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Brain Microglia Express Steroid-Converting Enzymes in the Mouse

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Abstract

In the CNS, steroid hormones play a major role in the maintenance of brain homeostasis and its response to injury. Since activated microglia are the pivotal immune cell involved in neurodegeneration, we investigated the possibility that microglia provide a discrete source for the metabolism of active steroid hormones. Using RT-PCR, our results showed that mouse microglia expressed mRNA for 17 β -hydroxysteroid dehydrogenase type-1 and steroid 5 α -reductase type-1, which are involved in the metabolism of androgens and estrogens. Microglia also expressed the peripheral benzodiazepine receptor and steroid acute regulatory protein; however, the enzymes required for *de novo* formation of progesterone and DHEA from cholesterol were not expressed. To test the function of these enzymes, primary microglia cultures were incubated with steroid precursors, DHEA and AD. Microglia preferentially produced delta-5 androgens (Adiol) from DHEA and 5 α -reduced androgens from AD. Adiol behaved as an effective estrogen receptor agonist in neuronal cells. Activation of microglia with pro-inflammatory factors, LPS and INF γ did not affect the enzymatic properties of these proteins. However, PBR ligands reduced TNF α production signifying an immunomodulatory role for PBR. Collectively, our results suggest that microglia utilize steroid-converting enzymes and related proteins to influence inflammation and neurodegeneration within microenvironments of the brain.

Keywords

LPS; cytokines; neurosteroids; steroidogenesis; PBR; DHEA; Adiol; estrogen

1. Introduction

The brain is under the constant influence of steroid hormones. Among these, sex hormones are known to affect multiple CNS processes such as the differentiation and maturation of specific brain regions, learning and behavior, neurogenesis, neurotransmission, synaptogenesis, and neuroprotection [1–7]. Sex hormones are derived from the steroid precursors dehydroepiandrosterone (DHEA) and androstenedione (AD), which originate from the

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metabolism of cholesterol and its further conversion to pregnenolone [8,9]. At one time sex steroids were thought to originate exclusively from the gonads and the adrenals; however, it is now accepted that local steroid synthesis occurs in a number of tissues [10], including the central nervous system [11]. It is further thought that steroids metabolized in the nervous system exert intracrine and paracrine effects on homeostasis, as suggested from studies in hippocampal neurotransmission [12–14], cerebellar development [15], or neuroprotection following neurotoxic challenge [16,17].

Certain types of brain cells are capable of metabolizing steroids [18], yet the role of microglia in this process has not been studied. Microglia are the resident brain macrophages, and as such have important immunological and pathological functions [19]. Through their production of cytokines and superoxides, activated microglia may underlie pathological processes such as Alzheimer's disease [20], Parkinson's disease [21], and multiple sclerosis [22].

Steroid metabolism in the CNS has been implicated as an adaptive coping mechanism following brain damage [3]. Additionally, neurodegenerative conditions have been negatively correlated with brain steroid levels [23]. As androgens and estrogens can exert pro- or anti-inflammatory actions [24–26], the steroid profile in a tissue may determine the extent of inflammation [27]. While microglia are critical mediators of brain inflammation, it is unknown if they participate in the steroid-metabolizing actions of the brain.

In previous studies we have reported that the microglia cell line, BV2, exhibits steroid-converting capacity [28–30]. Given the neuroprotective properties of steroids and the critical role that microglia play during brain damage responses, and in the steady state [31,32], we have characterized the expression of the main proteins and enzymes in the steroidogenic pathway by real-time PCR (RT-PCR) from *ex vivo* isolated microglia and primary cultured cells. The function of key enzymes involved in steroid metabolism was assessed by the metabolic conversion of the sex steroid precursors DHEA and AD and analysis of their metabolites by thin layer chromatography (TLC). Our results indicate that microglia express steroid-converting enzymes that contribute to the metabolism of steroids into active androgens and estrogens within the brain.

2. Materials and Methods

2.1. Animals

For these studies, the transgenic mouse line p7.2fms-EGFP (C57BL6/6 X CBA background) was used [33]. Enhanced green fluorescent protein (EGFP) expression is driven by the promoter and the regulatory elements of the *c-fms* gene that encodes the receptor for macrophage colony stimulating factor (CSF-1), resulting in EGFP expression in cells of the mononuclear phagocytic lineage, including microglia [33]. Animals were bred in Rockefeller University facilities under 12:12 light:dark cycle and free access to chow and water. All experimental procedures were approved by the Rockefeller University Animal Care and Use Committee. To induce inflammation, mice received a single injection with *Salmonella typhimurium* lipopolysaccharides (LPS; 1–5 mg/kg, i.p.; Sigma, L2262; St. Louis, MO).

2.2. Ex vivo Microglia Isolation by fluorescence activated cell sorting (FACS)

Previously reported methods to obtain a single population of microglia by FACS were used [34]. In brief, adult mice (2–3 months of age) were anaesthetized with pentobarbital (750mg/kg) and rapidly decapitated. Brains were removed and placed on ice in Hank's balanced salt solution (Gibco, Carlsbad, CA), and meninges, blood vessels and choroid plexus were carefully removed under a dissecting scope. Brain cell suspensions, obtained after incubation with type II-S collagenase (600U; Sigma) and DNase (450U; Invitrogen, Carlsbad, CA) for 30min at

37°C in 15ml HBSS supplemented with 90 mM CaCl₂, were homogenized by repetitive gentle pipetting with fire-polished Pasteur pipettes on ice, followed by filtering through a 40 mm cell strainer (BD, Franklin Lakes, NJ). Cells were washed by centrifugation and subject to Percoll gradient centrifugation as described previously [34]. Cells collected from the 30/70 interphase, were washed and re-suspended in 5% fetal calf serum (FCS)-PBS containing 100ng/ml propidium iodide (PI), before sorting in a FACS Vantage SE Flow Cytometer (BD), with smHighPurity precision. Post-sort analysis was performed to ensure the purity of the collection process.

2.3. Primary microglia (1 °MG) cultures and cell stimulation

Microglia cultures were prepared following standard protocols (23). Briefly, day 2-old mouse pup brains were dissected on ice, and the meninges were carefully removed. The forebrains were minced in 5% FCS-PBS, dissociated using fire polished Pasteur pipettes, and then passed through a 40mm nylon cell strainer (BD). Cells were washed once in buffer and seeded in culture media at a density of roughly two forebrains per 75mm flask. Culture media, 10%FCS DMEM (Gibco), was changed every 5 days and supplemented with 5ng/ml granulocyte-monocyte colony stimulating factor (GM-CSF; Sigma). After 2 weeks in culture at 37°C, 5 % CO₂, cells were shaken at 125rpm for 5hrs at 37°C to harvest detached microglia. Microglia were counted and seeded in 6-well plates at a density of 1 million cells/well (for RNA-PCR) or in 24-well assay plates at a density of 0.25–0.3 million cells/well (for hormone metabolism-TLC, and cytokine assays). After plating, microglia were allowed to adhere for 1hr and then rinsed to remove non-adherent glial cells, fed and incubated as described above. The following day, cells were rinsed and incubated in DMEM for 24hr at 37°C with vehicle or the various treatments: LPS, interferon- γ (INF γ) (Sigma), dibutyryl-cyclic adenosine monophosphate (db-cAMP) (Sigma), (see results). For PBR ligand stimulation, cultures were pre-treated for 10 minutes with Ro 5-4864 (Ro) and PK-11195 (PK) (Sigma), or corticosterone (Sigma) as positive control, before LPS+INF γ stimulation. INF γ was supplemented to LPS to obtain a robust nitric oxide (NO) response. LPS+INF γ conditioned media (LCM) was obtained by stimulating 1°MG cultures with 1%FCS DMEM plus 100ng/ml LPS+ 10ng/ml INF γ , collecting the supernatants 24hr later, and centrifuging at 2000rpm for 5min to clear any debris. LCM contains elevated levels of several inflammatory cytokines such as TNF α , IL-6 and NO, as well as IL1 β , IL-12, MCP-1, MCP-5, and RANTES (Gottfried-Blackmore, unpublished results). This LCM was used to stimulate fresh cultures of 1°MG.

2.4. FACS Analysis of 1 °MG

After shaking for 5hr, microglia were collected and washed in FACS buffer (5% FCS PBS). Cells were then blocked for 15 min at 4°C with 5% mouse serum. Cells were then stained for 15 min at 4°C with phycoerythrin (PE) conjugated anti-CD11b (1:200) (BD), or its corresponding PE-conjugated isotype, anti-rat IgG2b (1:200) (BD). Finally, cells were washed 3X in FACS buffer and then analyzed using a BD FACSCalibur system (BD) under the FITC and PE channels. Data was analyzed using FlowJo software (Tree Star Inc., OR).

2.5. Real-time PCR

Adult microglia were sorted by FACS into RNA lysis buffer (Absolute RNA Microprep kit (Stratagene, La Jolla, CA)), frozen in dry ice, and kept at -80°C until processing. RNA was isolated using the Absolute RNA Microprep kit. RNA from 1°MG cultures was obtained by rinsing the cell cultures with PBS and then lysing cells in 350 μ l RNA lysis buffer (RNeasy Mini kit (Qiagen, Valencia, CA)). RNA was extracted using the RNeasy Mini kit protocol (Qiagen). Both extraction methods included a step with DNase incubation (Qiagen). RNA was quantified with RiboGreen RNA Quantitation kit (Molecular Probes), following manufacturer instructions. 10ng of RNA were retrotranscribed with SuperScript II Reverse Transcriptase

(Invitrogen) and 3 μ l of a 1:3 dilution of the cDNA were amplified by real-time PCR using SYBR Green master mix (AB) in a 7900HT SDS thermal cycler (AB). The steroidogenic protein transcripts quantified were peripheral benzodiazepine receptor (PBR), steroidogenic acute regulatory protein (StAR), cytochrome p450 side chain cleavage enzyme (p450scc), cytochrome p450 21-hydroxylase (p450c21), cytochrome p450 17-hydroxylase (p450c17), aryl sulfatase (Arsa), DHEA sulfotransferase (Sult), 3 β -hydroxysteroid dehydrogenase (isoforms 1,2,4) (3 β HSD1,2,4), 17 β -hydroxysteroid dehydrogenase type 1 (17 β HSD1), cytochrome p450 aromatase (P450Arom), steroid 5 α -reductase type 1 (5 α R), 3 α -hydroxysteroid dehydrogenase (3 α HSD), and ribosomal protein L27A (L27A). Primers sequences were designed using Primer Express Software (ABI) and are indicated in Table 1; amplicons span at least one intron in order to avoid potential genomic DNA amplification. All primers were blasted on NCBI databases for target specificity and tested using appropriate positive control tissues such as ovary and adrenal glands. Each sample was run in triplicate and the average Ct (threshold cycle) was used to calculate the relative amount of product by the $-\Delta\Delta$ Ct-method (AB), using the ribosomal L27A as a housekeeping gene. The ratio of enzyme Ct to L27A Ct values was calculated as a way of assessing the relative expression levels of each enzyme comparing 1 $^{\circ}$ MG and *ex vivo* MG. In each experiment, both positive (1 μ g ovary mRNA/cDNA1:3) and negative (RT minus and water) controls were included to ensure that the PCR reaction was working properly.

2.6. Incubation of tritiated hormones and steroid extraction

The day after seeding, cells were rinsed and incubated in 0.2ml DMEM containing [1,2,6,7- 3 H] DHEA (60 Ci/mmol) or [1,2,6,7- 3 H] AD (105 Ci/mmol) (Perkin Elmer Life Science, Shelton, CT) for 22–24hr. All incubations were conducted in a 5% CO₂ atmosphere at 37 $^{\circ}$ C. The reaction was stopped by vortexing the supernatant with acetone (0.2 ml) and ethyl acetate (0.5ml). A 0.2 ml portion of the organic phase extract was evaporated to dryness at room temperature, the residue was re-dissolved in methanol, and the yield of metabolites was determined after separation by thin layer chromatography (TLC).

2.7. TLC Identification of Metabolites

Products of tritiated steroid incubations were separated by TLC on silica gel containing a fluorescent indicator on pre-coated aluminum sheets (Fisher, Carlsbad, CA) using chloroform/ethyl-acetate/xylene (68/23/9 % by vol. for [3 H] DHEA) and (62/21/17 % by vol. for [3 H] AD). Non-radioactive steroids of known identity were added to the TLC sheets on lanes adjacent to the putative metabolites and were visualized and identified by their chromogenic properties after spraying with 5% (by vol.) sulphuric acid in methanol and heating on a hot plate. Unlabeled steroids E2 and E1 were purchased from Sigma, and T, AD, 5 α AD, DHEA, Adiol from Steraloids Inc. Purity of [3 H]-DHEA and [3 H]-AD (<98 %) was determined by TLC.

2.8. Cytokine and Nitric Oxide detection

After 24hr stimulation with PBR ligands and LPS+INF γ , 1 $^{\circ}$ MG supernatants were collected, centrifuged (2,500rpm) for 5 minutes at 4 $^{\circ}$ C in a tabletop microfuge, and then frozen at -20 $^{\circ}$ C. For detection of TNF α and IL-6, ELISA was performed according to manufacturer's instructions (eBioscience, SanDiego, CA). Nitric Oxide (NO) was determined indirectly though the Greiss Assay (Promega, Madison, WI).

2.9. Transfection and Luciferase Assay

EtC.1 cells were seeded in 24-well plates, 2 \times 10⁴ cells/well, in DMEM containing 10% charcoal stripped serum. 24hr later cells were transfected using Lipofectamine Plus (Invitrogen), following manufacturer's instructions, with 0.4 μ g of plasmid DNA/well. The 3X ERE-Luciferase plasmid was a generous gift of Don McDonnell (Duke University Durham, NC),

and the β -Galactosidase plasmid was from Promega (pSV- β -Gal control vector). 24hr after transfection, cells were incubated with various concentrations of E2 or Adiol for another 24hr. 100nM ICI pre-treatment was done for 30 minutes. Cell lysates were prepared and luciferase activity measured using Promega Luciferase Assay System according to manufacturer's instructions (Promega). β -Gal activity was measured from cell lysates to normalize for transfection efficiency.

2.10. Statistical analysis

Statistical analysis was performed using StatView (SAS Institute Inc., Cary, NC). Experiments involving 2 groups were compared using a Student t-test. Experiments involving more than 2 groups were compared by Analysis of Variance (ANOVA), followed by posthoc analysis with Tukey Honestly Significant Difference (HSD) when variances were homogeneous (using Levene's test); or with nonparametric Games-Howell test. Graphs show the mean \pm the standard error of the mean (S.E.M.). $P < 0.05$ was considered statistically significant. *, $p < 0.05$, **, $p < 0.01$ and ***, $p < 0.0001$; (n.s.), non-significant.

3. Results

3.1. Purity of microglia assays

Exclusion of brain cells with high steroid-converting capacity, such as astrocytes [18], from our experiments with 1° MG cultures was verified by EGFP expression and by FACS analysis. Cultured 1° MG showed 99–98% purity by EGFP expression and staining with the macrophage-specific marker CD11b (Fig.1). Therefore, gene expression and enzymatic activity in our assays can be attributed mainly to microglia.

3.2. Gene expression of steroid-converting enzymes in microglia

To investigate the expression of the main steroid-converting enzymes in mouse microglia, gene specific primers were designed for Real Time PCR (RT-PCR) (Table 1). Some of these enzymes have multiple isoforms (i.e. 3β -hydroxysteroid dehydrogenase has 7 isoforms in mouse). Selection of isoforms in this study was based on reported predominance (common vs. rare) and activity (dehydrogenase vs. oxo-reductase). RT-PCR analysis of RNA from adult, FACS-sorted, microglia (*ex vivo* MG) and from primary microglia cultures (1° MG) revealed positive expression of the peripheral benzodiazepine receptor (PBR), steroid acute regulatory protein (StAR), steroid (StS) and aryl (Arsa) sulfatase, 17β -hydroxysteroid dehydrogenase type 1 (17β HSD1), and 5α -reductase type 1 (5α R) (Table 2). However, cytochrome p450 side chain cleavage enzyme (p450scc), cytochrome p450 21-hydroxylase (p450c21), cytochrome p450 17-hydroxylase (p450c17), DHEA sulfotransferase (Sultt), 3β HSD type 1, 2, and 4, cytochrome p450 aromatase (P450Arom), and 3α -hydroxysteroid dehydrogenase (3α HSD) were not detected (Table 2). The ratio of specific gene C_t values to the C_t value of L27A, a housekeeping gene, was calculated as a way of assessing the relative expression levels of each gene within *ex vivo* MG and 1° MG (Table 2). The most abundantly expressed genes were the PBR (ratio of 0.80; 0.94) and Arsa (ratio of 0.85; 0.83), while the lowest expression was found for StAR (ratio of 0.68; 0.62) (Table 2). No differences were found between the expression levels of *ex vivo* MG and 1° MG (Student T-test; $p = 0.16$ – 0.64), except for the PBR ($p < 0.0001$) and StS ($p < 0.01$), which were expressed at higher levels in 1° MG (Table.2). A summary of the expression of steroidogenic proteins in microglia is depicted in Figure 2.

3.3. Inflammatory LPS stimulation modulates expression of steroid-converting enzymes in microglia

Intraperitoneal injections (i.p.) of LPS into mice leads to the activation of brain microglia, as indicated by changes in morphology and the expression of pro-inflammatory cytokines [34,

35]. Adult *cfms*-EGFP male mice were injected i.p. with saline or 1mg/kg LPS to evaluate the expression of the principal steroid-converting enzymes in activated microglia. Twenty-four hour *in vivo* stimulation with LPS led to an overall decrease in the expression of steroid-converting enzymes (Fig.3). StAR expression was decreased by $38.8 \pm 9.3\%$ ($p < 0.05$), 17β HSD1 by $71.4 \pm 0.8\%$ ($p < 0.05$), and 5α R by $66.6 \pm 1.8\%$ ($p < 0.05$) (Fig. 3). One notable exception was the PBR, whose expression was significantly increased 7.7 \pm 2.2-fold ($p < 0.05$) (Fig.3). However, i.p. injection of LPS did not induce the expression of p450scc, p450c21, p450c17, Sulft, 3β HSD1,2,4, P450Arom, or 3α HSD in *ex vivo* microglia.

Since i.p. injection of LPS is known to induce high levels of circulating cytokines, we sought to assess whether the down regulation of steroid enzymes in microglia was due to a direct or indirect effect of LPS. 1° MG cultures were stimulated *in vitro* for 24hrs with different pro-inflammatory stimuli (LPS+INF γ , INF γ , and LPS-conditioned media (LCM), see methods) and cells were processed for RT-PCR analysis (Fig.4).

Stimulation with LPS+INF γ led to an increase in the expression of all steroid-converting enzymes expressed in 1° MG (Fig.4A–D). PBR expression was increased 9.6 \pm 1.0-fold ($p < 0.01$) (Fig.4A), StAR increased 4.5 \pm 0.7-fold ($p < 0.05$) (Fig.4B), 17β HSD1 increased 3.3 \pm 0.4-fold ($p < 0.01$) (Fig.4C), and 5α R was increased 5.1 \pm 0.6-fold ($p < 0.01$) (Fig.4D). INF γ stimulation alone had no significant effect (Fig.4) and LCM did not induce a down-regulation of steroid-converting enzymes. Instead, it mimicked the LPS+INF γ effect on 1° MG, but to a lesser extent (Fig.4). Finally, db-cAMP, a known inducer of steroidogenesis in various endocrine cells, was tested on 1° MG, yet it presented no effect on the expression of steroid-converting enzymes in 1° MG (Fig.4). None of the treatments in 1° MG induced the expression of p450scc, p450c21, p450c17, Sulft, 3β HSD1,2,4, P450Arom, or 3α HSD (data not shown).

3.4. Steroid-converting activity in microglia

The results from our RT-PCR analysis suggested that microglia express enzymes capable of metabolizing DHEA, androgens, and estrogens (Fig.2). The steroid-converting activity of microglia was evaluated by incubating 1° MG with the steroid intermediate dehydroepiandrosterone (DHEA) and measuring its metabolism. After 24hr incubation, 1° MG showed a 30 \pm 3.3% ($p < 0.0001$) conversion of H³-DHEA, with Adiol being the only product detected of this conversion, accounting for 15.5 \pm 1.3 ($p < 0.0001$) of H³-radioactivity (Fig. 5A). Gene expression experiments indicated that LPS+INF γ increased expression of steroid converting enzymes; however, microglia stimulation with LPS+INF γ had no effect on the conversion or profile of steroids produced from H³-DHEA (Fig.5B).

Androstenedione (AD) is the main intermediary product from DHEA in the synthesis of androgens and estrogens. 1° MG incubated with H³-AD for 24hr showed a 36.7 \pm 1.7% ($p < 0.0001$) conversion of this steroid (Fig.6A), with the main products of its conversion being T, 12.8 \pm 0.8% ($p < 0.0001$); 5α AD, 3.8 \pm 0.8% ($p < 0.01$); and DHT, 14.4 \pm 3.1% ($p < 0.01$) (Fig. 6A). Further, stimulation of the cells using LPS+INF γ did not affect the conversion or the profile of steroids produced from H³-AD (Fig.6B).

3.5. PBR role in microglia

Of the genes studied, PBR was the most abundant transcript detected in microglia; additionally, it showed a robust induction by LPS+INF γ . The absence of cholesterol metabolizing enzymes, i.e. low levels of StAR and absence of p450scc and p450c17, suggest that PBR may play an alternate role in microglia. Indeed, the PBR has been associated with numerous biological functions such as cellular proliferation, porphyrin transport, heme biosynthesis, anion transport, apoptosis, and immunomodulation [rev. by [36]]. Stimulation of microglia with two selective PBR ligands, Ro and PK, led to a specific reduction of LPS-induced production of

TNF α (16.2 \pm 10.8% decrease (p <0.05), and 40.9 \pm 8.9% (p <0.05) respectively), but had no effects on IL-6 and NO (Fig.7), except a 12.4 \pm 6.4% (p <0.05) increase by PK on NO (Fig.7). Despite the modulatory effects on these inflammatory products, PBR ligands had no effect the metabolism of DHEA or AD (data not shown).

3.6. Adiol is an effective estrogen receptor agonist

The sole product of DHEA metabolism in 1 $^{\circ}$ MG was Adiol. This steroid has been reported to have androgenic and estrogenic properties [37]. To determine whether Adiol could function as a specific estrogen receptor (ER) agonist, we utilized an ER-expressing neuronal cell line, EtC.1 [Gottfried-Blackmore, *et al.* submitted], transfected with a luciferase gene reporter coupled to 3 estrogen response elements (EREs). Incubation of these cells with Adiol induced the expression of the luciferase gene reporter at a dose of 1 μ M, but not at 10nM (Fig.8). Moreover, this induction was completely abrogated by pre-treatment of the cells with the specific ER-antagonist ICI-182,780 (100nM) (Fig.8). Microglia were not used for this assay because of their low expression of ERs and negligible responsiveness to E2 [Sierra *et al.* submitted].

4. Discussion

The involvement of sex hormones in neuroprotection is widely reported [38,39]. Additionally, steroid-converting enzymes are up-regulated following neuronal damage, and blocking such enzymes increases neuronal death in a number of injury models [40]. These studies indicate the critical role of steroid metabolism in the brain's responses to injury. Although the steroid-converting capacity of neurons, astrocytes and oligodendrocytes has been described to some extent, there are no reports addressing the contribution of microglia to brain steroidogenesis. In this study, microglia cells from adult and neonatal mice are reported to express steroid-converting enzymes and our data suggest microglia participate in the brain's steroid-converting capacity.

In the brain it has been hypothesized that neurosteroidogenesis occurs involving the *de novo* synthesis from cholesterol of steroid hormone precursors, such as pregnenolone, progesterone and DHEA. At present, it remains controversial that neurosteroidogenesis can occur given conflicting data on expression of p450c17 in the adult brain [41,42], required for synthesis of DHEA. Our results revealed that microglia do not express p450c17 nor p450scc, p450c21, or 3 β HSD1-2, and therefore, do not have the capacity to synthesize steroids *de novo* from cholesterol.

Alternate roles for steroidogenic proteins

Despite the lack of p450scc and p450c17, microglia expressed low levels of StAR mRNA, which is the main mediator for cholesterol transfer to the mitochondria for the initiation of steroidogenesis [43,44]. StAR expression in microglia may be playing a different role in these cells, such as cytosolic free sterol transfer, as has been suggested for other cholesterol binding proteins in macrophages [45]. Further studies will need to address this in microglia.

The PBR, like StAR, also participates in the transfer of cholesterol into the mitochondria [46]. In the brain, PBR expression is increased following nerve injury and can increase steroidogenesis locally [47]. In addition to its role in steroidogenesis [48], the PBR is reported to participate in other processes such as apoptosis, cancer, and immunomodulation [49–51]. This protein is widely expressed in monocytic cells [52], and has been identified as a marker of activated microglia [53]. Of the genes we studied, the most highly expressed in microglia was the PBR. In the CNS, PBR is mainly expressed in glial cells, and its expression levels increase following glial activation induced by inflammation or neuronal damage [rev. by

[36]]. 1°MG and *ex vivo* MG challenged with inflammatory stimuli responded with an increased expression of PBR, consistent with previous reports. However, PBR stimulation with specific ligands, which increase steroidogenesis [47,48], had no effect on DHEA or AD conversion in microglia. This may be due to the fact that these ligands affect the initial transfer of cholesterol into the mitochondria providing more substrate for p450_{scc}, which is lacking in microglia. However, in our study we found that PBR ligands selectively decreased the production of TNF α in response to LPS+INF γ stimulation in 1°MG, in accordance with previous reports [54–56]. Interestingly, PBR ligands did not affect IL-6 and only slightly affected NO indicating that PRB ligands may preferentially interfere with signaling pathways involved in the expression of TNF α . Our data further support an immunomodulatory role of PBR in microglia, which may account for the neuroprotective effects reported for its ligands [57,58].

Expression of steroid-converting enzymes and microglia LPS activation

Our results showed a significant reduction in all the steroid-converting enzymes expressed in *ex vivo* microglia after i.p. LPS stimulation. Systemic injection of LPS *in vivo* causes a rapid induction of circulating cytokines and inflammatory mediators [59], which can induce inflammatory genes in microglia [34,35]. *In vitro*, LPS+INF γ caused an increase in steroid-converting enzyme expression, while single cytokine stimulation (INF γ), or stimulation with a combination of cytokines (LPS-conditioned media) showed only marginal effects. Our results would suggest that other *in vivo* factors may be causing the down-regulation of steroid-converting enzymes in microglia. Following the systemic rise of cytokines, activation of the Hypothalamic-Pituitary-Adrenal axis causes a 3–4 fold increase in circulating glucocorticoid levels (rev. by [60]). Increased glucocorticoid levels can block steroidogenesis in testicular Leydig cells [61,62], and may have similar effects in brain microglia. We tested whether direct stimulation with corticosterone would decrease steroid-converting enzyme expression, however corticosterone failed to reduce StAR, 17 β HSD1, or 5 α R expression *in vitro* in both resting and LPS-activated microglia (data not shown). These results suggest other factors are involved in the *in vivo* reduction we observed. Another possibility for the discrepancy of the results is the developmental stage of the cells. Although primary microglia cultures are widely accepted to resemble adult cells, they may retain characteristics from developing cells. Such developmental differences could be pursued in future studies.

The LPS+INF γ activation of 1°MG caused increased enzyme expression *in vitro*, but did not affect the rate or metabolism of H³-DHEA and H³-AD conversion. The catalytic activity of several of these enzymes is determined by the redox potential set by the NADP⁺ to NADPH cofactor ratio [63]. Cofactor availability should be addressed in future studies comparing resting vs. activated microglia.

Macrophage secreted products, such as TNF α [64,65], IL1 β [66] and NO [67], can inhibit production of DHEA and androgens in the gonads via inhibition of p450_{c17} gene expression [68–70]. Our results indicate that 17 β HSD1 and 5 α R activity are not affected by inflammatory stimuli. This point is critical, considering that inflammatory conditions can dramatically reduce steroid metabolism in astrocytes co-cultured with microglia [41].

Steroid Metabolism in CNS Microglia

In this report, we show microglia expressed 17 β HSD1 and 5 α R, enzymes that are known to be involved in the metabolism of DHEA, androgens and estrogens. Conversion of H³-DHEA to AD did not occur in 1°MG consistent with absence of 3 β HSD; however these cells converted H³-AD into T, DHT, and 5 α AD, corroborating the activity of 17 β HSD1 and 5 α R in microglia. Similar findings have been reported in human alveolar macrophages where the principal metabolites of AD were 5 α AD and T [71]. 5 α R activity in microglia could be significant in

amplifying the actions of androgens, as DHT is three times more efficient than T [72] and is considered a non-aromatizable androgen. Microglia had no detectable expression of aromatase, in contrast to differentiated monocytes/myeloid cells that show low levels of estrogen formation [73,74].

In the CNS, DHEA has multiple effects reminiscent of sex hormones [75]. The absence of a specific DHEA receptor [76] has led investigators to suggest that DHEA is metabolized into active sex hormones that mediate these observed effects [28,30,74]. Neurons, astrocytes and oligodendrocytes can metabolize DHEA into sex hormones [18]. Recently we reported that the microglial cell line, BV2, is able to convert DHEA into Adiol and validated the identity of this product by high-performance liquid chromatography [30]. The current study corroborates these findings in 1°MG and shows the expression of 17 β HSD1, required for this conversion, both in *ex vivo* MG and 1°MG.

DHEA levels are higher in brain than in circulation and have been negatively correlated with aging and neurodegeneration [77]. Moreover, conversion of DHEA in the brain, into metabolites like Adiol, may be reduced in patients with neurodegenerative disease [78]. Our data indicate that both resting and activated microglia can metabolize DHEA and specifically convert it to Adiol. This delta-5 steroid has reported androgenic and estrogenic properties in peripheral tissues [37,79,80] and in the male rat pituitary [81]. In this study we present evidence confirming that Adiol is an effective estrogen receptor agonist in ER-expressing granule neuronal cells (EtC.1). These results are significant given that microglia are situated among ER-expressing neurons and astrocytes [Sierra *et al.* submitted] suggesting that DHEA metabolism by microglia may be a source of active estrogens in the brain.

Microglia also expressed StS and Arsa. DHEA-sulfate, a predominant sulfated steroid in the brain with neuro-excitatory properties [82], cannot be utilized for sex steroid production in its sulfated form. The expression of StS and Arsa in microglia, consistent with that found in macrophages [83], suggests that microglia could modulate the levels of locally available DHEA [84] for androgen and estrogen conversion. Further studies will be required to elucidate these roles of StS and Arsa in microglia.

In conclusion, this report demonstrates for the first time that microglia express steroid-converting enzymes. Our data suggest that microglia depend on available steroid precursors for the conversion of androgens, rather than presenting *de novo* synthesis of sex hormones from cholesterol. Microglia differ from neurons, astrocytes and oligodendrocytes in the expression of steroid-converting enzymes, particularly those involved in the *de novo* formation of progesterone and DHEA from cholesterol. It is likely that some of these proteins may have alternate roles in microglia, such as StAR, or PBR, as demonstrated by the ability of PBR ligands to reduce TNF α production. Although inflammatory microglia are implicated in neural pathology, particularly by their accumulation at neurodegenerative loci, their steroid-converting activity may add to the adaptive neuroprotective mechanisms of the brain following damage. Therefore, understanding the impact of microglia steroid metabolism and the factors involved in its regulation will allow for better interventions and therapeutic use of steroid hormones.

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Abbreviations

DHEA, 3 β -hydroxy-5-androstene-17-one (dehydroepiandrosterone)
 AD, 4-androstene-3,17-dione (androstenedione)
 5 α AD, 5 α -androstane-3,17-dione (5 α -Adione)
 Adiol, 5-androstene-3 β ,17 β -diol (Δ^5 -Adiol)
 T, 4-androsten-17 β -ol-3-one (testosterone)

DHT, 5 α -androstane-3-one-17 β -ol (dihydrotestosterone)
TLC, thin layer chromatography
RT-PCR, real-time polymerase chain reaction
PBR, peripheral benzodiazepine receptor
StAR, steroidogenic acute regulatory protein
P450scc, cytochrome p450 side chain cleavage enzyme
P450c21, cytochrome p450 21-hydroxylase
P450c17, cytochrome p450 17-hydroxylase
Arsa, aryl sulfatase
Sulft, DHEA sulfotransferase
3 β HSD1,2,4, 3 β -hydroxysteroid dehydrogenase isoforms 1,2,4
17 β HSD1, 17 β -hydroxysteroid dehydrogenase type 1
p450Arom, cytochrome p450 aromatase
5 α R, steroid 5 α -reductase type 1
3 α HSD, 3 α -hydroxysteroid dehydrogenase
L27A, ribosomal protein L27A
db-cAMP, dibutyryl cyclic AMP
LPS, lipopolysaccharide
INF γ , interferon gamma
NO, nitric oxide
ERE, estrogen response element
EGFP, enhanced green fluorescent protein

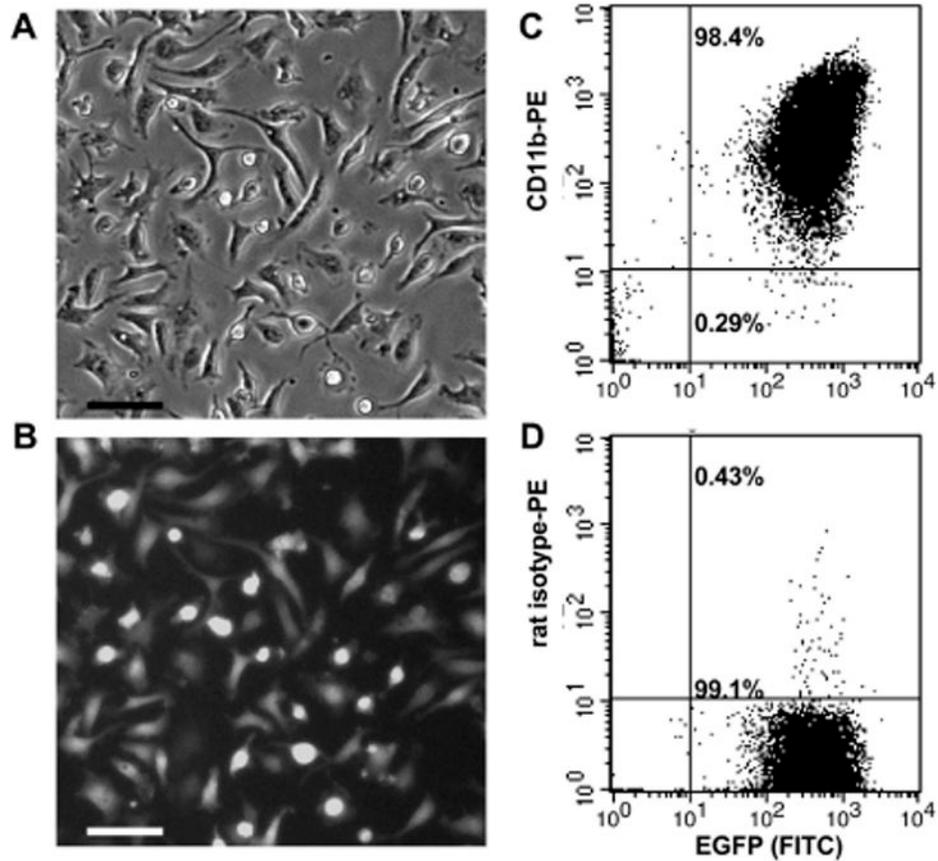


Fig. 1. Purity of *in vitro* cultures of primary microglia (1°MG) from *cfms*-EGFP mice
Purity of 1°MG cultures obtained from *cfms*-EGFP neonatal (day 2) mice was verified by fluorescence microscopy (A–B) for expression of EGFP, as well as by FACS analysis using EGFP expression and staining with the macrophage marker CD11b (C). Isotope control staining is shown in (D). Scale bar = 20 μ M.

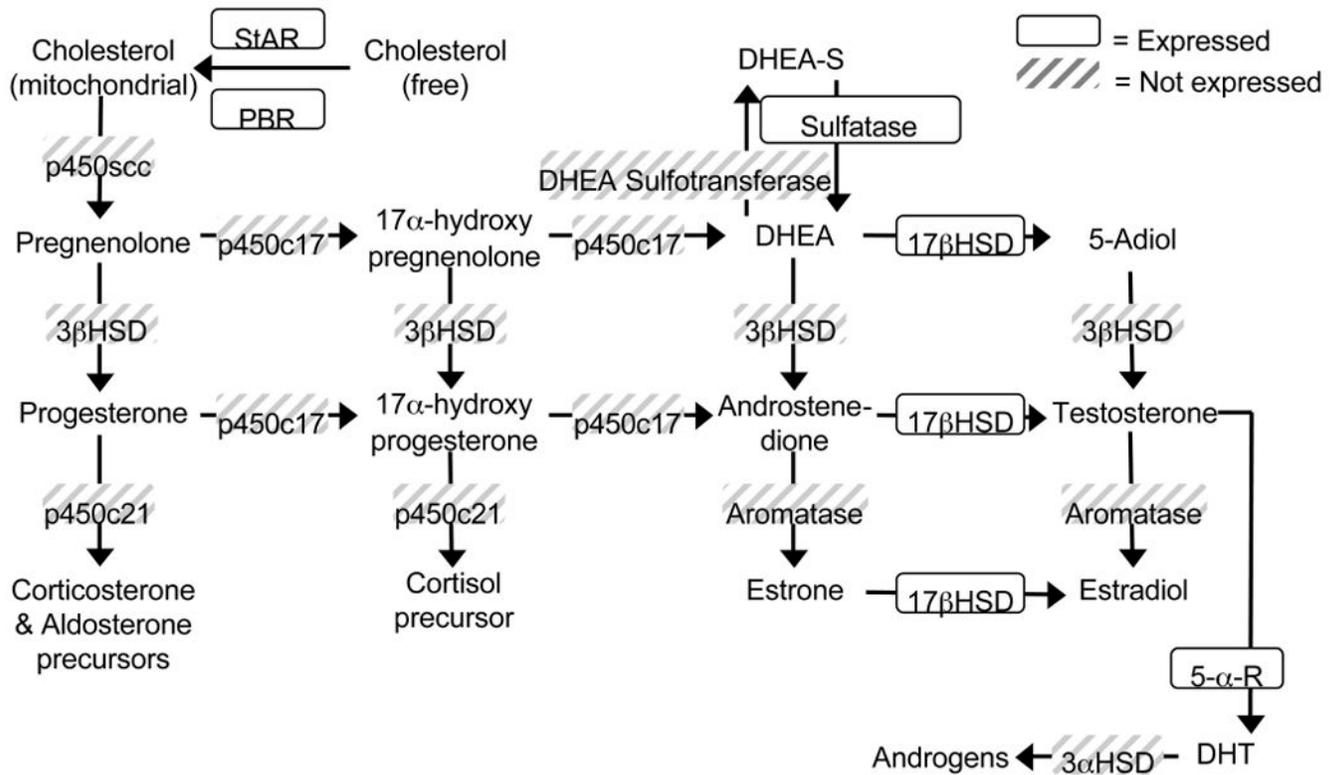


Fig. 2. Expression summary of steroid converting proteins in microglia

Diagram of the steroidogenic pathway showing the main steroidogenic enzymes and their steroid substrates. The boxed enzymes are those expressed in resting microglia, the shadowed ones were not expressed.

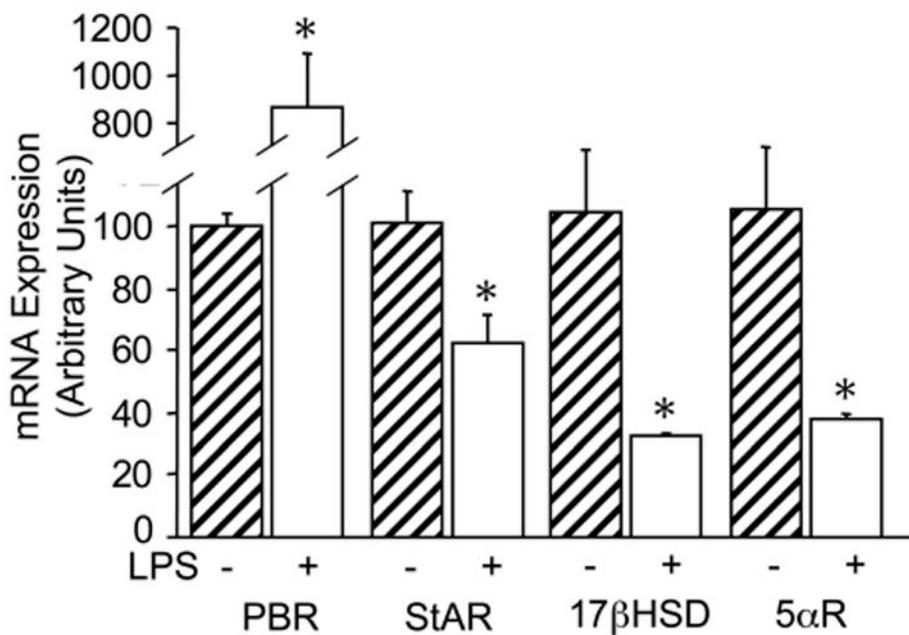


Fig. 3. *In vivo* LPS stimulation reduces steroidogenic enzyme expression in *ex vivo* microglia
 RT-PCR analysis of steroidogenic enzymes in FACS-sorted microglia from adult mice treated for 24hr with an IP injection of saline (hatched bars) or 1mg/kg LPS (open bars). Bars represent the % expression of each enzyme with respect to control samples and normalized to the housekeeping gene L27A. Values are the mean \pm the SEM of three animals. *, $P < 0.05$ vs. control.

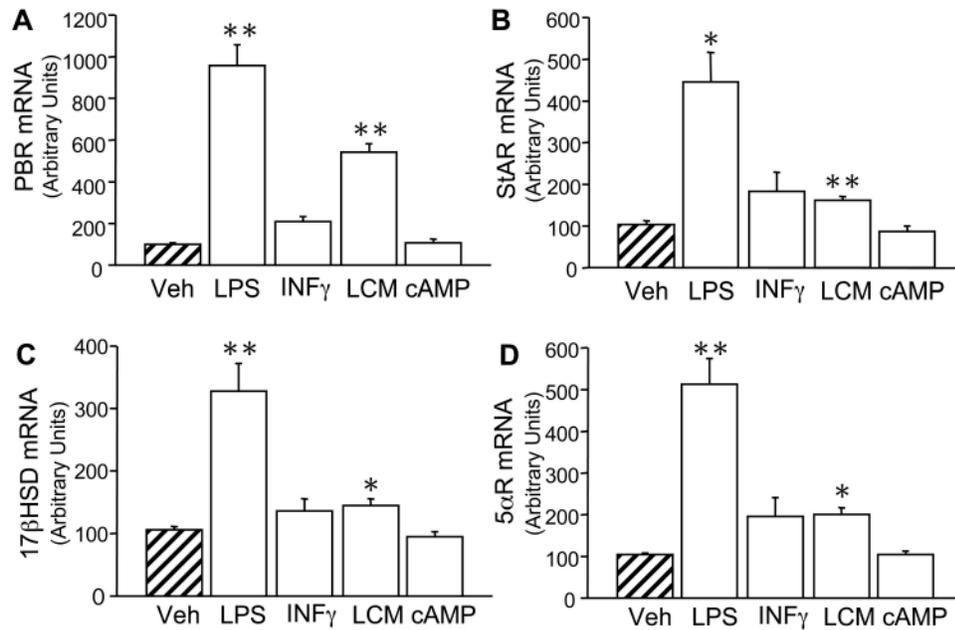


Fig. 4. LPS+INF γ increases the expression of steroidogenic enzymes in cultured 1°MG
 RT-PCR analysis of steroidogenic enzymes in cultured 1°MG. A–D) 1°MG cells were stimulated for 24hr with 100ng/ml LPS+ 10ng/ml INF γ (LPS), 10ng/ml INF γ (INF γ), LPS-conditioned media (LCM), 1mM db-cAMP (db-cAMP), or vehicle (Veh) (hatched bars). Bars represent the % expression of each enzyme with respect to vehicle and normalized to the housekeeping gene L27A. Values are the mean \pm the SEM of 2–3 independent experiments done in triplicate. *, **, P < 0.05, P < 0.001 vs. Veh, respectively.

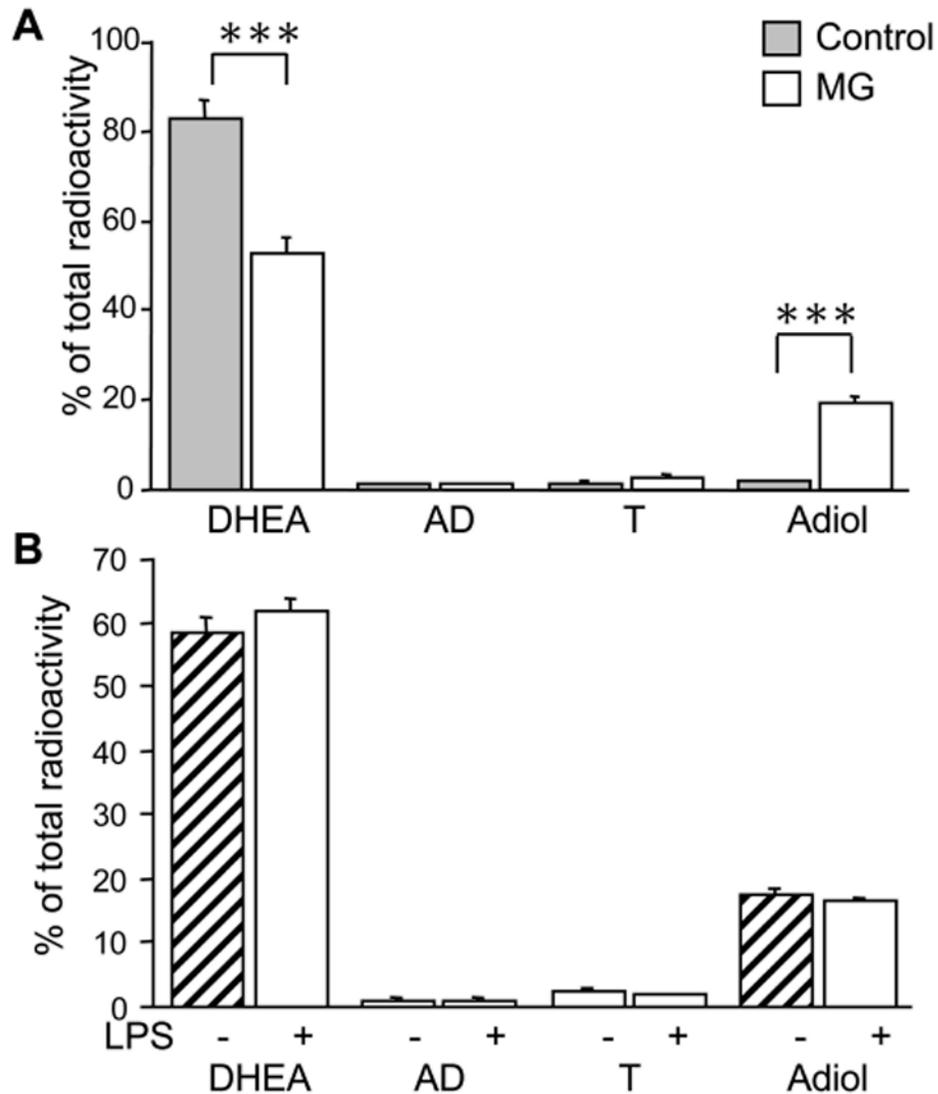


Fig. 5. Metabolism of H³-DHEA by 1°MG

TLC analysis of H³-DHEA metabolism in 1°MG. Cells were incubated for 24hr with H³-DHEA and metabolites extracted from culture supernatants were resolved by TLC and counted on a scintillation counter. A) Resting microglia (MG, open bars) showed a significant conversion of DHEA compared to the no-cell controls (Control, grey bars), with the only product being Adiol. B) LPS (100ng/ml LPS+ 10ng/ml INF γ) stimulation of the cells (hatched bars) didn't alter the metabolic activity or profile of steroids produced from DHEA in microglia. Bars represent the % of total H³ radioactivity collected from the TLC assay. Values are the mean \pm the SEM of 2–3 independent experiments done in triplicate. ***, $P < 0.0001$ vs. MG.

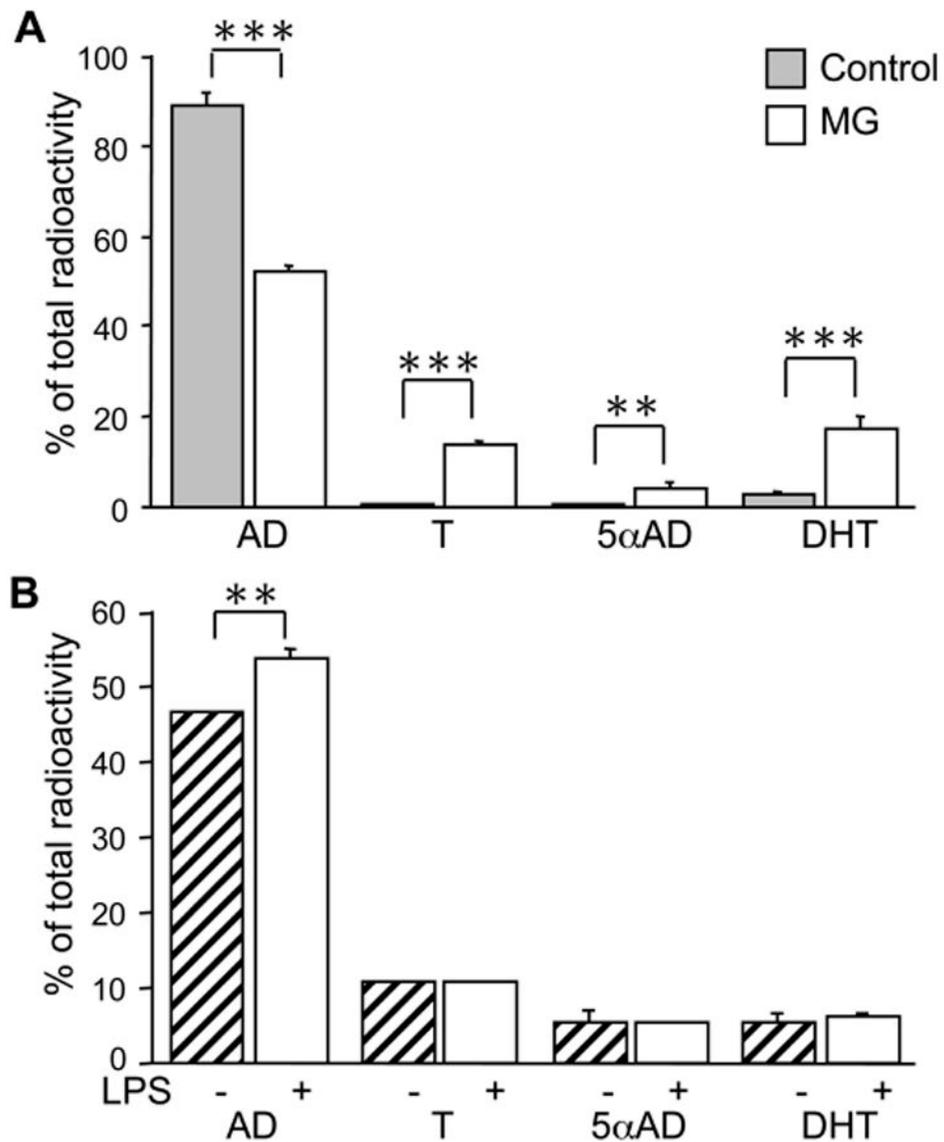


Fig. 6. 1°MG convert H³-AD into downstream steroids

TLC analysis of H³-AD metabolism in 1°MG. Cells were incubated for 24hr with H³-AD and metabolites extracted from culture supernatants were resolved by TLC and counted on a scintillation counter. A) Resting microglia (MG, open bars) showed a significant conversion of AD compared to the no-cell controls (Control, grey bars). The main products of AD were testosterone (T), 5αAndrostanedione (5αAD), and dihydrotestosterone (DHT). B) LPS (100ng/ml LPS+ 10ng/ml INFγ) stimulation (hatched bars) slightly decreased the conversion of AD compared to vehicle treated cells (open bars), but not the production of downstream steroids. Bars represent the % of total H³ radioactivity collected from the TLC assay. Values are the mean ± the SEM of 2–3 independent experiments done in triplicate. **,***, P < 0.001, P < 0.0001 vs. MG, respectively.

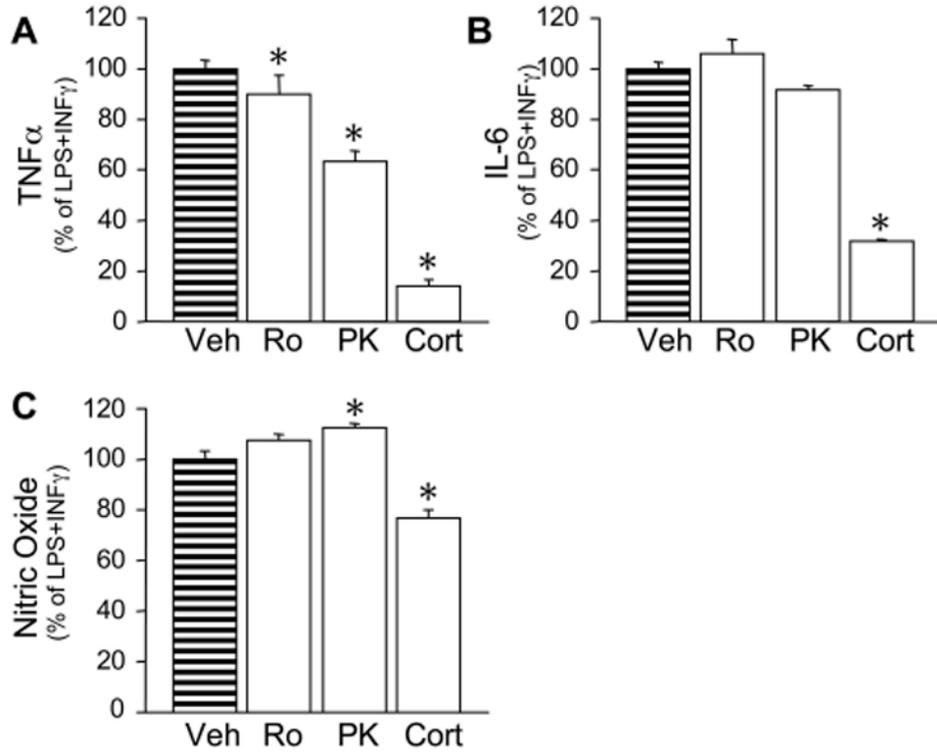


Fig. 7. PBR ligands selectively modulate TNF α production in 1 $^{\circ}$ MG

Cells were incubated for 24hr with LPS (100ng/ml LPS+ 10ng/ml INF γ) with a 10-minute pre-treatment of the PBR ligands Ro (10pM) and PK (10pM), or 1 μ M corticosterone as positive control. Culture supernatants were assayed for TNF α (A), IL-6 (B), and nitric oxide (NO) by ELISA and Greiss assay, respectively. Bars represent the % of total cytokines detected in the LPS controls. Mean cytokine levels measured for the LPS+INF γ controls were: TNF α (32ng/ml), IL-6 (70ng/ml) and NO (35mM). Un-stimulated cells had no detectable cytokine production. Bars represent the mean \pm S.E.M. of 2-3 independent experiments done in triplicate. *, P < 0.05 vs. LPS+INF γ .

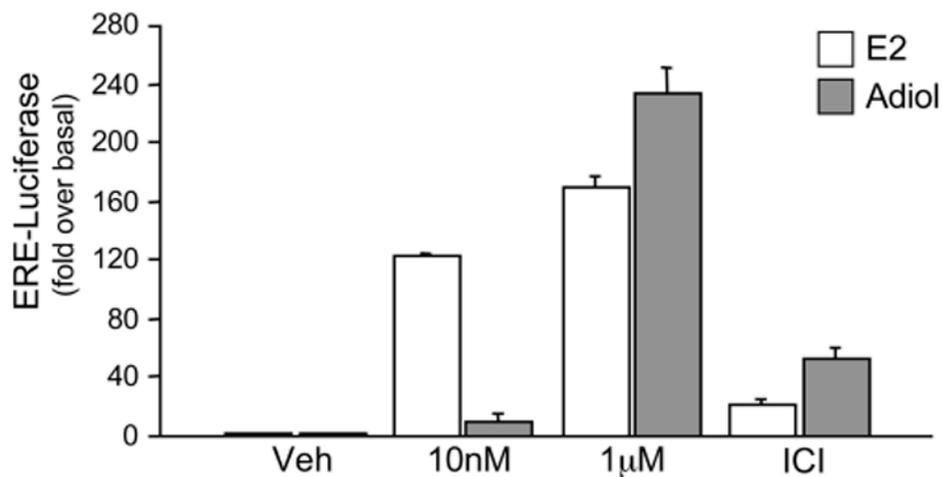


Fig. 8. Adiol is an effective estrogen receptor agonist

Estrogen receptor luciferase reporter assay reflected by luciferase expression from an ERE-Luciferase containing plasmid transfected into the neuronal cell line EtC.1. Bars represent luciferase activity determined by luminometry in cells treated with increasing doses of estrogen (E2) (open bars) or Adiol (grey bars). The third group of columns is from cells pre-treated 30 min with 100nM ICI 182,780 and then treated with the high doses of E2 and Adiol (ICI). Values are given in fold over basal \pm the SEM, from one representative experiment done in triplicate.

Table 1**Oligonucleotide Primers designed for RT-PCR Amplification of Steroidogenic Enzymes**

Primers for RT-PCR used to determine the expression of steroidogenic enzymes. The table shows the name of the protein analyzed (name of gene in parenthesis), specific oligonucleotide sequence, and the exons spanned by each amplicon.

Protein(Gene)		Sequence (5'-3')	Bases	Exons
StAR (<i>Star</i>)	Fwd	GAGCTCTGCTTGGTTCTCAA	218-239	2-3
	Rev	TTGAGTATGCCCAAGGCCTT	325-316	
PBR (<i>Tspo</i>)	Fwd	TGCAGAAACCTCTTGGCATC	173-193	2-3
	Rev	TGAAACCTCCCAGCTCTTTCC	286-266	
p450scc (<i>Cyp11a1</i>)	Fwd	CCTATTCCGCTTTTCTTTGAGTCC	636-660	3-4
	Rev	CGTCCCAAATATAACACTGCTG	686-663	
p450c21 (<i>Cyp21a1</i>)	Fwd	TGCCCATCGTGCAACTAGG	1064-1083	8-9
	Rev	AGCCGGAGATGCTGTAGCC	1102-1083	
p450c17 (<i>Cyp17a1</i>)	Fwd	TCGGCCCCAGATGGTGACTC	416-435	1-2
	Rev	TGGTCCGACAAGAGGCCTAGAG	454-433	
Sts (<i>Sts</i>)	Fwd	CCACTACTGCAACGCCTACCT	<i>Isles, et al. 2004</i> <i>Hum. Mol. Gen.</i>	
	Rev	CGTGAAGTAGAAGGCCTTCCA		
Arsa (<i>Arsa-1</i>)	Fwd	TTCACTGCAGATAACGGTCCTG	2038-2059	5-6
	Rev	AGGAGTAATGTGACCTGGCCA	2178-2158	
Sulft (<i>Sult2a2</i>)	Fwd	ACAGCTCTTCCAAGCCATGA	614-634	5-6
	Rev	TCCCCAGTTGTGCCTTTTCT	722-703	
3βHSD1 (<i>Hsd3b1</i>)	Fwd	TCTGAAAGGTACCCAGAACCTATTGG	433-458	3-4
	Rev	TTGCTTGAACACAGGCCTCCA	476-456	
3βHSD2 (<i>Hsd3b2</i>)	Fwd	AAAGGTACCCAGAACTTATTGGAGGC	417-442	3-4
	Rev	GGCACACTGGCTTGGATACAGG	463-442	
3βHSD4 (<i>Hsd3b4</i>)	Fwd	GGTCGAAAACAGGAAGAGGAATTGTC	118-144	1-2
	Rev	TGGTCTTGTCTGCAGCTTGGAC	163-140	
17βHSD1 (<i>Hsd17b1</i>)	Fwd	AGTGTGGGAGGCTTGTATGGGA	458-478	3-4
	Rev	CACTTCGTGGAATGGCAGTCC	496-479	
Arom (<i>Cyp19a1</i>)	Fwd	CTTTGGAGAACAATTCGCCCTTTC	418-441	3-4
	Rev	GCCCCGCAGAGCTTTCATAAAGAA	462-439	
5αR (<i>Srd5a1</i>)	Fwd	TGTTTCTGACAGGCTTTGCC	436-457	2-3
	Rev	CCATGCCCACTAACACAGGG	475-455	
3αHSD (<i>Hsd3a1</i>)	Fwd	GCCATCGTGAAAAACAATGG	692-711	6-7
	Rev	AATCAGCGCAGGAGTTCGA	795-777	
L27A (<i>Rpl27a</i>)	Fwd	TGTTGGAGGTGCCCTGTGTTCT	442-462	1-1
	Rev	CATGGAGAGAAGGAAGGATGC	542-522	

Table 2

Relative comparison of steady-state microglia mRNA levels of proteins involved in steroid metabolism

Expression levels of steroid converting proteins were evaluated by real-time PCR in microglia obtained from FACS-sorted cells from adult mouse brains (*ex vivo* MG), or from primary cultures (1°MG). Ovary tissue was used as a positive control. Values represent the mean (n=3) ± the SEM of the ratio of enzyme expression to the housekeeping ribosomal gene L27A. Cycle threshold values (C_t) are shown for L27A for reference.

	PBR	Star	p450scc	p450c21	p450c17	Sis
ExVivo MG	0.80 ± 0.03	0.68 ± 0.01	ND	ND	ND	0.67 ± 0.02
1°MG	0.94 ± 0.02	0.62 ± 0.01	ND	ND	ND	0.83 ± 0.00
Ovary	0.79 ± 0.02	0.79 ± 0.03	0.95 ± 0.0	0.55 ± 0.02	0.85 ± 0.0	0.66 ± 0.07
	Arsa	Sulft	3βHSD1	3βHSD2	3βHSD4	
ExVivo MG	0.85 ± 0.04	ND	ND	ND	ND	ND
1°MG	0.83 ± 0.02	ND	ND	ND	ND	ND
Ovary	0.69 ± 0.02	0.54 ± 0.01	0.73 ± 0.03	0.69 ± 0.01	0.69 ± 0.02	
	17βHSD	Arom	5αR	3αHSD	L27A(C _t)	
ExVivo MG	0.69 ± 0.01	ND	0.69 ± 0.01	ND	1.00 (21.5)	
1°MG	0.70 ± 0.02	ND	0.70 ± 0.01	ND	1.00 (23.9)	
Ovary	0.77 ± 0.03	0.67 ± 0.04	0.64 ± 0.0	0.54 ± 0.01	1.00 (14.6)	