

Dysregulation of neural activity and microglia function following exposure to the global environmental contaminant perfluorooctane sulfonate (PFOS)

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ABSTRACT

Pollution is a driving force in climate change and an important modifier of human health. Humans are chronically exposed to complex chemical mixtures and, correspondingly, researchers are disentangling the contribution of different contaminants to human neuropathologies. Per- and polyfluoroalkyl substances (PFAS) are biopersistent pollutants and, due to their diverse applications, have become global contaminants. Perfluorooctane sulfonate (PFOS), a prevalent PFAS congener, impairs humoral immunity; however, its impact on innate immunity is unclear. Given the critical roles of innate immune cells, namely microglia, in brain development and homeostasis, we asked whether exposure adversely affects microglial function. Herein, we demonstrate developmental PFOS exposure produces microglial activation and upregulation of the microglia activation gene *p2ry12*. PFOS-induced microglial activation heightened microglial responses to brain injury, in the absence of increased cell death or inflammation. Use of the photoconvertible calcium indicator CaMPARI revealed PFOS exposure heightened neural activity, while optogenetic silencing of neurons was sufficient to normalize microglial responses to injury. Exposure to perfluorooctanoic acid, an immunotoxic PFAS, did not alter neuronal activity or microglial behavior, further supporting a role for neural activity as a critical modifier of microglial function. Together, this study reveals how contaminant-induced changes in brain activity can shape brain health.

INTRODUCTION

Pollution poses a global threat to environmental, ecologic, and economic health and stability. As international geologic committees consider whether we have now entered the “Anthropocene Age,” it is of increasing importance to understand the broad impacts of human-made pollutants. It is also worth emphasizing that the burden of pollution is disproportionately shared, with minority and low-income communities being more susceptible to the physical consequences of insufficient regulatory laws, industrial waste disposal, and air pollution (Bullard 1994, Muller, Sampson and Winter 2018, Liu et al. 2021). If we

are to protect our health, society, and environment, more research efforts are needed to address the costs of our past and present environmental negligence.

One major class of chemicals of increasing concern are per- and polyfluoroalkyl substances (PFAS). While PFAS were only introduced in industrial manufacturing in the mid-20th century, the expansive and international utilization of PFAS has led to near universal exposure in humans (Kannan et al. 2004, Paul, Jones and Sweetman 2009). PFAS congeners are found in fire-fighting foams, commercial household products such as water-repellant fabrics, food packaging, and non-stick finishes, and are used in a variety of applications in the aerospace, aviation, and automotive industries (USEPA). Structurally, PFAS consist of fully or polyfluorinated aliphatic substances of varying carbon chain length and head groups, with longer chain length tending to be associated with increased toxicity (Buck et al. 2011, Gomis et al. 2018, Chambers, Hopkins and Richards 2021). The strength of carbon-fluorine bonds lends most PFAS to be biopersistent, bioaccumulative, and resistant to degradation. As such, PFAS have acquired the disconcerting moniker “forever chemicals.”

Of the 4,700+ PFAS congeners (OECD 2018), one of the most environmentally prevalent and extensively studied is perfluorooctane sulfonate (PFOS). PFOS has an 8-carbon chain with a sulfonic acid head group and adversely impacts the functioning and health of several major organ systems (Sunderland et al. 2019, Zeng et al. 2019), and is considered a metabolic, endocrine, and immune disruptor (DeWitt, Copeland and Luebke 2009a, DeWitt et al. 2012, DeWitt et al. 2009b, DeWitt et al. 2016, Stein et al. 2016, Braun 2017, Chang et al. 2016). *In utero* and developmental exposure to PFOS has particularly concerning consequences on adaptive immunity. Grandjean et al. demonstrated that elevated PFOS levels in infancy and early childhood significantly attenuate adaptive immune responses, curb antibody production, and limit vaccine efficacy (Grandjean et al. 2012, Grandjean et al. 2017). Such studies reinforce the necessity of understanding the impact of pollution on population health, especially considering the need to vaccinate individuals against emerging or evolving pathogens. PFOS-induced humoral immune suppression (DeWitt et al. 2012, Peden-Adams et al. 2008) prompted the National Toxicology Program to classify PFOS as a “presumed immune hazard to humans” in 2016 (NTP 2016). However, the effects of PFOS on innate

immunity are still largely inconclusive, with some groups finding that PFOS dampens innate cell infiltration, gene expression, or activity (Castano-Ortiz, Jaspers and Waugh 2019, Keil et al. 2008, Ryu et al. 2014), and others describing heightened inflammation or immune function (Dong et al. 2012, Qazi et al. 2009, Wang et al. 2021). Even less understood is the tissue-specific impact of PFOS on local immune populations, namely tissue-resident macrophages. Tissue-resident macrophages are largely yolk-sac or fetal liver-derived, heterogenous, and most notably carry out discrete, non-canonical tissue-specific functions essential for development and homeostasis (Davies et al. 2013). Determining whether tissue-resident macrophages are PFAS targets, especially during development, is essential for our understanding of long-term consequences of macrophage dysregulation following PFAS exposure.

In this work, we focus on microglia, the resident immune population of the central nervous system (CNS), as a potential PFOS target. Beyond effector immune responses, microglia have a highly dynamic and diverse repertoire of homeostatic functions, including developmental pruning of extra-numerary synapses (Paolicelli et al. 2011, Schafer et al. 2012), facilitating synaptogenesis and maturation (Hoshiko et al. 2012), and regulation of neural plasticity and dendritic spine density through frequent synaptic monitoring (Parkhurst et al. 2013, Szepesi et al. 2018, Wake et al. 2009). As sentinels of the CNS, these persistent, self-renewing cells are also highly sensitive and rapidly adaptable to any changes in their environment. As such, microglia are often categorized into two broadly termed polarization states: homeostatic and ramified versus activated and reactive. Homeostatic microglia have highly motile processes that continuously extend and retract to survey the surrounding environment, including active synapses in an activity-dependent manner (Dissing-Olesen et al. 2014, Eyo et al. 2014, Nimmerjahn, Kirchhoff and Helmchen 2005, Tremblay, Lowery and Majewska 2010, Wake et al. 2009). Conversely, activated microglia, which are actuated by pathogenic or local endogenous stimuli, exhibit an amoeboid-like morphology, are migratory, proliferative, phagocytic, and secrete immunomodulatory and neuroactive factors (Kettenmann et al. 2011).

Microglia activation has been well documented in disease models of various neuropathological states, including Down Syndrome (Pinto et al. 2020), Alzheimer's disease (Mancuso et al. 2019), and epilepsy (Hiragi, Ikegaya and Koyama 2018). Due to the critical regulatory roles of microglia in neural health and

development, there has been a concerted effort to clarify the mechanisms of bidirectional cross-talk between neurons and microglia (Szepesi et al. 2018). However, the impact of environmental exposures on neuron-microglia interactions during developmentally sensitive periods of life, including gestation, has been largely overlooked. Some epidemiological studies suggest correlations between developmental PFOS exposure and ADHD incidence, while others found no such relationship (Fei et al. 2008, Lien et al. 2016, Liew et al. 2015, Ode et al. 2014). Meanwhile in mice, a single neonatal PFOS exposure is sufficient to alter proteins required for neuronal growth and synaptogenesis, as well as cause spontaneous behavior and hyperactivity in adults (Johansson, Eriksson and Viberg 2009, Johansson, Fredriksson and Eriksson 2008). Developmental exposure to PFOS was also shown to cause hyperactive locomotor activity in larval zebrafish (Gaballah et al. 2020, Huang et al. 2010, Jantzen, Annunziato and Cooper 2016). *In vitro* experiments using cultured neurons revealed that PFOS interacts with inositol 1,4,5-triphosphate receptors (IP₃Rs) and ryanodine receptors (RyRs), leading to the release of intracellular calcium stores, which suggests a potential role for calcium in PFOS-induced neurotoxicity (Liu et al. 2011). However, it is still not known whether the induced locomotor activity in zebrafish is attributed to neuronal hyperactivity or skeletal muscle calcium utilization (Christou et al. 2020). It is also not known if potential changes in the neuronal environment following PFOS exposure can impact microglia function, and vice versa.

Herein, we employed functional neuroimaging in larval zebrafish to provide the first *in vivo* account of PFOS-induced disruption of neuronal activity. We further observed the direct effects of PFOS-induced neuronal hyperactivity on microglia activation and function by using an established brain injury model for zebrafish (Schmidt et al. 2014). By utilizing optogenetics to independently modulate the neuronal or microglial electrical state, we demonstrate that PFOS-induced neuronal hyperactivity and microglia hyperresponsiveness could be reversed, respectively, further supporting the role of neuronal activity on

microglia function. Lastly, by exposing larvae to a non-excitatory PFAS congener, we demonstrate that structurally similar compounds can have significantly distinct effects on CNS health and neural cell function.

RESULTS

PFOS exposure activates microglia and renders them hyperresponsive to minor brain injury.

Studies utilizing immortalized microglia cell lines suggest PFOS exposure decreases microglial viability, mitochondrial stability, and increases reactive oxygen species production *in vitro* in a concentration-dependent manner (Ge et al. 2016, Zhang et al. 2011). However, whether PFOS affects microglia activation and function *in vivo* is not yet known. Given the critical developmental and modulatory roles of microglia in the CNS, we sought to determine whether embryonic exposure to PFOS had a direct effect on microglia colonization of the developing brain. We selected a concentration exposure range based on previously published work demonstrating the adverse effects of PFOS exposure on pancreatic development in larval zebrafish (Sant et al. 2017, Chen et al. 2014). Transgenic zebrafish embryos with macrophage-specific GFP expression under the *mpeg1* promoter (*Tg(mpeg1:EGFP)*) (Ellett et al. 2011b) were chronically exposed to either a control solution (0.1% DMSO) or 28 μ M PFOS from 4 hours post-fertilization (hpf) until 3 days post-fertilization (dpf) (Figure 1A). Using liquid chromatography with high-resolution mass spectrometry (LC-HRMS), we validated our exposure solutions and determined that 28 μ M PFOS resulted in a total body burden of 32.22 ± 2.57 ng/embryo at 48 hpf and 70.46 ± 2.72 ng/embryo at 72 hpf (Table 1). Next, we examined microglial number and morphology at 72 hpf, a time point at which no significant increase in mortality was observed across the tested concentration range (Table 2). Non-parenchymal, parenchymal, and total microglia number was not changed at 72 hpf between the control and 28 μ M PFOS-exposed larvae (Figure 1—figure supplement 1A-C). While microglia number was unchanged, cell morphology was significantly altered. PFOS-exposed microglia acquired a less ramified and more amoeboid cell shape, resembling an activated phenotype (Figure 1D-E" vs 1I-J"). This corresponded to a significant reduction in cell area (Figure 1N), cell perimeter (Figure 1O), and the acquisition of a more

circular cell shape as calculated by a decrease in the perimeter-to-area ratio (Figure 1P). Relative mRNA analysis for the gene *p2ry12*, a G-protein coupled receptor directly involved in microglia activation and migration behavior (Badimon et al. 2020, Davalos et al. 2005, Yu et al. 2019), was also significantly upregulated in the brains of PFOS-exposed larvae (Figure 1Q), further supporting the activation of microglia. To determine whether the observed phenotypic changes were accompanied by functional changes, we tested whether pollutant-induced microglia activation affected the ability of microglia to respond to minor brain injury using an established zebrafish injury model (Schmidt et al. 2014, Sieger et al. 2012). At 72 hpf, larvae were injured at the right hemisphere of the telencephalon using a pulled glass needle (OD 9 μ m; Figure 1B) and microglia recruitment to the injury site was monitored for the first 4.5 hours post-injury (hpi; Figure 1C). PFOS-exposed larvae had a significant increase in microglial recruitment in the first 4.5 hpi compared to sibling controls (Figure 1R; Figure 1—figure supplement videos 1 versus 2). Additionally, the area of microglial response in exposed larvae was significantly expanded at 4 hpi (Figure 1G vs 1L; Figure 1S), which persisted up to 12 hpi (Figure 1H vs 1M; Figure 1T). A major function of microglia following brain injury is the removal of damaged cells and cellular debris to prevent excessive inflammation, which is deleterious to brain health (Donat et al. 2017). Using the nucleic acid stain acridine orange in live larvae, we asked whether an increased incidence of cell death was driving the microglia response to injury in pollutant-exposed larvae; however, PFOS-exposure did not result in an increase in cell death in the whole brain nor at the injury site at 1 or 4 hpi (Figure 1—figure supplement 1D-H). Moreover, relative mRNA expression of the proinflammatory cytokines *il1 β* , *tnfa*, and *il6* were unchanged in larval brains following exposure (Figure 1—figure supplement 1I). While these data suggest microglial hyperresponsiveness in PFOS-exposed larvae is not directed by canonical damage signaling, the

morphological and transcriptional shift of microglia toward an activated phenotype (Figure 1N-Q) suggest that microglia are activated following PFOS exposure and may be targets of PFOS toxicity.

Optogenetic Reversal of Microglia Activation State Normalizes the Microglial Response to Injury

To determine the extent to which microglial activation state contributed to the heightened microglial response to injury, we next asked whether PFOS-induced microglia activation could be reversed *in vivo*, and if this reversal of activation state was sufficient to normalize the injury response. Since the homeostatic microglia membrane potential is largely maintained by chloride channel currents (Newell and Schlichter 2005), we utilized transgenic zebrafish expressing the light-gated chloride pump halorhodopsin (eNpHR3.0) specifically in microglia (*Tg(mpeg1:Gal4FF;UAS:eNpHR3.0-mCherry)*) (Figure 2A). This third-generation opsin exhibits reliable membrane trafficking for uniform surface expression, is resistant to inactivation, and has step-like, potent photocurrents that are stable over long timescales (Gradinaru et al. 2010). Larvae with halorhodopsin⁺ microglia were exposed to 28 μ M PFOS to induce microglia activation by 72 hpf, at which point the larvae were injured and exposed to 589 nm light for 4 hours (Figure 2B,C). We confirmed the morphological shift toward an activated phenotype in unstimulated PFOS-exposed larvae (Figure 2D-E'') and compared the microglia morphology of unstimulated larvae to larvae stimulated by 589 nm light. Halorhodopsin stimulation of PFOS-exposed microglia produced a significantly more ramified morphology (Figure 2H-I''). Additionally, while stimulation had no effect on microglia area (Figure 2L), it significantly increased the cell perimeter (Figure 2M) and perimeter-to-area ratio (Figure 2N), supporting the transition from amoeboid to ramified. We also confirmed 589 nm stimulation alone had no impact on microglia morphology by stimulating and assessing microglia from PFOS-exposed larvae lacking halorhodopsin expression, *Tg(mpeg1:Gal4FF;UAS:nfsb-mCherry)* (Figure 2—figure supplement 1). We next tested whether optogenetic modulation of microglia activation state was sufficient to rescue their responses to brain injury. Indeed, while unstimulated larvae had exacerbated microglia responses to injury (Figure 2F vs 2G), stimulation of halorhodopsin in PFOS-exposed microglia significantly reduced the microglia response (Figure 2G vs 2K; area of response quantified in Figure 2O). These data suggest that modulation of

microglia activity state alone is sufficient to normalize their functions following toxicant exposure. Additionally, given that microglia are highly responsive to their microenvironments, it suggests that microglia hyperresponsiveness in PFOS-exposed larvae may be influenced by changes in neuronal communication.

Developmental PFOS exposure alters global and regional neuronal network activity

While evidence suggests PFOS exposure can impact developing and adult brain health, the direct effects of exposure on neuronal network activity, whole brain metabolome, and neurotransmitter release are not well understood. To understand the effect of PFOS on global brain activity *in vivo*, we assessed neuronal calcium activity by driving the fluorescent calcium sensor CaMPARI (Calcium-Modulated Photoactivatable Ratiometric Integrator) under the pan-neuronal promoter *elavl3* (*Tg(elavl3:CaMPARI(W391F+V398L))^{tg}*). CaMPARI is a permanent photoconvertible calcium sensor that undergoes allosteric chromophore modulation from green-to-red in response to ultraviolet light, but only upon simultaneous binding of free intracellular calcium (Fosque et al. 2015). Therefore, inactive neurons at the time of photoconversion are green, while active neurons convert to red (Figure 3A-D). In this study, free swimming larvae were subjected to 135 mW/cm² of 405 nm light for 1 minute inside our behavioral unit (Figure 3—figure supplement 1A-D). We validated our CaMPARI photoconversion, imaging, and analysis pipeline by exposing larvae to pentylentetrazol (PTZ), a GABAA receptor antagonist known to cause neuronal hyperactivity and seizures in zebrafish (Baraban et al. 2005) (Figure 3—figure supplement 1E-H versus 1E'-H'). Indeed, 10 mM PTZ led to a significant ratio-metric increase of neuronal intracellular calcium in the optic tectum and cerebellum, as well as the whole brain (Figure 3—figure supplement 1J). To determine whether PFOS exposure affected brain activity, we applied this validated pipeline to 3 dpf larvae exposed to 28 μ M PFOS, the same concentration used for the injury model. PFOS exposure caused a notable increase in neuronal activity in the forebrain, optic tectum, and hindbrain, with a significant increase in the cerebellum and whole brain collectively (Figure 3G,I). We also examined whether lower concentrations of PFOS were able to induce changes in neuronal activity. While there were no significant changes in network activity at 3 dpf following 7 μ M PFOS (Figure 3F,H), this group had significant increases in regional and global activity at 5 dpf (Figure 3K,M), as did the 14 μ M exposed group (Figure 3L,N). We also assessed gross brain morphology of PFOS-

exposed larvae and found that while whole brain area was slightly reduced at 3 dpf, brain area is unchanged by 5 dpf, suggesting that the effects of PFOS on morphology are nominal and temporary (Figure 3—figure supplement 2). Together, these data reveal that PFOS has a time and concentration dependent effect on network activity, exclusive of effects on gross morphology.

To better understand the neurochemical changes in the PFOS exposed brain, we performed an untargeted metabolome wide association study (MWAS) on isolated heads of 3 dpf control and 28 μ M PFOS exposed larvae. Several features were significantly upregulated or downregulated between the control and exposed groups (Figure 3—figure supplement 3A). Significantly changed metabolites from the MWAS were further analyzed for pathway enrichment using MetaboAnalystR (Chong and Xia 2018). Several of the significantly enriched pathways, including glutamate metabolism, aspartate and asparagine metabolism, tyrosine metabolism, and phosphoinositide metabolism, are involved in neuronal excitation, catecholamine synthesis, and neurotransmitter release, uptake, and recycling (Figure 3—figure supplement 3C; Table 3). Together with the CaMPARI analysis, these data suggest the brain is a PFOS target and significant neurochemical imbalances occur following exposure.

We next sought to determine whether the neurochemical imbalances were predictors of abnormal behavioral activity. A single exposure event to PFOS is sufficient to drive spontaneous behavior and hyperactivity in adult mice (Johansson et al. 2008) and can lead to tonic convulsions in response to an ultrasonic stimulus (Sato et al. 2009). Additionally, developmental exposure to PFOS decreases the frequency but increases the intensity of spontaneous locomotor activity in both mice and larval zebrafish in a concentration-dependent manner (Spulber et al. 2014). To first determine whether our exposure paradigm replicates previously reported behavioral hyperactivity, we exposed embryos to either 7 μ M, 14 μ M, or 28 μ M PFOS at 4 hpf, and performed a behavioral photomotor response assay at 3 dpf, 4 dpf, and 5 dpf (Figure 4—figure supplement 1A,B). Consistent with previous locomotor assays at 6 dpf (Gaballah et al. 2020), PFOS-exposed larvae exhibited significantly increased photomotor responses to light changes, designated by increased distance traveled within the well at 3 dpf following 28 μ M exposure (Figure 4—figure supplement 1E-H) and 5 dpf following 7 μ M and 14 μ M exposure (Figure 4—figure supplement 1N-

R). In addition to distanced traveled, behavior videos were analyzed to assess the frequency of center avoidance within the wells of a 24-well plate. Similar to the mammalian open-field test, center avoidance in a well is an indication of zebrafish anxiety, while willingness to cross the center suggests a reduced anxiety level (Colwill and Creton 2011). As such, we examined the time each larva spent along the well's edge (anxious) versus center (exploratory) throughout each photomotor assay (Figure 4 A,B). PFOS exposure resulted in a notable increase in anxiety-like behavior in all dosed groups at 3, 4, and 5 dpf (Figure 4 C-E). Since anxiety, fear, and aversive behaviors are regulated by the medial habenula in zebrafish (Mathuru and Jesuthasan 2013, Okamoto, Agetsuma and Aizawa 2012), we assessed neuron activity at the habenula using CaMPARI. Only 14 μ M PFOS exposed larvae showed a significant difference in habenular activity at 3 and 5 dpf (Figure 4—figure supplement 2), suggesting that habenular dysregulation alone may not be an accurate predictor of anxiety in larval zebrafish. Interestingly, while 14 μ M PFOS-exposed larvae exhibited the most pronounced anxiety responses at 3, 4, and 5 dpf, they did not have an increase in swim activity at 3 dpf (Figure 4—figure supplement 1H), and only showed an increase in activity when the behavior box light was “On” at 4 dpf (Figure 4—figure supplement 1K). This suggests that anxiety behaviors are separable from swim hyperactivity and support the need to more thoroughly understand the independent behavioral changes associated with PFOS exposure.

Microglia mutants have reduced swim behavior, but are more reactive to light transitions following PFOS exposure Neuron-microglia bi-directional signaling enables the reciprocal regulation of microglia and neuronal functions, including microglial modulation of neuronal activity and circuit refinement (Marinelli et al. 2019, Szepesi et al. 2018). Considering that microglia become activated following PFOS exposure, we sought to determine whether or not PFOS-induced microglial dysfunction during early larval development influenced neuronal excitation and behavior. To do this, we exposed PFOS to zebrafish with a null mutation for *irf8* (*irf8*^{st96/st96}), a gene required for macrophage formation during primitive and transient definitive hematopoiesis (Shiau et al. 2015). *irf8* mutant larvae lack the earliest embryonic-derived macrophage populations, including microglia (Shiau et al. 2015). Control *irf8* mutants had a consistent reduction in overall swim behavior and were significantly less reactive to the light-dark transitions (Figure 5A,B). Exposure to 8 μ M PFOS significantly increased swim behavior of 5 dpf wildtype and *irf8* mutant

larvae (Figure 5A). However, unlike the control-treated *irf8* mutants, PFOS-exposed mutants were not consistently less active during the photomotor assay. PFOS-exposed mutants also exhibited the most dramatic behavioral response to the light-dark transitions, especially during the second dark cycle (Figure 5B; “First Minute”). In addition, while wildtype control larvae and wildtype PFOS-exposed larvae had a 14.9% and 23.8% reduction in swim activity by the end of the second dark cycle, respectively, PFOS-exposed *irf8* mutants had a 47.6% reduction (Figure 5B). These data suggest that microglia may attempt to modulate the PFOS-exposed brain, such that the absence of microglia may further promote PFOS-induced behavioral reactivity. Heightened reactivity may also result in behaviors that impair swim ability, such as seizing or convulsions. Microglia-deficient larvae with transgenic expression of neuron-driven CaMPARI also had significant increases in regional and global calcium activity over controls, though these were not significant between genotypes (Figure 5C).

Neuronal activity modulates microglia responsiveness

While microglia are not wholly responsible for the behavioral or neuronal phenotypes seen with PFOS-exposure, PFOS-induced neuronal hyperactivity may still be an important modulator of microglia. To test this hypothesis, 3 dpf larvae were treated with 5 mM PTZ to induce neuronal activity, then were injured in the right telencephalon as described. PTZ-induced neuronal hyperactivity significantly increased the microglia response at 4 hpi, similar to that of PFOS-exposed larvae (Figure 6—figure supplement 1). We next asked whether inhibiting neuronal activity using optogenetics could normalize the hyperresponsive microglial phenotype seen with PFOS exposure. Previous studies implementing halorhodopsin for neuronal hyperpolarization in zebrafish demonstrated significant suppression of coiling behavior, swim activity, and silencing of single neuron spiking (Antinucci et al. 2020, Arrenberg, Del Bene and Baier 2009). Larvae expressing *Tg(elavl3:Gal4;cryaa:RFP;UAS:eNpHR3.0;mpeg1:EGFP)*, which have pan-neuronal expression of halorhodopsin as well as GFP-labeled macrophages, were exposed to either a control solution or 28 μ M PFOS at 4 hpf. After brain injury at 3 dpf, larvae were either left unstimulated or stimulated for 4 hours with 589 nm light to silence neuronal activity. Unstimulated PFOS-exposed larvae exhibited a significant increase in microglia response to injury, as demonstrated earlier (Figure 6E,F,I). Conversely,

neuronal hyperpolarization rescued microglia hyperresponsiveness (Figure 6G,H,I), further supporting that pollutant-induced changes in the neural signaling environment significantly influences microglia behavior, independent of cell death or inflammation.

Exposure to PFOA, an immunotoxic PFAS congener, does not result in microglia hyperresponsiveness

To further assess if microglia hyperresponsiveness following PFOS exposure is a result of neuronal hyperactivity, we exposed larval zebrafish to a structurally similar PFAS congener, perfluorooctanoic acid (PFOA). Whereas PFOS has a sulfonic acid head group, PFOA is an 8-carbon PFAS with a carboxylic acid head group. PFOA has been shown to cause significant health effects, including immunotoxicity, elevated cholesterol, dysregulated liver metabolism, kidney dysfunction, thyroid disease, among many others (Reviewed in (Post, Cohn and Cooper 2012)). Previous reports using larval zebrafish have found that PFOA concentrations ranging from 4.4 μ M to 80 μ M did not result in a significant change in 6 dpf swim behavior during photomotor response assays (Gaballah et al. 2020). Therefore, we hypothesized that PFOA may also have unchanged neuronal network activity, making it an ideal candidate to address the capacity of neuronal activity to influence microglial function. Following the same exposure paradigm used for PFOS exposure (Figure 1A), zebrafish embryos were dosed with either a control solution (0.1% DMSO) or 64 μ M PFOA at 4 hpf. In concordance with previous findings, we found that exposure to 64 μ M PFOA did not result in altered swim behavior during photomotor response assays at 5 dpf (Figure 7—figure supplement 1A-D). While PFOA-exposed larvae had shorter body length, there were no observable gross morphological effects on the spine that would affect swimming ability (Figure 7—figure supplement 2). Using neuronally-driven CaMPARI, PFOA exposure did not result in any regional or global changes in neuronal network activity in 5 dpf larvae (Figure 7A-C), unlike the PFOS-exposed groups at this timepoint (Figure 3). Given that PFOA did not result in neuronal hyperactivity, we asked whether PFOA exposure affected microglial responses to brain injury. Indeed, PFOA exposure did not result in altered microglial responses to brain injury (Figure

7D-F). Together, these data further support that neuronal hyperactivity is a key driver of microglia response, and that structurally similar PFAS congeners can have distinct impacts on the CNS.

DISCUSSION

For more than two decades, researchers have revealed the many biologic, ecologic, and environmental ramifications of PFOS toxicity, as well as the toxicity profiles for a subset of other ‘forever chemical’ congeners. Here, we used a combination of *in vivo* imaging of cellular behavior, functional neuroimaging, optogenetic modulation, and behavioral assays to address the impact of PFOS exposure on microglia-neuron interactions in larval zebrafish. Our data demonstrate that developmental PFOS exposure activates microglia and induces hyperactivity in response to injury independent of cell death or inflammation. Since the homeostatic microglia membrane potential is largely maintained by chloride channel currents (Newell and Schlichter 2005), we asked whether activating the optogenetic chloride channel halorhodopsin (eNpHR3.0) could rescue microglia hyperresponsiveness. Indeed, electrical modulation of microglia was sufficient to revert microglia from an activated to a homeostatic morphology and normalize their responses to injury, suggesting electrical dysfunction as a previously underappreciated pathway worth interrogating when studying microglial activation and immunotoxicity. PFOS-exposed larvae also exhibited global and regional increases in neuronal activity and anxiety-like behaviors, a previously unidentified neurodevelopmental phenotype of PFOS exposure in zebrafish, with only nominal and temporary impacts on regional brain morphology. We demonstrated that PFOS-induced neuronal hyperactivity was a key mediator of microglia reactivity and that optogenetic silencing of neurons was sufficient to normalize microglia responses to injury. Further, exposure to PFOA did not result in neuronal hyperactivity nor microglia hyperresponsiveness. Together, this study provides the first detailed account of the effects of PFOS exposure on the developing brain *in vivo* and adds neuronal hyperactivity as an important endpoint to assess when studying the impact of toxicant exposures on microglia function.

While the immunotoxic impact of PFOS on adaptive immunity has been well documented (DeWitt, Blossom and Schaidler 2019), the effects on the innate immune arm are less understood and, at times, contradictory. Studies showing either innate immune activation or suppression following PFOS exposure

are likely attributed to varying exposure paradigms, PFOS concentration, animal model, age, or cell line used (Castano-Ortiz et al. 2019, Mollenhauer et al. 2011, Qazi et al. 2009, Ryu et al. 2014, Wang et al. 2021). The innate immune system is highly sensitive to both endogenous and xenobiotic stimuli and reacts rapidly, creating a signaling cascade that informs all downstream immune functions (Clark and Kupper 2005). Macrophages in particular have the ability to dynamically polarize from a homeostatic state to be pro- or anti-inflammatory depending on the environmental needs. As antigen-presenting cells, macrophages also have the vital role of instructing adaptive immune cells on their responses (Clark and Kupper 2005). The dependency of the adaptive immune system on the innate emphasizes the need to clarify the immunotoxic mechanisms of pollutants like PFOS.

This work constitutes the first *in vivo* analysis of microglia, the resident innate immune population of the CNS, following PFAS exposure. Previous *in vitro* studies using immortalized microglia cell lines suggest PFOS exposure decreases microglial viability, mitochondrial stability, and increases ROS production *in vitro* in a concentration-dependent manner (Ge et al. 2016, Zhang et al. 2011). We found that PFOS exposure activates microglia, demonstrated by their morphological transition from ramified to amoeboid shaped, as well as by the upregulation of the microglia activation gene *p2ry12*. P2ry12 is also a purinergic receptor that responds to ATP. It has been demonstrated that microglia response to injury is mediated by glutamate-evoked calcium waves and ATP release from the injury site (Sieger et al. 2012); therefore ATP released by the injury site and/or through high neuronal activity could be contributing to the microglial hyperresponsiveness in PFOS-exposed larvae. It is likewise still unclear to what extent PFOS is acting directly on the microglia or indirectly through neuronal activation. Microglia activation and injury responses were attenuated by both microglia and neuronal optogenetic silencing, suggesting that the neuronal environment has a significant influence on the microglia. While we are providing the first demonstration of microglia activation in response to toxicant exposure, microglia activation has been well documented in disease models of various neuropathological states. In addition, pharmacological prevention of microglia activation or microglia depletion has led to cognitive and functional improvements in some neurological disease models (Goldfarb et al. 2021, Mancuso et al. 2019, Pinto et al. 2020). However, it is worth noting that activated microglia are not inherently pathological. For example, microglia depletion

following stroke significantly increased infarct size, neuronal cell death, and caused calcium overload (Szalay et al. 2016). Microglia depletion during acute seizures also exacerbated excitotoxicity and seizure sensitivity (Liu et al. 2020). Additionally, the regional specificity of microglia activation in a mouse model of chronic stress was considered a protective and/or adaptive response (Tynan et al. 2010).

Due to the developmental and homeostatic roles microglia have on maintaining the excitatory/inhibitory balance (Henstridge, Tzioras and Paolicelli 2019), as well as the situation-specific adverse effects that microglia activation may have on the CNS, we asked whether aberrant microglial activation following PFOS exposure was contributing to the behavioral or neuronal hyperactivity. Using microglia-deficient zebrafish with a mutation in the gene *irf8*, we found that control-treated *irf8* mutants had a consistent reduction in swim behavior that was not seen the PFOS-treated mutants. Additionally, PFOS-exposed mutants had the greatest response to the light-dark transition, suggesting that microglia may actually attempt to repress PFOS-induced behavioral hyperactivity. While microglia loss did not affect baseline neuronal calcium signaling, it is worth noting that microglia loss is distinct from microglia dysfunction, and thus does not rule out the contribution of microglia to neuronal hyperactivity in this context. In addition, microglia increasingly accumulate in synaptic regions between 7 and 28 dpf in zebrafish (Silva et al. 2021), suggesting that any significant death-dependent or activity-dependent neuronal pruning may occur developmentally later than the timepoints investigated in this study. Lastly, the concentrations of PFOS used in this study might increase neuronal activity too substantially or even irreversibly, such that electrical modulation or loss of microglia was not sufficient to influence the activity in a measurable way.

To our knowledge, this is the first report of *in vivo* neuronal hyperactivity caused by embryonic PFOS exposure. We show that embryonic PFOS exposure increased intracellular calcium concentrations across multiple brain regions at various concentrations (7 μ M, 14 μ M, and 28 μ M) and time points (3 dpf & 5 dpf). Both gestational and adult PFOS exposures impact calcium dependent signaling molecules important for memory, including Ca^{2+} /calmodulin dependent kinase II (CaMKII) and cAMP response element-binding protein (CREB) in rats (Liu et al. 2010), suggesting that disrupted calcium signaling is an important mediator of PFOS-induced neurotoxicity. *In vitro* studies also link PFOS toxicity to disrupted calcium homeostasis (Liao et al. 2008, Liu et al. 2011). While the underlying mechanisms remain unknown,

possible explanations include increased influx through activation of L-type Ca^{2+} channels (Liao et al. 2008) and release of intracellular calcium stores through interaction with ryanodine and inositol 1,4,5-trisphosphate receptors (Liu et al. 2011). Intracellular calcium excess in neurons promotes excitotoxicity and can cause brain damage leading to various neurological and neurodegenerative disorders (Armada-Moreira et al. 2020, Olloquequi et al. 2018), emphasizing the need to further understand the influence of PFAS on neuronal function. PFAS compounds have also been shown to activate peroxisome proliferator-activated receptors (PPARs) *in vitro* as well as in zebrafish (Rosen et al. 2017, Takacs and Abbott 2007). PPAR- γ in particular is expressed in neurons, microglia, and astrocytes and mediates inflammatory responses in the CNS (Reviewed in (Villapol 2018)). Interrogating PPAR activation in the CNS following PFOS exposure could also provide important information about the specific pathways impacted by exposure.

The susceptibility of the developing human brain to PFOS remains a contentious point due to conflicting data. While some studies have demonstrated significant correlations between developmental PFOS exposure and ADHD incidence (Hoffman et al. 2010, Lenters et al. 2019), others found no such relationship (Fei et al. 2008, Lien et al. 2016, Liew et al. 2015, Ode et al. 2014). In the mouse brain, PFOS concentrations have been shown to increase over time and lead to tonic convulsions, despite a lack of morphological phenotypes (Sato et al. 2009). The increase in regional and global network activity demonstrated in this study warrant an assessment of convulsive phenotypes and seizure activity. While outside the scope of this study, determining the threshold and timescale at which neuronal hyperactivity increases incidence of convulsion would be an interesting to pursue

Similar to previous reports (Gaballah et al. 2020, Jantzen et al. 2016, Huang et al. 2010), we observed hyperactive swim behavior in PFOS-exposed larvae during photomotor response assays. However, we are the first to report PFOS-induced hyperactivity during the light and dark phases prior to 6 dpf. Hyperactive behavioral changes corresponded well with increased intracellular neuronal calcium concentrations. Of note, increased anxiety-like behaviors were separable from increased swim activity at 3dpf. This suggests that there may be further disruptions in neuronal communication beyond just increased calcium concentrations influencing swim activity. In-depth regional brain activity analyses, possibly through

the integration of developed analytical pipelines (Randlett et al. 2015), could provide further insight into how PFOS induced neural activation is linked to behavioral abnormalities in the larval zebrafish.

We identify increased anxiety-like behaviors as a novel phenotype of PFOS neurotoxicity in the larval zebrafish using an adapted open-field test model (Kalueff and Stewart 2012, Ahmad and Richardson 2013, Richendrfer et al. 2012). Anxiety is an associated symptom of several neurobehavioral disorders linked to PFAS exposures, including autism spectrum disorder and ADHD (Ode et al. 2014, Skogheim et al. 2021, Oh et al. 2021, Shin et al. 2020). PFOS-induced center avoidance has been observed in mice exposed during adulthood (Fuentes et al. 2007), but developmental anxiety has not been previously reported. Of note, previous published research reporting PFOS-induced larval hyperactivity utilized 96-well plates for higher throughput (Jantzen et al. 2016, Gaballah et al. 2020); however, smaller well sizes may not be conducive to conducting anxiety-like behavioral analyses. We therefore conducted behavioral experiments using 24-well plates, which allowed us to define regions within the well to quantify where fish spent their time swimming. Because dysregulation of habenular activity has been associated with increased anxiety, depression, and fear across species, including the zebrafish (Browne, Hammack and Lucki 2018, Mathuru and Jesuthasan 2013, Okamoto et al. 2012), we assessed whether neuronal calcium was affected in the habenula following PFOS exposure. While only the 14 μ M PFOS group showed neuronal dysregulation at the habenula, all groups displayed anxiety like behaviors. This suggests that the habenula may be but one region dictating anxiety-like responses in larval zebrafish, and/or that global neuronal excitation is enough to trigger anxiety responses independent of habenular activity.

Not only does this study provide the first *in vivo* assessment of neuronal calcium activity following PFOS exposure, but it is also the first to evaluate PFOA exposure in this context. The chemical structure of PFOA is very similar to PFOS, differing only by the presence of a carboxylic acid head group rather than sulfonic acid, respectively. At the concentrations and timepoints tested, PFOA exposure did not result in increased larval swim behavior or neuronal network activity. However, others have reported that 5 dpf larvae exposed to 1 μ M PFOA (Rericha et al. 2021) as well as 14 dpf larvae exposed to 2 μ M PFOA (Jantzen et al. 2016) did result in swim hyperactivity. In addition, while neonatal exposure to PFOA in mice has been shown to result in abnormal expression of proteins important for brain growth and neuron function

(Johansson et al. 2009), PFOA exposure of rat primary cortical neurons did not affect spontaneous neuronal activity or burst duration (Tukker et al. 2020). Further analysis is needed to clarify the extent by which PFOA, and other shorter chain length or carboxylic group PFAS congeners, could be neurotoxic or contribute to neural dysregulation. Nevertheless, that PFOA-exposed larvae had normal microglia responses to injury, compared to those exposed to PFOS, highlights the role of neuronal activity on microglia function.

This study provides the first *in vivo* analysis of how developmental PFOS exposure disrupts larval brain health and function. It also highlights the relevance of understanding pollution-induced effects on innate immune cells, in both a non-canonical developmental and homeostatic context, as well as considering the long-term consequences of potential antigen-presentation dysfunction. In summary, the complexity of neural cell communication, especially during the sensitive period of brain development, emphasizes the importance of studying pollutant exposure in non-isolated systems.

MATERIALS & METHODS

Zebrafish husbandry

Zebrafish (*Danio rerio*) maintenance and experimental procedures were approved by the Brown University Institutional Animal Care and Use Committee (IACUC; 19-12-0003) adhering to the National Institute of Health's "Guide for the Care and Use of Laboratory Animals." Zebrafish colonies were maintained in an aquatic housing system (Aquaneering Inc., San Diego, CA) maintaining water temperature ($28.5 \pm 2^\circ\text{C}$), filtration, and purification, automatic pH and conductivity stabilization, and ultraviolet (UV) irradiation disinfection. Adult and larval zebrafish were sustained in a 14:10 hour light-dark cycle (Westerfield 2000).

Adult zebrafish were placed into 1.7 L sloped spawning tanks (Techniplast, USA) 15-18 hours prior to mating. Sexes were separated by a transparent partition. Within 2 hours of light cycle onset, the partition was removed, and zebrafish were allowed to spawn for 1 hour. Embryos were collected in fresh egg water (60 mg/L Instant Ocean Sea Salts; Aquarium Systems, Mentor, OH) and placed into 100 mm non-treated

culture petri dishes (CytoOne, Cat. No. CC7672-3394) until time of toxicant exposure. Embryonic and larval zebrafish were maintained at $28.5 \pm 1^\circ\text{C}$ in an incubator (Powers Scientific Inc., Pipersville, PA) up to 120 hours post-fertilization (hpf).

Zebrafish lines

The following zebrafish lines were used in this study, either independently or in combination: *Tg(mpeg1:EGFP)* (Ellett et al. 2011b); *Tg(UAS:eNpHR3.0-mCherry)* (Arrenberg et al. 2009, Gradinaru et al. 2010); *Tg(elavl3:CaMPARI(W391F+V398L))^{if9}* (Fosque et al. 2015); *Tg(elavl3:Gal4-VP16)* (Kimura, Satou and Higashijima 2008); *Tg(mpeg1:Gal4FF)^{gl25}* (Ellett et al. 2011a); *Tg(HuC:Kaede)* (Sato, Takahoko and Okamoto 2006); *Tg(UAS:nfsB-mCherry)* (Davison et al. 2007); *irf8* mutant (TALE-NT 2.0; st96 allele) (Shiau et al. 2015).

Perfluorooctanesulfoinc Acid (PFOS) and Perfluorooctanoic Acid (PFOA) exposure

Perfluorooctanesulfonic acid (Sigma-Aldrich, Cat. No. A-5040) and Perfluorooctanoic acid (Sigma-Aldrich, Cat. No. 171468) were prepared by dissolving the powdered compounds in 100% DMSO. PFOS stock concentrations were verified using a LC-HRMS (Table 1) (method details provided below). PFOS stock solution was diluted by a factor of 5000x in a mixture 1:1 methanol:water and 2 mM ammonium acetate to accommodate the detection range of the LC-HRMS. LC/MS grade water and methanol were purchased from Honeywell (Muskegon, MI 49442). Ammonium acetate solution (5 M) was purchased from Millipore Sigma (Burlington, MA 01803).

Timed spawns of relevant transgenic zebrafish crosses were performed for 1 hr. Embryos were collected and screened for embryo quality at 4 hpf. Healthy embryos were placed in 24-well plates at a density of 3 embryos per well. Prior to treatment, PFAS compounds were diluted in egg water to the final concentration of interest. Egg water containing 0.1% DMSO was used as vehicle control. Embryos were dosed with 2 mL of diluted PFOS solution or vehicle control at 4 hpf. The 24-well plates were sealed with parafilm to limit

evaporative loss and placed in an incubator ($28.5 \pm 1^\circ\text{C}$). Embryos were dechorionated at 24 hpf and statically exposed until the experimental timepoint of interest.

Brain injury model

At 72 hpf, larval zebrafish were anesthetized in 0.02% tricaine-s solution (Syndel, Ferndale, WA) and restrained dorsal-side-up in 2% agarose (Fisher Scientific, Cat. No. BP160-100). Larvae were injured anteriorly at the right telencephalon with a 9 μm OD pulled glass capillary needle. This method was an adaptation for larval fish (Kishimoto, Shimizu and Sawamoto 2012). For time-lapse imaging, larvae were immediately mounted dorsally on a 35 mm glass bottom microwell dish (MatTek, Part No. P35G-1.5-14-C) in 2% low-melting agarose (Fisher Scientific, Cat. No. BP160-100) surrounded by egg water. Multi-slice projection images of the forebrain and optic tectum at various timepoints post-injury, as well as time-lapse videos composed of 10-minute imaging intervals, were captured using a Zeiss LSM 880 confocal microscope at 20x magnification. Area of microglia response was measured using Zen Blue (Zeiss).

Microglial cell quantification

Adult transgenic zebrafish expressing *Tg(HuC:Kaede)* and *Tg(mpeg1:EGFP)* were crossed to generate *Tg(HuC:Kaede; mpeg1:EGFP)* fish, which had pan-neuronal expression of the photoconvertible protein kaede as well as green macrophages. Embryos were exposed to PFOS as described. At 3 dpf, larvae were fixed in 4% paraformaldehyde (PFA, Sigma-Aldrich, Cat. No. P6148) for 18-24 hours at 4°C . Post-fixation, larvae were washed 3 times in PBS-T (phosphate buffered solution + 0.6% Triton-X 100) (Sigma-Aldrich, Cat. No. X100). PBS-T was removed and VECTASHIELD Mounting Media (Vector Labs, Cat. No. H-1000) was added to the samples. Samples incubated for 5 minutes at room temperature, then were gently mixed and placed in -20°C until imaging. The larvae were removed from VECTASHIELD and mounted in 2% low-melting agarose (Fisher Scientific, Cat. No. BP160-100) in 35mm glass bottom microwell dishes (MatTek, Part No. P35G-1.5-14-C). Prior to image acquisition, the kaede fluorophore was photoconverted from green to red fluorescence using 405 nm light (about 1-minute exposure) on the Zeiss LSM 880 confocal microscope. Microglia were defined as macrophages (*mpeg1*⁺ cells) amongst and in contact with

differentiated neurons (HuC+ cells) in the forebrain, optic tectum, and hindbrain. The entirety of the zebrafish brains were imaged at 20x magnification (2 tile panels with 10% overlap). Microglia were counted by panning through confocal z-stacks using FIJI/ImageJ. Immune cells in contact with HuC+ differentiated neurons were counted as a parenchymal cells (i.e. microglia). Surface cells in contact with differentiated neurons that failed to extend into the parenchyma were defined as non-parenchymal cells (i.e. macrophage). Location and proximity of mpeg1+ cells were confirmed using z-stack orthogonal views. Each treatment group was compared to controls (0.1 % DMSO) using a One-way ANOVA in GraphPad Prism (Dotmatics, San Diego, CA).

Microglial morphology quantification following PFOS exposure

Microglial morphology was performed on high-resolution confocal micrographs of the optic tectum in fixed transgenic zebrafish larvae expressing either *Tg(mpeg1:EGFP)*, *Tg(mpeg1:Gal4FF;UAS:eNpHR3.0-mCherry)*, or *Tg(mpeg1:Gal4FF;UAS:nfsb-mCherry)*. Individual microglia were isolated based on their specific z-stack range, to prevent analysis of overlapping cells, using FIJI/ImageJ. Z-slices were also processed to rescale intensity (compensating for loss of intensity in deeper tissue) and modestly smoothed using Gaussian blur with a sigma radius of 1 to better define cellular extensions and reduce signal noise. Each cell was individually outlined, and morphological parameters measured with CellProfiler software (version 3.1.9) or FIJI/ImageJ. Area, perimeter, and cell shape (perimeter-to-area ratio) were quantified. Unpaired t-tests with Welch's correction were used to evaluate statistical significance using GraphPad Prism (Dotmatics, San Diego, CA).

Quantitative RT-PCR

Larval zebrafish heads were collected at 3 dpf. To collect, larvae were placed on ice for 4 minutes to immobilize swim activity but prevent pain or distress during collection (Wallace et al. 2018). The heads were removed by cutting at the base of the hindbrain at a 45° angle, limiting collection of the heart and pericardium. Heads were pooled (n = 10) and flash frozen in liquid nitrogen. RNA isolation and purification was carried out using the RNeasy Plus Kit (QIAGEN). cDNA synthesis was achieved using the SuperScript

IV Reverse Transcriptase First-Strand Synthesis System kit (Invitrogen, Cat. No. 18091050). qRT-PCR for genes of interest was performed with 7.5 ng/uL of cDNA using the ViiA7 Real Time PCR System (Applied Biosystems). Gene targets were detected by using either pre-designed TaqMan probes (Thermo Fisher Scientific) or custom primers with Power Track SYBR Green Master Mix (Thermo Fisher, Cat. No. A46012). See list of probes and primers for qRT-PCR and genotyping below.

Gene	TaqMan ID or Primer Sequences
<i>il6 (m17)</i>	Dr03098117_g1
<i>il1b</i>	Dr03114368_m1
<i>tnfa</i>	Dr03126850_m1
<i>actb1 (b-actin)</i>	Dr03432610_m1
<i>pr2y12</i>	F: 5'-CTTCAGGTCGTCGCTGTTTA-3' R: 5'-AGTGCGTTTCCCTGTTGAT-3'
<i>b-actin</i>	F: 5'-CGAGCAGGAGATGGGAACC-3' R: 5'-CAACGGAACGCTCATTGC-3'

Genotyping *irf8* mutants

irf8^{st96/st96} were generated by Shiau et al. via TALEN-targeting (Shiau et al. 2015). After swim behavior testing or CaMPARI imaging, whole larvae were placed in DNA lysis buffer to be used for genotyping. The *irf8* exon 1 fragment was amplified using the following primers: 5'-ACATAAGGCGTAGAGATTGGACG-3' and 5'-GGATGAGGACCGCACTATGTTTC-3'. As this *irf8* mutant has a frameshift mutation at an Aval site, the PCR product was used for restriction digest with Aval (New England BioLabs, Cat. No. R0152L) to identify the presence of the mutation.

Acridine orange stain

Cell death was determined using the vital dye acridine orange (AO) in live zebrafish embryos. AO is a cell permeable stain that selectively intercalates uncoiled nucleic acid present in apoptotic cells (Delic et al. 1991). Zebrafish embryos were injured at 72 hpf and placed in 5 ug/mL acridine orange (ThermoFisher, Cat. No. A1301) diluted in egg water for 20 minutes. Zebrafish were then washed in egg water for 15 minutes, refreshing the solution every 5 minutes. Live zebrafish were immediately imaged on a Zeiss LSM 880 confocal microscope. For 1 hour-post injury (hpi) analysis, staining took place from 40 minutes to 1 hpi. For 4 hpi analysis, staining took place from 3 hours and 40 minutes to 4 hpi.

CaMPARI Photoconversion and Image Acquisition

Tg[elavl3:CaMPARI(W391F+V398L)]jf9 embryos were screened and the brightest (highest expressing) were selected for toxicant exposure and subsequent photoconversion. Individual embryos were placed into a modified 1-well dish (15 mm diameter) containing PFOS or 0.1% DMSO. The dish was placed onto a constructed pedestal within a DanioVision Observation Chamber (Noldus, Wageningen, The Netherlands) adapted with optogenetics components (Prizmatix, Southland, MI). Use of the pedestal decreased working distance from the LED light source to the free-swimming larva, concordantly increasing light intensity. Cells expressing CaMPARI with high calcium content will photoconvert (green-to-red) in the presence of 405 nm light. Photoconversion was performed by exposing zebrafish to 405 nm (135 mW/cm²) wavelength light for 1 minute. Live zebrafish larvae were then anesthetized in 0.02% tricaine-s (MS-222) and mounted in 2% low-melting agarose in 35mm glass bottom microwell dishes. Confocal z-stacks were acquired on a Zeiss LSM 880 confocal microscope and maximum intensity projections generated in Zen Black (Zeiss) representing a global snapshot of neural activity. Image parameters were set during acquisition of the first image and maintained for the duration of each experiment.

CaMPARI analyses (R/G ratio calculations)

CaMPARI photoconversion from green-to-red during exposure to 405 nm wavelength light was used to determine neuronal calcium levels following PFOS exposure. To assess the level of photoconversion, maximum intensity projections were imported into FIJI/ImageJ. Brain regions of interest including forebrain (FB), habenula (H), optic tectum (OT), cerebellum (Ce), hindbrain (HB), and whole brain (WB) were manually selected and integrated density of red and green was measured. Measurements were blank corrected by selecting 3 separate background regions of each image (Equation 1). The ratios of corrected integrated density were then used to determine R/G ratio for each respective image (Equation 2). R/G ratios were then normalized to the average control R/G ratio within a particular experiment (Equation 3). Unpaired t-tests with Welch's correction were performed on the normalized R/G ratios to determine difference between PFOS treated and control groups.

Equation 1: Corrected Integrated Density

b = background integrated density

d = integrated density of region of interest

Corrected Integrated Density (c) = $d - (\Sigma(b1 + b2 + b3)) \div 3$

Equation 2: R/G Ratio

R/G Ratio (r) = $C_{Red} \div C_{Green}$

Equation 3: Normalized R/G Ratio

\bar{X} = mean of control R/G ratios (for each experimental day)

Normalized R/G Ratio = $r \div \bar{X}$

Pentylentetrazol (PTZ) Exposure

For validation of CaMPARI functionality, 3 dpf *Tg(elavl3:CaMPARI(W391F+V398L))^{if9}* larvae were exposed to 10 mM pentylentetrazol (PTZ) for 10 minutes. PTZ was washed out 3 times with egg water immediately followed by CaMPARI photoconversion, as described. For assessment of microglia response to injury following PTZ exposure, larvae were exposed to control solution, 28 μ M PFOS, or 5 mM PTZ at 72 hpf. Larvae were injured in the right telencephalon, as described, and placed back in their original dosing solutions for 4 hours. At 4 hpi, multi-slice projection images of the forebrain and optic tectum were captured using a Zeiss LSM 880 confocal microscope at 20x magnification. Area of microglia response was measured using Zen Blue (Zeiss). Statistics were performed using unpaired t-tests with Welch's correction using GraphPad Prism (Dotmatics, San Diego, CA).

Photomotor Behavioral Response

Behavioral assessments were performed in 24-well plates (ThermoFisher, Cat. No. 144530) using a DanioVision Observation Chamber with EthoVision XT live-tracking software (Noldus, Wageningen, The Netherlands). Larval zebrafish were transferred into individual wells of the same dosing solution

approximately 18-24 hours prior to behavioral assessment. All photomotor response assays were performed between 8 AM and 12 PM to prevent behavioral changes attributable to circadian differences. Briefly, zebrafish were placed into the DanioVision observation chamber and acclimated during a 15-minute dark cycle, followed by a 5-minute light cycle (Light 1), a 5-minute dark cycle (Dark 1), another 5-minute light cycle (Light 2), and completed with a 15-minute dark cycle (Dark 2). Behavioral experiments lasted a total of 45 minutes.

Total distance moved (mm) during light/dark cycles was quantified using the EthoVision XT software as a measure for hyperactivity. Anxiety following PFOS exposure was assessed by monitoring larval well location (center vs. edge) during experimentation. Defined center and edge regions each constituted 50% of total well area. Edge versus center preference was quantified using EthoVision XT. Anxiety data was normalized relative to controls for each experimental replicate. Statistical analyses were performed using GraphPad Prism (Dotmatics, San Diego, CA).

Optogenetic Manipulation in Noldus Behavioral Unit

To stimulate the optogenetic channel halorhodopsin in the transgenic lines *Tg(elavl3:Gal4;UAS:eNpHR3.0-mCherry)* and *Tg(mpeg1:Gal4ff;UAS:eNpHR3.0-mCherry)*, screened larvae were placed a DanioVision observation chamber (Noldus, Wageningen, The Netherlands) outfitted with a 570 nm wavelength laser (Prizmatix, Southland, MI). Halorhodopsin⁺ neurons or microglia were stimulated with 570 nm light for 4 hours then immediately imaged for cell morphology or injury response.

Targeted Analysis of PFOS Using Liquid Chromatography High Resolution Mass Spectrometry

Chemicals: Certified PFOS and isotope labelled reference standards were purchased from Waters (Milford, MA 01757). LC/MS grade water and methanol were purchased from Honeywell (Muskegon, MI 49442). Ammonium acetate solution (5 M) was purchased from Millipore Sigma (Burlington, MA 01803).
Sample Extraction: Larvae were stored in -80°C freezer until extracted and defrosted at 20°C. 1 mL methanol was added to the centrifuge tube containing the embryos. The samples were sonicated for 90

minutes, vortex mixed 1 minute, and allowed to reach equilibrium for 3 hours at 20°C. Samples were then centrifuged at 3,000 rpm for 10 minutes. The following was added to an LC analysis vial: 50 µL of the methanol extract, 10 µL labelled PFOS internal standard, and 440 µL of a mixture containing 1:1 methanol:water and 2 mM ammonium acetate.

Targeted analysis of PFOS was performed using a Thermo Liquid Chromatography (LC) Orbitrap Q Exactive HF-X mass spectrometer (MS) equipped with a Thermo Vanquish UHPLC system. Mobile phase A contained 2 mM ammonium acetate in water and mobile phase B contained 2 mM ammonium acetate in methanol. 20 µL of each sample extract was injected in triplicate and separated in a Thermo Hypersil Gold Vanquish C18 column (50 mm X 2.1 mm x 1.9 µm) at a constant temperature of 60°C. PFOS was eluted from the column at a constant flow rate of 0.4 mL/min using a mobile phase gradient as follows: equilibration with 10% B for 1 minute, followed by a gradient ramp from 10% B to 95% B over 4 minutes and held for 2 minutes, and back to 10 % B over 1 minute and held for 2 minutes (total run time 9 minutes, data collected from 0.6 to 8.5 minutes). The MS was operated in full scan dd-MS² mode (70 NCE) with an inclusion list for PFOS. Ionization was performed in negative mode with an ionization window of 1.0 m/z, sheath gas flow rate of 40, auxiliary gas flow rate of 10, sweep gas flow rate of 2, spray voltage of 2.7 kV, 310°C capillary temperature, funnel RF level of 35, and 320°C auxiliary gas heater temperature. Ions were further fragmented in the HCD collision cell filled with N₂ (produced by a Peak Scientific Nitrogen Generator, Genius NM32LA). For the full-scan, the Orbitrap was operated with a resolution of 120,000, automatic gain control (AGC) of 3e6, and maximum dwell time of 100ms. For dd-MS², the Orbitrap was operated with a resolution of 15,000, AGC of 2e5, and maximum dwell time of 400ms.

Four ions were monitored for PFOS, including 498.9302 m/z (quantifying), 79.9573 m/z (confirming), 98.9556 m/z (confirming), and 82.9607 m/z (confirming). The retention time of PFOS was 5.18 min. Quantification was performed in TraceFinder 5.0 General with external seven-point calibration curve prepared by serial dilutions of the calibration standard. Limits of detection (LOD) were determined by injecting a calibration 7 standard seven times and using Equation 4:

Equation 4: Limits of Detection (LOD) Calculation

$$\text{LOD}=(3*s)/m$$

where s is the sample standard deviation and m is the calibration curve slope. The resulting PFOS LOD was 33.87 ppt.

Untargeted Metabolome Wide Association Study

High resolution metabolomics: The sample extracts were re-analyzed using LC-HRMS to collect untargeted metabolomics data. A 10 µL volume was injected in triplicate onto on the Thermo LC-Orbitrap system described above. Two chromatography separation methods were used, normal and reverse-phase. The normal-phase LC was performed with a HILIC column (Thermo Synchronis HILIC 50 mm X 2.1 mm x 3 µm) at a constant temperature of 25°C. Mobile phase A contained 2 mM ammonium acetate in acetonitrile and mobile phase B contained 2 mM aqueous ammonium acetate. Metabolites were eluted from the column at a constant flow rate of 0.2 mL/minute using a solvent gradient as follows: equilibrate with 10% B for 1 minute, increase to 65% B for 9 minutes and hold for 3 minutes, decrease to 10% over 1 minute and hold for 1 minute. The reverse-phase LC was performed with a C18 column (Thermo Hypersil Gold Vanquish, 50 mm X 2.1 mm x 1.9 µm) at a constant temperature of 60°C. Mobile phase A contained 2 mM aqueous ammonium acetate and mobile phase B contained 2 mM ammonium acetate in acetonitrile. Metabolites were eluted from the column at a constant flow rate of 0.5 mL/minute using a mobile phase gradient as follows: equilibration with 2.5% B for 1 minute, increase to 100% B over 11 minutes and held for 2 minutes, and back to 2.5% B over 1 minute and held for 1.5 minutes (total run time 16.5 minutes, data were collected from 0.05 to 12.5 minutes). For both normal and reverse-phase LC, the MS was operated in full scan mode with 120,000 resolution, automatic gain control of 3×10^6 , and maximum dwell time of 100 ms. Electrospray ionization was conducted in positive mode for normal-phase and negative mode for reverse phase LC. Ionization was performed at a sheath gas flow of 40 units, auxiliary gas flow of 10 units, sweep gas flow of 2 units, spray voltage of 3.5 kV, 310°C capillary temperature, funnel radio frequency (RF) level of 35, and 320°C auxiliary gas heater temperature.

Metabolomics data analysis: Data files were converted from *.raw files to *.cdf files using XCalibur file Converter, and then processed in R packages apLCMS (Yu et al. 2009) and xMSanalyzer (Uppal et al. 2013) to produce m/z feature tables. Feature intensities were normalized by log2 transformation. Association of metabolite feature with PFOS exposure was assessed using a t-test ($P < 0.05$) in comparison to the control samples. The P -values were adjusted using Benjamini and Hochberg with a false discovery rate (FDR) $\leq 20\%$ to control for Type I errors in multiple comparisons. Significant metabolites were analyzed for pathway enrichment using MetaboAnalystR (Chong and Xia 2018) using the zebrafish mummichog curated model, which includes the KEGG, BiGG, and Edinburgh maps. All metabolomics data analysis was performed in R (version 4.0.2).

Statistical analyses and reproducibility

As noted in the Methods, control, PFOS, and PFOA stock solutions were validated. Each experiment was carried out in at least three independent experimental replicates. An experimental replicate was considered a cohort of zebrafish that were spawned on separate days and, when applicable, dosed with separate freshly prepared dosing solutions. When applicable, dosing groups for each experimental replicate were composed of siblings, such that sibling controls could be compared to dosed siblings. Statistical analyses for each figure are listed in the accompanying figure legends. When comparing two groups, unpaired t-tests with Welch's correction were performed, which is used when two samples have unequal variance and sample size and therefore does not assume equal standard deviations. When comparing three or more groups, One-Way ANOVA was performed. All statistics were performed using GraphPad Prism (Dotmatics, San Diego, CA).

Data availability

All data is available in the main text and supplementary materials. All targeted and untargeted metabolomics files will be deposited into Metabolomics Workbench.

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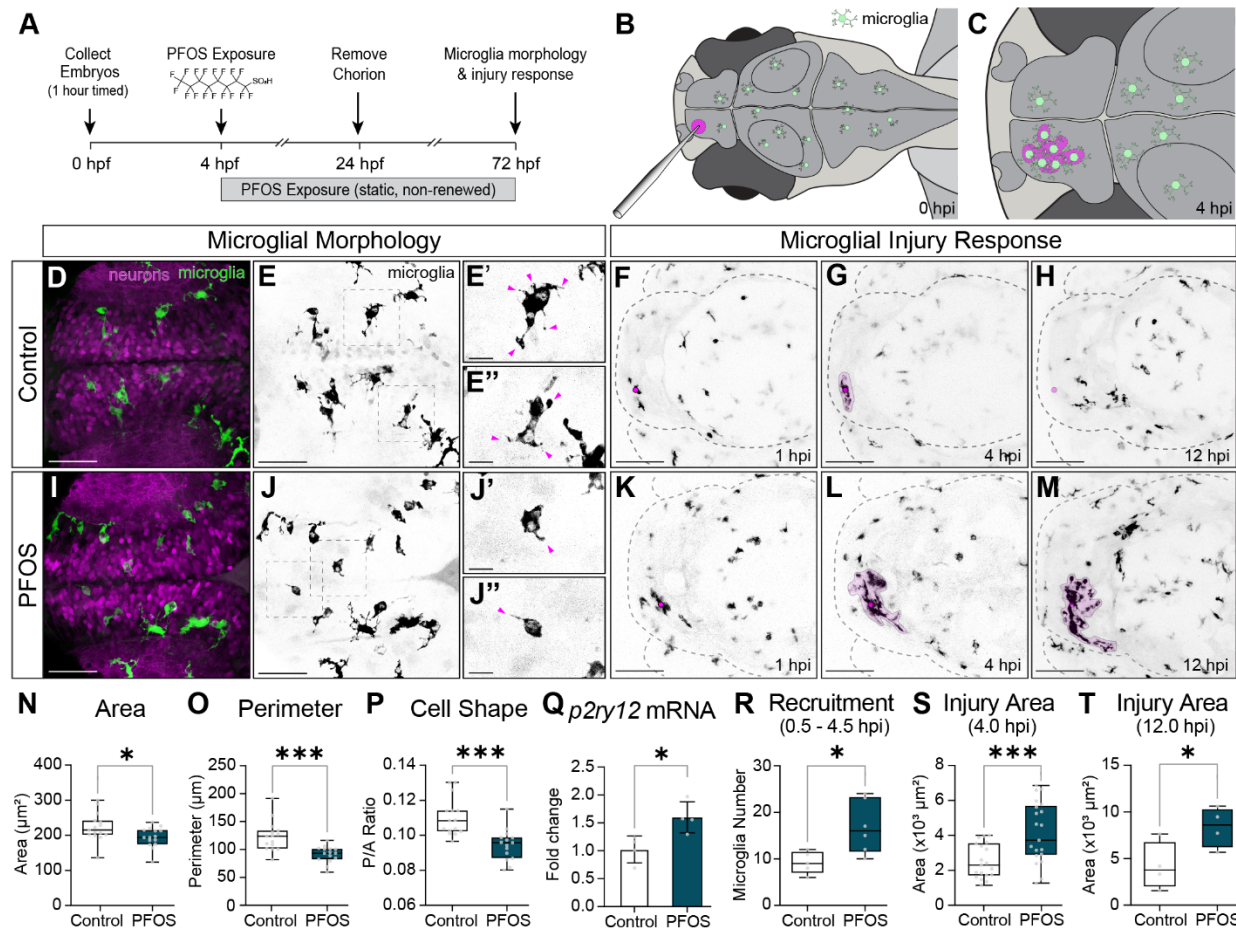


Figure 1. Microglia become activated and hyperresponsive to minor brain injury following developmental exposure to PFOS. (A) Exposure paradigm: zebrafish embryos from *Tg(mpeg1:EGFP)* adults were collected after a 1-hour timed spawn. At 4 hpf, embryos were dosed with 0.1% DMSO (Control) or 28 μ M PFOS. Treatment solutions were static and not renewed. Embryos were dechorionated at 24 hpf, and imaging and injury experiments were conducted at 72 hpf. (B) Schematic of larval brain injury: a pulled glass needle (OD 9 μ m) was used to puncture the right telencephalon of larval zebrafish. (C) Hypothetical schematic of microglia response 4 hpi. (D-E) Homeostatic, non-activated microglia of 3 dpf control larvae have several projections emanating from their cell bodies. (F-H) Control microglia respond to brain injury, with visible accumulation at the injury site at 4 hpi that mostly clears by 12 hpi. (I-J) Microglia of 3 dpf larvae exposed to 28 μ M PFOS become phenotypically activated, with rounder cell bodies and reduced projections. (K-M) In PFOS-exposed larvae, microglia robustly respond to brain injury by 4 hpi and are still maintained at the site at 12 hpi. Quantifications of the images reveal that PFOS-exposed microglia (N) area ($P = 0.0342$) and (O) cell perimeter ($P = 0.0006$) are significantly reduced. (P) The perimeter-to-area ratio ($P = 0.0002$) is also significantly decreased, indicative of a rounder cell morphology. $n = 15$ fish per group (3-18 cells counted per fish). (Q) qRT-PCR for the microglia activation gene, *p2ry12*, was significantly upregulated in isolated heads of PFOS-exposed larvae ($P = 0.0205$; $n = 10$ pooled heads per sample). PFOS exposure resulted in a significant increase in (R) microglia recruitment in the first 4.5 hpi ($P = 0.0216$; $n = 5-6$ per group), as well as increased response area at (S) 4 hpi ($P = 0.0006$; $n = 19-20$ per group) and (T) 12 hpi ($P = 0.0467$; $n = 4$ per group). Confocal micrographs at 40x magnification (D-E' & I-J') or 20x magnification (F-H; K-M). Unpaired t-test with Welch's correction.

Table 1. PFAS body burden in zebrafish larvae.

PFAS Body Burden (ng/embryo)								
Chemical	Formula	MW (g/mol)	Control		28 μ M PFOS		56 μ M PFOS	
			48 hpf	72 hpf	48 hpf	72 hpf	48 hpf	72 hpf
PFOS	C ₈ HF ₁₇ O ₃ S	500.13	0.12 \pm 0.17	0.12 \pm 0.16	32.22 \pm 2.57	70.46 \pm 2.72****	46.04 \pm 3.75	109.15 \pm 5.99****
PFNS	C ₉ HF ₁₉ O ₃ S	550.14	0.01 \pm 0.02	0.00 \pm 0.00	0.19 \pm 0.05	0.39 \pm 0.07****	0.23 \pm 0.04	0.69 \pm 0.20***
PFHpS	C ₇ HF ₁₅ O ₃ S	450.12	0.00 \pm 0.00	0.00 \pm 0.00	0.35 \pm 0.17	0.45 \pm 0.13	0.40 \pm 0.17	0.55 \pm 0.11*
PFHxS	C ₈ HF ₁₃ O ₃ S	400.12	0.006 \pm 0.012	0.008 \pm 0.014	0.008 \pm 0.014	0.007 \pm 0.011	0.008 \pm 0.012	0.008 \pm 0.011

*p<0.05; ***p<0.001; ****p<0.0001

Table 2. Survival rate of zebrafish larvae exposed to 14 μ M, 28 μ M, or 56 μ M PFOS over time.

Survival Rate Percentage				
Age	PFOS Concentration			
	Control	14 μ M	28 μ M	56 μ M
24 hpf	97.93 \pm 2.94	97.67 \pm 1.75	98.15 \pm 1.73	97.22 \pm 2.84
48 hpf	97.74 \pm 2.76	97.14 \pm 1.21	97.09 \pm 2.10	92.49 \pm 2.84
72 hpf	97.43 \pm 3.25	96.18 \pm 1.81	95.62 \pm 2.43	81.54 \pm 13.23
96 hpf	96.41 \pm 5.82	95.72 \pm 1.20	50.65 \pm 10.06*	25.00 \pm 9.89**
120 hpf	95.93 \pm 5.85	89.14 \pm 3.65	7.73 \pm 3.03****	5.33 \pm 2.67****

*p<0.05; **p<0.01; ****p<0.0001

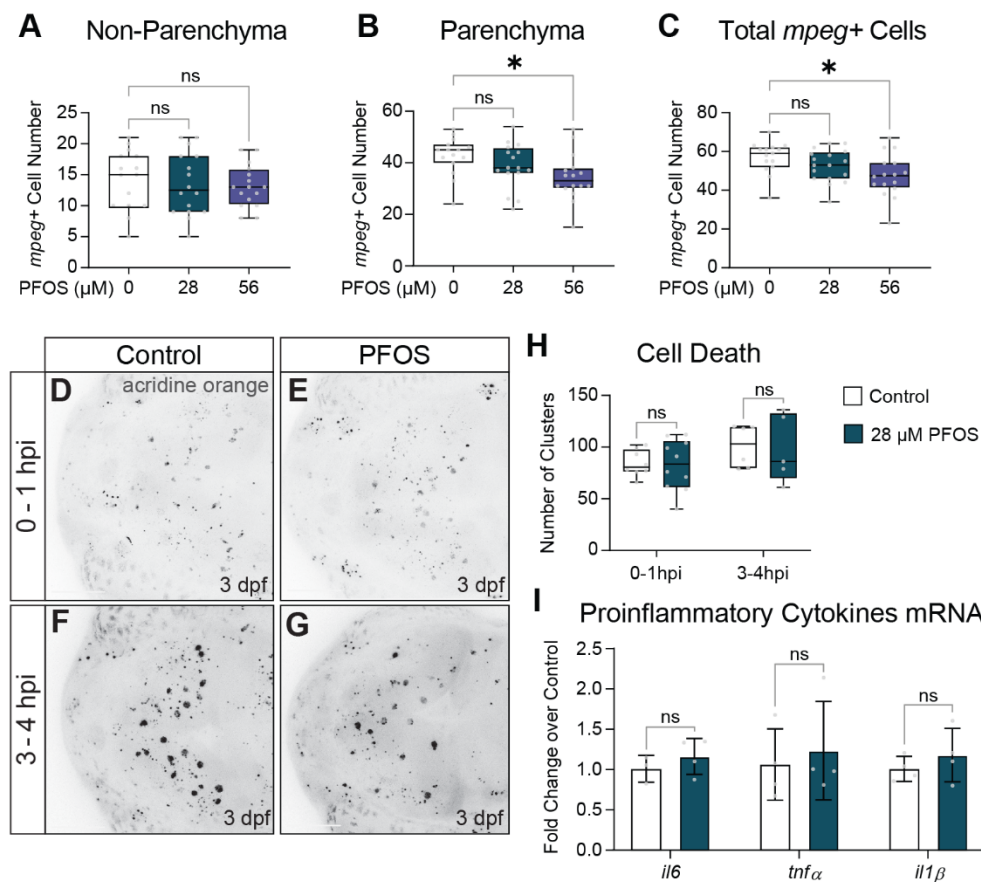
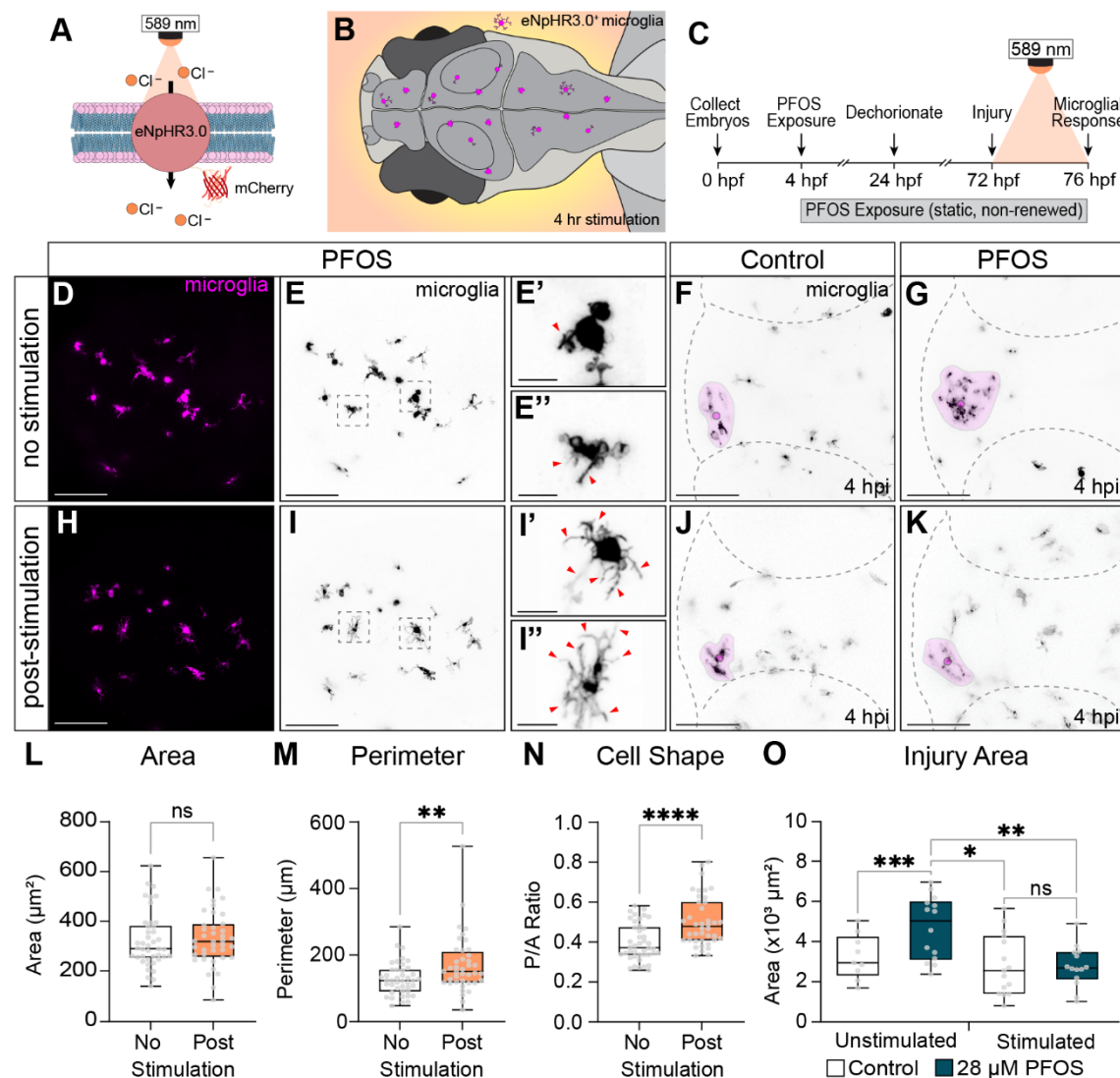


Figure 1-Supplement 1. Microglia hyperresponsiveness is not attributed to changes in microglia number, cell death, or brain inflammation. (A) Non-parenchymal, (B) parenchymal, and (C) total microglia on were quantified in 3 dpf control (n = 15), 28 μ M PFOS (n = 16), or 56 μ M PFOS (n = 16) exposed larvae. There was no significant difference in parenchymal or non-parenchymal microglia number in 28 μ M PFOS-exposed larvae compared to controls, while 56 μ M PFOS larvae had significantly fewer parenchymal ($P = 0.0084$) and total microglia ($P = 0.0105$). (D,E) To assess cell death, live 3 dpf larvae were stained with 5 μ g/mL acridine orange during the first hour post injury or (F,G) between 3-4 hpi. (H) Quantification of acridine orange-positive clusters in 3 dpf control and 28 μ M PFOS-exposed larvae show no significant change in cell death (n = 5-9 per group). (I) qRT-PCR for the inflammatory genes *il6*, *tnf α* , and *il1 β* were not significantly changed in isolated heads of 3 dpf control or 28 μ M PFOS-exposed larvae (n = 10 pooled heads per sample). Confocal micrographs at 20x magnification. Unpaired t-tests with Welch's correction. * $P < 0.05$.



824

825 **Figure 2. PFOS-induced microglia activation can be reverted by electrical modulation.** (A) Schematic
826 of halorhodopsin: Optogenetic modulation of microglia electrical state is achieved via photo-stimulation of
827 the light-gated chloride pump, halorhodopsin (eNpHR3.0). eNpHR3.0 is most responsive to 589 nm
828 wavelength light. (B) eNpHR3.0 was driven under a pan-macrophage promoter
829 (*Tg(mpeg1:GalFF;UAS:eNpHR3.0-mCherry)*) to achieve optogenetic control of microglia in zebrafish
830 larvae. (C) Experimental paradigm: At 72 hpf, injured or uninjured zebrafish were stimulated for 4 hours
831 with 589 nm light in the enclosed Noldus DanioVision Behavior Unit. (D-E'') Unstimulated halorhodopsin⁺
832 microglia of 3 dpf 28 μM PFOS-exposed larvae were rounded with few projections. As shown in Figure 1,
833 (F) unstimulated control microglia were responsive to minor injury at 4 hpi, though (G) unstimulated PFOS-
834 exposed microglia had a significantly heightened response. (H-I'') Following 4-hour stimulation of
835 halorhodopsin⁺ in PFOS-exposed larvae, microglia became more ramified. (J, K) Stimulation of
836 halorhodopsin⁺ microglia following PFOS exposure also normalized the microglia response to injury. (L)
837 Stimulated microglia area was unchanged, but (M) they had significantly increased cell perimeter ($P =$
838 0.0079) and (N) an increased perimeter-to-area ratio ($P < 0.0001$), indicative of a more ramified cell shape
839 ($n = 40-42$ cells per group from 3 independent experiments). Unpaired t-tests with Welch's correction. (O)
840 The injury response area was significantly increased between unstimulated control and 28 μM PFOS

exposed larvae ($P = 0.0177$); however, there was no significant difference in response area between 4-hour stimulated control and PFOS-exposed larvae. Additionally, the stimulated PFOS-exposed area was significantly decreased compared to the unstimulated PFOS group ($P = 0.0008$; $n = 9-15$ per group). One-way ANOVA.

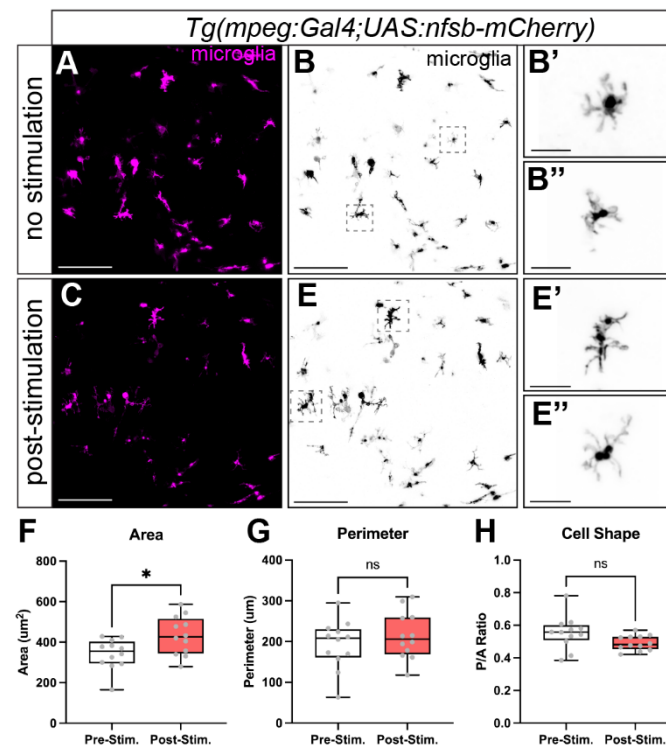


Figure 2-Supplement 1. Light stimulation alone does not impact microglia morphology. To ensure that 589 nm light alone does not alter microglia morphology, PFOS-exposed transgenic larvae expressing macrophage-driven mCherry (*Tg(mpeg1:gal4FF;UAS:nfsb-mCherry)*) were either (A-B'') unstimulated or (C-E'') subjected to the 589 nm light for 4 hours. (F) While microglia area was increased following light exposure (* $P < 0.05$), (G) the light stimulation did not result in significant changes in perimeter or (H) the perimeter-to-area ratio. Therefore, light stimulation alone does not result in the ramification of microglia. $n = 12$ cells. Unpaired t-test with Welch's correction.

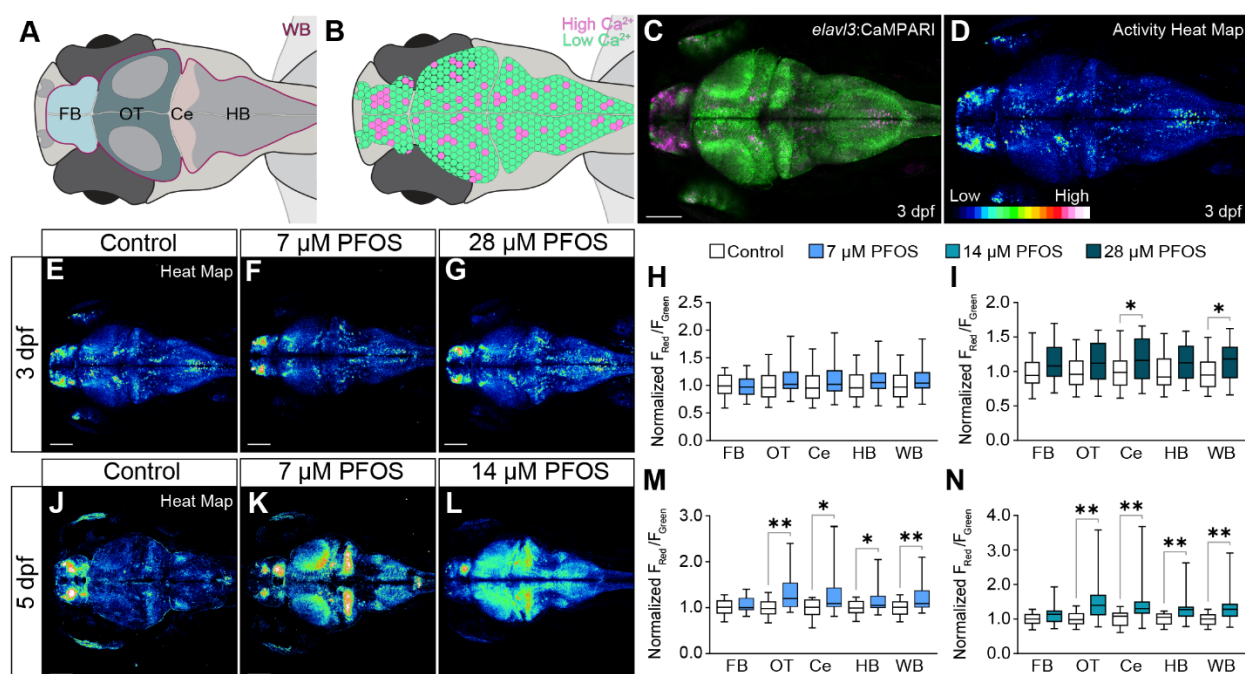


Figure 3. Developmental PFOS exposure alters global and regional neuronal network activity. (A) Illustrative representation of a larval zebrafish brain with anatomical regions outlined: forebrain (FB), optic tectum (OT), cerebellum (Ce), hindbrain (HB), and whole brain (WB). (B) Illustrative representation of neuron-driven CaMPARI: neurons with low intracellular calcium (low Ca^{2+}) remain green following 1 minute exposure to 405 nm light, while neurons with high intracellular calcium (high Ca^{2+}) are photoconverted to red. (C) Confocal micrograph of a 3 dpf larvae expressing neuron-specific CaMPARI (*Tg(elav1:CaMPARI)*) following 1 minute photoconversion. (D) Generated high intensity LUTs heat map of the red, photoconverted channel in C depicting high activity neurons. Low to high Intracellular calcium is depicted by a blue-red-white spectrum. (E) Micrographs of active neurons at 3 dpf in control, (F) 7 μM PFOS, and (G) 28 μM PFOS larvae following 1-minute photoconversion. Neuronal activity can be quantified by determining the ratio of fluorescent intensity in the red versus green channels ($F_{\text{Red}}/F_{\text{Green}}$). (H) 7 μM PFOS does not result in a regional or global (WB) change in neuron activity in 3 dpf larvae. However, (I) 28 μM PFOS-exposed 3 dpf larvae demonstrate a modest increase in regional brain activity, and a significant increase in the Ce ($P = 0.0483$) and globally (WB; $P = 0.0429$). (J) Brain activity was also determined in 5 dpf larvae exposed to control, (K) 7 μM PFOS, or (L) 14 μM PFOS. At 5 dpf (M) 7 μM PFOS-exposed larvae have significant increases in brain activity in the OT ($P = 0.0016$), Ce ($P = 0.0221$), HB ($P = 0.0294$), and globally ($P = 0.0075$). (N) Larvae exposed to 14 μM PFOS also had increases in the OT ($P = 0.0025$), Ce ($P = 0.0050$), HB ($P = 0.0038$), and globally ($P = 0.0036$). Confocal micrographs at 10x magnification. $n = 21$ -23 fish per group. Unpaired t-test with Welch's correction.

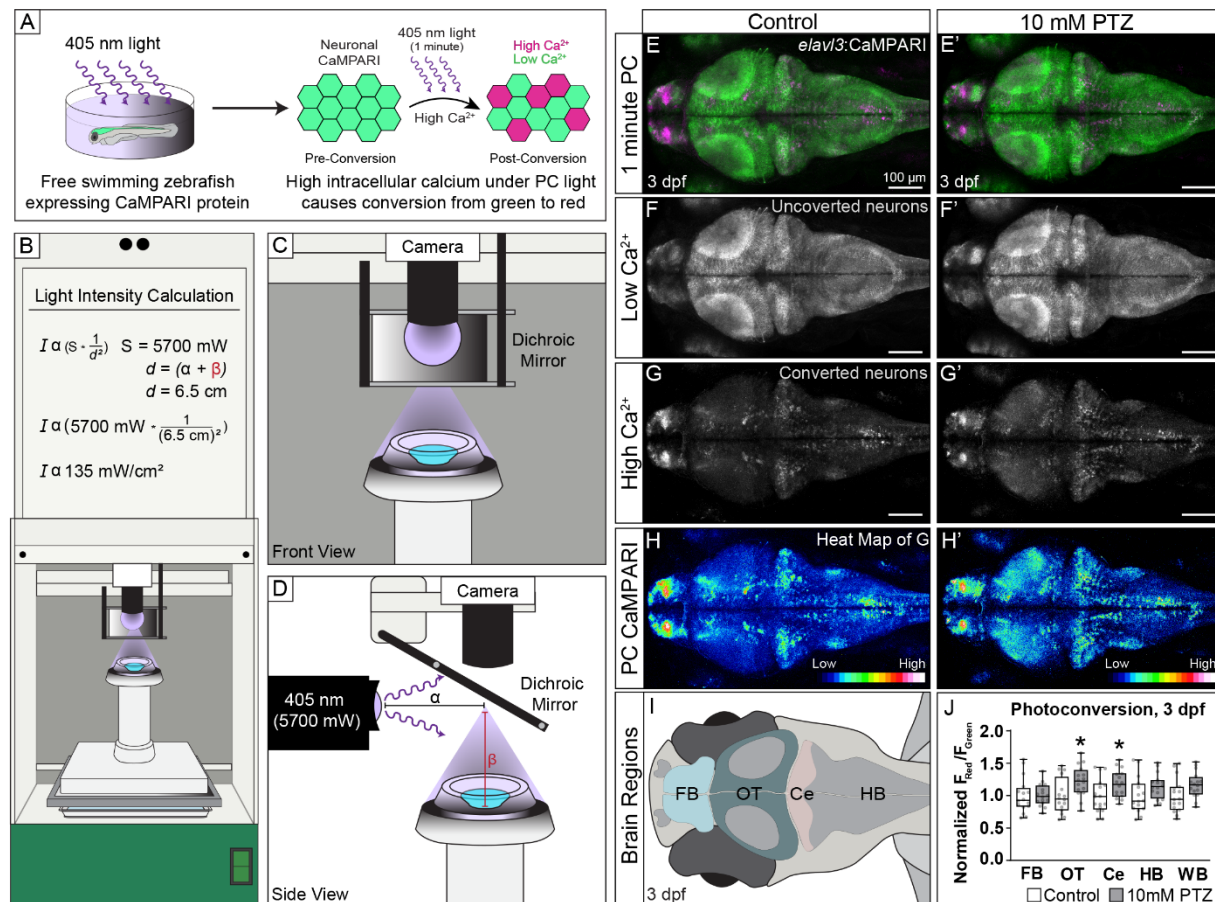
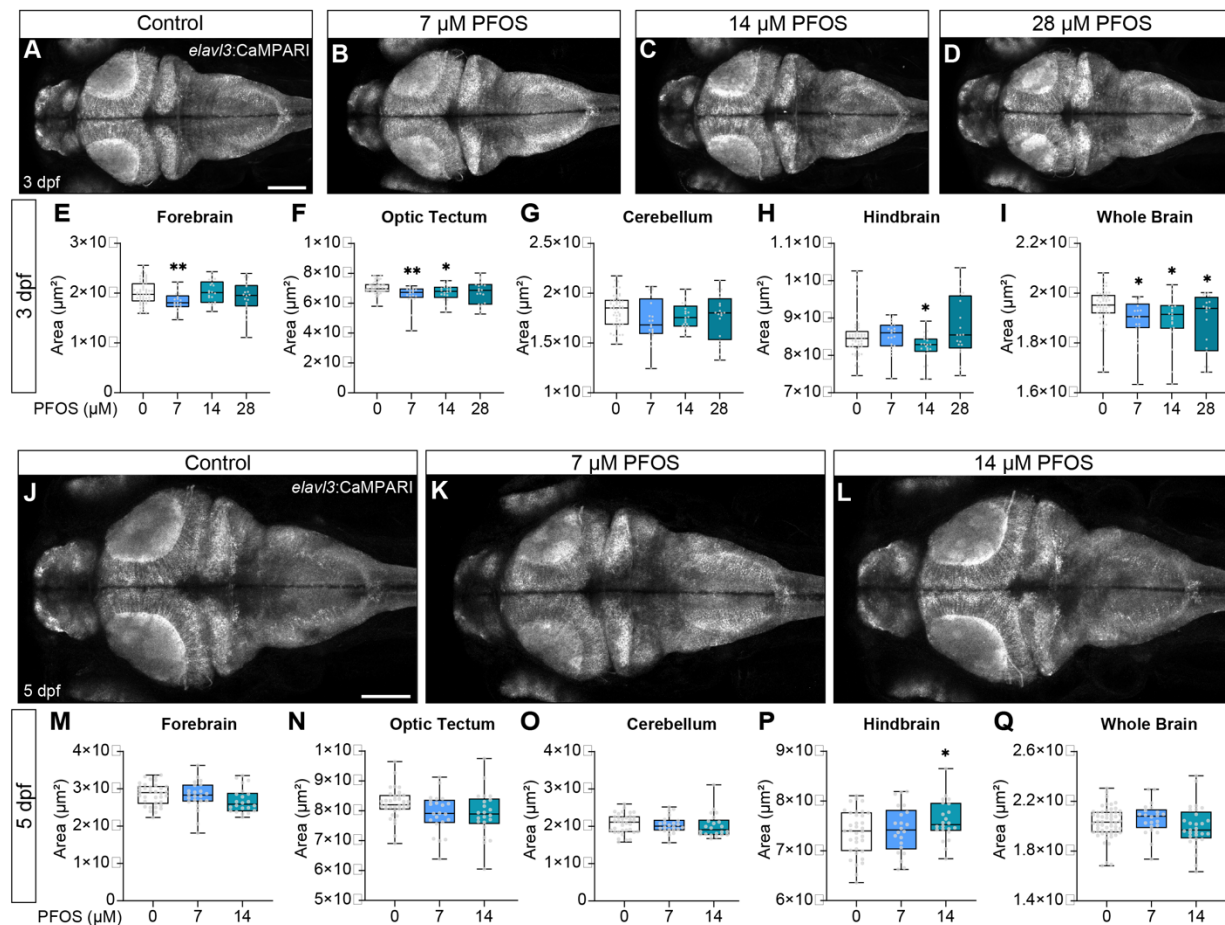


Figure 3- Supplement 1. Functional neuroimaging to understand the effects of toxicant exposure on neuronal activity. (A) Schematic overview of CaMPARI photoconversion: free-swimming zebrafish larvae with neuron-specific expression of the genetically encoded calcium indicator CaMPARI protein (*Tg(elav13:CaMPARI)*) are subjected to 405 nm light for 1 minute. Exposure to blue light photoconverts any neurons with high intracellular calcium from green to red. (B-D) We modified our Noldus DanioVision Behavioral unit, outfitted with LEDs for optogenetic manipulation, to efficiently photoconvert CaMPARI in free-swimming zebrafish. (B) A 3D printed pedestal is used to reduce the working distance between the light source and zebrafish larvae. Applying the inverse-square law, the light intensity at the apex of the pedestal was calculated to be 135 mW/cm². (C) Illustration of the apex of the pedestal containing a single well dish (OD 15 mm). (D) The 405 nm light has an intensity of 5700 mW at the source. The light travels horizontally then is reflected off a dichroic mirror within the unit. The working distance was calculated as the distance from the light source to the dichroic mirror (α) plus the distance from the dichroic mirror to the apex of the pedestal (β). (E) Confocal micrographs of neuronal calcium following 1-minute photoconversion of CaMPARI in 3 dpf larvae treated with egg water or (E') after 10 minutes in 10 mM pentylenetetrazol (PTZ), a GABA_A-inhibitor known to induce hyperactivity. (F, F') Green, non-active neurons with low intracellular calcium from E & E'. (G, G') Photoconverted, active neurons with high intracellular calcium, pseudo-labeled as magenta, from E & E'. (H, H') An intensity LUTs applied to the active neurons in G & G' spectrally maps the regional increases of neuronal activity on a scale from less active (blue) to more active (red-white). (I&J) Regional and global neuronal activity can be quantified by determining the ratio of fluorescent intensity in the red versus green channel ($F_{\text{Red}}/F_{\text{Green}}$). Exposure to 10 mM PTZ significantly increased brain activity in the optic tectum (OT; $P = 0.0200$), cerebellum (Ce; $P = 0.0386$), and nearly in the

whole brain (WB; $P = 0.0501$), but not in the forebrain (FB) or hindbrain (HB). Red/Green ratios were normalized relative to vehicle control. $n = 16$ fish per group. Unpaired t-test with Welch's correction.



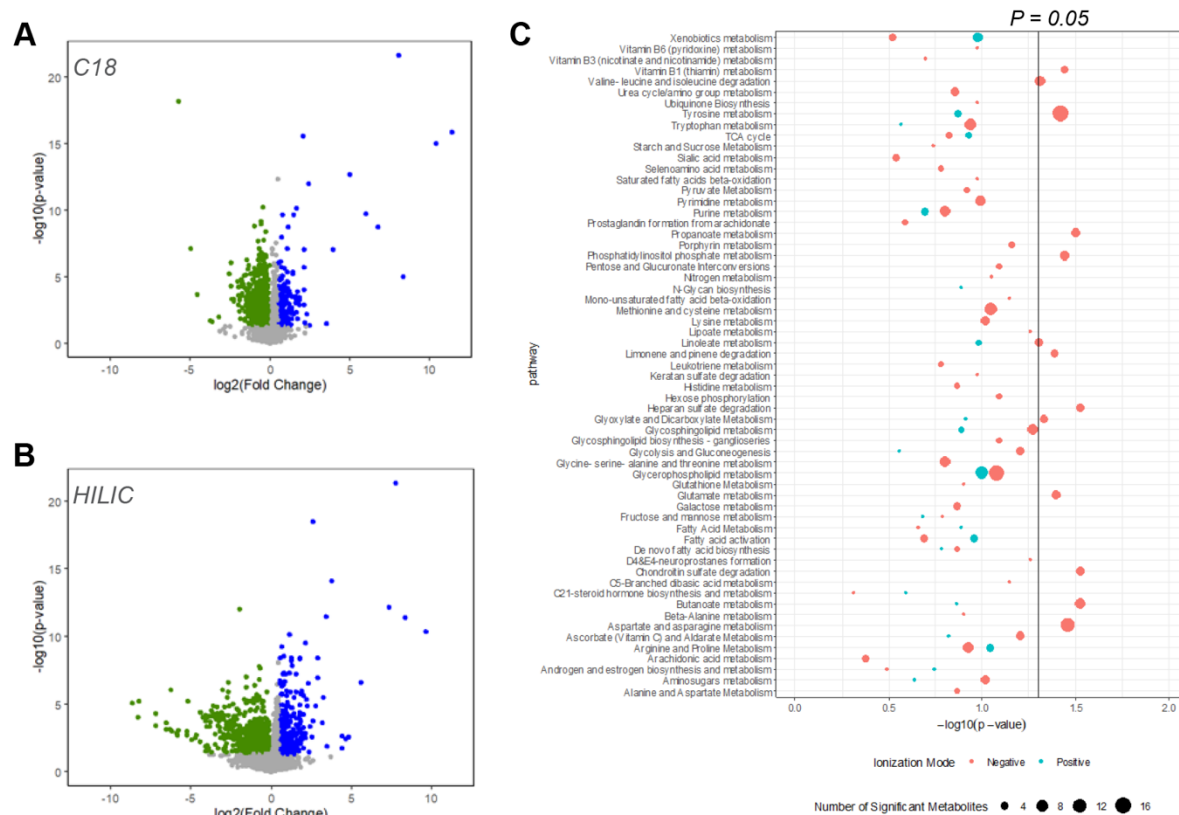


Figure 3-Supplement 3. MWAS of heads collected from control versus PFOS-exposed larvae. Heads were collected from 3 dpf control or 28 μ M PFOS exposed larvae for an untargeted metabolome wide association study (MWAS). (A) Volcano plots of metabolites detected using a C18 column with negative ionization and (B) a HILIC column with positive ionization reveal several significantly down- or up-regulated metabolites in the PFOS-exposed brain (Green = downregulated; Blue = upregulated). Statistical significance was set at $P = 0.05$. (C) There are several significantly enriched pathways following both negative ionization (pink) and positive ionization (teal). The vertical line is at $P = 0.05$. Size of each dot represents the number of significant metabolites in that pathway. HILIC, hydrophilic interaction liquid chromatography column.

Table 3. Significantly enriched metabolic pathways following PFOS exposure in 3 dpf larvae

Significantly Enriched Pathway	Physiological Relevance in CNS	Source
<i>Vitamin B1 (thiamin) metabolism</i>	<ul style="list-style-type: none"> - Microglia regulation - Neuronal K⁺ channels regulation - Myelinogenesis - Improves cognitive function 	(Ba 2008, Mkrtchyan et al. 2015)
<i>Valine-leucine and isoleucine degradation</i>	<ul style="list-style-type: none"> - Neurotransmitter metabolism, including glutamate - Protein synthesis and energy production - Nitrogen homeostasis and neurotransmitter cycling 	(Polis and Samson 2020, Salcedo et al. 2021)
<i>Tyrosine metabolism</i>	<ul style="list-style-type: none"> - Catecholamine synthesis - Regulator of neuronal longevity 	(Kobayashi et al. 1995) (Parkhitko et al. 2020)
<i>Propanoate metabolism</i>	<ul style="list-style-type: none"> - Modulates glutamine synthetase in glia - Inhibits histone deacetylase in GABAergic neurons, increasing GABA levels 	(Morland et al. 2018, Nguyen et al. 2007)
<i>Phosphatidylinositol phosphate metabolism</i>	<ul style="list-style-type: none"> - Neurotransmitter receptor expression regulation - Synaptic vesicle regulation and recycling - Neurite and dendrite morphogenesis - Clathrin-dependent membrane trafficking - Ion channel and transporter activity regulation 	(Clayton, Minogue and Waugh 2013, Raghu et al. 2019)
<i>Linoleate metabolism</i>	<ul style="list-style-type: none"> - Modulates astrocyte inflammatory response - Stimulates axonal growth of cortical neurons 	(Saba et al. 2019, Hennebelle et al. 2020)
<i>Limonene and pinene degradation</i>	<ul style="list-style-type: none"> - Represses neuronal cell death - Decreases activated glial cell number - Involved in neuroprotection - Anti-inflammatory and analgesic roles 	(Eddin et al. 2021)
<i>Heparan sulfate degradation</i>	<ul style="list-style-type: none"> - Regulator of axon guidance and synapse development and specificity - Involved in regulating the dopamine system - Part of the core synaptic organizing complexes neurexin and neuroligin 	(Condomitti and de Wit 2018, De Risi et al. 2021)
<i>Glyoxylate and dicarboxylate metabolism</i>	<ul style="list-style-type: none"> - Important for carbohydrate metabolism and energy availability - Altered following traumatic brain injury 	(Baker et al. 2018)

<i>Glutamate metabolism</i>	<ul style="list-style-type: none"> - Involved in nitrogen trafficking and ammonia homeostasis in brain - Excitatory neurotransmitter and immediate precursor for the neurotransmitter GABA 	(Schousboe et al. 2014)
<i>Chondroitin sulfate degradation</i>	<ul style="list-style-type: none"> - Role in development, plasticity, and regulation of cortical circuitry - Modulates ion channel properties - Inhibits structural plasticity (scar after injury, axonal pathfinding, and synapse formation during development) - Neuronal excitability modulation 	(Hudson et al. 2015)
<i>Butanoate metabolism</i>	<ul style="list-style-type: none"> - Neuromodulator largely produced by gut microbiota - Attenuates neuronal apoptosis - Inhibits amyloidogenesis - Protects neurons from ischemic damage - Improves long-term memory - Neurodegenerative attenuation 	(Kim et al. 2020, Xu et al. 2021, Zhou et al. 2021)
<i>Aspartate and asparagine metabolism</i>	<ul style="list-style-type: none"> - Secondary excitatory neurotransmission - Amino acid involved in glutamate synthesis - Influences NMDAR-mediated transmission - Can evoke presynaptic release of endogenous l-glutamate release in selective brain regions - Synaptic strength and connectivity modulation 	(Pardo et al. 2011, Errico et al. 2018)

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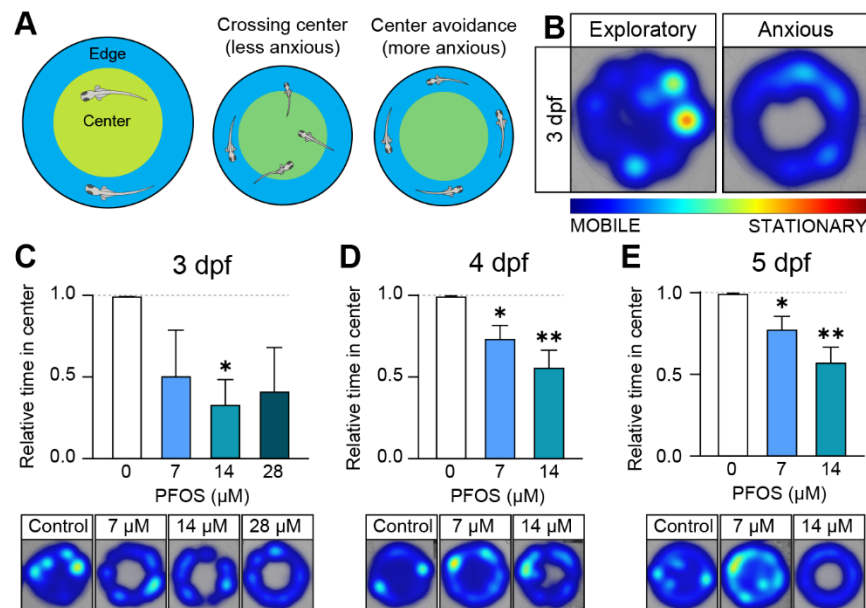


Figure 4. Developmental PFOS exposure increases anxiety-like behaviors. (A) Larval swim behavior during a 30-minute photomotor response assay can be interrogated to determine time spent in the well's center (crossing center; less anxious) versus time spent along the well's edge (center avoidance; more anxious). (B) Heat maps were generated to indicate mobile (blue) versus stationary (red) swim activity within the well, as well as the areas traversed within the well. (C) At 3 dpf, larvae exposed to 7 μM and 28 μM PFOS spend considerably less time in the center of the wells, and 14 μM PFOS spend significantly less time in the center ($P = 0.0232$). (D) At 4 dpf, both 7 μM ($P = 0.0410$) and 14 μM ($P = 0.0018$) PFOS-exposed larvae spend significantly less time in the center compared to control larvae. (E) This is also true in 5 dpf larvae exposed to 7 μM ($P = 0.0440$) and 14 μM PFOS ($P = 0.0013$). $n = 76$ -101 fish per group. Unpaired t-test with Welch's correction. Error bars represent SEM.

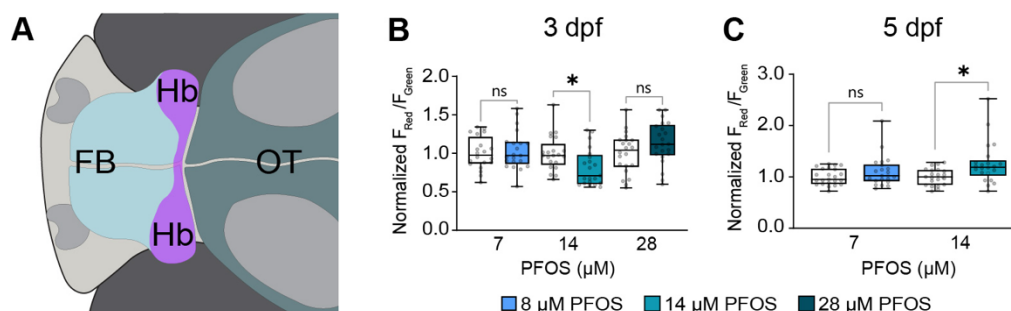


Figure 4- Supplement 2. (A) Illustrative representation of a larval zebrafish brain with anatomical regions outlined: forebrain (FB), habenula (Hb), and the optic tectum (OT). Activity quantification of neuron-driven CaMPARI was performed in the entire developing habenula following 1-minute exposure to 405 nm light. (B) At 3 dpf, 14 μM PFOS exposure caused a significant decrease in habenular Ca^{2+} activity ($P = 0.0138$), but not 7 μM nor 28 μM PFOS. (C) At 5 dpf, 14 μM PFOS-exposed larvae had a significant increase in habenular Ca^{2+} activity ($P = 0.0101$), but not 7 μM PFOS. $n = 20$ -22 fish per treatment. Unpaired t-test with Welch's correction.

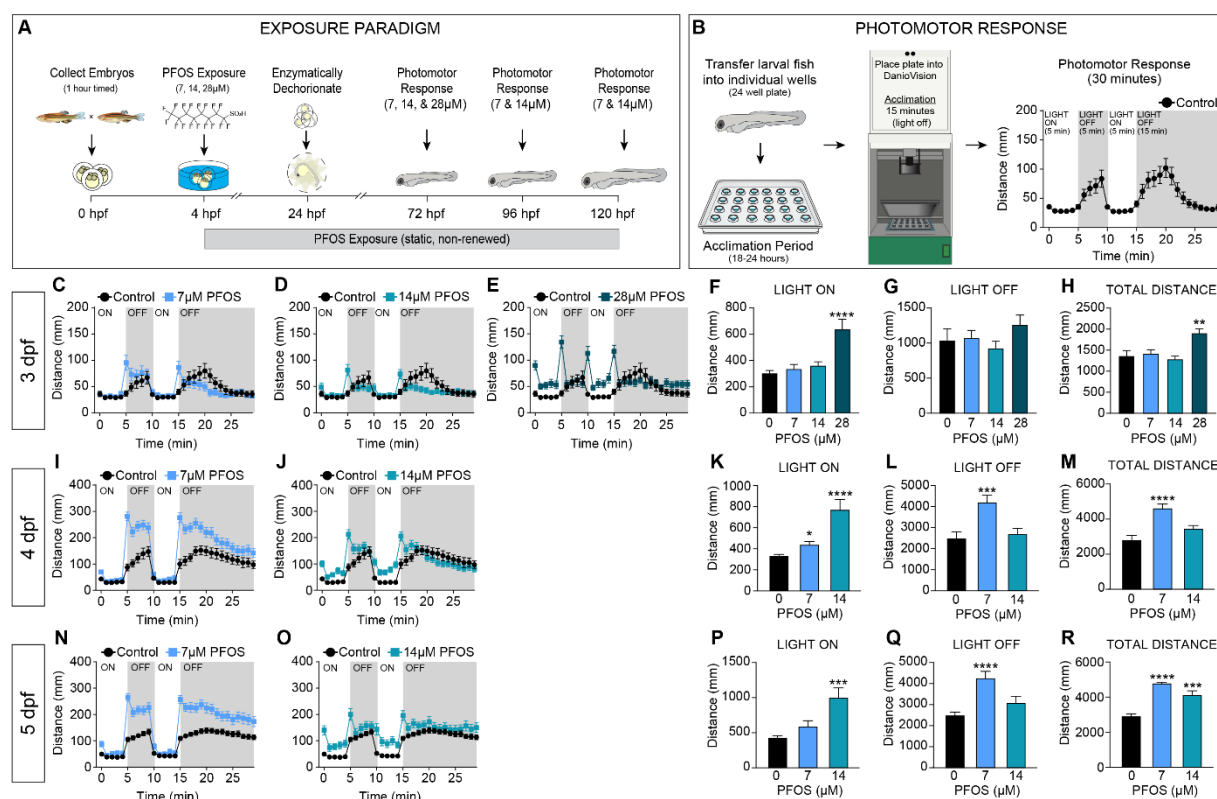


Figure 4-Supplement 1. Developmental PFOS exposure increases larval zebrafish behavior. (A) Exposure paradigm: at 4 hpf, zebrafish embryos were statically exposed to 0.1% DMSO (Control) or varying concentrations of PFOS (7, 14, or 28 μM). At 24 hpf, embryos were enzymatically dechorionated. Behavior was captured at 72, 96, and 120 hpf. (B) Photomotor response assay: Zebrafish larvae were placed in individual wells of a 24-well plate 18-24 hours prior to the behavior assay to allow for well acclimation. A photomotor assay was performed using the Noldus DanioVision Behavioral Unit. The photomotor assay was as follows: 15-minute acclimation in the unit with the light off, 5 minutes with the light on, 5 minutes with the light off, 5 minutes with the light on, and 15 minutes with the light off. (C-H) At 3 dpf, larvae exposed to 28 μM PFOS had a significant increase in swim activity during the (F) light on cycles ($P < 0.0001$) and (H) over the course of the assay ($P = 0.0023$). (K) At 4 dpf, 7 μM PFOS larvae had a significant increase in swim distance when the light was on ($P = 0.0103$), as did 14 μM PFOS larvae ($P < 0.0001$). (L) At 4 dpf, 7 μM PFOS-exposed larvae also had a significant increase in swim behavior when the light was off ($P = 0.0006$) and (M) in total over the course of the assay ($P < 0.0001$). (P) At 5 dpf, 14 μM PFOS had increased swim behavior when the light was on ($P = 0.0002$), and (Q) 7 μM PFOS larvae had increased swim behavior when the light was off ($P < 0.0001$). (R) Both 7 μM ($P < 0.0001$) and 14 μM PFOS ($P = 0.0001$) larvae had a significant increase in total distance moved. $n = 130-190$ for controls, $n = 88-99$ for PFOS-exposed. One-way ANOVA. Multiple comparisons were performed using Tukey's multiple comparisons test.

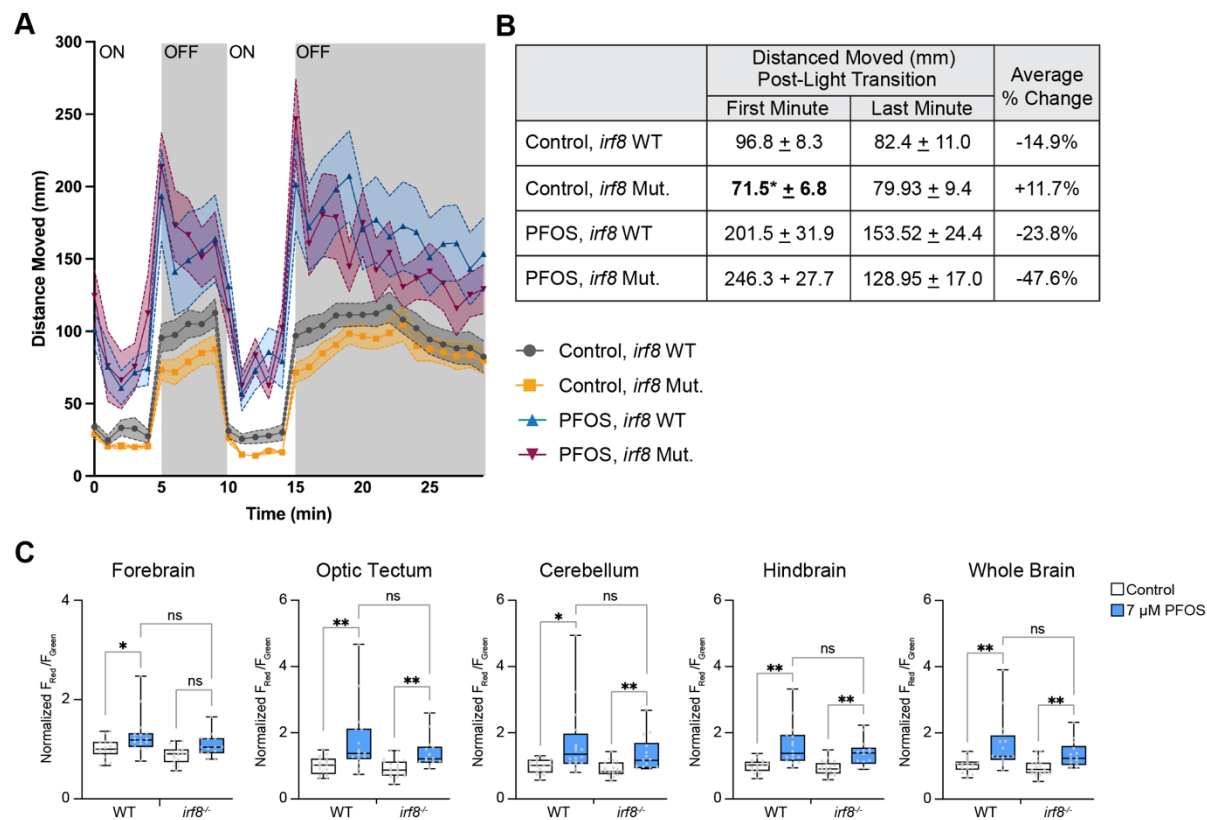


Figure 5. Microglia may influence larval behavior but are not responsible for PFOS-induced behavioral or neuronal phenotypes. Wildtype and microglia-deficient *irf8*^{st96/st96} larvae dosed with either control or 7 μ M PFOS were assessed for potential changes in swim behavior and neuronal activity. (A) Photomotor response assay of 5 dpf wildtype or *irf8* mutant control or PFOS-exposed larvae. (B) Assessing the distance moved following the first minute of the second light “off” cycle (i.e., minute 15 of the assay), control-treated *irf8* mutant larvae responded significantly less than wildtype controls ($P = 0.0237$). While not significant, PFOS-exposed *irf8* mutants had a heightened response to the first minute of the dark cycle compared to PFOS-exposed wildtype larvae. PFOS mutants also had twice the percent recovery in swim activity by the last minute of the assay (i.e., minute 15 versus minute 30). (C) There were no significant differences in regional or global PFOS-induced neuronal hyperactivity between the wildtype or mutant larvae expressing *Tg(elav13:CaMPARI)*. Unpaired t-test with Welch’s correction. $n = 14$ -22 per group for behavior; $n = 13$ -16 per group for neuroimaging.

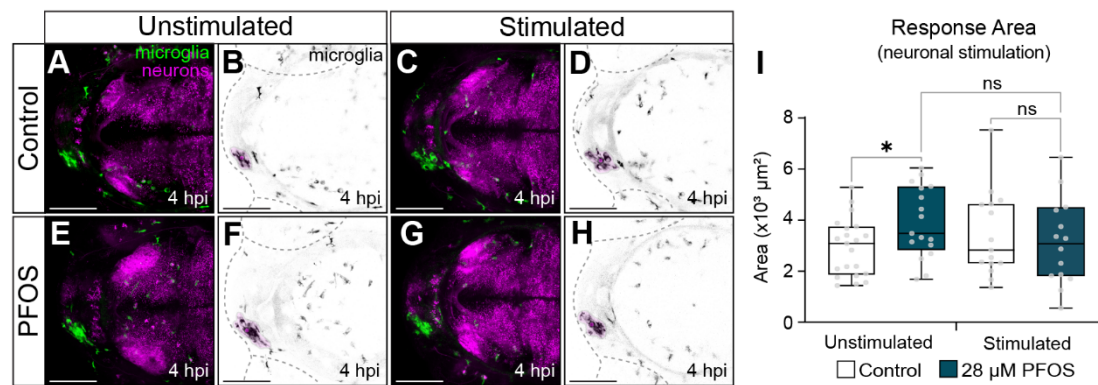


Figure 6. Neuronal silencing normalizes microglia responsiveness in PFOS-exposed larvae. Larvae expressing *Tg(elavl3:Gal4;cryaa:RFP;UAS:eNpHR3.0;mpeg1:EGFP)* were dosed with (A-D) control or (E-H) 28 μM PFOS and injured at the right telencephalon at 3 dpf and imaged 4 hpi. (E&F) PFOS-exposed unstimulated larvae had a significant increase in microglia response compared to controls ($P = 0.0256$). (C&D) Optogenetic stimulation of halorhodopsin with 570 nm light for 4 hpi did not affect microglia response to injury in control larvae. (G&H) However, neuronal silencing in PFOS-exposed larvae normalized the microglia response, such that the response was equivalent to the stimulated controls. (I) Quantification of microglia response area. Unpaired t-test with Welch's correction. $n = 14-20$ per group.

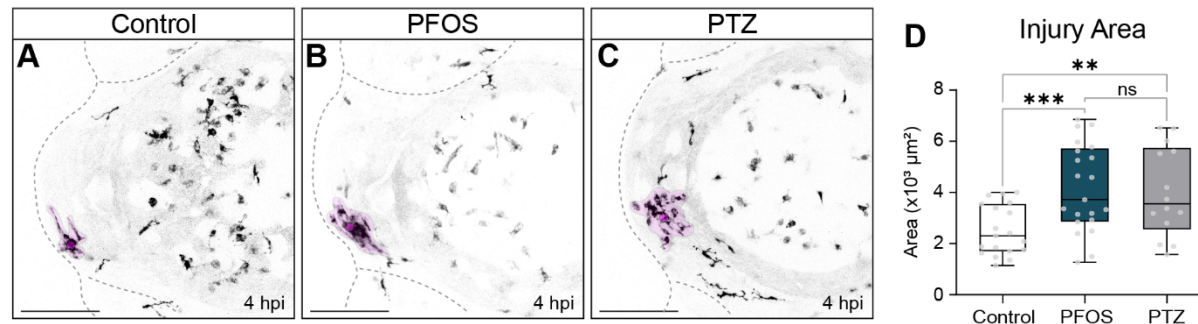


Figure 6-Supplement 1. Neuronal excitation recapitulates microglia hyperresponsive phenotype seen in PFOS-exposure. Larvae with transgenic expression of macrophages (*Tg(mpeg1:EGFP)*) were dosed with either (A) control or (B) 28 μM PFOS at 4 hpf, or (C) 5 mM PTZ at 72 hpf, followed by brain injury. (D) At 4 hpi, PFOS-exposed larvae had a significant increase in microglia response area ($P = 0.0006$). In addition, neuronal excitation by PTZ also resulted in a significant increase in microglia response ($P = 0.0092$). Unpaired t-test with Welch's correction. $n = 14-21$ per group.

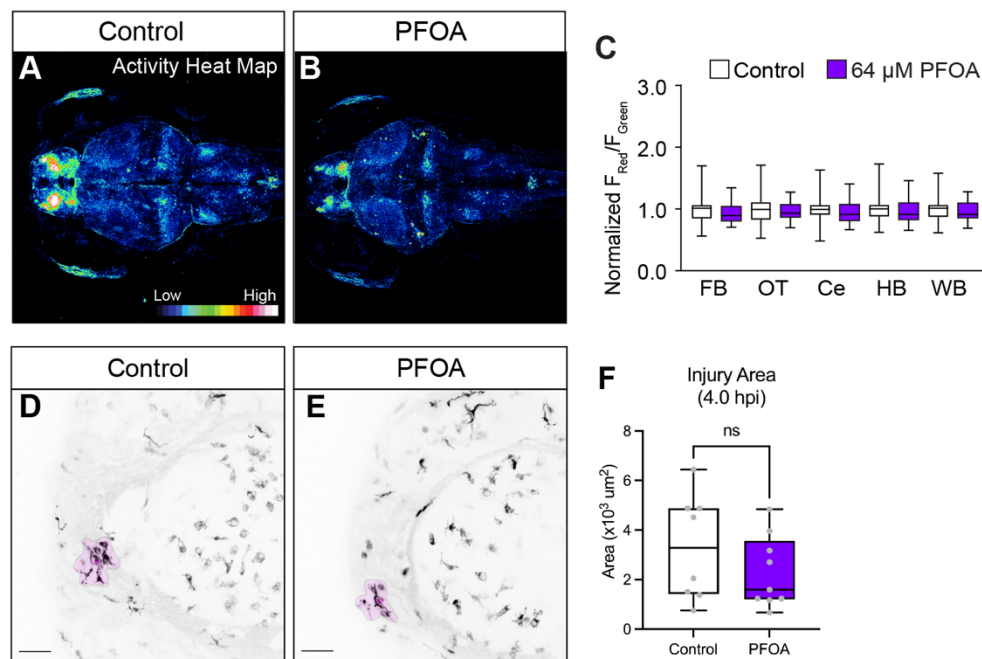


Figure 7. Exposure to a non-excitatory PFAS does not result in microglia hyperresponsiveness. PFOA is an 8-carbon PFAS compound with a carboxylate head group. To observe the effects of PFOA on neuronal activity, we again used larvae of the Tg(*elavl3:CaMPARI*) background. (A) At 5 dpf, control and (B) 64 μM PFOA-exposed larvae show no significant changes in regional or global neuronal activity, quantified in (C). (D) Larvae expressing Tg(*mpeg1:EGFP*) were also dosed with control solution or (E) 64 μM PFOA to determine if this non-excitatory PFAS compound resulted in microglia response differences. (F) At 4hpi, 64 μM PFOA-exposed larvae did not have a significant change in microglia response to injury compared to sibling controls. n = 8-9 per group. Unpaired t-test with Welch's correction.

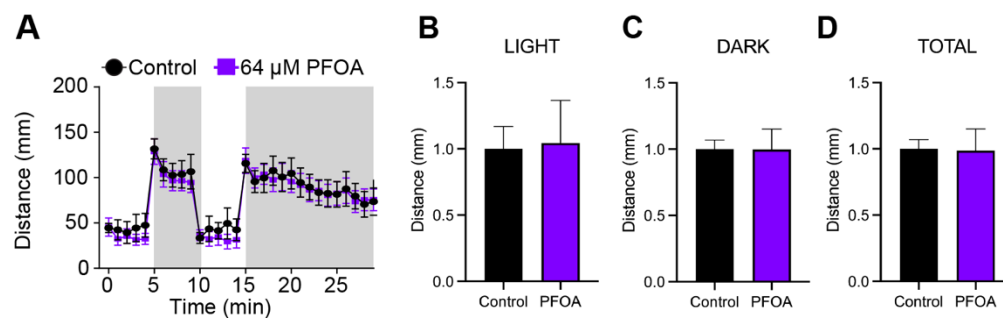


Figure 7-Supplement 1. PFOA exposure does not increase zebrafish swim behavior. Zebrafish larvae were subjected to a 30-minute photomotor response assay at 5 dpf, as previously described. (A) There were no apparent changes in distance moved during the 30-minute assay. (B) Total distance moved during light cycles, (C) dark cycles, and (D) in total throughout the assay were not significantly changed. n = 24 for controls, n = 46 for PFOA exposed. Unpaired t-test with Welch's correction.

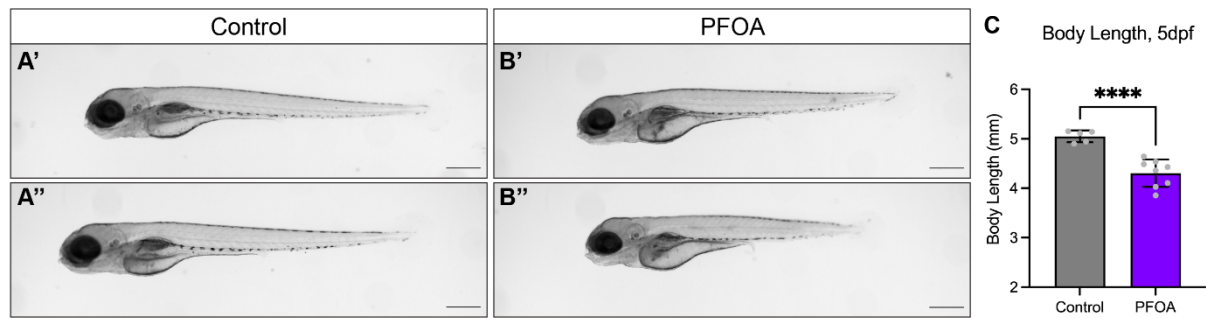


Figure 7-Supplement 2. PFOA exposure does not result in morphological changes that would impact swim ability. (A'-A'') Images of 5 dpf control-treated larvae. (B'-B'') Images of 5 dpf larvae chronically treated with 64 μM PFOA. (C) Body length measurements of control versus PFOA treatment at 5 dpf ($P < 0.0001$). Unpaired t-test with Welch's correction. $n = 5-8$ per group. Scale = 500 μm.

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