Neurobiology of Disease

The Soluble Isoform of CX3CL1 Is Necessary for Neuroprotection in a Mouse Model of Parkinson's Disease

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The chemokine CX3CL1/fractalkine is expressed by neurons as a transmembrane-anchored protein that can be cleaved to yield a soluble isoform. However, the roles for these two types of endogenous CX3CL1 in neurodegenerative pathophysiology remain elusive. As such, it has been difficult to delineate the function of the two isoforms of CX3CL1, as both are natively present in the brain. In this study we examined each isoform's ability to regulate neuroinflammation in a mouse model of Parkinson's disease initiated by the neurotoxin 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP). We were able to delineate the function of both CX3CL1 isoforms by using adenoassociated virus-mediated gene therapy to selectively express synthetic variants of CX3CL1 that remain either permanently soluble or membrane bound. In the present study we injected each CX3CL1 variant or a GFP-expressing vector directly into the substantia nigra of CX3CL1 — mice. Our results show that only the soluble isoform of CX3CL1 is sufficient for neuroprotection after exposure to MPTP. Specifically, we show that the soluble CX3CL1 isoform reduces impairment of motor coordination, decreases dopaminergic neuron loss, and ameliorates microglial activation and proinflammatory cytokine release resulting from MPTP exposure. Furthermore, we show that the membrane-bound isoform provides no neuroprotective capability to MPTP-induced pathologies, exhibiting similar motor coordination impairment, dopaminergic neuron loss, and inflammatory phenotypes as MPTP-treated CX3CL1 — mice, which received the GFP-expressing control vector. Our results reveal that the neuroprotective capacity of CX3CL1 resides solely upon the soluble isoform in an MPTP-induced model of Parkinson's disease.

Introduction

A key hallmark of Parkinson's disease (PD) is the dramatic depletion of dopaminergic (DA) neurons in the substantia nigra pars compacta (SNpc). While the exact etiology of the disease remains elusive, recent evidence suggests that microglial activation is a key component contributing to DA degeneration in the SNpc (Wu et al., 2002; Tieu et al., 2003; Cardona et al., 2006; Kim and Joh, 2006; Yasuda et al., 2008). Recent literature has identified the chemokine fractalkine (CX3CL1, FKN) as a neuroimmune regulatory protein signaling through its sole receptor (CX3CR1) exclusively on microglia with anti-inflammatory and thus neuroprotective effects (Harrison et al., 1998; Ransohoff et al.,

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2007). Accordingly, multiple reports have begun to characterize CX3CL1-CX3CR1 interactions as important mechanisms of microglial regulation (Harrison et al., 1998; Zujovic et al., 2000, 2001; Mizutani et al., 2007; Wynne et al., 2010; Bachstetter et al., 2011; Pabon et al., 2011). Functionally, exogenous CX3CL1 peptide has been shown to inhibit microglial activation by downregulating the production of cytotoxic mediators (TNF α , IL-1 β , iNOS, and IL-6) through multiple transduction pathways (Maciejewski-Lenoir et al., 1999; Garcia et al., 2000; Zujovic et al., 2000, 2001; Ré and Przedborski, 2006; Ransohoff et al., 2007). Endogenous expression of CX3CL1 produces a membraneanchored isoform that is proteolytically cleaved by ADAM10/17, yielding a soluble isoform. Both isoforms of CX3CL1 have been shown to ligate CX3CR1 (Chapman et al., 2000; Garton et al., 2001; Ré and Przedborski, 2006; Hundhausen et al., 2007; Ludwig and Weber, 2007). However, the individual signaling functions of each CX3CL1 isoform in a neurodegenerative condition are not fully understood.

In the present study, we investigated the neuroprotective capacity of each CX3CL1 isoform in a 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP) mouse model of PD. MPTP has been reliably used to induce DA cell loss in the SNpc of mice and non-human primates akin to what is observed in the human condition (Przedborski et al., 2001). To delineate the functional significance of each CX3CL1 isoform independently (soluble versus membrane bound), we constructed two recombinant adenoassociated viruses (rAAVs) that selectively expressed either

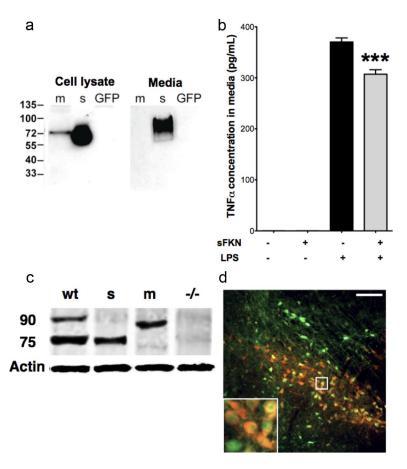


Figure 1. Generation and validation of rAAV-FKN variants. a, Western blot analysis of sFKN (s), mFKN (m), and GFP expressed in HEK293 cells using an anti-hemagglutinin antibody. Only sFKN was detected in the medium, while both FKN variants were found in the crude cell lysate. b, ELISA quantification of BV2 secreted TNF α . Pretreatment with sFKN-conditioned medium attenuates (****p < 0.05) the secretion of TNF α from BV2 cells when exposed to 100 ng of LPS for 4 h. Importantly, sFKN-conditioned media alone does not induce the secretion of TNF α . c, Western blot analysis of in vivo expression of wild-type (wt), sFKN, mFKN, and $CX3CL1^{-/-}$ (-/-) tissue lysates using an anti-CX3CL1 antibody. Both sFKN and mFKN produce a single band consistent in molecular weight with previous literature and relative to WT. Importantly, mFKN yields a single band, indicating that constitutive proteolytic cleavage by ADAM10/17 is abated. d, ln vivo expression pattern of rAAV-GFP, which uses the same promoter as the rAAV-FKN variants and CED vector delivery to the SNpc, resulted in transduction of numerous neurons of the SN. Green, GFP; red, TH $^+$ neurons; yellow-orange, colocalized expression. Scale bar, 200 μ m. Data are presented as mean \pm SEM.

soluble CX3CL1 or a mutant isoform, which is proteolytically resistant to cleavage and remains membrane bound, as well as a control vector expressing green fluorescent protein (GFP). Each of the three vectors was injected into the SNpc of *CX3CL1*^{-/-} mice such that the only CX3CL1 present was either solely soluble or membrane bound.

We therefore examined each isoform's ability to dampen the neurotoxic effects of MPTP, which were assessed through common behavioral, immunological, and biochemical assays. Our results show a significant reduction in the classic neurotoxic pathology associated with MPTP exposure in CX3CL1-null mice that received the vector expressing only the soluble isoform. Furthermore, CX3CL1-null mice, which expressed only the membrane-bound isoform of CX3CL1 in the SNpc, had pathologies akin to CX3CL1-null mice expressing the GFP control vector. These data suggest that the neuroprotective capacity of CX3CL1 is exclusively reliant on the soluble isoform in an MPTP model of PD.

Materials and Methods

AAV production. Recombinant AAV serotype 9 (rAAV9) vectors expressing the two different forms of CX3CL1 (GI:114431260) were cloned using PCR

from mouse brain cDNA. The soluble form of fractalkine (sFKN; amino acids 1-336) was generated with primers Frac5' (GAGACCGGTCCA CCATGGCTCCCTCGCCGCTCGCG) and Frac3' (CTCGCTAGCTCACATGGCATAGTC AGGCACGTCATAAGGATAGCTAGAAGCC ATTGTGGCTGCCTGGGTGTCGGGGAC). The mutant membrane-bound form of fractalkine (mFKN; mutations to stop ADAM10/17 cleavage into the soluble form, R337A+R338A) was generated by cloning two PCR fragments with primers Frac5' plus Fracma (GTAGCCCCACTGCCTGGGCAGCTGTGGC TGCCTGGGTG) and Frac3' (CTCGCTAGCT CACATGGCATAGTCAGGCACGTCATAAG GATAGCTAGAAGCCATCACTGGCACCAG GACGTATGAGTTAC) and Fracms (CACCCA GGCAGCCACAGCTGCCCAGGCAGTGGGG CTAC). sFKN and mFKN were cloned into the pTR2-MCS at the AgeI and NheI cloning sites. This vector contains the AAV2 terminal repeats and the hybrid cytomegalovirus—chicken β -actin promoter for CX3CL1 mRNA transcription. A hemagglutinin (HA) tag was added to the C terminus of both sFKN and mFKN with the PCR primers. rAAV9 vectors were generated and purified as described previously (Carty et al., 2010). rAAV particles are expressed as vector genomes (vg)/ml. Vector genomes were quantitated using a modified version of the dot plot protocol described by Zolotukhin et al. (2002). The protocol was modified to use a nonradioactive biotinylated probe for fractalkine generated by PCR. Bound biotinylated probe was detected with IRDye 800CW (Li-Cor Biosciences) and quantitated on the Li-Cor Odyssey. UF11 plasmid was used to generate rAAV9-expressing GFP as described previously (Carty et al., 2008).

In vitro cell culture experiments. sFKN conditioned medium was generated by transfection of HEK293 cells with Lipofectamine 2000 (Invitrogen) and the pTR2-sFKN plasmid. Cells were incubated in Opti-Mem (Invitrogen; containing 100U/ml penicillin and 100 μ g/ml streptomycin) for 48 h. Following incubation, medium was harvested and cleared by

centrifugation at 2000 \times g. The cleared sFKN conditioned medium was used for the proceeding cell culture experiment. BV2 cells (Blasi et al., 1990) were seeded at 5 \times 10 5 cells per well in Opti-Mem (Invitrogen; supplemented containing 100U/ml penicillin and 100 μ g/ml streptomycin) of a 24 well plate and allowed to adhere for 16 h. Immediately before treatment, cells were washed with fresh medium to remove any dead or nonadhered cells. Cells were treated with conditioned media, conditioned media plus LPS, LPS, or medium alone for 4 h. Following incubation, medium was removed from each of the four groups and analyzed for TNF α using a commercial ELISA kit (R & D Systems) following the manufacturer's protocol.

Animals. All experiments were conducted in accordance with the National Institutes of Health Guide for the Care and Use of Laboratory Animals and were approved by the Institutional Animal Care and Use Committee of the University of South Florida College of Medicine (Tampa, FL). CX3CL1^{-/-} mice (Merck Sharp & Dohme) were backcrossed to the C57BL/6J (WT) background (The Jackson Laboratory) for >10 generations. Colonies of wild-type (WT) and CX3CL1^{-/-} littermates were maintained at the University of South Florida, and genotyping was outsourced using a commercially available service (Transnetyx). Twelve- to sixteen-week-old WT and CX3CL1^{-/-} littermates were used in the proceeding experiments. Mice were pair housed in environmen-

tally controlled conditions (12:12 h light:dark cycle at 21 \pm 1°C) and provided food and water *ad libitum*.

Surgical procedure. Mice for the treatment groups were randomly selected and age stratified to ensure equal representation. Three groups of 12- to 16-week-old male $CX3CL1^{-/-}$ mice (n=30 per group) were injected with mFKN, sFKN, or a vector expressing GFP for viral and protein control, while WT mice were injected with sterile saline (n=60). Briefly, mice from each rAAV group received 1.5 μ l at 1×10^{12} vg/ml (2.5 μ l/min) bilateral direct injections into the SNpc (anteroposterior, -2.8; mediolateral, ± 1.4 ; dorsoventral, -4.6) using a convectionenhanced delivery (CED) 26 gauge needle (Carty et al., 2010). WT mice and an additional group of $CX3CL1^{-/-}$ mice (n=15) received sterile saline via the same injection parameters.

Acute MPTP administration. Six weeks following SNpc injections, sFKN-, mFKN-, and GFP-treated CX3CL1 -/- mice were weighed and serially injected with MPTP. Additionally, the group of WT mice were weighed and randomly split such that half received either serial injections of MPTP (WT-MPTP) or sterile saline to serve as a double sham for both surgery and injection paradigms (WT-Sham). Lastly, the group of CX3CL1 ^{-/-} mice that received intracranial saline also received i.p. saline as well (FKN $^{-/-}$ Sham). This group served to examine whether there was a genotype effect in the baseline of all endpoints. MPTP injections were administered as previously described (Cardona et al., 2006). Briefly, MPTP (Sigma Aldrich; 10 mg per kg body weight) was diluted in sterile saline and injected i.p. four times with a 1 h injection interval. Following injections, animal cages were placed on flexible, water-heating pads (37°C) for \sim 24 h postinjection in an effort to maintain body temperature and combat excessive mortality. After 5 days the mice were weighed, and mice that lost >5% body mass were excluded from the study. Mortality ranged from 25-28% and was uniform across all groups of mice exposed

Behavioral analysis. Five days after MPTP injections, mice were tested for overall balance and motor coordination. This test was performed on an accelerating rotarod apparatus (Ugo Basile) with a 3 cm diameter rod starting at an initial rotation of 4 rpm accelerating to 40 rpm over 5 min. Mice were tested for the time spent on the rod during each of four trials with a 30 min intertrial interval. Each trial was completed when the mouse fell off the rod (distance of 12 cm) onto a spring-cushioned lever (Rogers et al., 2011). To assess motor coordination deficits and not motor learning, only the first three trials (day 1) were analyzed. Following behavioral analysis, animals were randomly divided for either perfusions or biochemical analyses.

Immunohistochemistry. Mice were euthanized with pentobarbital (50 mg/ kg, i.p.) and transcardially perfused with PBS, followed by 4% paraformaldehyde in PBS. The brains were postfixed in 4% paraformaldehyde for 12 h, after which they were transferred into 30% sucrose in PBS for at least 16 h at 4°C. Exhaustive coronal sections were made at 40 μm using a Microm cryostat (Richard-Allan Scientific) and stored in cryoprotectant at 4°C. Labeling of dopaminergic neurons was performed using an antibody against tyrosine hydroxylase (TH; 1:4000, Immunostar). Neurons were labeled using NeuN (Millipore, 1:500), which is a pan marker for neuronal nuclei. Activated microglia as well as T cells, were visualized with antibodies against CD68, CD11b, and CD3 (Serotec 1:200). Standard staining procedures were conducted on free-floating sections using every sixth section for the entire SNpc and included sections before and after to ensure that the entire structure was sampled. Following incubations (24-48 h) with appropriate primary and biotinylated secondary antibodies, sections were treated with Vectastain ABC reagent (Vector Labs) and visualized with DAB reaction (Sigma). As a control measure, the specificity of all antibodies was confirmed by omitting primary antibodies.

Stereological quantification and imaging. To determine numbers for TH $^+$ and CD3 $^+$ cells, the optical fractionator method of unbiased stereological cell counting techniques (West et al., 1991) was used with a Nikon Eclipse 600 microscope and quantified using Stereo Investigator software (MicroBrightField). The virtual grid (150 \times 150 μ M) and counting frame (75 \times 75 μ M) were optimized to count at least 200 cells per animal with error coefficients <0.07. Outlines of the anatomical structures were done using $10\times/0.45$ objective, and cell quantification was conducted using $60\times/1.40$ objective.

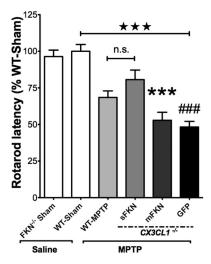


Figure 2. MPTP-induced impairment of motor coordination is rescued by sFKN. Five days after the last injections of MPTP or saline, mice were subjected to behavioral analysis to analyze balance and coordination on an accelerating rotarod. Motor coordination was assessed by the average of three trials during training. FKN $^{-/-}$ sham group performed similarly to WT-Sham animals. As a whole, exposure to MPTP induced significant deficits in motor coordination compared to WT-Sham mice, which received sterile saline (three-way ANOVA; $F_{(1,145)}=42.11$, ***p< 0.001). $CX3CL1^{-/-}$ mice that received sFKN performed significantly better than those which received mFKN (Tukey's HSD; ***p= 0.002) or GFP (Tukey's HSD; *##p< 0.001). Additionally, treatment with sFKN recovered the MPTP-induced motor coordination deficit akin to WT-Sham mice (Tukey's HSD; p= 0.380). Data are presented as mean time in seconds as a percentage of WT-Sham \pm SEM. n.s. = not significant.

Cell density quantification and imaging. Cell density was measured using a Zeiss Mirax slide scanner, as described previously (Rogers et al., 2011). Briefly, slides with NeuN-, CD68-, and CD11b-immunolabeled sections were scanned and converted into digital image files. Following scanning, the digital images were analyzed for positive area using Zeiss NeuroQuant IAE analysis software. An investigator unaware of sample treatment conditions defined the anatomical regions comprising the SNpc for each tissue section. Positive cells were defined using the analysis software based upon their threshold to include both the cell body and processes, while eliminating background values. Data are represented as the average area ratio of positive cells relative to the total measurement area.

Biochemical analysis tissue preparation. Mice used for biochemical studies were euthanatized by rapid decapitation. Brain tissues encompassing the ventral mesencephalon (VM) and striatum were separately dissected and rapidly frozen in liquid nitrogen before storage at -80° C. Tissues were homogenized using an electric tissue homogenizer in 1:10 weight-to-volume ratio of ice-cold RIPA buffer (Millipore) containing protease inhibitors and EDTA (Pierce). Following homogenization, sample lysates were centrifuged at $10,000 \times g$ at 4°C for 15 min, and supernatant was collected. Protein concentration was determined using BCA assay (Pierce). Tissue lysates were used for subsequent Western blot and ELISA analyses.

Western blot analysis. Fifty micrograms of total protein per lane was loaded onto a 4–15% SDS-polyacrylamide gel (Bio-Rad) for electrophoresis. Proteins were transferred onto a nitrocellulose membrane for immunodetection. Membranes were subsequently blocked for 1 h in 5% nonfat dry milk (NFDM; Bio-Rad) in PBS-Tween (PBS-T; 0.1% Tween 20). Anti-HA (1:500; Roche), anti-CX3CL1 (1:500, directed against N terminus; R & D Systems), anti-CX3CR1 (1:800; Abcam), anti-TH (1: 500; Millipore), and mouse β-actin (1:3000; Sigma) were incubated overnight at 4°C in 1% NFDM in PBS-T. Following washes, appropriate secondary antibodies (Li-Cor) were incubated for 1 h at room temperature in 1% NFDM in PBS-T. Membranes were scanned using Li-Cor Odyssey near-infrared imager and raw intensity for each band was measured using Li-Cor Odyssey image analysis software. Scanned pseudocolor images were converted to black and white.

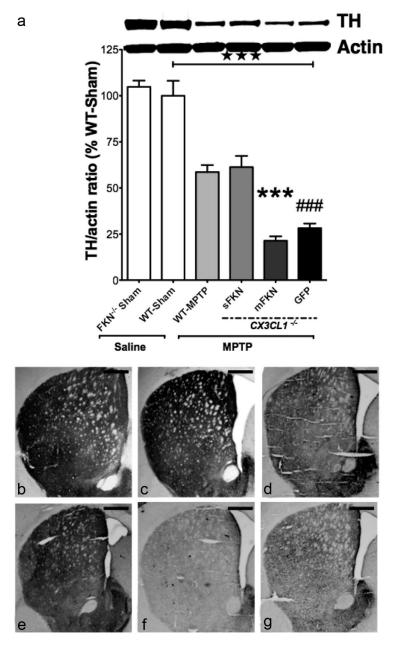


Figure 3. MPTP-depleted TH in the striatum is recovered by sFKN. Both hemispheres were rapidly dissected 7 days following the last MPTP or saline injections for biochemical analysis. \boldsymbol{a} , Western blot analysis revealed that MPTP exposure significantly reduced the availability of TH in the striatum (three-way ANOVA; $F_{(1,23)}=34.42$, ***p<0.001). Additionally, FKN $^{-/-}$ sham animals had no distinguishable difference in TH production compared to WT-Sham mice. $CX3CL1^{-/-}$ mice expressing sFKN in the SNpc had an attenuated loss of TH compared to mFKN (Tukey's HSD; ***p<0.001) and GFP (Tukey's HSD; ***p<0.001) expressing mice. There were no significant differences between $CX3CL1^{-/-}$ mice expressing either mFKN or GFP (Tukey's HSD; p=0.627). $\boldsymbol{b}-\boldsymbol{g}$, Representative immunostaining in the striatum shows progressive loss of TH for FKN $^{-/-}$ sham (\boldsymbol{b}) , WT-Sham (\boldsymbol{c}) , WT-MPTP (\boldsymbol{d}) , sFKN (\boldsymbol{e}) , mFKN (\boldsymbol{f}) , and GFP (\boldsymbol{g}) groups. Data are presented as the mean band density ratio (n=5 per group) of TH to actin as a percentage WT-Sham \pm SEM. Scale bars, 200 μ m.

ELISA quantification. CX3CL1, TNFα, and IL-1 β concentrations were quantified using standard ELISA technique. VM lysates were run in triplicate at a concentration of 100 μ g per well and incubated overnight at 4°C. Following incubation, the manufacturer's (RayBiotech) suggested protocol was followed. Optical density values for each ELISA plate were measured on a plate reader (BioTek), and sample concentrations were calculated based upon the supplied standard curve. Values were converted from pg/ml to pg/ μ g of protein loaded. Antibodies for CX3CL1 were directed against the N terminus.

Data analysis. All data were analyzed using SPSS software (IBM, version 20) and are presented as the mean \pm SEM. Statistical analyses were

performed using three-way ANOVA accounting for genotype, MPTP, and rAAV treatments. Group differences were assessed by Tukey's honestly significant difference (HSD) *post hoc* multiple comparisons test, with p values of <0.05 considered significant.

Results

Generation and validation of rAAV9-FKN variants

Both rAAV9-FKN variants produced a murine version of the CX3CL1 gene driven by the hybrid chicken β -actin cytomegalovirus promoter and were tagged with a HA peptide sequence for ease of detection. Before virus production, the rAAV vector was tested in HEK293 cells to evaluate the expression cassette. Cell lysate and conditioned media from transfected cells were examined using Western blot analysis to determine sFKN and mFKN expression. An anti-HA tag Western blot of transfected HEK 293 cells revealed that sFKN is readily secreted into the medium, while mFKN remains only in the cell lysate fraction (Fig. 1a). We next wanted to confirm that the synthetic FKN variants were biologically active. Briefly, conditioned medium from HEK 293 cells transfected with the sFKN plasmid was used to confer bioactivity by measuring the amount of secreted TNF α from BV2 microglia-like cells. It has been well documented that the induction of cytokine production from microglia as a result of LPS exposure can be blunted if the cells are pretreated with exogenous soluble CX3CL1 peptide (Zujovic et al., 2000; Mizuno et al., 2003; Lyons et al., 2009; Wynne et al., 2010). In accordance with previous literature, we show that BV2 cells preconditioned with sFKN-conditioned medium had an attenuated response to LPS (100 ng/ml) as measured by TNF α secretion (Fig. 1b). Furthermore, addition of sFKN-conditioned medium alone did not induce TNF α production. This attenuated response is comparable to previous literature (Mizuno et al., 2003; Lyons et al., 2009) given the dilute concentration of CX3CL1 found in sFKN-conditioned medium. As we were able to confer the functionality and bioactivity in vitro, we next examined the expression of the rAAV9-

FKN vectors *in vivo*. Using CX3CL1-null mice, we injected each vector bilaterally into the SNpc. Six weeks postinjection the harvested brain tissues were analyzed for sFKN and mFKN expression by Western blot. Using an anti-mouse CX3CL1 antibody, *in vivo* expressions of each of the rAAV-FKN constructs created their specific variants of CX3CL1 consistent with predicted molecular weights as we had previously documented (Bachstetter et al., 2011). Importantly, mFKN produces only a single band, indicating that the dibasic arginine mutation effectively inhibits

cleavage by ADAM10/17 (Fig. 1c). Furthermore, direct injection using the CED method (Carty et al., 2010) with rAAV variants transduced numerous neurons throughout the SNpc (Fig. 1d). Viral transduction was not limited to DA neurons of the SNpc, as other nondopaminergic neurons were transduced. Of these populations, we found >90% transduction of neurons (data not shown). Additionally, as we have documented previously (Carty et al., 2010), parenchymal injections of rAAV9 vectors via CED resulted in neuronal transduction with minimal astrocytic expression (data not shown).

MPTP-induced impairment of motor coordination is ameliorated by sFKN but not mFKN

We next assessed the effects of each rAAV-FKN treatment on MPTP-induced motor deficits using an accelerating rotarod. FKN^{-/-} Sham mice displayed no motor coordination impairment relative to WT-Sham (Fig. 2). Five days after the last MPTP injection, all MPTP-treated groups displayed a significant performance impairment $(F_{(1,145)} = 42.11, p < 0.001)$ as measured by average latency to fall across three trials compared to WT-Sham mice (Fig. 2). Treatment with sFKN significantly improved MPTP-induced behavioral deficits compared to mFKN (p = 0.002) and GFP (p < 0.001) viral control. Following rotarod performance assessment, mice were killed and striatal tissues were dissected for biochemical analysis. In agreement with each group's rotarod performance, Western blot analysis revealed significant loss of striatal TH as a result of MPTP administration (p < 0.001), ranging from 40-75% (Fig. 3a). Furthermore, sFKN treatment was able to attenuate exhaustive striatal TH loss by 35 and 30% compared to mFKN (p < 0.001) or GFP groups (p < 0.001), respectively.

MPTP-induced neurodegeneration of SNpc is rescued by sFKN

It has been previously documented that disruption of CX3CL1 signaling via genetic ablation of CX3CR1 in mice increases MPTP neurotoxicity in SNpc (Cardona et al., 2006). Seven days after the last MPTP injection, mice were transcardially perfused with a buffered paraformaldehyde solution and brain tissues were removed. We evaluated the level of MPTP-induced neuro-degeneration in the SNpc through stereological quantification of TH (Fig. 4a) and by measuring the density of NeuN (Fig. 4b) staining. Additionally, FKN -/- Sham mice showed no significant reduction in the number of TH-positive neurons or the density of NeuN-stained neurons relative to WT-Sham, indicating that genetic disruption of CX3CL1 signaling alone does not produce a PD-like phenotype (Fig. 4a,b). Compared to WT-

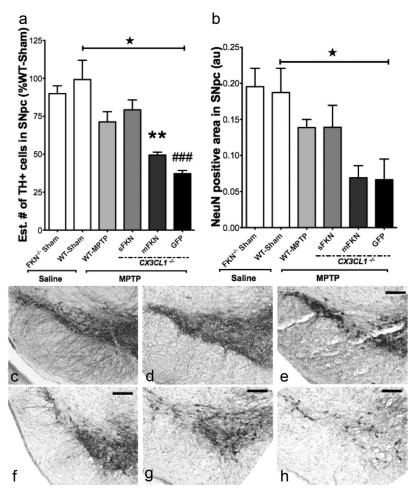


Figure 4. Stereological analysis of TH-positive neurons in the SNpc. TH immunoreactivity was analyzed in the SNpc of coronal sections made from mice 7 days after the final MPTP or saline injections. a, Exposure to MPTP resulted in a marked reduction in the estimated number (Est. #) of TH-positive cells in the SNpc (three-way ANOVA; $F_{(1,12)} = 4.733$, *p = 0.050) compared to WT-Sham. Again, FKN $^{-/-}$ sham animals exhibited no phenotypic differences in TH $^+$ cell counts relative to WT-Sham mice. Furthermore, sFKN provided the most significant TH neuroprotection relative to mFKN (Tukey's HSD; **p = 0.003) and GFP (Tukey's HSD; **p < 0.001). Paralleling our data in the striatum, there was no significant difference in CX3CL1 $^{-/-}$ mice expressing either mFKN or GFP (p = 0.324). b, Graph of NeuN $^+$ cell density in the SNpc. Although injections with MPTP did reduce the immunoreactivity of NeuN $^+$ cells compared to WT-Sham (three-way ANOVA; $F_{(1,18)} = 6.81$, *p = 0.018), there were no significant differences revealed among group interactions (two-way ANOVA; $F_{(3,12)} = 3.14$, p = 0.065). Akin to TH $^+$ cell counts, there were no differences between FKN $^{-/-}$ sham and WT-Sham animals for NeuN density. c-h, Representative immunostaining for TH in the SNpc for FKN $^{-/-}$ sham (c), WT-Sham-(d), WT-MPTP-(e), sFKN-(f), mFKN-(g), and GFP-treated (f) groups. Data are presented as mean estimate of TH $^+$ cells or mean density ratio as a percentage of WT-Sham \pm SEM. Scale bars, 200 μ m.

Sham, MPTP exposure produced a marked reduction in the estimated number of TH-positive cells across all groups ($F_{(1, 12)} = 4.733$, p = 0.050). As shown previously by Cardona et al. (2006), MPTP produced a larger lesion in the $CX3CL1^{-/-}$ mice. sFKN treatment attenuated MPTP-induced dopaminergic cell loss compared to mFKN- (p = 0.003) and GFP-treated (p < 0.001) animals. Furthermore, exposure to MPTP reduced the NeuN-positive cell density in the SNpc ($F_{(1, 12)} = 4.733$, p = 0.050), although there were no significant differences when groups were compared among each other ($F_{(1,12)} = 3.14$, p = 0.065).

Administration of MPTP does not alter CX3CL1–CX3CR1 signaling

Recent literature has suggested that acute inflammatory insults can downregulate both transcript and protein levels of CX3CR1 on microglia (Kremlev and Palmer, 2005; Wynne et al., 2010).

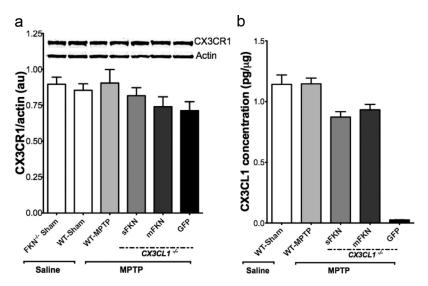


Figure 5. MPTP does not alter CX3CL1—CX3CR1 signaling in the VM. Tissue lysates from the VM of each group were analyzed for protein levels of either CX3CR1 by Western blot and CX3CL1 by ELISA. \boldsymbol{a} , Representative bands for CX3CR1 and actin for each group from Western blot analysis of VM tissue lysate using an anti-mouse CX3CR1 antibody. There were no significant alterations of CX3CR1 protein levels as a result of either MPTP or saline injections. Furthermore, neither the presence nor absence of CX3CL1 (regardless of isoform) affected protein levels of CX3CR1 (three-way ANOVA; $F_{(1,19)} = 0.66$, p = 0.428). \boldsymbol{b} , VM lysates were analyzed for CX3CL1 concentrations by standard ELISA. Exposure to MPTP did not alter the production of CX3CL1 compared to WT-Sham (three-way ANOVA; $F_{(1,23)} = 3.74$, p = 0.066). Furthermore, each rAAV-FKN variant sufficiently restored the levels of CX3CL1 in the VM to that resembling WT-Sham. Importantly, each variant expressed approximately the same amount of CX3CL1 (sFKN vs mFKN; Tukey's HSD; p = 0.703). Data are presented as the mean band density ratio of CX3CR1 to actin and the mean concentration of CX3CL1 in pq/ μ q \pm SEM. au, Arbitrary units.

We analyzed VM tissue lysate using Western blot analysis for CX3CR1 (Fig. 5a). Acute exposure to MPTP failed to affect ($F_{(1,19)} = 0.66$, p = 0.428) the protein levels of CX3CR1 *in vivo*. Additionally, neither the presence of the individual rAAV-FKN variants nor the complete absence of CX3CL1 in the GFP group altered CX3CR1 protein levels in the VM ($F_{(3,13)} = 1.47$, p = 0.268). These data would suggest that expression of CX3CR1 is not only constitutive (Harrison et al., 1998) but also independent of the CX3CL1 isoform. Additionally, we quantified the production of CX3CL1 from each rAAV-FKN variant in the VM. Each rAAV-FKN variant was able to restore CX3CL1 levels resembling WT-Sham, and importantly sFKN and mFKN produced similar concentrations of CX3CL1 (Fig. 5b). Furthermore, here we show that MPTP does not affect the production of CX3CL1 (WT-Sham vs WT-MPTP; p > 0.05).

Induction of CD68- and CD11b-reactive microglia is attenuated by sFKN

Microglial activation has been implicated in the propagation of SNpc neurotoxicity in multiple animal models of PD. Recent studies further report that an intact CX3CL1 to CX3CR1 signaling axis dampens the neurotoxic effects of both 6-OHDA and MPTP (Cardona et al., 2006; Pabon et al., 2011). Postmortem analysis of idiopathic PD patients has revealed strong immunoreactivity for CD68, a marker of phagocytic microglia (Croisier et al., 2005; Vroon et al., 2007). Furthermore, administration of MPTP has been reliably shown to induce this phagocytic microglial phenotype in the SNpc of mice (Vroon et al., 2007; Chung et al., 2010, 2011). Quantification of CD68-positive cell density revealed a significant induction of microglial reactivity (Fig. 6a) as a result of MPTP injections compared to WT-Sham ($F_{(1, 26)} = 4.25$, p = 0.046). Furthermore, loss of CX3CL1 (FKN $^{-/-}$ Sham) is not Sham) is not sufficient to induce significant upregulation of CD68 immunoreactivity. However, CD68 immunoreactivity was most profound

in both the mFKN-treated (p = 0.001) and GFP-treated (p < 0.001) groups compared to sFKN. Furthermore, sFKNtreated mice resembled those of WT-MPTP (p = 0.971) in that there were very few CD68-positive cells. Here our data also show that mice treated with mFKN have a CD68 phenotype similar to that of mice with control AAV-GFP vector (Fig. 6k,l; p = 0.913). We next analyzed a broader marker for microglial activation by using an antibody for CD11b. Immunolabeled CD11b microglia have previously been shown to increase reactivity following administration of the dopaminergic neurotoxin MPTP in the SNpc (Wu et al., 2003; Chung et al., 2011). Mice exposed to MPTP had a significant increase in CD11b-reactive microglia (Fig. 6*b*) compared to WT-Sham ($F_{(1, 27)} = 7.35$, p = 0.021). Again, FKN^{-/-} Sham mice displayed no induction phenotype for this marker relative to WT-Sham mice. Moreover, mice that received sFKN injections again presented a similar CD11b expression phenotype as that of WT-MPTP (p = 0.792). Furthermore, a strong induction of CD11b density by MPTP exposure was observed in the SNpc

of both mFKN-treated (Fig. 6m, p = 0.007) and GFP-treated (Fig. 6n, p < 0.001) mice compared to sFKN-treated (Fig. 6j) CX3CL1-null mice. Additionally, CD11b immunoreactivity was similar following MPTP exposure between mFKN and GFP-treated CX3CL1-null mice (p = 0.499). Recent literature has identified a subset of cytotoxic (CD4 $^+$ CD28 $^-$) T cells that express CX3CR1, which is thought to mediate T cell entry to brain parenchyma in an animal model of multiple sclerosis (Broux et al., 2012). As others have shown (Brochard et al., 2009), we did observe an increase in CD3 $^+$ cells in the SNpc following MPTP exposure; however we did not find any differences between sFKN- and mFKN-treated mice (data not shown).

Expression of MPTP induced proinflammatory cytokines is diminished by sFKN

Postmortem analysis of human PD tissue has shown that microglia are immunoreactive for multiple proinflammatory cytokines, including TNF α and IL-1 β (McGeer and McGeer, 2004). Furthermore, mice that are genetically altered to inhibit cytokine production or are deficient in receptors for these cytokines provide neuroprotection in the SNpc following MPTP exposure (Klevenyi et al., 1999; Sriram et al., 2002). Accordingly, as CX3CL1 signaling has been shown to attenuate the production of these cytokines in vitro and in vivo following various inflammatory stimuli, we examined the anti-inflammatory properties of each rAAV-FKN variant in response to MPTP exposure in the CX3CL1-null mice. Tissue lysates of the VM were analyzed using standard ELISA techniques for the proinflammatory cytokines TNF α and IL-1 β . There was not a significant change in these cytokines for FKN^{-/-} Sham mice relative to WT-Sham mice. MPTP treatment significantly upregulated the production of TNF α (Fig. 7*a*; $F_{(1, 23)} = 18.36$, p < 0.001) and IL-1 β (Fig. 7*b*; $F_{(1, 23)} = 11.97, p = 0.002$) compared to all treatment groups. These findings are consistent with our data showing an in-

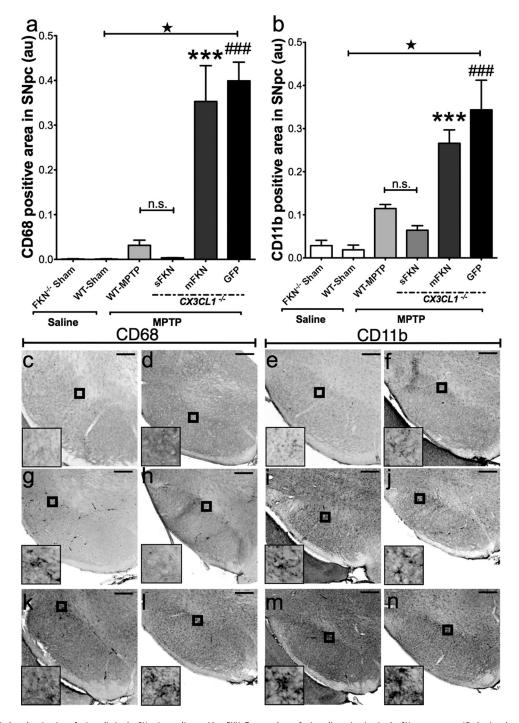


Figure 6. MPTP-induced activation of microglia in the SNpc is ameliorated by sFKN. Two markers of microglia activation in the SNpc were quantified using densitometric analysis. a, MPTP exposure as a whole induced a significant immunoreactive profile for CD68 compared to saline injections (three-way ANOVA; (F(1, 26) = 4.25, $\star p$ = 0.046). No difference was observed in CD68 staining for FKN $^{-/-}$ sham mice compared to WT-Sham mice. Interestingly, CD68 immunoreactivity was noticeably absent in sFKN-treated mice, akin to levels found in WT-Sham. Furthermore, sFKN-expressing mice had a dramatic reduction in CD68 immunoreactivity relative to $CX3CL1^{-/-}$ mice expressing either mFKN (Tukey's HSD; ****p < 0.001) or GFP (Tukey's HSD; ****p < 0.001). Although there was a visual induction of the CD68 phenotype in WT-MPTP mice, this was not significant compared to sFKN mice (Tukey's HSD; p = 0.971). b, Densitometric analysis of CD11b immunoreactivity in the SNpc. As a whole, MPTP exposure induced significant upregulation of CD11b immunoreactivity compared to saline injections (three-way ANOVA; $F_{(1,27)} = 7.35$, *p = 0.021). There was a slight increase in CD11b reactivity for FKN $^{-/-}$ sham mice, but this was not significant compared to the WT-Sham group. $CX3CL1^{-/-}$ mice expressing sFKN in the SNpc had a blunted response to MPTP compared to mFKN (Tukey's HSD; ***p = 0.007) and GFP (Tukey's HSD; p = 0.092). c-n, Representative immunoreactivity of CD68- and CD11b-reactive microglia in the SNpc for FKN $^{-/-}$ sham (c, e), WT-Sham (d, f), WT-MPTP (g, f), sFKN (f), f), mFKN (f), f), mFKN

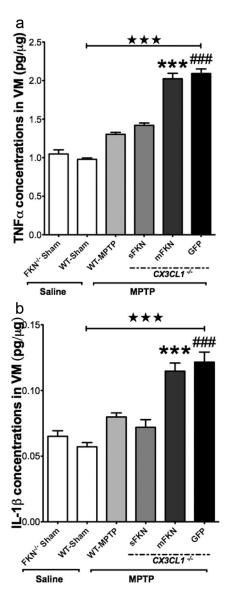


Figure 7. Induction of proinflammatory cytokines is attenuated in $CX3CL1^{-/-}$ mice expressing sFKN. TNF α and IL-1 β concentrations were measured using standard ELISA techniques for VM lysates. **a**, TNF α concentrations were upregulated following MPTP administration (three-way ANOVA; $F_{(1, 23)} = 18.36$, ***p < 0.001). Comparatively, $CX3CL1^{-/-}$ mice expressing sFKN in the SNpc had significantly lower concentrations of TNF α relative to mFKN (Tukey's HSD; ***p < 0.001) and GFP (Tukey's HSD; *#*p < 0.001) expressing mice. There were no significant differences between sFKN and WT-MPTP (Tukey's HSD; p = 0.384) or mFKN and GFP (Tukey's HSD; p = 0.773). **b**, The IL-1 β concentrations in the VM were significantly upregulated for mice exposed to MPTP (three-way ANOVA; $F_{(1, 23)} = 11.97$, ***p = 0.002). Similar to the pattern of TNF α , IL-1 β concentrations in $CX3CL1^{-/-}$ mice expressing sFKN were significantly blunted compared to both mFKN (Tukey's HSD; ***p = 0.001) and GFP (Tukey's HSD; ***p = 0.001). As was observed with TNF α , there were no significant differences in IL-1 β concentrations between WT-MPTP and sFKN (Tukey's HSD; p = 0.785) or mFKN and GFP (Tukey's HSD; p = 0.845) expressing mice. Data are presented as the mean \pm SEM.

crease in activated microglia (CD11b $^+$ and CD68 $^+$) following exposure to MPTP. Interestingly, sFKN attenuated the production of both TNF α (p < 0.001) and IL-1 β (p < 0.001) compared to mFKN-treated and GFP-treated mice. Comparably, the production levels of both proinflammatory cytokines in sFKN mice were similar to the levels found in the WT-MPTP group.

Discussion

Disruption of CX3CL1–CX3CR1 signaling in the brain increases MPTP-induced DA neurodegeneration (Cardona et al., 2006). Functionally, this signaling pair has been shown across a variety of studies to dampen microglial activation and subsequent toxicity. Until now, it has been difficult to isolate the individual effects of the two forms of CX3CL1. We have developed a novel method to examine the individual properties of the soluble versus the uncleaved membrane-anchored isoforms of CX3CL1 through the use of rAAV gene therapy in CX3CL1-null mice. Our study demonstrates that only sFKN attenuates the neurotoxic effects of MPTP on the nigrostriatal network and that the membraneanchored mFKN isoform had no effect in this model. Here, we show that treatment with sFKN reduces impairment of motor coordination, prevents DA neuron loss, and ameliorates microglial activation and proinflammatory cytokine release, all of which are commonly associated with the MPTP mouse model of PD.

These important discoveries provide novel insight into the role for CX3CL1 signaling in an acute inflammatory state. We have previously reported that an exogenous truncated peptide of CX3CL1 was able to reduce 6-OHDA-associated neurotoxicity in a rat model of PD (Pabon et al., 2011). However, this model still left unaccounted the actions of the full-length membrane-bound CX3CL1 isoform, as native CX3CL1 (both soluble and membrane bound) was still present in these animals. Additionally, Lyons et al. (2009) concluded that membrane-bound CX3CL1 possesses the same anti-inflammatory abilities as its soluble counterpart. However, Lyons et al. (2009) neglected to inhibit the constitutive cleavage mechanism of ADAM10/17 in their experimental paradigm. Here we have addressed this concern by mutating the known cleavage site of CX3CL1, thus preventing the soluble variant from being produced. Furthermore, by using CX3CL1-null mice, we were able to isolate the effects of each rAAV-FKN variant in response to MPTP.

In the present study we showed that the MPTP-induced motor deficit was significantly ameliorated by treatment with sFKN but not mFKN. Accordingly, mice expressing sFKN had significantly more TH immunoreactivity in the striatum compared to both mFKN-expressing and GFP-expressing mice. Furthermore, the depletion of TH-positive cells in the SNpc in response to MPTP was ameliorated by treatment with sFKN but not mFKN. Expression of TH has been shown to be somewhat plastic in response to MPTP, and therefore a decline in TH is not necessarily an indicative marker of degenerating DA neurons (Jackson-Lewis et al., 1995). To accurately assess whether neuronal loss was occurring, we measured the amount of NeuN staining in the SNpc. Our data show that the neuronal loss associated with MPTP administration shows an inclination of recovery by administration of sFKN but not mFKN.

CX3CL1-to-CX3CR1 communication is one way that neurons are thought to directly interact with microglia. Several studies have indicated that an acute inflammatory insult such as LPS exposure is sufficient to disrupt this dialog via downregulation of CX3CR1 transcript and protein levels. Here we show that exposure to acute administration of MPTP does not, under these circumstances, affect the signaling capabilities of this pair, as both CX3CR1 and CX3CL1 protein levels remain unchanged in response to MPTP exposure. Importantly, each of our rAAV-FKN variants restores CX3CL1 back to levels seen in WT mice.

The aberrant activation of microglia has been shown to increase the pathophysiology associated with PD. Moreover, in

vitro, a truncated CX3CL1 peptide has been shown to reduce the production of multiple proinflammatory molecules (Zujovic et al., 2000, 2001; Biber et al., 2007; Mizutani et al., 2007). Our current data show that only sFKN attenuates the production of both IL-1 β and TNF α in the VM of MPTP-injected mice and reduces microglial expression of CD68 and CD11b. These data suggest that treatment with mFKN was not efficacious in reducing the histological and molecular proinflammatory phenotype.

There has been a recent surge in the literature regarding the role for CX3CL1 signaling in both physiological and pathological conditions. We have recently shown that genetic disruption of CX3CL1 signaling is sufficient to impair cognitive function (Rogers et al., 2011), while others have demonstrated both neuroprotective (Bhaskar et al., 2010) and neurotoxic (Fuhrmann et al., 2010; Lee et al., 2010) effects of CX3CL1 signaling. Here we report a novel interaction wherein CX3CL1-mediated anti-inflammatory control of microglia is accomplished solely through the soluble isoform in a MPTP mouse model of PD. In this model, through the exclusive interactions of sFKN and CX3CR1, we demonstrate that amelioration of impaired motor coordination is mediated through a reduction in microglial activation and subsequent proinflammatory response. These effects are completely absent in mice expressing only mFKN. A role of the membrane-bound form has been demonstrated in the periphery in mice that only express either the membrane-bound or soluble forms of CX3CL1 (Kim et al., 2011). In these mice the soluble form is shown to be important for intestinal macrophages to form transepithelial dendrites; however, only the membrane-bound form was able to rescue Ly6C lowCX3CR1 high monocytes. As such, a role for different types of CX3CL1 may depend upon the site of action.

Given that only the soluble form of CX3CL1 provides an antiinflammatory and therefore neuroprotective function in this model of PD, we suggest that ligation with the soluble but not the membrane-bound isoform allows for CX3CR1 internalization. Recent literature has indicated that multiple chemokine receptors actively retreat through internalization to the cytoplasm following ligation to affect downstream signaling pathways (Neel et al., 2005; Borroni et al., 2010). Although ligation of mFKN with CX3CR1 is possible through the chemokine domain, its permanent attachment to the membrane may prevent subsequent internalization to the cytoplasm of microglia, thereby altering any downstream effectors. Therefore, only the soluble domain of CX3CL1 can be readily internalized with CX3CR1 ligation. Recent literature corroborates this hypothesis in that CX3CR1 actively removes soluble CX3CL1 from the surrounding milieu, as CX3CR1 -/- mice have substantially more circulating CX3CL1 (Cardona et al., 2008). Moreover, this justifies the need for the constitutive and endogenous proteolytic cleavage by ADAM10/17 to yield soluble CX3CL1 in an effort to maintain a constant neuroprotective dialog between neurons and microglia.

We have created two very effective tools to examine the signaling characteristics of the two native isoforms of CX3CL1. Although we used an acute inflammatory model to examine these effects, it is worth noting that the relationships we detailed in this study may not apply to all inflammatory-related neurodegenerative conditions. Further examination of both of the CX3CL1 isoforms in various animal models of neuroinflammation and degenerative disease models is necessary to expand our understanding of this unique chemokine signaling pair.

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