tumor injection in the lung. Monocyte tracks transposed to a common origin from representative 20 min movies (scale bar=100  $\mu$ m, representative tracks shown from one mouse, median speed calculated from tumor areas analyzed in 3 separate mice per group,  $*=p<0.001$ . (D) Percentage of Ly6C− PMo containing LLC-RFP tumor material in the lung 3 hrs after IV injection of tumor into representative wild-type (WT),  $Cx3cr1^{-/-}$ , or  $Tlr7^{-/-}$  mice (left), and quantification of tumor material uptake (right) (n=3 per group;  $*p<0.001$  versus WT). (E) Percentage of CD31<sup>+</sup> CX3CL1<sup>+</sup> lung endothelial cells isolated from Untreated (UN) or 24 hrs or 7 days after IV injection of B16F10-YFP tumor cells into CX3CL1-mCherry mice. (F) Representative imaging of CX3CL1-mCherry (Red) expression in lung 24 hrs after IV injection of B16F10-YFP tumor cells (Green) in CX3CL1-mCherry mice. CD45<sup>+</sup> immune cells are labeled in blue. (G) Relative chemokine mRNA expression in Ly6C<sup>+</sup> or Ly6C<sup>−</sup> monocytes isolated from lung by FACS 24 hrs after IV B16F10 tumor injection (monocyte

populations isolated from three separate mice; \*=p<0.01; experiment repeated 3 times). (H) Percentage of natural killer (NK) cells in the lungs of CSF1R-Cre<sup>−</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>−</sup>) or CSF1R-Cre<sup>+</sup>Nr4a1<sup>fl/fl</sup> (CSF1R-Cre<sup>+</sup>) mice 7 days after IV injection of  $3\times10^5$  B16F10luciferase tumor cells (n=6 mice per group,  $*=p<0.01$ ).