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RESEARCH ARTICLE

DYNAMICS OF NEUROSTEROIDS IN THE BRAIN: SPECIAL REFERENCE TO STEROIDOGENIC ENZYME 3α -HSD GENE EXPRESSION IN THE BRAIN OF AN INDIAN MAJOR CARP *LABEO ROHITA* (HAM.)

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ABSTRACT

3α -Hydroxy steroid dehydrogenase (3α -HSD) is a member of the aldo-keto reductase family. The enzyme is used in the synthesis of neuroactive steroids in steroidogenesis. The 3α -HSD catalyzes the conversion of 5α -dihydroprogesterone (5α -DHP) and 5α -dihydrotestosterone (5α -DHT) into $3\alpha,5\alpha$ -tetrahydroprogesterone ($3\alpha,5\alpha$ -THPROG) and $3\alpha,5\alpha$ -tetrahydrotestosterone ($3\alpha,5\alpha$ -THT) respectively. The progesterone metabolite of neuroactive steroid, $3\alpha,5\alpha$ -THPROG is synthesized in different regions of brain. In the present study, the gene expression of 3α -HSD in the brain of *Labeo rohita* was identified. The gene was observed by PCR amplification with a specific primer. The result showed has formed a basis for future investigations on the regulation and function of these enzymes in the fishes.

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INTRODUCTION

The steroids synthesised in the brain tissues or neural tissues are generally called as 'neurosteroids'. Pregnenolone (PREG) and progesterone (PROG) can be synthesized in the nervous system. A crucial biochemical reaction in vertebrates is the progesterone conversion into neuroactive metabolites such as 5α -dihydroprogesterone (5α -DHP) and $3\alpha,5\alpha$ -tetrahydroprogesterone ($3\alpha,5\alpha$ -THPROG), which regulate several neurobiological processes, including stress, depression, neuroprotection, and analgesia. $3\alpha,5\alpha$ -THPROG is a potent stimulator of type A receptors of γ -amino butyric acid (GABA), the main inhibitory neurotransmitter (Belelli and Herd, 2003; Belelli and Lambert, 2005; Pinna, *et al.*, 2006; Belyaeva, *et al.*, 2007). GABAergic neurons of the reticular thalamic nucleus express high levels of 5α -RI and 3α -HSD and their nerve endings may secrete allopregnanolone and release GABA in the proximity of postsynaptic GABA_A receptors located on the dendrites and somata of glutamatergic thalamocortical output neurons (Pinault, 2004). Similar considerations are also expected for allopregnanolone synthesized by medium spiny GABAergic neurons in the caudate or putamen and also very likely by Purkinje cells that modulate GABA receptors expressed postsynaptically on cell bodies or dendrites of deep cerebellar nuclei neurons (Agis-Balboa *et al.*, 2006). The aldo-keto reductase superfamily is involved in the metabolism of endogenous substrate, such as

steroid hormones, prostaglandins, bile acids and xenobiotics, such as drugs and environmental carcinogens (Khanna *et al.*, 1995). $3\alpha,5\alpha$ -THPROG with a steroid chemical structure and low molecular weight of 318.49, easily penetrates the blood-brain barrier to induce central nervous system (CNS) effects, including anxiolytic and sedative hypnotic properties (Gee *et al.*, 1998; Brinton, 1994). Functional analyses indicate that $3\alpha,5\alpha$ -THPROG induces myelin formation in both the CNS and the peripheral nervous system (Baulieu and Schumacher 2000; Schumacher *et al.*, 2003) and promotes neuron survival. In humans, recent advances in functional brain imaging have identified critical neural circuits in cortico-limbic structures involved in the modulation of fear responses, aggressiveness, anxiety, and sexual behaviors that appear to be affected in mood disorders (Carlson *et al.*, 2006). These circuits include the amygdala, hippocampus, and the medial prefrontal cortex (mPFC) (Carlson *et al.*, 2006; Nelson and Trainor, 2007). In rodents, the olfactory system has also been implicated (Mandiyani *et al.*, 2005). 3α -HSD gene expression has been reported in some other vertebrates like humans, rat, and cattle (Penning *et al.*, 1984; Qin *et al.*, 1993; Lin *et al.*, 1997).

Recently it has been reported that both allopregnenolone and isopregnenolone are also synthesized from PROG in CNS tissues throughout the embryonic and early postnatal period of development in the rat (Pomata *et al.*, 2000). However, there is no evidence for the identification of 3α -HSD gene in fishes. This is the first attempt to know the 3α -HSD enzyme in brain of fish, particularly among the Indian carp. *Labeo rohita*

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(commonly called as Rohu) is an animal model for the present research work and it is an economically important species among the Indian major carps.

MATERIALS AND METHODS

Sample collection

Fish samples of *Labeo rohita* were collected from the Lake Sevilimedu located at Kanchipuram. Fishes were caught in live condition and they were dissected to collect brain samples. The brain tissues were fixed in a sterilized vials containing RNAlater and they were stored at 4°C until analysis.

Total RNA isolation

Total RNA was isolated by homogenizing the whole brain sample (200mg) with 500 μ l of TriReagent (Sigma) and 200 μ l of DEPC water. After homogenizing the sample, it was then incubated at -20°C for 5 minutes. 0.2ml of chloroform was added and incubated at -20°C for 5 minutes. It was then centrifuged at 12,000rpm (4°C) for 15 minutes. The supernatant was collected in a fresh tube. RNA was precipitated by adding equal volume of isopropanol and it was stored at -20°C for 45 minutes. The sample was centrifuged again at 12,000 rpm (4°C) for 15 minutes. Total RNA was obtained as a pellet and 75% ethanol (7.5ml of Ethanol was mixed with 2.5ml of DEPC water) was added. It was centrifuged at 12,000 rpm for 5 minutes and air dried and dissolved by adding 40 μ l of DEPC water. This was frozen and stored at -20°C for half-an-hour. The sample was then tested with the 1.2% agarose gel for its purity.

Synthesis of first strand cDNA

Total RNA was isolated from the brain of *L. rohita*. The separated RNA has been reverse transcribed into cDNA using RT-PCR method. A clean PCR tube was taken to this 1 μ l of the sample, 1 μ l of Oligo (dT)₁₈ primer, 9.5 μ l of de-ionized water was added and it was spun gently for few seconds in a micro centrifuge. This mixture was incubated at 70°C for 5 minutes. After incubation, 4 μ l of 5X reaction buffer, 0.5 μ l of Ribonuclease inhibitor (40 μ l/dl), 2 μ l of 10mM dNTP mix were added. This was mixed gently, centrifuged and incubated at 42°C for 5 minutes. After incubation, it was added with M-MuLV reverse transcriptase (20U/ μ l) to make it around 20 μ l volumes. This was incubated finally at 42°C for 60 min and 25°C for 10 min.

The reaction was stopped by heating at 70°C for 10 min and chilled on ice. PCR products were then tested with agarose gel electrophoresis.

Polymerase Chain Reaction

2 μ l of cDNA was taken in a sterilized PCR tube with 1 μ l of 3 α -HSD primers of sense and antisense. The following primers were used at different reactions. Sense- 5'CTGTGCCTGAGAAGGTTGCT3', antisense-5'CATGTGTCACAGATATCCAC3', sense-5' GGAGGCCATGGAGAAGTGTAT3', antisense 5'CACCCATGTTTTGTCTCGTG3'and sense- 5'ACAAGCGATGGATTCCATAT3' 25 μ l PCR master mix consists of all basic components : Taq DNA Polymerase, dNTPs and reaction buffer(1.5mM Magnesium chloride) were added. The PCR amplification was used to check the expression of the mRNAs of 3 α -HSD enzyme in the brain sample. The temperature adopted in the amplification is as follows: 94°C for 2 minutes in 1 cycle, 94°C for 1 minute, 56°C for 1 minute and 72°C for 1 minute in 35cycles, the finally holding temperature was 4°C. The PCR products of gene specific primer of 3 α -HSD along with 100bp DNA Ladder were then subjected to 1.2% agarose gel electrophoresis. After running the gel, the image of specific bands were captured by using UV trans illumination under the JH Bio geldoc system.

Sequencing

The PCR product was sequenced and the mRNA sequences of 3 α -HSD obtained for *L. rohita* were shown in the result. The mRNA sequences of 3 α -HSD of *L. rohita* were multiple aligned with vertebrate organisms to know the similarity, relationship and evolutionary significance of *L. rohita* with other organisms.

RESULTS

Total RNA was isolated from the entire male brain of *L. rohita*. Two RNA bands as 28S and 18S were obtained and shown in Fig.1A. The isolated RNA was reverse transcribed into cDNA using RT-PCR. The single strand cDNA was obtained and it appeared as a smear. The PCR was made to run at specific temperature to analyze the gene expression of 3 α -HSD by adding the gene specific primer to the cDNA. The PCR product shows a single band which is represented in Fig.1B and Fig.1C. The PCR product and the 100bp DNA Ladder

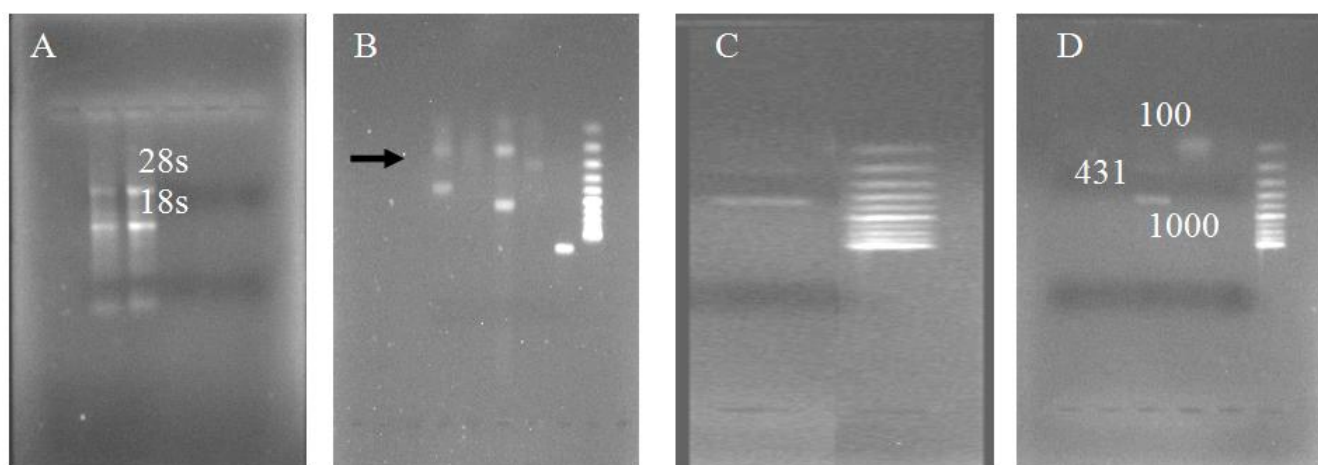


Fig.1. A. Shows the Two RNA bands as 28S and 18S were obtained from brain tissue of *Labeo rohita*, B and C. Shows the PCR products from cDNA amplified along with specific primer of 3 α -HSD, D. Shows the 100bp marker DNA and 431bp expression of 3 α -HSD gene in the brain of Indian major carp, *Labeo rohita*

were loaded in separate lanes in the gel. Single band at 431bp was obtained and is shown in Fig.1D. The mRNA sequence of 3 α -HSD obtained for *L. rohita* are shown below. The obtained mRNA sequence of 3 α HSD of *L. rohita* were multiple aligned with different species of vertebrates to know the similarity of *L. rohita* with other organisms. Relative score table and multiple sequence alignment of complete mRNA sequence of 3 α -HSD of vertebrates along with the obtained sequence of *Labeo rohita* showing the percentage of similarity shown in Fig. 2.

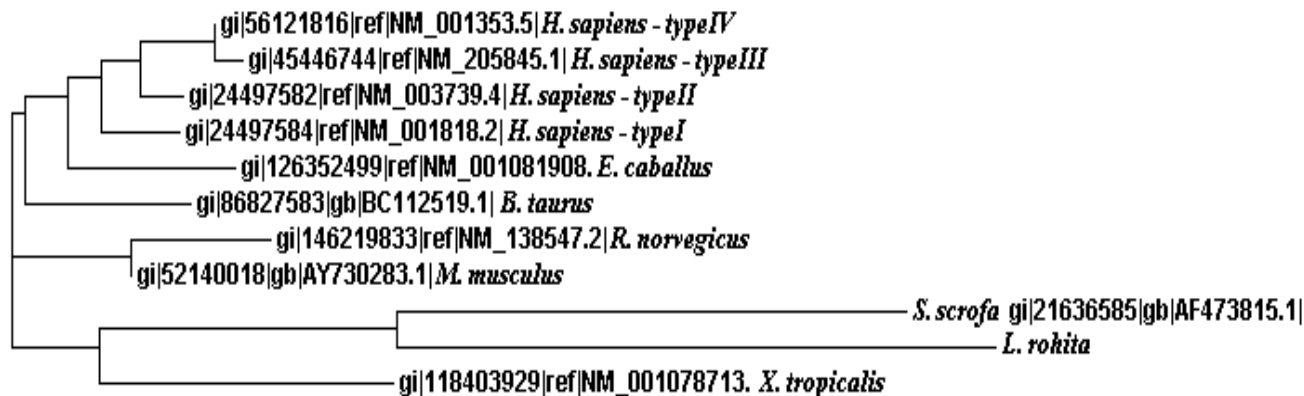


Fig. 2. Phylogenetic tree of mRNA sequences of 3 α -HSD of different classes of vertebrates along with the obtained sequences from *Labeo rohita* showing the relationship and evolutionary distance among the organisms when compared with each other

The partial mRNA Sequences of 3 α -HSD of *Labeo rohita* obtained from the present study

>gi|326325828|gb|JF683819.1| *Labeo rohita* 3-alpha hydroxy steroid dehydrogenase (AKR1C) mRNA, complete sequence
 GCGATGGTGCTGTGCTCTTGGATACCGTCATTTTTTCAT
 CGTCGCGAGCGTACTTAGGTGACCTGCCTGGCCACTT
 GCATCTGCACGGCATTGCTTGGCTGGTCTGTCCAATA
 TATTTGTGGGATTTTGAGAGTGAAGTAGCTCCTCTGTT
 GGCTGCATTTGTAAGTCCATTCTGGCTGCAAATGAA
 CCTGCTGGGAATGATTTCTAGTCTGCAAGTGTGCGA
 TCTAAGCTGCATGCGTTCATGTCTATTTGCATCGAAGC
 TACGAGAGTGGTGTGGAACCATGTTTGGCAACAGCGT
 TTCTGACATGATTCTACCAGGTAGAGGTCCGTCCTT
 CCGTTGTCTCGATCTAACCTTGGCCTACCTCTCCTGCC
 GGTACTTACTCTACCTCGTTGAGAATGTATGGACAGT
 GGATATCTGGTGACACATGA

DISCUSSION

Our results confirm the presence of 3 α -HSD in the brain of an Indian Major Carp, *L. rohita*. Results from the present experiment indicates the expression of the mRNAs of 3 α -HSD, the enzyme involved in the conversion of 5 α -DHPROG into 3 α , 5 α -THPROG in the brain of *L. rohita*. Aldo-keto reductases interconvert weak androgens, estrogens, progesterin, mineralocorticoids and glucocorticoids to their more potent counterparts by catalyzing the reduction and oxidation of keto- and hydroxysteroids, respectively, thereby regulating a wide range of physiological processes involved in development, homeostasis and reproduction (Bauman *et al.*, 2004). In this manner, AKRs regulate the occupancy and transactivation of several steroid receptors in target tissues leading to

transcription of hormone-responsive genes (Penning *et al.*, 2004). The products of AKR activity have been implicated in prostate disease, breast cancer, obesity, poly cystic ovary disease and delay in the onset of puberty in humans (Stanbrough *et al.*, 2006; Rittner *et al.*, 1997; Nonneman *et al.*, 2006). They reported the AKR1C gene expression in different tissues of pig. From the report it is noted that spleen, ovary, lung, adrenal, kidney, and endometrial expressed all the five genes. AKR1C2 and AKR1C4 were the most widely expressed genes. AKR1C2 was expressed in all tissues except pancreas and brain. Unlike human AKR1C4, pig AKR1C4

expression was not specific to the liver but was expressed in all tissues and AKR1C2 was the only other AKR1C gene expressed in brain. Because of their location in the swine genome and their implication in reproductive physiology, this gene cluster was characterized and evaluated for effects on reproductive traits in swine. Results of the present experiment confirm the presence of the gene AKR1C in the brain tissue of *L. rohita*. The mRNA sequence obtained for *L. rohita* was partial and it was multiple aligned with *B. Taurus*, *M. musculus*, *R. norvegicus*, *X. tropicalis* and *H. sapiens*. The phylogenetic analysis revealed that the tree was very well bifurcated into two branches with *H. sapiens* at the top and *L. rohita* at the bottom. *H. sapiens* and *B. tarus* arise from a same point of origin whereas *R. norvegicus* and *M. musculus* have arisen from a different branch which reveals that they share quite conserved regions. *L. rohita*, *S. scrofa* and *X. tropicalis* have risen from a common point which reveals that *L. rohita* is more closely related to *S. scrofa* and distantly related to *H. sapiens*. The present identified gene might be the AKR1C2 which is specific to brain of Indian major carp. The identification of other genes AKR1C1, AKR1C3, AKR1C4 and AKR1C5 need to be studied in this carp to state whether the 3 α -HSD expression is tissue specific. Further studies are required to understand the expression of the enzyme in the brain which will highlight the presence or absence of AKR1C2 or AKR1C1.

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