

Centrally Synthesized Estradiol Is a Potent Anti-Inflammatory in the Injured Zebra Finch Brain

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In homeotherms, injury to the brain, such as a penetrating wound, increases microglial cytokine expression and astroglial aromatase (estrogen synthase). In songbirds, injury-induced synthesis of estrogens is neuroprotective as aromatase inhibition and replacement with estradiol (E_2) exacerbates and mitigates the extent of damage, respectively. The influence of induced aromatization on inflammation, however, remains unstudied. We hypothesized that injury-induced aromatization, via E_2 synthesis, may affect neuroinflammation after a penetrating brain injury. Using adult zebra finches, we first documented an increase in the transcription of cytokines but not aromatase, 2 hours after the injury. Twenty-four hours after the injury, however, aromatase was dramatically elevated and cytokine expression had returned to baseline, suggesting that aromatization may be involved in the decrease of cytokines and neuroinflammation. In two subsequent experiments, we tested the influence of the inhibition of induced aromatization and aromatase inhibition with concomitant central E_2 replacement on the transcription of the cytokines TNF- α , IL-1 β , and IL-6, the enzyme cyclooxygenase-2 (cox-2), and its product prostaglandin E2 (PGE2). Administration of fadrozole, an aromatase inhibitor, caused a sustained elevation of IL-1 β in females and TNF- α , cox-2, and PGE2 in both sexes. This prolonged neuroinflammation appears to be due to a failure to synthesize E_2 locally because intracranial E_2 replacement lowered IL-1 β in females, TNF- α in males, and cox-2 and PGE2 in both sexes. IL-6 was not affected by injury, aromatase inhibition, or E_2 replacement in either sex. These data suggest that E_2 synthesis after a penetrating brain injury is a potent and inducible anti-inflammatory signal, with specific modulation of discrete cytokine signaling. (*Endocrinology* 157: 2041–2051, 2016)

Estrogens like 17 β -estradiol (E_2) organize and activate the vertebrate brain (1–3) and are potent regulators of many neural processes across vertebrates (4–6). E_2 also regulates peripheral physiology with dramatic effects on metabolism, muscle function, and the immune system (7, 8). Indeed, premenopausal women are at a lower risk for several diseases including cardiac disease, osteoporosis, and stroke, compared with age-matched men and postmenopausal women not on hormone replacement therapy, suggesting that E_2 may be protective centrally and peripherally (9, 10).

E_2 is synthesized in multiple tissues including but not limited to the ovary, adipose, and liver (8). Several vertebrates can also synthesize E_2 in the brain, via expression of

the enzyme aromatase (*estrogen synthase*) in neurons at discrete neural loci (11–15). However, in songbirds and mammals, after an injury to the brain, such as a penetrating stab wound, aromatase is induced in reactive astrocytes at the site of damage (16–19). In songbirds, the induction of aromatase in astroglia is particularly dramatic (19–21), with demonstrated neuroprotective effects such as decreases in injury-induced apoptosis (22–24), and increases in cyto- and neurogenesis (25).

Injury to the homeotherm brain also causes neuroinflammation, characterized in part by the rapid synthesis and secretion of proinflammatory cytokines like TNF- α , and interleukins, IL-1 β and IL-6 (26–30). The secretion of these cytokines induces cyclooxygenase-2 (cox-2), which

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Abbreviations: CNS, central nervous system; cox-2, cyclooxygenase-2; Ct, detection threshold; δ Ct, δ -threshold cycle number; E_2 , 17 β -estradiol; EIA, enzyme immunoassay; ER, estrogen receptor; GAPDH, glyceraldehyde-3-phosphate dehydrogenase; PGE2, prostaglandin E2; qPCR, quantitative PCR; SSV, steroid suspension vehicle.

synthesizes the prostanoid prostaglandin E₂ (PGE₂) (31, 32). PGE₂ is responsible for classic signs of tissue damage, including but not limited to edema, erythema, and pain (33). PGE₂ also can recruit macrophages and neutrophils, activate microglia, and increase inflammatory processes by amplifying proinflammatory cytokine and chemokine secretion (31, 33, 34). Inflammatory signaling can have beneficial and detrimental effects that may differ between the acute and chronic periods after trauma. Specifically, although the inflammatory response can initially promote phagocytosis and repair, the chronic expression of inflammatory signaling can activate toxic cascades and, in the central nervous system (CNS), can promote further neurodegeneration (35–38).

Previous studies have revealed a strong interaction between estrogens and the innate immune system. Specifically, TNF- α and IL-1 β increase in women with low circulating E₂ due either to natural (39) or surgical menopause (40). Resident macrophages isolated from female mice are more plentiful and express higher levels of toll-like receptors, compared with males (41), perhaps suggesting a higher sensitivity of the female immune system. Indeed, ovariectomized mice have higher neural cytokine expression after peripheral endotoxin treatment relative to sham controls (42), suggesting an anti-inflammatory role for circulating E₂.

Whereas interactions between the endocrine and immune system have been studied in songbirds (43–45), surprisingly little is known about how E₂ affects the innate immune system. Given the interactions among a penetrating brain injury, cytokine expression and E₂-dependent neuroprotection, we hypothesized an influence of injury-induced aromatization on neuroinflammation. Because zebra finches (*Taeniopygia guttata*) of both sexes demonstrate rapid increases in cytokines (20, 46) and a robust, sustained, and reliable induction of glial aromatase (21, 46) after a penetrating brain injury we hypothesized that E₂ synthesis, via aromatization, may affect neuroinflammation in males and females of this species. First, we assessed the time course of cytokine and aromatase induction. Using time points established from this experiment, we tested the effect of aromatase inhibition and E₂ replacement on the transcription of the proinflammatory cytokines TNF α , IL-1 β , and IL-6. Toward understanding the downstream effects of cytokine expression, in these same subjects, we also assessed the effects of injury-induced aromatization and E₂ on the expression of *cox-2* and the central levels of PGE₂.

Materials and Methods

Adult male and female zebra finches (>100 d after hatch) were obtained from a commercial breeder and housed in same-sex

aviaries at The American University, under a 12-hour light, 12-hour dark cycle, with food and water provided ad libitum. The American University Institutional Use and Animal Care Committee approved all animal procedures.

Surgery and tissue preparation

In all experiments, subjects served as their own controls (within animal design) with experimental treatments administered to contralateral brain hemispheres. Within each experiment (see below), all treatments were counterbalanced between the left and the right telencephalic hemispheres. As previously published (20, 22–25), animals were anesthetized and positioned in a stereotaxic apparatus with the head angled at 45°. The cranium was exposed, a craniotomy was performed, and a 22-gauge Hamilton syringe (Hamilton Co) was targeted toward the entopallial nucleus, 2 mm anterior to the pineal gland, 3 mm lateral to the midline, and 3 mm ventral to the brain surface (see 22, 23). This form of injury is similar to penetrating brain injury used in other laboratories (47–51). Two or 24 hours after the injury, birds were decapitated, and the telencephalon was detached from the rest of the brain. To concentrate the contribution of the injury tract to the tissue processed, the anterior telencephalon approximately 4 mm anterior to the pineal (52) was discarded. The injury site was always in the posterior lobes and, based on previous studies, is estimated to span about 10%–20% of the remaining telencephalic volume (23–25). The posterior telencephalon was separated across the midline, weighed, and then stored at –80°C. Just prior to the procedures described below, samples were homogenized in 1 mL of 0.1M phosphate buffer.

Experiments

Experiment 1

Time course of injury-induced changes in the expression of TNF- α , IL-1 β , IL-6, and aromatase mRNA after a neural injury

To determine the time course of cytokine and aromatase induction, 20 adult zebra finches (10 per sex) received a unilateral brain injury as described above. A 5- μ L injection of a steroid suspension vehicle (SSV) (22, 25) was administered through the syringe over 60 seconds. Control hemispheres were left uninjured. Birds were either killed at 2 hours ($n = 10$; five per sex) or 24 hours after the injury ($n = 10$; five per sex), and the tissue was collected as described above. Tissue was subsequently processed for quantitative PCR (qPCR) to measure the expression of TNF- α , IL-1 β , IL-6, and aromatase (see below).

Experiment 2

*Expression of TNF- α , IL-1 β , IL-6, *cox-2*, and PGE₂ after aromatase inhibition*

Ten adult zebra finches (five per sex) received bilateral injuries as described above. Subjects received either 5 μ L of a 10-mg/mL solution of the aromatase inhibitor fadrozole suspended in SSV (50 μ g of fadrozole per injection) (23) or 5 μ L of SSV into contralateral hemispheres. Animals were killed 24 hours after the injury, a time when injury-induced aromatase expression is robust and reliable (23–25). The expression of TNF- α , IL-1 β , IL-6, and *cox-2* was assessed using qPCR, and neural levels of PGE₂ were measured using a validated enzyme immunoassay (EIA) (see below).

Experiment 3

Expression of TNF- α , IL-1 β , IL-6, cox-2, and PGE2 after E₂ replacement

Ten adult zebra finches (five per sex) received bilateral injuries through which either 5 μ L of 10-mg/mL fadrozole suspended in SSV alone or 5 μ L of the same solution containing 1 μ g of E₂ was administered (24). Animals were killed 24 hours after the injury, and qPCR and EIA were used to analyze the expression of TNF- α , IL-1 β , IL-6, cox-2, or PGE2 as described below.

Quantitative PCR

Total RNA was isolated from 500 μ L of homogenate using the RNeasy mini extraction kit (QIAGEN) according to the manufacturer's instructions. The purity and concentration were analyzed on a ND-1000 spectrophotometer (NanoDrop), and only extracts that exceeded a 260:280 ratio of 1.9 were used.

Primers for aromatase, TNF- α , IL-1 β , IL-6, cox-2, and the housekeeping gene, GAPDH (see Table 1) were designed based on the known zebra finch sequence for each gene. Amplicons were run on an agarose gel to confirm the presence of a single band of the expected size (approximately). Because the size of small sequences (~100bp) is often difficult to determine unequivocally, the bands were sequenced (Genewiz; Beckman Coulter Genomics) and compared with the zebra finch genome. We also performed a serial dilution by varying template concentration. Efficiency was assessed at greater than 100%. Melt curves of the PCR products were evaluated to confirm the specificity of primer annealing.

The expression of target transcripts was analyzed using 96-well optical plates and the SuperScript Sybr Green one-step quantitative RT-PCR kit (Invitrogen). Each sample was amplified in triplicate, and each well (reaction) had 75 ng total RNA in a 15- μ L reaction. In each experiment, samples for one target gene were run on a single plate. To permit and assess for plate-to-plate variation (53) an interrun calibrator was used on each plate, and glyceraldehyde-3-phosphate dehydrogenase (GAPDH) expression was analyzed using a one-way ANOVA to ensure that GAPDH did not differ between groups (data not shown). A no-template control and a no-reverse transcriptase control were run on each plate to eliminate the possibility of contaminants.

Prostaglandin E₂ enzyme immunoassay

A combined ether and solid-phase extraction were performed on 300 μ L of homogenate to maximize the recovery of steroids from the zebra finch brain as previously described (54, 55).

Ether extraction

Homogenate was extracted with 2 mL of diethyl ether (Sigma-Aldrich), vortexed on a low setting for 30 seconds, and centrifuged at 2100 rpm for 5 minutes at 4°C. After the centrif-

ugation, the mixture was placed in a MeOH/dry ice bath to freeze the aqueous phase, and the organic phase was poured into a clean glass tube (Kimble Chase). This was repeated for a total of three times. After the third extraction, samples were dried down under a stream of air until fully evaporated. Then approximately 50 μ L of 1:1 mixture of MeOH/CH₂Cl₂ was dripped down the side of the tubes and evaporated under air. Fully dried down samples were stored at -80°C until solid-phase extraction.

Solid-phase extraction

Ether-extracted samples (see above) were resuspended in 300 μ L of 1.0 M acetate buffer according to the manufacturer's instructions (Arbor Assays). Solid-phase extraction was performed using a Visiprep SPE vacuum manifold (Supelco) and 1 mL per 500 mg C18 cartridges (Agilent). Columns were conditioned under vacuum pressure by adding 500 μ L ethanol, followed by 750 μ L of ultrapure water. Cartridges were not allowed to dry. Samples were then applied to an SPE cartridge and eluted into a waste container and then were washed with 250 μ L of ultrapure water. At this time, the waste bin was replaced with the final collection glass tubes (Kimble Chase), and 300 μ L of ethyl acetate containing 1% methanol was applied to the column and was eluted by gravity. Samples were then stored at -80°C.

PGE2 enzyme immunoassay

PGE2 was measured using a commercial kit developed by Cayman Chemical. Cayman Chemical reports a high specificity for PGE2 and low cross-reactivity (<0.01%) with similar prostaglandins or prostaglandin metabolites and a detection range from 7.8 to 1000 pg/mL. On the day of the assay, samples were dried down under air stream, according to the manufacturer's instructions (Arbor Assays). Samples were then resuspended in 300 μ L EIA buffer (Cayman Chemical) and assayed at a 1:5 dilution in triplicate according to the manufacturer's instructions.

PGE2 immunoassay validation

To validate the specificity of the assay in the zebra finch, a serial dilution was first performed on zebra finch brain samples. Three dilutions (1:1, 1:3, 1:5 dilution) were chosen and the measured concentrations decreased by expected proportion. Second, the accuracy of the kit was assessed by spiking brain tissue samples with known amounts of PGE2. Before ether and solid-phase extraction, spiked samples were also run to estimate recovery. Two samples were spiked with radioinert PGE2 to 125 pg/mL. There was a high correlation between expected and observed values, and the recovery rate was assessed at 87%. Finally, as part of a different study and as an in vivo assay validation, injections of a cox-2 inhibitor significantly decreased brain levels of PGE2 compared with injected controls ($P = .04$) (56).

Table 1. List of Primers Used for Amplification for qPCR

Gene Symbol	Accession Number	Forward Primer (5'-3')	Reverse Primer (5'-3')
Aromatase	AH008861.2	CTCCACCGACAAAAATCCAC	GGGTTTCCAGGACCATCT TT
GAPDH	NM_001198610.1	TGCTGCTCAGAACATTATCCC	TTTCCACAGCCTAGCAGCT
TNF- α	XM_002197321.2	TGTCCCATCTGC ACCACCTTCTTA	ATTCCCTTCCATCTGGCTTCTGT
IL-1 β	XM_002195564.2	TTCCGGTGCATCAGAGGCAGTTAT	GCACGAAGCACTTGTGGTCAATGT
IL-6	XM_002191284.2	CGTCTGCCAGAACAGCATGGAAAT	TATCCTCATTAAGCCGGCGAGCA

Statistical analyses

Quantitative PCR Data

The delta threshold cycle number (δCt) method was used for quantification using the detection threshold (Ct) for each target gene less the Ct for the housekeeping gene, glyceraldehyde 3-phos-

phate dehydrogenase (GAPDH). All statistical analyses were performed on δCt , and are presented as means \pm SEM in both the text of the results and in the figures (see Figures 1, 3, and 6). Some data are shown as fold change + SEM using the formula $2^{-\delta(\delta\text{Ct})}$, and the normalization of the control condition δCt value is set to 1 (20, 46, 57). In this case, each individual experimental data point is expressed as a function of the control condition mean, resulting in a distribution (with variability). Whereas this permits an intuitive appreciation of the difference in expression between treatment and control hemispheres, the reader is reminded that it obscures any baseline differences across control conditions and groups (see Figures 2, 4, and 7).

δCt -values for TNF- α , IL-1 β , IL-6, and aromatase were analyzed separately using a three-way ANOVA with treatment, time, and sex (experiment 1) and a two-way ANOVA with treatment and sex (experiments 2 and 3) as the main variables. In all experiments, treatment was coded as a within-subject variable. The source of significant main effects was queried using Tukey-Kramer post hoc analysis, and significant interactions were assessed with Fisher least significant differences pairwise comparisons.

To account for brain weight, data obtained in picograms per milliliter were converted to picograms per milligram based on the wet weight of samples collected at the time the animals were killed. A two-way ANOVA was run with variables of treatment and sex, and treatment was coded as a within-subject variable, and post hoc analysis was conducted as described above.

PGE2 enzyme immunoassay

To account for brain weight, data obtained in picograms per milliliter were converted to picograms per milligram based on the wet weight of samples collected at the time the animals were killed. A two-way ANOVA was run with variables of treatment and sex, and treatment was coded as a within-subject variable, and post hoc analysis was conducted as described above.

Results

Experiment 1: Time course of injury-induced changes in the expression of TNF- α , IL-1 β , IL-6, and aromatase mRNA after neural injury

Tumor necrosis factor- α

Statistical analysis revealed the main effects of treatment, time, and sex. Overall, TNF- α expression was higher in the injured relative to the uninjured lobe ($F [1, 8] = 38.51, P < .01$), at 2 hours relative to 24 hours after the injury (time [$F [1, 8] = 70.30, P = .01$], and in females relative to males ($F [1, 8] = 8.715, P =$

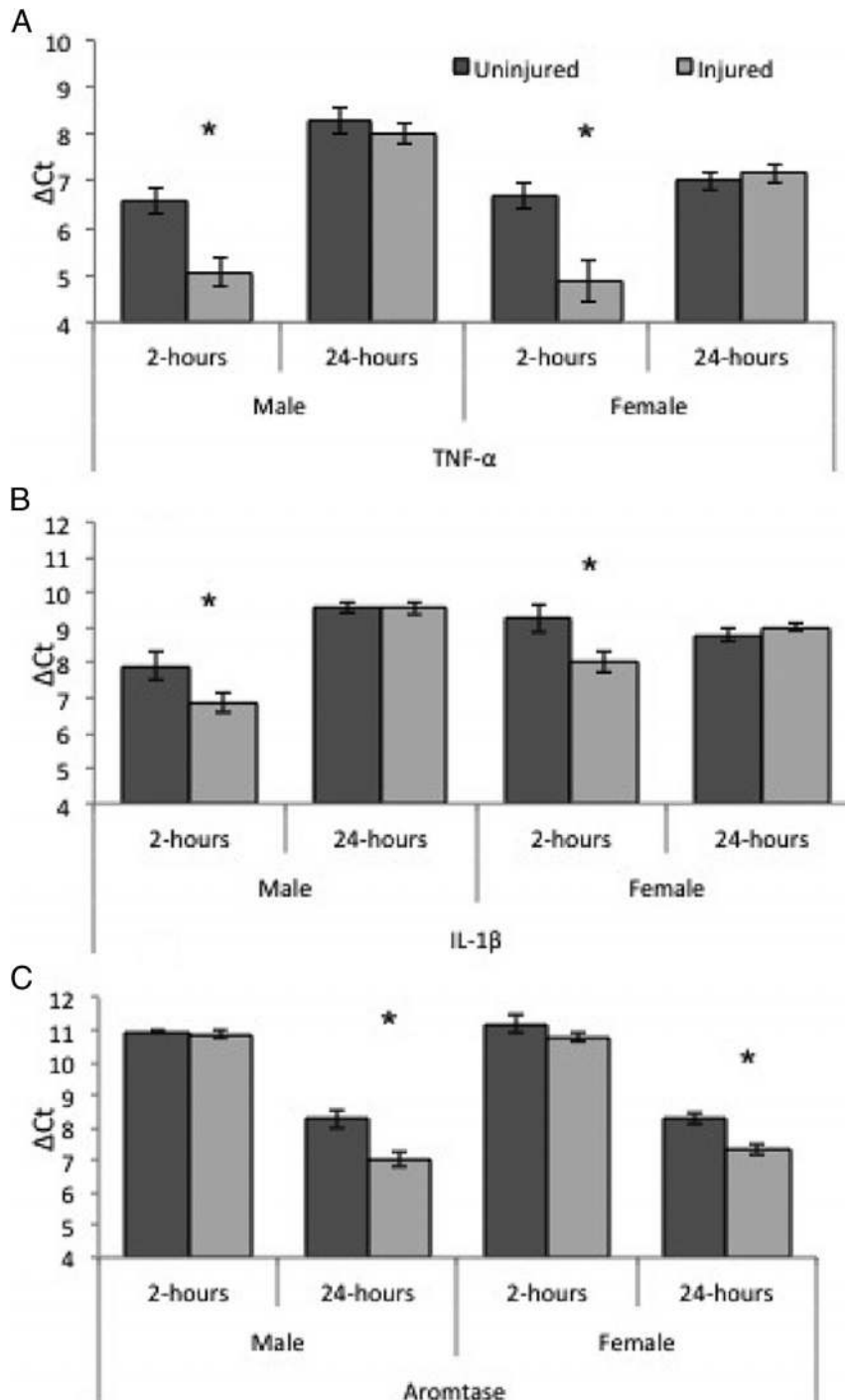


Figure 1. Levels of TNF- α , IL-1 β , and aromatase mRNA relative to GAPDH (δCt values \pm SEM) in adult male and female zebra finches from experiment 1. In both males and females, TNF- α (A) and IL-1 β mRNA (B) is increased 2 hours after the injury but not at 24 hours after the injury. At 24 hours after the injury, females have higher TNF- α expression relative to males (A). At 2 hours, males have increased IL-1 β mRNA relative to injured females (B). Aromatase mRNA is increased at 24 hours but not at 2 hours after the injury (C). *, $P < .05$. Note: y-axes do not begin at zero.

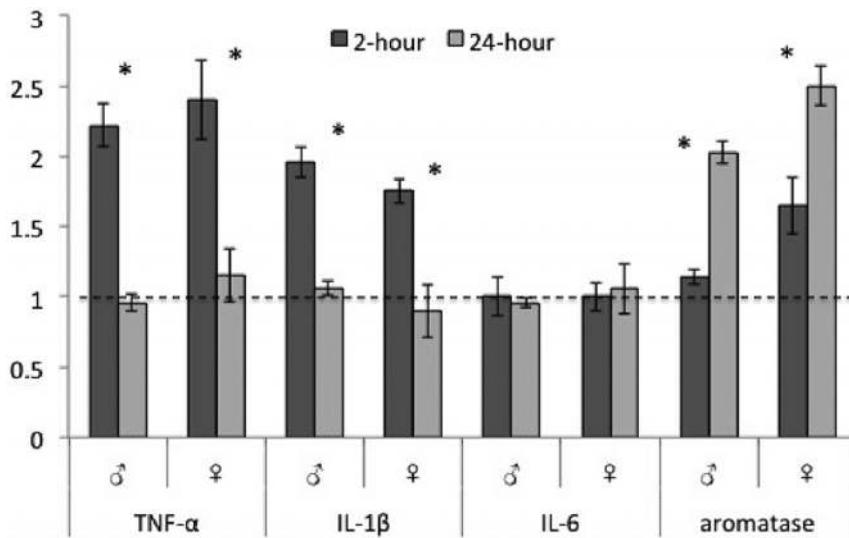


Figure 2. Levels of cytokines and aromatase mRNA relative to GAPDH (fold change values \pm SEM) in adult male and female zebra finches. TNF- α and IL-1 β mRNA are up-regulated at 2 hours after the injury in both males and females; however, all cytokines are at control levels 24 hours after the injury. At this time point, aromatase transcript is increased in both males and females. Dashed line represents uninjured controls (controls set to 1 for fold change calculation). *, $P < .05$.

.01). We also found a significant interaction between treatment \times time ($F [1, 8] = 36.167, P < .01$). This interaction reflects a higher expression in the injured relative to uninjured lobes at 2 (4.95 ± 0.26 vs 6.67 ± 0.19 , respectively; $P < .01$) but not at 24 hours after the injury (7.71 ± 0.24 vs 7.68 ± 0.24 , respectively; $P = .88$). There was also a significant interaction of sex \times time ($F [1, 8] = 8.76, P < .01$). This interaction reflects similar levels of expression across sexes at 2 hours (5.81 ± 0.32 vs $5.82 \pm 0.40, P = .96$; males vs females) but a lower expression in the male relative to the female brain at 24 hours (8.27 ± 0.28 vs $7.16 \pm 0.19, P = .01$; males vs females). There were no other significant interactions (treatment \times sex: $F [1, 8] = 1.34, P = .23$; treatment \times sex \times time: $F [1, 8] = 0.064, P = .80$). Means \pm SEM of δ Ct (in which a lower number equals a higher expression) are shown in Figure 1A. Fold change data are depicted in Figure 2.

Interleukin-1 β

In a largely similar pattern, IL-1 β transcription varied systematically over treatment ($F [1, 8] = 12.45, P < .01$) and time ($F [1, 8] = 16.49, P < .01$) but not across sex ($F [1, 8] = 1.01, P = .33$). We also found a significant interaction of treatment \times time ($F [1, 8] = 16.23, P < .01$). As above, this interaction reflects a higher expression in the injured relative to uninjured lobe at 2 hours (7.61 ± 0.31 vs 8.59 ± 0.35 , respectively; $P < .01$) but not at 24 hours (9.19 ± 0.72 vs 9.05 ± 0.21 , respectively; $P = .81$) after the injury. There also was a significant interaction of sex \times time ($F [1, 8] = 24.24, P < .01$). This interaction

reflects a higher expression in males relative to females at 2 hours after the injury (7.3 ± 0.29 vs $8.39 \pm 0.28, P < .01$; males vs females) but not 24 hours after the damage (9.56 ± 0.11 vs $9.6 \pm 0.19, P = .64$; males vs females). No other significant interactions were found (treatment \times sex: ($F [1, 8] = 0.51, P = .48$; treatment \times sex \times time: ($F [1, 8] = 0.12, P = .91$). Means \pm SEM of δ Ct (in which the lower number equals a higher expression) are shown in Figure 1B and fold change data are depicted in Figure 2.

IL-6

In contrast to the other cytokines investigated, no significant main effects or interactions on IL-6 expression were detected (treatment: $F [1, 8] = 1.008, P = .33$; time: $F [1, 8] = 0.24, P = .64$; sex: $F [1, 8] = 0.009, P = .93$) or interactions (treatment \times time: $F [1, 8] = 1.28, P = .28$; treatment \times sex: $F [1, 8] = 0.033, P = .89$; sex \times time: $F [1, 8] = 0.30, P = .59$; treatment \times sex \times time ($F [1, 8] = 0.57, P = .46$; see Figure 2).

Aromatase

Three-way ANOVA revealed significant effects of treatment ($F [1, 8] = 39.238, P < .01$) and time ($F [1, 8] = 381.09, P < .01$) but not sex ($F [1, 8] = 0.57, P = .46$). We also detected a significant interaction of treatment \times time ($F [1, 8] = 15.35, P < .01$) but no other interactions (treatment \times sex: $F [1, 8] = 0.21, P = .89$; sex \times time: $F [1, 8] = 0.59, P = .81$; treatment \times sex \times time: ($F [1, 8] = 1.91, P = .19$). The interaction of treatment \times time is driven by higher aromatase in injured lobes at 24 hours but not at 2 hours after the injury (7.16 ± 0.14 vs $10.81 \pm 0.08, P < .01$; 24 hours vs 2 h). Means \pm SEM of δ Ct (in which a lower number equals a higher expression), and fold change data are depicted in Figures 1C and 2, respectively.

Experiment 2: Expression of TNF- α , IL-1 β , IL-6, cox-2, and PGE2 after aromatase inhibition

Tumor necrosis factor- α

At 24 hours after the injury, a time when aromatase, but not cytokines, is elevated (experiment 1), we found significant effects of aromatase inhibition ($F [1, 8] = 29.72, P = .01$) and sex ($F [1, 8] = 73.83, P < .01$) but no interaction (treatment \times sex: $F [1, 8] = 2.847, P = .13$).

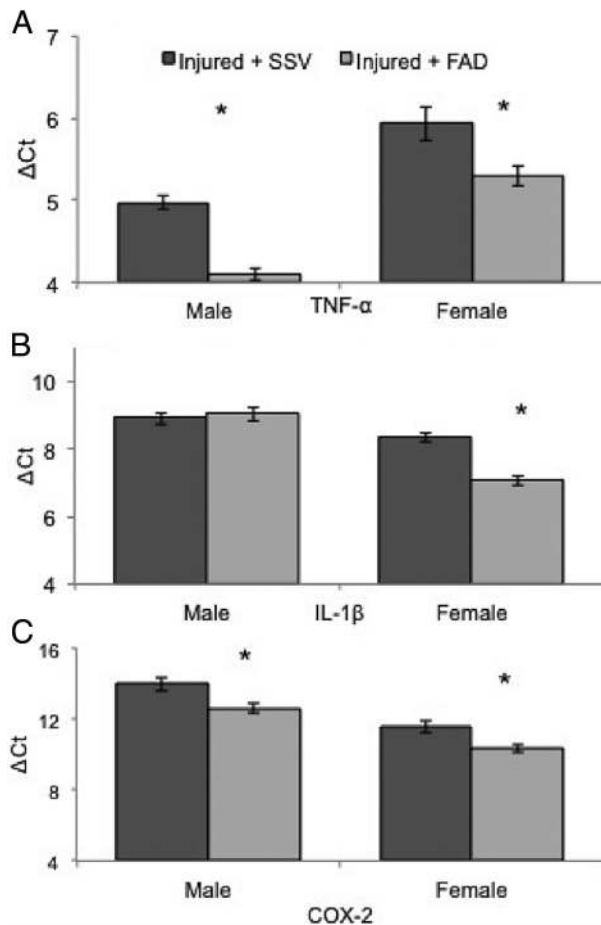


Figure 3. Levels of TNF- α , IL-1 β , and cox-2 mRNA relative to GAPDH (δCt values \pm SEM) in adult male and female zebra finches from experiment 2. Only significant gene targets are shown. Fadrozole (FAD) administration increased TNF- α mRNA in both males and females; however, males have more TNF- α , regardless of treatment (A). Fadrozole treatment after the injury increased IL-1 β mRNA only in females (B). Fadrozole administration increased cox-2 mRNA in both males and females; however, females have higher cox-2 levels, regardless of treatment (C). *, $P < .05$. Note: y-axes do not begin at zero.

Fadrozole administration increased TNF- α mRNA expression 24 hours after injury compared with control SSV treatment (4.71 ± 0.18 vs 5.45 ± 0.168), independent of sex. Overall, males had higher TNF- α than females, regardless of treatment (4.56 ± 0.13 vs 5.61 ± 0.15 ; males vs females). Means \pm SEM of δCt and fold changes are presented in Figures 3A and 4, respectively.

Interleukin-1 β

Two-way ANOVA indicated a main effect of treatment (treatment ($F[1, 8] = 24.52, P < .01$), sex ($F[1, 8] = 7.71, P < .01$), and their interaction ($F[1, 8] = 27.45, P < .01$). IL-1 β expression was higher in injured lobes treated with fadrozole compared with vehicle, independent of sex (8.04 ± 0.32 vs 8.63 ± 0.12). Furthermore, females had a higher IL-1 β expression than males independent of treat-

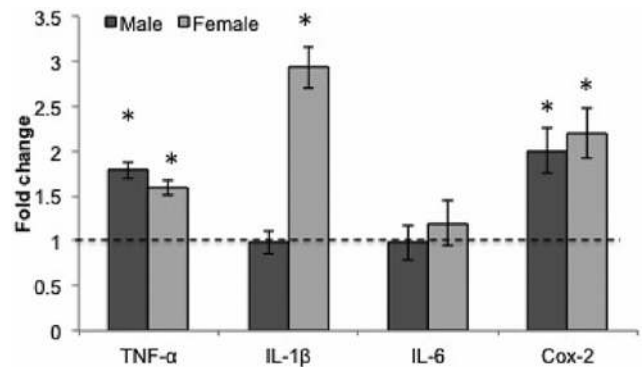


Figure 4. Levels of cytokine mRNA relative to GAPDH (fold change values \pm SEM) after injury with fadrozole administration in adult male and female zebra finches. Fadrozole administration resulted in sustained elevation of TNF- α and cox-2 in both males and females compared with injured controls, with males having a more pronounced increase of TNF- α . Aromatase inhibition also resulted in a sustained elevation of IL-1 β but is evident only in females. Dashed line represents injured controls (controls set to 1 for fold change calculation). *, $P < .05$.

ment (7.69 ± 0.21 vs 8.98 ± 0.11). The treatment \times sex interaction reflects higher IL-1 β expression in the lobe treated with fadrozole compared with SSV in females (7.05 ± 0.13 vs 8.34 ± 0.13 ; $P = .01$; fadrozole vs SSV) but not in males (9.03 ± 0.20 vs 8.93 ± 0.13 , $P = .74$; fadrozole vs SSV). Means \pm SEM of δCt and fold changes are presented in Figures 3B and 4, respectively.

Interleukin-6

As in experiment 1, we failed to detect any differences in IL-6 expression. Two-way ANOVA indicated no significant effects (treatment: $F[1, 8] = 0.69, P = .43$; sex: $F[1, 8] = 0.59, P = .46$) or interactions (treatment \times sex: $F[1, 8] = 2.09, P = .19$; see Figure 4).

Cyclooxygenase-2

We found significant effects of treatment ($F[1, 8] = 17.0, P < .01$) and sex ($F[1, 8] = 27.5, P < .01$) on cox-2 with no interaction of treatment \times sex ($F[1, 8] = 0.83, P = .83$). Cox-2 expression was higher in lobes treated with fadrozole compared with SSV (11.49 ± 0.42 vs 12.76 ± 0.45) independent of sex. Overall, females had higher cox-2 expression compared with males independent of treatment (10.95 ± 0.31 vs 12.30 ± 12.51). Means \pm SEM of δCt and fold changes are presented in Figures 3C and 4, respectively.

Prostaglandin E2

Inhibition of aromatase increased PGE₂ levels, with a main effect of treatment ($F[1, 8] = 33.8, P < .01$) and its interaction with sex ($F[1, 8] = 9.30, P < .01$) but not sex alone ($F[1, 8] = 1.09, P = .32$). PGE₂ levels were higher in lobes treated with fadrozole compared with SSV

(27.53 ± 3.47 vs 7.59 ± 1.16 , respectively), independent of sex. The interaction of treatment \times sex is due to lower PGE2 expression after an injury with SSV in females compared with males (3.97 ± 0.48 vs 11.21 ± 0.34 pg/mg, $P < .01$; females vs males) but higher expression in females compared with males after an injury with fadrozole (34.38 ± 6.17 vs 20.69 ± 1.95 , $P = .05$; females vs males). Data, presented in picograms per milligram, are shown in Figure 5.

Experiment 3: Expression of TNF- α , IL-1 β , IL-6, cox-2, and PGE2 after E₂ replacement

Tumor necrosis factor- α

Replacement with E₂ did not affect TNF- α expression ($F[1, 8] = 4.148$, $P = .76$), but significant effects of sex ($F[1, 8] = 18.82$, $P < .01$) and its interaction with treatment were found ($F[1, 8] = 7.87$, $P = .02$). Overall, TNF- α expression was higher in males relative to females (4.63 ± 0.24 vs 6.16 ± 0.42 , respectively), independent of treatment. The interaction of treatment and sex reflects that E₂ replacement compared with just aromatase inhibition lowered TNF- α expression in males (4.99 ± 0.29 vs 4.01 ± 0.33 ; $P = .02$; E₂ vs fadrozole) but had no such

effect in females (6.11 ± 0.27 vs 6.22 ± 0.11 ; $P = .63$; E₂ vs fadrozole). Means \pm SEM of δ Ct and fold changes are presented in Figures 6A and 7, respectively.

Interleukin-1 β

IL-1 β expression showed no effect of E₂ replacement ($F[1, 8] = 4.30$, $P = .07$) but an effect of sex ($F[1, 8] = 20.09$, $P < .01$) and its interaction with treatment ($F[1, 8] = 6.12$, $P = .03$). Overall, females had higher IL-1 β expression compared with males independent of treatment (8.07 ± 0.18 vs 9.84 ± 0.17). The interaction of treatment \times sex is driven by females (8.46 ± 0.08 vs 7.59 ± 0.26 , $P = .03$; E₂ vs fadrozole) but not males (9.98 ± 0.29 vs 9.81 ± 0.22 ,

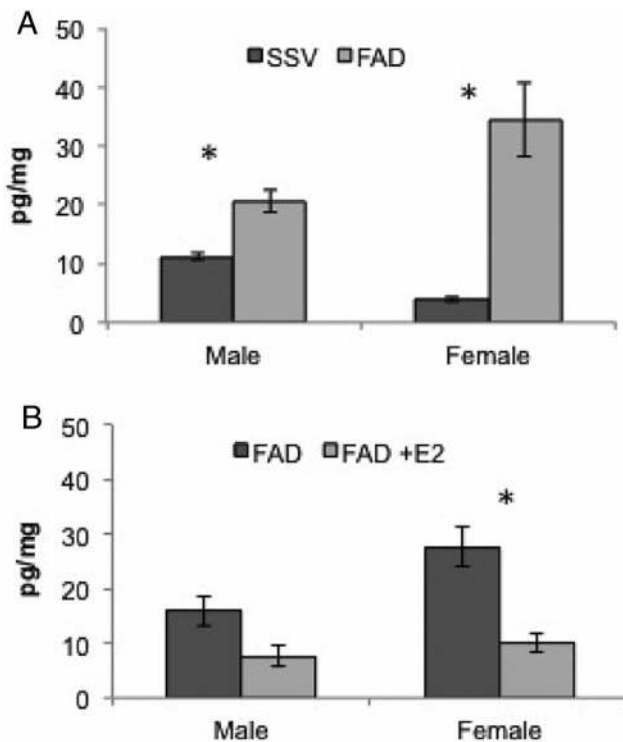


Figure 5. Levels of PGE2 after injury with fadrozole (FAD) administration in adult male and female zebra finches (A) or after injury with fadrozole administration and central E₂ replacement (B) in adult male and female zebra finches. Fadrozole administration resulted in increased levels of PGE2 in both male and females compared with injured controls. Injury with central E₂ replacement rescued PGE2 levels in females, and there was a trend for males. *, $P < .05$.

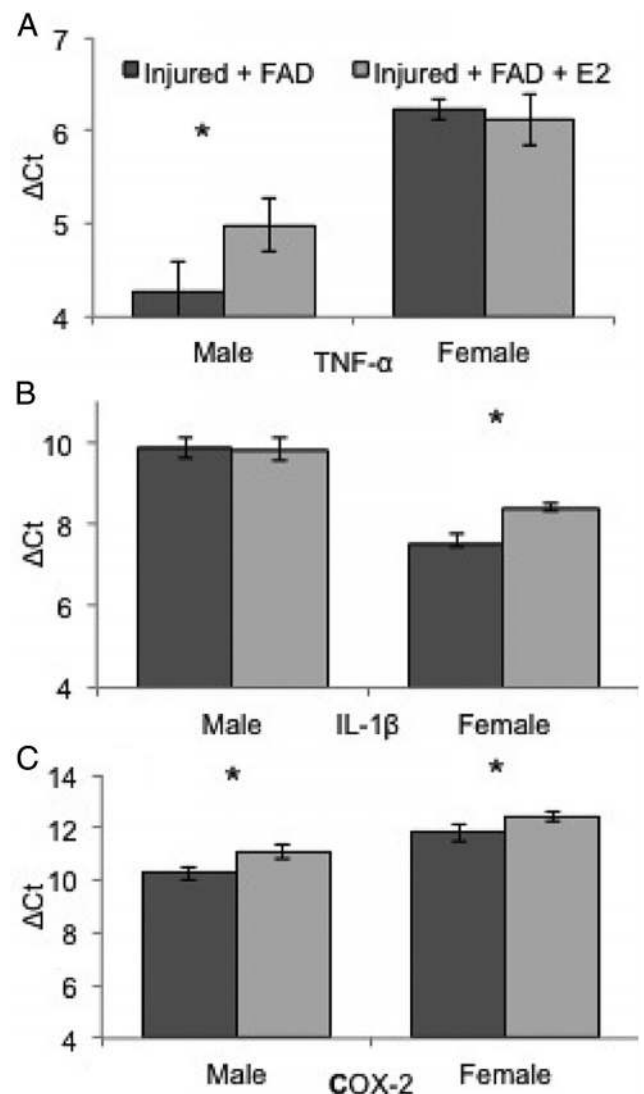


Figure 6. Levels of TNF- α , IL-1 β , and cox-2 mRNA relative to GAPDH (δ Ct values \pm SEM) in adult male and female zebra finches from experiment 3. Only significant gene targets are shown. E₂ replacement decreased TNF- α in males (A) and IL-1 β in females (B), and cox-2 in both sexes (C). In addition, males have higher overall levels of TNF- α (A) and cox-2 (C). Females had higher levels of and IL-1 β , regardless of treatment (B). *, $P < .05$. Note: y-axes do not begin at zero. FAD, fadrozole.

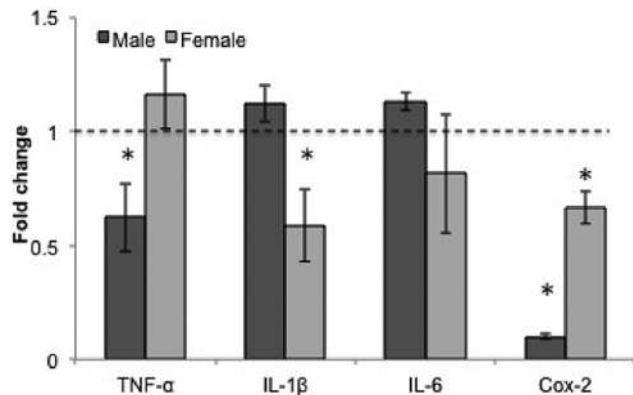


Figure 7. Levels of cytokine mRNA relative to GAPDH (fold change values \pm SEM) after an injury with fadrozole administration and central E₂ replacement in adult male and female zebra finches. Injury with central E₂ replacement rescued sustained TNF- α elevation after fadrozole in males and sustained IL-1 β in females. It also rescued sustained cox-2 elevation in both sexes. Dashed line represents injured aromatase inhibited fadrozole controls (controls set to 1 for fold change calculation). *, $P < .05$.

$P = .78$; E₂ vs fadrozole), showing decreases in IL-1 β mRNA after an injury with E₂ replacement, compared with fadrozole-injured birds. Means \pm SEM of δ Ct and fold changes are presented in Figures 6B and 7, respectively.

Interleukin-6

Once again, no significant effects on IL-6 expression were found (treatment: $F [1, 8] = 0.94$, $P = .36$; sex: $F [1, 8] = 2.65$, $P = .14$; treatment \times sex: $F [1, 8] = 0.27$, $P = .13$; see Figure 7).

Cyclooxygenase-2

Replacement with E₂ after an injury affected cox-2 expression with significant effects of treatment ($F [1, 8] = 52.8$, $P < .01$), sex ($F [1, 8] = 77.5$, $P < .01$), and a significant interaction of treatment \times sex ($F [1, 8] = 25.2$, $P < .01$). Both sexes showed decreases in cox-2 levels after an injury with E₂ treatment compared with fadrozole alone (11.75 ± 0.26 vs 10.63 ± 0.43 , respectively). In addition, males had higher cox-2 expression compared with females independent of treatment (10.27 ± 0.31 vs 11.11 ± 0.20). The interaction of these variables reflects a marginally stronger decrease following E₂ replacement compared with fadrozole in males (11.07 vs 10.26 , respectively; $P = .05$; E₂ vs fadrozole) relative to that seen in females (12.42 ± 0.3 vs 11.9 ± 0.17 , respectively, $P = .07$; E₂ vs fadrozole). Means \pm SEM of δ Ct and fold changes are presented in Figures 6C and 7, respectively.

Prostaglandin E2

In partial agreement with the data above, PGE₂ levels were affected by E₂ replacement ($F [1, 8] = 29.5$ $P < .01$)

and sex ($F [1, 8] = 6.58$ $P < .03$) but not by the interaction of these variables ($F [1, 8] = 3.78$ $P < .08$). PGE₂ levels were lower in lobes receiving E₂ replacement compared with just fadrozole (8.80 ± 1.27 vs 21.79 ± 2.28 , respectively) independent of sex. Additionally, independent of treatment, females had higher levels of PGE₂ compared with males (18.81 ± 3.45 vs 11.77 ± 2.12). Data, presented in picograms per milligram, are shown in Figure 5.

Discussion

In adult zebra finches of both sexes, we found that postinjury elevation of TNF- α and IL-1 β was low when aromatase was elevated (experiment 1), remained elevated upon pharmacological inhibition of aromatase activity (experiment 2), and did so in an E₂-dependent manner (experiment 3). In addition, inhibition of induced aromatase sustained the expression of, whereas replacement with E₂ appeared to lower, cox-2 and PGE₂ expression, respectively. Most effects were apparent in both sexes except that E₂ replacement decreased PGE₂ levels in females, but its effect in males was not significant ($P = .09$). These data strongly support the idea that E₂, synthesized in response to brain damage, is a potent anti-inflammatory signal that may prevent the CNS from the effects of chronic neuroinflammation.

In the zebra finch, there is extensive documentation of injury and inflammation-dependent aromatase expression in cells of astrocytic morphology (18, 19, 23) and those that coexpress the astrocytic marker vimentin (20–22, 24). Importantly, these changes in astrocytic aromatase expression are in excellent agreement with penetrating brain injury and inflammation-induced increases in aromatase transcription (20, 21). Thus, we have good reason to believe that the observed changes in aromatase transcription and the manipulation of its activity in the current set of studies reflect the injury-dependent expression of this enzyme in reactive astrocytes around the site of damage that has been reported over a decade (20, 21, 46). However, because the present studies rely on homogenates from telencephalic regions that express abundant neuronal aromatase, we cannot exclude this source of E₂ as a contributor to the effects described. Further studies that restrict the samples to just the injured entopallium (where aromatase is exclusively glial) are necessary to unequivocally test the specific effects of astrocytic aromatase to neuroinflammation.

Similarly, whereas measurement of cytokine expression in the current study was limited to transcript, we hypothesize that these cytokines are indeed translated into protein and secreted. Administration of fadrozole in-

creased, and replacement with E_2 lowered, the expression of *cox-2* and *PGE2*, respectively. There is excellent agreement between the expression of cytokines and that of *cox-2* in the present studies. Given these similarities and because *cox-2* expression and activity represents a point farther downstream in signaling, (31–33), it is very likely that the changes in cytokine transcription measured in the current studies do result from the activity of multiple elements of the inflammatory signaling cascade. Indeed, the measured changes in *PGE2* protein are strong evidence of robust cytokine secretion and activation of the inflammatory cascade including the expression and activity of *cox-2*. Taken together, the data strongly suggest that injury-induced E_2 synthesis is a potent modulator of neuroinflammation associated with brain damage from a penetrating brain injury in the finch brain.

There is substantial evidence supporting a role for circulating E_2 as an anti-inflammatory in vertebrates. Indices of inflammation are higher in postmenopausal women and ovariectomized mice compared with premenopausal, age-matched controls and intact animals, respectively. Specifically, expression and secretion of $TNF-\alpha$, $IL-1\beta$, and $IL-6$ are higher at times of low circulating E_2 relative to controls, as is the expression of their cognate receptors (39). The present data extend these findings to the brain by demonstrating a role for injury-induced aromatization within the CNS, one that involves a potent inhibition of multiple components of the inflammatory cascade within neural tissue.

Interestingly, although similar in general, we found some intriguing differences in the anti-inflammatory effects of E_2 across sexes. When aromatase is inhibited, females have a prolonged elevation of $TNF-\alpha$ and $IL-1\beta$, whereas only $TNF-\alpha$ remains high in males (Figure 4). In partial agreement, E_2 administration lowers $TNF-\alpha$ in males and $IL-1\beta$ in females but not vice versa (Figure 7). Previous studies have hypothesized that cytokines may serve different biological functions in men and women (58), and the rate of injury-induced aromatase expression differs between sexes in songbirds (21). Thus, it is likely that E_2 manipulation may affect inflammation in a sexually differentiated manner. Experiments varying severities of injury and time points of injury need to be explored to increase confidence in this interpretation. Furthermore, no experiments in the current study detected differences in $IL-6$ in either sex. Although we did not see changes in $IL-6$, previous studies have shown that $IL-6$ protein peaks 4 hours after penetrating brain injury (20). It is possible that $IL-6$ would have been affected by injury or treatment manipulation at a different time point. Alternatively, there may be specific, highly regulated mechanisms promoting E_2 's anti-inflammatory properties in the zebra finch brain.

It is difficult, however, to extract overall patterns regarding sex differences, response to injury or E_2 -dependent signaling, from these studies because only one experimental group is shared across experiments. Specifically, although the injured lobe 24 hours after the damage and the SSV lobe are similar between experiments 1 and 2, it is the fadrozole group that is similar between experiments 2 and 3. In general, however, interactions between injury-induced aromatization and inflammation appear to demonstrate strong effects on $TNF-\alpha$ expression in the male brain but stronger effects on $IL-1\beta$ expression in the female brain. First, 24 hours after the injury, $TNF-\alpha$ appeared to return to lower levels in the male brain relative to the female brain (experiment 1), and the effect of E_2 replacement affected $TNF-\alpha$ only in males (experiment 3). In contrast, the female brain responded to aromatase inhibition with higher levels of $IL-1\beta$ expression compared with males (experiment 2); and $IL-1\beta$, but not $TNF-\alpha$, was affected by E_2 replacement (experiment 3).

Because these studies did not contain any measures of unmanipulated birds, it is difficult to know whether these differences are due to baseline differences in cytokine expression or differences in responsiveness of the inflammatory cascade to injury and/or aromatization. The latter hypothesis is supported by a report of a more rapid induction of aromatase in the female brain after an injury (21), suggesting that E_2 -dependent effects may be temporally distinct between sexes. Potential differences in the temporal patterns of response to injury and inflammation may prove even more important in terms of the interaction between *PGE2* and aromatase. In rodents, aromatase is sensitive to *PGE2* both developmentally and in adulthood (59, 60), perhaps via a distinct binding site on the aromatase promoter (61). This interaction may prove key in the dynamics of injury-induced aromatase expression, the feedback effects of locally synthesized E_2 , and their modulation by *PGE2*. Current work in our laboratory is testing these hypotheses (56).

Nevertheless, in both sexes, aromatization and consequent E_2 synthesis affects signaling cascades downstream of cytokine expression, as evidenced by changes in the expression of *cox-2* and central *PGE2*. Thus, independent of the particular cytokine(s) affected, the present data suggest that the inflammatory cascade is potentially affected by aromatization in both sexes. Current studies in our laboratory focus on the mechanisms whereby E_2 may affect microglial cytokine synthesis, and furthermore, how these cytokines may impact inflammatory cascades in neurons and other cells within the CNS.

Whereas E_2 appears to be a potent, inducible anti-inflammatory after an injury, the mechanism by which it decreases cytokine activity is unknown. Microglia, the

producers of cytokines after injury, contain classical estrogen receptors (ER)- α and ER- β and also the nonclassical receptor G protein-coupled receptor-30. These receptors are up-regulated after neural damage in mammals (62, 63), which may help E₂ reduce cytokine production and therefore limit neural damage. However, whether passerine microglia express ERs is unclear and requires further study.

In summary, these experiments suggest that induced E₂ synthesis via astrocytic aromatization is a potent anti-inflammatory in the brain. E₂ synthesis may impact inflammatory expression and secretion after injury to the brain, thereby reducing the deleterious effects of chronic neuroinflammation. We believe this is the first demonstration of induced steroid synthesis acting as an anti-inflammatory in the vertebrate brain.

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