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Reduced microglial CX3CR1 expression delays neurofibromatosis-1 glioma formation

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Abstract

While traditional models of carcinogenesis have largely focused on neoplastic cells, converging data have revealed the importance of non-neoplastic stromal cells in influencing tumor growth and progression. Leveraging a genetically-engineered mouse model of NF1-associated optic glioma, we now demonstrate that stromal microglia express the CX3CR1 chemokine receptor, such that reduced CX3CR1 expression decreases optic nerve microglia. Moreover, genetic reduction of *Cx3cr1* expression in *Nf1* optic glioma mice delays optic glioma formation. Coupled with previous findings demonstrating that microglia maintain optic glioma growth, these new findings provide a strong preclinical rationale for the development of future stroma-directed glioma therapies in children.

Keywords

genetically-engineered mice; monocyte; astrocytoma; brain tumor; fractalkine

Introduction

Neurofibromatosis type 1 (NF1) is the most common inherited tumor predisposition syndrome in which affected children develop low-grade glial neoplasms (pilocytic astrocytomas). These gliomas primarily form in the optic pathway, such that tumor progression is often associated with reduced visual acuity.¹ Histological analysis reveals a heterogeneous cellular composition, including neoplastic cells (astroglial cells) lacking *NF1* gene expression and nonneoplastic cells (microglia and endothelial cells) with reduced *NF1* gene expression. Since NF1-optic pathway gliomas (OPGs) are rarely resected, mechanistic insights have largely derived from studies using *Nf1* genetically-engineered mouse (GEM) strains.

One of the important observations to emanate from the analysis of *Nf1* GEM models is that *Nf1* inactivation in astroglial progenitors is insufficient for gliomagenesis unless coupled with reduced *Nf1* gene expression (*Nf1*^{+/-} microenvironment).² The requirement for *Nf1*^{+/-} stromal cells in gliomagenesis, together with similar findings using *Nf1* GEM models of plexiform neurofibroma,³ establishes a critical role for a supportive microenvironment in NF1-associated tumorigenesis.

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Based on the abundance of microglia in human sporadic and NF1-associated pilocytic astrocytomas,⁴ we have previously demonstrated that optic glioma proliferation (maintenance) is attenuated by pharmacologic or genetic inhibition of microglia function.^{5,6} In the current study, we apply a genetic approach using *Nf1* GEM strains to show that CX3CR1⁺ microglia are required for optic glioma formation *in vivo*.

Methods

Mice

All mice were maintained on a C57BL/6 background in accordance with established Washington University Animal Studies protocols. The mice employed in this study were generated by successive intercrossing (**Supplementary Table 1**).

Optic nerve measurements

Following paraformaldehyde fixation, optic nerves/chiasmata were microdissected and photographed.⁷ Optic nerve diameters were measured at the chiasm and at 150, 300, and 450 μm anterior to the chiasm, and the volumes of the three 150 μm cones summed to calculate total optic nerve volume.

Immunohistochemistry and immunofluorescence

Following anesthetization, mice were transcardially perfused with Ringer's solution and then 4% paraformaldehyde in 0.1M sodium phosphate buffer (pH 7.4). Optic nerves with intact chiasmata were post-fixed overnight and processed for either paraffin or OCT embedding. Immunohistochemistry was performed using appropriate antibodies (**Supplementary Table 2**) for 3,3'-diaminobenzidine development or immunofluorescence microscopy. Images were acquired on a Nikon Eclipse TE300 microscope (Tokyo, Japan) equipped with an optical camera (Optronics, Goleta, CA). Image J image analysis (<http://rsbweb.nih.gov/ij/>; National Institute of Mental Health, Bethesda, MD) and MetaMorph Microscopy Automation & Image Analysis (Molecular Devices, LLC, Sunnyvale, CA) software were used to obtain single collapsed fluorescence images.

Apoptosis was assessed by terminal deoxynucleotidyl transferase-mediated biotinylated UTP nick end labeling (TUNEL) staining according to the manufacturer's instructions (Roche Diagnostics, Indianapolis, IN).

Flow cytometry (FACS)

Microglia from pools of 7-10 mice per experiment were collected and processed for antibody-mediated flow cytometry (**Supplementary Table 2**) using appropriate controls for gating.⁶

Statistical analyses

Investigators were blinded to mouse genotypes and treatments, and percentages of positive cells were calculated using the total number of cells in each image (hematoxylin or DAPI nuclear staining). Data were analyzed with Graphpad Prism 5 software using a Student's t-test, and outliers were excluded using Grubbs' test. Data are displayed as mean \pm S.E.M.

Results

Previous studies from our laboratory have demonstrated that short-term pharmacologic (minocycline) inhibition or genetic ablation (ganciclovir treatment of CD11b-TK-expressing mice) of microglia at 3 months of age was sufficient to reduce tumor proliferation in

Nfl^{+/-}-GFAP^{CKO} mice.^{5,6} To determine whether microglia are also critical for tumor formation (gliomagenesis), we used *Nfl*^{+/-}-GFAP^{CKO}-TK mice to reduce CD11b⁺ cells (microglia) following ganciclovir (GCV) treatment during glioma formation (6 weeks to 3 months of age; **Supplementary Fig 1**). While we found that long-term GCV treatment reduced optic glioma proliferation, interpretation was confounded due to the anti-proliferative properties of GCV in *Nfl*^{+/-}-GFAP^{CKO} mice lacking the CD11b-TK transgene (data not shown). Thus, we sought an alternative method to target microglia.

Microglia infiltration and function is partly driven by chemokines acting on their cognate receptors, including the CX3CL1 receptor, CX3CR1.⁸ Consistent with previous studies, CX3CL1 is found in abundant levels in the optic nerve and retina, both in wild-type and *Nfl*^{+/-}-GFAP^{CKO} mice (**Supplementary Fig 2A**). Using FACS, CD45^{low} microglia in the mouse optic nerve and brainstem co-express CX3CR1 and CD11b (**Fig 1A and 1B**). Employing a mouse strain in which one copy of the *Cx3cr1* gene is inactivated following germline insertion of a green fluorescent protein (GFP) gene,⁸ double-labeling immunofluorescence reveals that all microglia identified with Iba1 or CX3CR1 antibodies in the optic nerves of *Cx3cr1*^{+gfp} mice also co-express GFP (**Fig 1C-D, Supplementary Fig 2B-D**), and that GFAP⁺ cells (astrocytes) and retinal ganglion cells lack GFP expression (**Fig 1E-1F, Supplementary Fig 2E**). Importantly, no reduction in optic nerve proliferation or microglia (%Iba1⁺ cells) was observed following targeted reduction of CX3CR1 expression in 3-month-old wild-type or *Nfl*^{+/-} mice (**Supplementary Fig 3A and 3B**).

Next, we generated *Nfl*^{+/-}-GFAP^{CKO} mice bearing one inactive *Cx3cr1* allele (*Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice). Using a specific CX3CR1 antibody, *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 optic nerves exhibited a 52% decrease in the percent of CX3CR1⁺ cells relative to *Nfl*^{+/-}-GFAP^{CKO} mice (**Fig 1G**). To determine whether reduced CX3CR1 expression impaired microglia-mediated optic glioma formation, *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice were examined at 6 weeks, 3 months, and 4 months of age. At 6 weeks of age (prior to obvious glioma development), the increased percent of microglia present in the *Nfl*^{+/-}-GFAP^{CKO} mouse optic nerve⁵ was reduced in *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice concomitant with decreased optic nerve proliferation (%Ki67⁺ cells, n=9 mice; **Fig 2A**). Similarly, at 3 months of age, *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice (n=10 mice) also had reduced optic nerve volumes, proliferation, and microglia (**Fig 2B, Supplementary Fig 4**) relative to *Nfl*^{+/-}-GFAP^{CKO} mice. Importantly, unlike *Nfl*^{+/-}-GFAP^{CKO} mice, there was no evidence of optic glioma formation in *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice, demonstrating that CX3CR1⁺ microglia are critical for optic gliomagenesis.

Since optic gliomas are found in the majority of *Nfl*^{+/-}-GFAP^{CKO} mice by 3 months of age,² we sought to determine whether gliomagenesis was delayed, rather than prevented, by reduced CX3CR1 expression. Examination of *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice (n=6 mice) at 4 months of age revealed optic nerve volumes, proliferation and microglia content indistinguishable from *Nfl*^{+/-}-GFAP^{CKO} mice (**Fig 3**). In contrast, retinal ganglion cell death (TUNEL⁺ cells) in *Nfl*^{+/-}-GFAP^{CKO}-CX3CR1 mice was not significantly different from *Nfl*^{+/-}-GFAP^{CKO} mice at any age (**Supplementary Fig 5A-C**). Collectively, these observations establish microglia as essential drivers of gliomagenesis, such that reducing CX3CR1⁺ microglia function delays *Nfl* murine optic glioma formation.

Discussion

Accumulating evidence from other solid tumors has revealed the importance of non-neoplastic cells in the tumor microenvironment for the maintenance of cancer growth.⁹ In brain cancer (glioma), microglia represent one of the main stromal cells.⁴ Consistent with previous immunohistochemistry and FACS studies demonstrating monocyte/microglia

infiltration using Iba1, CD11b, CD163, CX3CR1, CD45, and CD68 antibodies, *Nfl* GEM optic gliomas exhibit increased numbers of Iba1⁺, CD11b⁺ and CX3CR1⁺ cells with low CD45 expression. Based on these findings, we utilized a genetic targeting strategy to impair microglia function (reduced CX3CR1 expression) and delay the formation of *Nfl* optic gliomas. These new observations have broad implications for glioma biology.

First, microglia are emerging as key stromal cells important for dictating both glioma formation and maintenance. We have previously shown that genetic or pharmacologic inhibition of microglia function reduces tumor proliferation, supporting a role for microglia in glioma maintenance. Similarly, others have reported that microglia ablation impairs glioma growth by limiting apoptosis and metalloproteinase-mediated tumor invasion.^{10,11} In addition, our new results using *Nfl*^{+/-}-GFAPCKO-CX3CR1 mice, coupled with earlier reports demonstrating that targeted disruption of IL-6 expression in a GFAP-v-src glioma model prevents glioma formation¹² and monocyte chemoattractant protein-1-induced monocyte infiltration increases transplanted glioma growth,¹³ argue that microglia also serve as essential mediators of glioma formation.

Second, CX3CL1 is one of the most abundantly expressed chemokines in the nervous system and regulates communication between neurons, glia, and microglia through activation of the CX3CR1 receptor. CX3CR1 function mediates pathology in many GEM models of neurological disease. In this regard, CX3CR1 loss in Alzheimer Disease (AD) mouse models attenuates neuronal loss¹⁴ and beta-amyloid deposition through increased microglial phagocytosis.¹⁵ Additionally, reduced CX3CR1 expression results in smaller infarct volumes following brain ischemia.¹⁶ The observation that CX3CR1 reduction alone in *Nfl* GEM delayed optic glioma formation establishes that CX3CL1/CX3CR1 axis function is also critical for microglia function relevant to brain tumor biology. While one report showed that high-grade glioma growth following implantation was not affected by host CX3CR1 loss,¹⁷ single nucleotide polymorphisms within the human *CX3CR1* locus are associated with increased survival of individuals with high-grade glioblastoma and decreased microglial infiltration.¹⁸ CX3CR1 variants have been found to contribute to defective adhesive function, fractalkine binding, signaling¹⁹, and impaired migration.²⁰ Taken together, these findings support a model in which chemokine-regulated microglia function represents a prime determinant of glioma biology.

As we enter an era of rational therapeutics, these data provide a scientific rationale for the design of future treatments that consider stromal cell (microglia) and stroma-derived factor (chemokines) targets. The combination of neoplastic and non-neoplastic cell therapies offers the potential of more durable outcomes with reduced toxicity to the normal cells within the developing brains of children.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

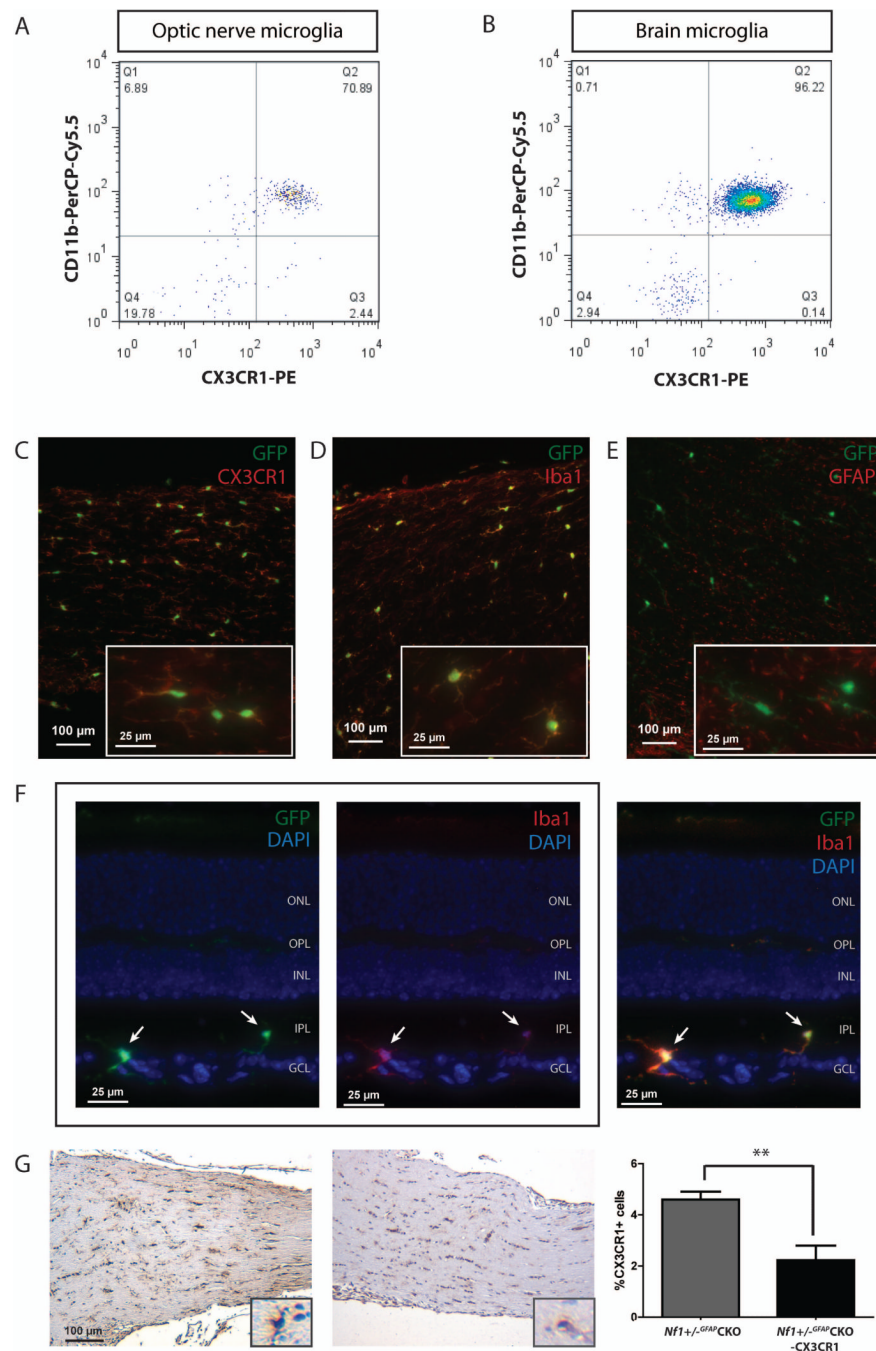
Acknowledgments

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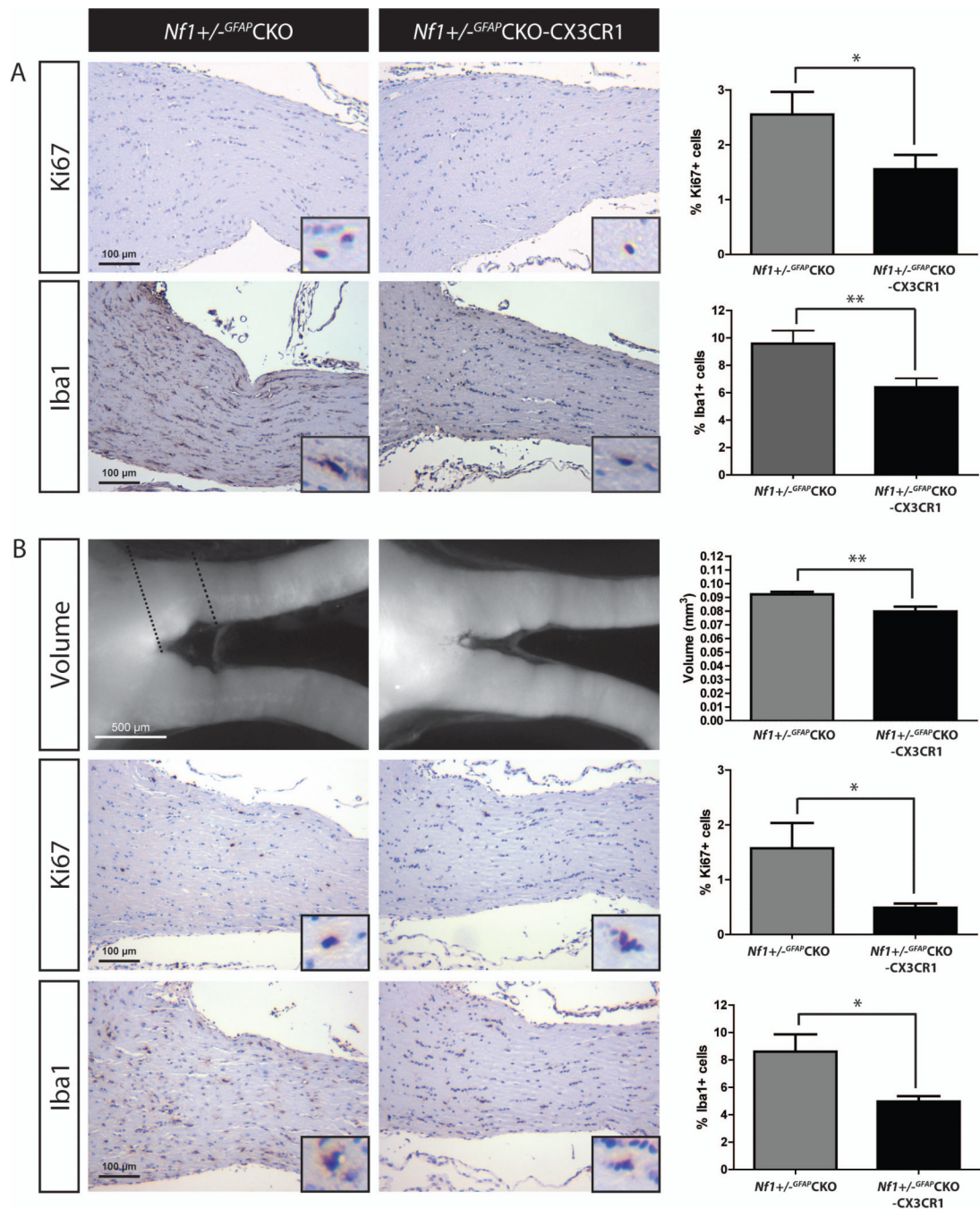
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**FIGURE 1.**

CX3CR1 is expressed by microglia in the optic nerve. Flow cytometry demonstrates that nearly 100% of monocytes in wild-type (A) optic nerves (7-10 pooled optic nerves) or (B) brainstem are double-positive for both CD11b and CX3CR1 expression.

Immunofluorescence analysis of paraformaldehyde-fixed tissue cryosections using CX3CR1 (C, red) or Iba1 antibodies (D, red) and endogenous GFP (green) shows that GFP is expressed by microglia in *Cx3cr1*^{+/-gfp} mouse optic nerves, but not by GFAP-expressing cells (E, red). Insets show representative labeled cells. (F) GFP (green) is also only expressed in microglia (Iba1⁺ cells, red) in the *Nf1*^{+/-GFAP}CKO-CX3CR1 mouse retina (outer nuclear layer, ONL; outer plexiform layer, OPL; inner nuclear layer, INL; inner

plexiform layer, IPL; ganglion cell layer, GCL). The two left panels show GFP or Iba1 immunofluorescence in paraformaldehyde-fixed tissue cryosections with DAPI (blue) nuclear staining, while the right panel shows the merged triple channel fluorescence. The white arrows denote GFP⁺ and Iba1⁺ double-labeled cells. **(G)** CX3CR1 immunohistochemistry reveals a 52% reduction in the percent of CX3CR1⁺ cells in the *Nf1*^{+/-}GFPCKO-CX3CR1 mouse optic nerve at 3 months of age (n=9, p = 0.0012). **, p<0.01.

**FIGURE 2.**

Targeted reduction of CX3CR1 expression reduces microglia content and optic glioma proliferation. **(A)** Optic nerves from *Nf1*^{+/-GFAP}CKO-CX3CR1 mice (n=9) exhibit a 39% reduction in the percent of Ki67⁺ cells relative to *Nf1*^{+/-GFAP}CKO (n=9) at 6 weeks of age (p = 0.0463) as well as a 33% reduction in the percent of Iba1⁺ cells (p = 0.0086). **(B)** 3-month-old *Nf1*^{+/-GFAP}CKO-CX3CR1 mice (n=10) have smaller optic nerve volumes relative to *Nf1*^{+/-GFAP}CKO mice (n=10, 13% reduction, p = 0.0039) and exhibit a 65% reduction in proliferation (%Ki67⁺ cells; p = 0.0358) as well as a 42% reduction in the percentage of Iba1⁺ microglia (p = 0.0114), relative to *Nf1*^{+/-GFAP}CKO mice. Dashed lines

delineate the representative area of the optic nerves used to calculate volume. Insets show representative positively-labeled cells. *, $p < 0.05$; **, $p < 0.01$.

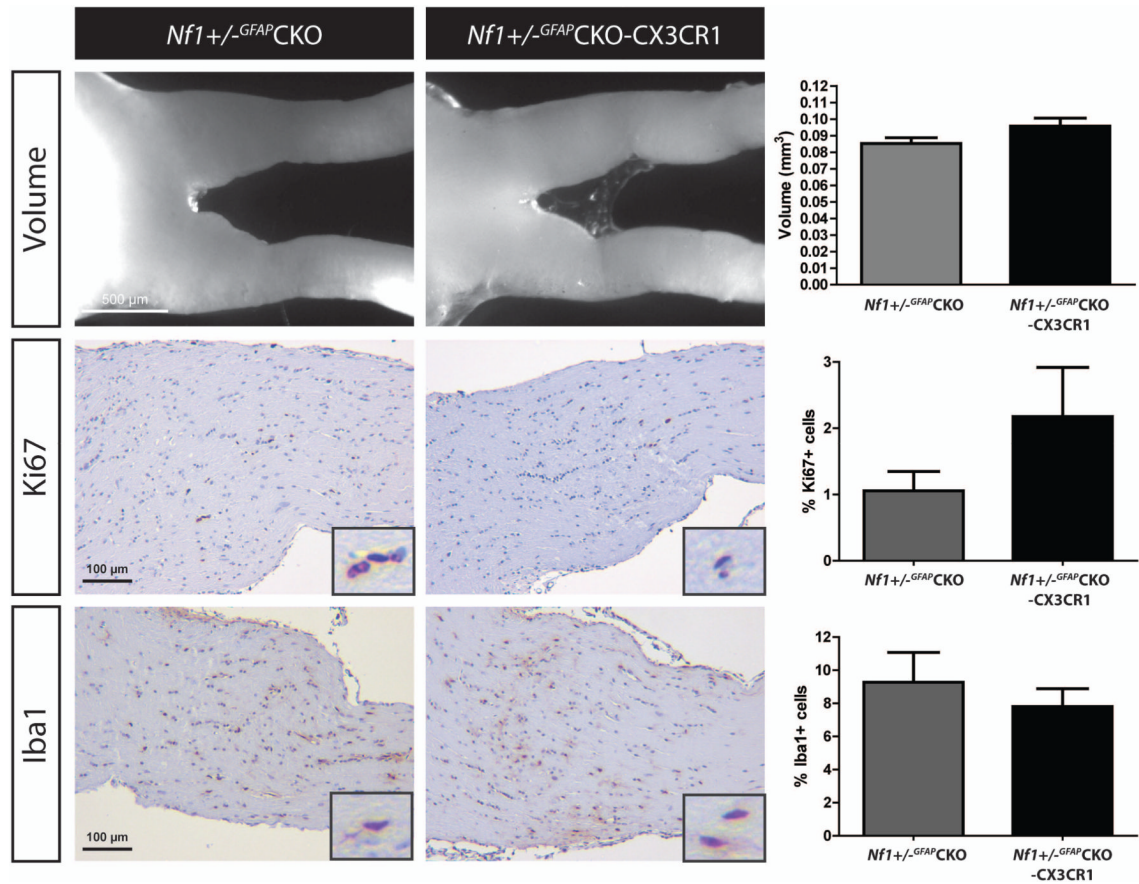


FIGURE 3.

Nf1 optic glioma formation is delayed by targeted reduction of CX3CR1 expression. At 4 months of age, *Nf1*^{+/-GFAP}CKO-CX3CR1 mice (n=6) have optic nerves with similar volumes, proliferation (%Ki67⁺ cells), and microglia (%Iba1⁺ cells) as *Nf1*^{+/-GFAP}CKO mice. Insets depict representative positively-labeled cells.