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Gene expression of cyclooxygenase-1 and Ca²⁺-independent phospholipase A₂ is altered in rat hippocampus during normal aging

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Abstract

Brain aging is associated with inflammatory changes. However, data on how the brain arachidonic acid (AA) metabolism is altered as a function of age are limited and discrepant. AA is released from membrane phospholipids by phospholipase A_2 (PLA₂) and then further metabolized to bioactive prostaglandins and thromboxanes by cyclooxygenases (COX) -1 and -2. We examined the phospholipase A_2 (PLA₂)/cyclooxygenase (COX)-mediated AA metabolic pathway in the hippocampus and cerebral cortex of 4, 12, 24 and 30 month-old rats. A 2-fold increase in brain thromboxane B_2 level in 24 and 30 months was accompanied by increased hippocampal COX-1 mRNA levels at 12, 24, and 30 months. COX-2 mRNA expression was significantly decreased only at 30 months. Hippocampal Ca^{2+} -independent iPLA₂ mRNA levels were decreased at 24 and 30 months without any change in Ca^{2+} -dependent PLA₂ expression. In the cerebral cortex, mRNA levels of COX and PLA₂ were not significantly changed. The specific changes in the AA cascade observed in the hippocampus may alter phospholipids homeostasis and possibly increase the susceptibility of the aging brain to neuroinflammation.

Keywords

aging; cyclooxygenases; calcium-independent phospholipase A₂; hippocampus; thromboxane; prostaglandin

1. Introduction

Aging is associated with increased inflammatory responses and vulnerability of neurons to degeneration [17,18,22,39]. Some authors have raised the possibility that inflammation may occur during normal aging and increase the vulnerability to neurodegenerative disorders such as Alzheimer's disease (AD) [7,20,26]. Since the arachidonic acid (AA) cascade plays a key role in neuroinflammation [37], we thought it of interest to identify the changes that occur in this pathway during physiological aging.

AA (20:4n-6) is released from membrane phospholipids by phospholipase A_2 (PLA₂) enzymes. The PLA₂ family can be generally divided into two groups: the Ca²⁺-independent PLA₂

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(iPLA₂) and the Ca²⁺-dependent PLA₂, which includes secretory (sPLA₂) and cytosolic PLA₂ (cPLA₂) [24]. iPLA₂ is thought to mediate homeostatic phospholipid remodeling through fatty acid deacylation/reacylation reactions [49] and may also be involved in cellular signaling [5]. In contrast to cPLA₂, which releases predominantly AA, iPLA₂ has been suggested to selectively release docosahexaenoic acid (DHA; 22:6n-3), a n-3 polyunsaturated fatty acid highly concentrated in brain membranes [42,43]. Once released, AA is then converted to bioactive prostaglandins and thromboxanes by cyclooxygenase (COX) enzymes. Two isoforms of COX have been described: COX-1, constitutively expressed in most tissues [30], but also induced by certain inflammatory stimuli in peripheral tissues [46], and COX-2, generally induced by inflammatory stimuli including cytokines, hormones, and mitogens [44], but also constitutively expressed in the central nervous system, especially in pyramidal neurons of hippocampal and cortical circuits [50].

Learning and memory deficits have been well documented in aged F344 rats [14,21,41,45]. Activated microglia and astrocytes [16] and elevated levels of pro-inflammatory cytokines [10,18,51] have been described in the rodent brain during normal aging. However, the involvement of COX-mediated AA metabolism in this "pro-inflammatory-status" remains unclear [4,23,38]. Aging-related changes in COX-mediated AA metabolism are suggested by age-dependent spatial memory deficits and increased neuronal apoptosis and astrocytic activation in transgenic mice overexpressing COX-2 [2]. We have previously shown in the Rhesus monkey an age-dependant decrease in the levels of cPLA₂ protein in the cerebellum and of COX-2 protein in the frontal pole [47]. Taken together, these data suggest that AA metabolism may be disturbed in normal aging. We chose two morphologically and functionally distinct regions, the hippocampus and the cerebral cortex, that have been shown to be affected by the aging process [14,35]. We assessed age-related changes in the metabolism of AA, with a particular focus on the PLA₂/COX pathway across a range of 4 ages (4, 12, 24, and 27–30 months).

2. Materials and methods

2.1 Animals

The study was approved by the National Institutes of Health (NIH) Animal Care and Use Committee in accordance with NIH guidelines on the care and use of laboratory animals. Male Fischer-344 rats, 4, 12, 24 and 27–30 month-old (NIA-sponsored colony at Harlan Sprague-Dawley, Indianapolis, IN) were housed at least one week in our animal facility, maintained at 25° C with a 12 hr light/dark cycle, with free access to food and water. All rats were killed with an overdose of sodium pentobarbital (100 mg/kg, *i.p.*). Then, rats used to measure prostaglandin levels were subjected to high-energy head-focused microwave irradiation (4.8 kW, 3.4 sec, Cober Electronics, Stanford, CT, USA) to stop metabolism, as reported [8,9,33]. Rats used for mRNA analysis were decapitated, hippocampus and cerebral cortex were rapidly dissected out on ice, immediately frozen in -50° C 2-methylbutane, and stored at -80° C until used.

2.2 Determination of brain prostaglandin E₂ (PGE₂) and thromboxane B₂ (TXB₂) concentration

After microwave fixation, the amount of the brain extract required to perform analysis did not allow measuring prostaglandin levels in individual brain regions without pooling samples from different animals. Therefore, we measured PGE₂ and TXB₂ concentration in the whole brain. Microwaved brains were weighed and extracted with 18 volumes of hexane: 2-propanol (3:2, by volume) using a glass Tenbroeck homogenizer. The prostanoids were purified from the lipid extract as described by Radin [34] and the concentrations of PGE₂ and TXB₂ were determined using specific enzyme-linked immunosorbent assay kits (Oxford Biomedical, Oxford, MI,

USA), as previously reported [8]. According to the manufacturer, the sensitivity of the PGE_2 and TXB_2 assays are 0.2 ng/ml and 0.009 ng/ml at 80% B/Bo, respectively.

2.3 Real-Time Quantitative PCR

Total RNA was extracted from hippocampus or cerebral cortex using RNeasy Lipid Tissue Midi Kit (Qiagen, Valencia, CA, USA) as directed by the manufacturer. Five g of total RNA was reverse transcribed using a High Capacity cDNA Archive kit (Applied Biosystems, Foster City, CA, USA). Five g of each RNA sample was incubated similarly in the absence of reverse transcriptase to ensure that PCR products resulted from amplification from the specific mRNA rather than from genomic DNA contamination.

The levels of gene expression of COX-1, COX-2, cPLA₂, iPLA₂, glial fibrillary acidic protein (GFAP) were measured by real-time quantitative RT-PCR, using the ABI PRISM 7000 Sequence Detection System (Applied Biosystems). Specific primers and probes were purchased from the available Assays-on-Demands (Applied Biosystems). We used glyceraldehyde-3-phosphate dehydrogenase (GAPDH) as an endogenous control, as its expression has been shown to remain unchanged during aging [40]. Data were analyzed using sequence detection systems software (Applied Biosystems). Results were expressed as the amount of target gene normalized to the endogenous control (GAPDH) and relative to the 4 month-old rats using the $\Delta\Delta C_{T}$ method [25].

2.4 Statistical analysis

Results are expressed as mean \pm SEM. Statistical analysis was performed using One-Way ANOVA followed by post-hoc Bonferroni test. *P* < 0.05 was considered statistically significant.

3. Results

3.1 TXB₂ levels are specifically increased in the aged brain

Levels of TXB₂, a stable product of the very short half-lived TXA₂, were increased by 2 fold [F(3,19) = 4.89, p = 0.010] in 24 and 30 month-old group compared to the 4 month-old group (Fig. 1A). PGE₂ levels (Fig. 1B) showed a trend towards an increase at 24 months, which did not reach statistical significance among the different groups [F(3,19) = 2.71, p = 0.072].

3.2 COX-1 mRNA expression is upregulated whereas COX-2 mRNA expression is downregulated in the hippocampus during aging

Hippocampal COX-1 mRNA expression (Fig. 2A) was increased by 20% at 12 months and by 27% at 24 and 30 months compared to the 4 months group [F(3,19) = 28.03, p < 0.0001]. In contrast, hippocampal COX-2 mRNA expression (Fig. 2B) was statistically reduced by 15% in the 30 months group compared to the 4 months group [F(3,19) = 4.47, p = 0.015]. COX mRNA expressions (Table 1) were not changed in the cerebral cortex of all age groups examined [COX-1: F(3,19) = 0.37, p = 0.773; COX-2: F(3,19) = 2.97, p = 0.058].

3.3 Hippocampal iPLA₂ (VI) mRNA expression is downregulated in aged rats

Hippocampal iPLA₂ mRNA expression (Fig. 3A) was decreased by 15% at 24 and 30 months compared to the 4 months group [F(3,19) = 4.67, p = 0.013]. Hippocampal cPLA₂ mRNA expression level (Fig. 3B) was not changed in all age groups examined [F(3,19) = 0.32, p = 0.81], neither did cortical cPLA₂ [F(3,19) = 1.89, p = 0.165] and iPLA₂ mRNA expression levels [F(3,19) = 1.37, p = 0.283] (Table 1).

3.4 GFAP mRNA expression is increased in hippocampus and cerebral cortex during aging

To confirm aging-related astrocytic activation, we measured the expression of GFAP, a specific marker of astrocytes. Hippocampal GFAP mRNA expression (Fig. 4) was increased by 46% at 24 months and by 72 % at 30 months compared to the 4 months group [F(3,19) = 31.21, p < 0.0001]. Cortical GFAP mRNA expression (Table 1) was increased by 2 fold at 24 months and by 2.4 fold at 30 months compared to the 4 months group [F(3,19) = 60.74, p < 0.0001].

4. Discussion

In this study, we found that brain TXB_2 level was increased by 2 fold in 24 and 30 month-old rats compared to 4 month-old rats and we also showed that parts of the AA cascade are specifically altered in the hippocampus during normal aging: namely 1) COX-1 expression is increased, 2) COX-2 expression is decreased, and 3) iPLA₂, but not cPLA₂, expression is decreased. In contrast, we did not find any significant change in the expression of COX or iPLA₂ in the cerebral cortex from aged rats.

The upregulation of COX-1 mRNA expression appeared quite early in adulthood (12 months) and therefore can be viewed as part of the normal aging process. In contrast, COX-2 mRNA level was downregulated only at 30 months. Previous studies have examined hippocampal and cortical COX-2 mRNA expression in different strains of rats aged from 3 to 27 months and reported no change [4,23]. Therefore, COX-2 gene expression appears to become affected only in the advanced phase of the aging process. COX-1 upregulation as a function of age may have different implications. First, it may be responsible for the increased brain levels of TXB₂, an AA-derived prostaglandin, via the subsequent metabolism of COX and thromboxane synthase. Indeed, evidence indicates that thromboxane synthase preferentially couples with COX-1 for TXB₂ production [11]. Although brain microvessels also contribute to COX-1 derived TXB₂ synthesis, since microvessels represent only 0.1 % of the whole brain [48] the increase in TXB₂ that we observed in this study is unlikely significantly contributed by the cerebral circulation. However, since in this study TXB₂ and PGE₂ levels were determined in the whole brain, it would be interesting to further examine the regional distribution of the changes in prostaglandin during normal aging. An increase in COX activity has been reported in the cerebrum of F344 rats at 24 months versus 6 month-old [4], corresponding approximately to the age at which we found increased levels of TXB_2 in this study. Since COX-1 is mainly localized in glia, as opposed to COX-2 which is mainly in neurons [13,50,52], the upregulation of COX-1 could be associated with the aging-related glial activation. In particular, the agerelated increase of GFAP mRNA that we show has been well documented in human and rodent brain [19,29,32]. Glial activation may contribute to neuronal dysfunction as both microglia and astrocytes become activated early during physiological aging [12,16]. It remains unclear why only the hippocampus is responsive to the age-related astrocytic activation which occurred in several brain regions. It is unlikely that COX-1 upregulation in the hippocampus alone could account for the increase of TXB_2 level in the whole brain. It is possible that other cerebral regions that we did not examine may have an upregulated COX-1 expression. Another possibility is that increased hippocampal COX-1 mRNA expression is accompanied by an upregulation in thromboxane synthase activity in the hippocampus and/or in other regions. Even though age-related changes occur in the whole brain, the aging process can exhibit regional specificity, the hippocampus being especially sensitive [6,14].

Only few reports have addressed changes in brain prostanoid production with aging. Using brain microwaving to stop lipid metabolism *in vivo* [3,9], we demonstrated an increased level of TXB_2 in the brains of 24 and 30 month-old animals compared to 4 month-old rats. The same tendency was observed for PGE₂ levels, although it did not reach statistical significance. Prostaglandins have been shown to have numerous effect on astrocytes *in vitro* including upregulating GFAP expression [31]. Thus, it should be further clarified if the astrocytic

activation observed during aging is either the consequence or the cause of altered AA metabolism. COX-1 upregulation with aging might play a role in altering the neuroinflammatory response and increasing vulnerability to neurodegenerative diseases with a marked inflammatory component [28].

We also found a decrease in the Ca²⁺-independent iPLA₂ mRNA levels in the hippocampus of 24 and 30 month-old rat. In contrast to cPLA₂, which preferentially cleaves AA from membrane phospholipids, iPLA2 is thought to show selectivity towards docosahexaenoic acid (DHA) [42]. DHA, the major n-3 polyunsaturated fatty acid found in cerebral membranes, plays a crucial role in physiological functions such as neurotransmission, membrane fluidity, ion channel and regulation of enzyme activity and gene expression [1], and it may protect against cognitive decline [27]. The decrease in iPLA₂ expression observed in this study is consistent with evidence that the DHA content in cerebral phospholipids is decreased during normal aging [15] and that cortical iPLA₂ is downregulated in DHA-deprived rats [36]. Considering the role of iPLA₂ in regulating homeostatic phospholipid levels [49], the downregulation of iPLA₂ expression during aging might alter phospholipid homeostasis by decreasing the release of DHA in the hippocampus. Therefore, the age-related decrease of AA and DHA in membrane phospholipids [15], coupled with the downregulation of iPLA₂, could contribute to increase the ratio of free AA/DHA and consequently alter the brain prostanoid profile. These changes may exert profound effects in the hippocampus, a region highly vulnerable to the aging process [6,14].

In summary, our results indicate that the mRNA expression of three important enzymes in the AA metabolic pathway, COX-1, COX-2, and iPLA₂, is altered in the hippocampus during normal aging. These changes may alter brain phospholipids homeostasis and possibly increase the susceptibility of the aging brain to neuroinflammation. Future studies should address the functional consequences of the changes in gene expression described here by examining protein expression and activity of the AA cascade enzymes, as well as regional distribution of COX-1 within all brain regions during aging.

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B.



Fig 1.

Thromboxane B₂ (A) and prostaglandin E₂ (B) levels in the microwaved brain of 4, 12, 24 and 30 month-old F344 rats. Data (mean \pm SEM, n= 5–6) are expressed as the percentage of the 4 month-old rats. *p* values are expressed as the comparison of aged rats with 4-month-old rats using One-Way ANOVA (**p*<0.05, ***p*<0.01; post-hoc Bonferroni test).

A.



Fig 2.

COX-1 and COX-2 mRNA expression in the hippocampi (A-B) of 4, 12, 24 and 30 month-old F344 rats. Data (mean \pm SEM, n=5–6) are normalized to the level of the internal control, GAPDH and are expressed as the percentage of the 4 month-old rats. *p* values are expressed as the comparison of aged rats with 4-month-old rats using One-Way ANOVA (**p*<0.05, ****p*<0.001; post-hoc Bonferroni test).

A.



Fig 3.

iPLA₂ (VI) and cPLA₂ mRNA expression in the hippocampi (A-B) of 4, 12, 24 and 30 monthold F344 rats. Data (mean \pm SEM, n=5–6) are normalized to the level of the internal control, GAPDH and are expressed as the percentage of the 4 month-old rats. *p* values are expressed as the comparison of aged rats with 4-month-old rats using One-Way ANOVA (**p*<0.05; posthoc Bonferroni test).



Fig 4.

GFAP mRNA expression (B) in the hippocampi of 4, 12, 24 and 30 month-old F344 rats. Data (mean \pm SEM, n=5–6) are normalized to the level of the internal control, GAPDH, and are expressed as the percentage of the 4 month-old rats. *p* values are

Table 1

mRNA expression of genes involved in the PLA₂/COX pathway in the rat cerebral cortex during aging using quantitative real-time PCR.

Genes ¹	4 months (n=6)	12 months (n=6)	24 months (n=6)	30 months (n=5)
COX-1 COX-2 cPLA ₂ iPLA ₂ GFAP	$100 \pm 2 \\ 100 \pm 5 \\ 101 \pm 5 \\ 100 \pm 3 \\ 101 \pm 6$	$100 \pm 5 \\ 126 \pm 12 \\ 113 \pm 6 \\ 96 \pm 4 \\ 122 \pm 4$	$105 \pm 9 \\ 100 \pm 3 \\ 118 \pm 6 \\ 90 \pm 6 \\ 206 \pm 12 \\ ***$	$108 \pm 797 \pm 8105 \pm 788 \pm 6241 \pm 11 ***$

 I Data (mean ± SEM, n=5–6) are normalized to the level of the internal control, GAPDH, and are expressed as the percentage of values in the 4 monthold rats. *p* values are expressed as the comparison of aged rats with 4-month-old rats using One-Way ANOVA

*** *p*<0.01; post-hoc Bonferroni test.