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Identification of zebrafish histamine H₁, H₂ and H₃ receptors and effects of histaminergic ligands on behavior

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ABSTRACT

Neuronal histamine regulates several functions in the vertebrate brain. The zebrafish brain contains a widespread histaminergic system and H₃ receptor ligand binding has been reported. In this study we provide evidence for the existence of histamine H₁, H₂ and H₃ receptor genes in zebrafish. Single copies of putative histamine H₁, H₂ and H₃ receptors were identified and cloned from the zebrafish brain. Expression analysis suggested that they are expressed in the brain and a few other tissues. Widespread distribution of zebrafish H₂ receptor binding sites was detected with [¹²⁵I]iodoaminopotentidine in brain sections. Zebrafish larvae were exposed to 1, 10 or 100 μM of the H₁ ligand pyrimidine, the H₂ ligand cimetidine and the H₃ ligands thioperamide and imipip for 5 days. Significant decreases in swimming distance were observed with the highest dose of all ligands, whereas cimetidine gave a significant decrease also with 1 and 10 μM doses. These results provide the first molecular biological evidence for the presence of histamine receptors in zebrafish. These histamine receptors resemble those of higher vertebrates and they provide a useful model for pharmacological and behavioral studies for characterizing the functions of histamine in more detail.

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1. Introduction

Histamine is involved in several regulatory mechanisms in the brain, including alertness and sleep, seizure threshold, hormone secretion and pain [1–4], but the detailed mechanism behind these functions are poorly established. It acts through at least four types of characterized G-protein-coupled receptors (GPCR) in mammals: histamine H₁, H₂, H₃ and H₄ receptors. Of these receptors only the H₄ receptor has

not been consistently found in the brain [5]. Mammalian histaminergic neurons are located in the tuberomammillary nucleus (TMN) and their fibers innervate the whole brain [6–8], supporting the concept that this is a phylogenetically well preserved system.

The histaminergic system in zebrafish resembles that of other vertebrates [9,10]. Expression of L-histidine decarboxylase has been identified only in the caudal part of the hypothalamus in areas corresponding to the TMN in the

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Abbreviations: α-FMH, α-fluoromethylhistidine; [¹²⁵I]APT, [¹²⁵I]iodoaminopotentidine; dpf, day post fertilization; GPCR, G-protein-coupled receptor; hpf, hour post fertilization; IC, intracellular loop; TM, transmembrane region; TMN, tuberomammillary nucleus 0006-2952/\$ – see front matter © 2007 Elsevier Inc. All rights reserved.

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mammalian hypothalamus [9]. Histamine H₃-like receptor binding and H₃-related G-protein activation have been described in the zebrafish brain [11]. However, no molecular characterization of this receptor has been done in zebrafish or any other fish species. The H₁ receptor has earlier been mapped using in vivo [¹⁴C]2-deoxyglucose autoradiography and in vitro receptor-binding methods in *Tilapia nilotica* [12]. Recently it was shown that both zolantidine and chlorpheniramine give inhibitory avoidance responses in goldfish *Carassius auratus*, suggesting the potential presence of both H₁ and H₂ receptor in fish [13]. To date there is no evidence of the molecular identity of these receptors in fish, including the most commonly used species, zebrafish (*Danio rerio*). The zebrafish brain histamine content can be reduced by α -fluoromethylhistidine (α -FMH), and this reduction is associated with changes in the exploratory behavior and in T-maze performance. These changes might be due to reduced anxiety and some memory-related mechanisms after histamine depletion [14].

The gene for H₁ receptor, as for several other GPCR, lacks introns [15–18]. It has long been known that, in addition to the well known, important function in allergic and inflammatory conditions [19], the H₁ receptor is connected to the sleep-wakefulness cycle. More recent reports suggest that the arousal effect of orexin A is due to activation of histaminergic neurotransmission mediated by H₁ receptor [20–22]. The H₁ receptor expression pattern in the CNS is rather widespread [23,24]. The expression in the hypothalamus suggests the involvement in regulation of feeding, and a role in mediating the effects of leptin has been documented [25]. Activation of the H₁ receptor is coupled to the stimulation of phospholipase C via G_{q/11} G-proteins [26] the important transcription factor NF- κ B [27].

The canine H₂ receptor [28] was the first cloned intron-lacking histamine receptor. It was soon followed by the cloning of the H₂ receptor from several species including human, rat and mouse [29,30]. The H₂ receptor signals through the G_s-protein, leading to stimulation of adenylyl cyclase and activation of CREB [31,32]. The H₂ receptor binding pattern has been well characterized in guinea pig brain using [¹²⁵I]iodoaminopotentidine ([¹²⁵I]APT) [33]. The involvement of H₂ receptors in defensive/escape behavior as a response to fear has been suggested [34–36].

The histamine H₃ receptor was first characterized as an autoreceptor which mediates inhibition of histamine synthesis and release [37,38]. It is a modulator of the release of several transmitters [39–42]. The cloning of H₃ receptor by Lovenberg et al. [43] was followed by identification of receptor subtypes [44–46]. The H₃ receptor signaling couples to the inhibitory G_{i/o} type of G-protein [43,47]. The activation of H₃ receptor causes an inhibition of adenylyl cyclase [43,45]. Distinct pharmacology and expression pattern has been described for some of the isoforms of H₃ receptors. The stimulation of isoforms H_{3A}, H_{3B} and H_{3C} induces phosphorylation of mitogen activated protein kinase (MAPK) [45]. Recently the isoforms H_{3D}, H_{3E} and H_{3F} with an alternative extracellular C terminus from rat were described [48].

The interest in the histamine receptors as drug targets has been increasing, particularly for the H₃ receptor [49,50], because histamine regulates important central functions

and the receptors are abundantly expressed in important brain areas. Existing data on the zebrafish histaminergic system indicates that histamine is mostly present in the CNS of the zebrafish [9], and manipulation of histamine levels alters behavior [14]. Earlier binding studies indicate that the zebrafish has a H₃-like receptor, however, no molecular biological evidence for the presence of any of the histamine receptors is available. The purpose of this study was to identify the genes for the histamine receptors, and find out if histaminergic ligands produce significant behavioral effects in developing zebrafish.

2. Methods

2.1. Experimental animals

Zebrafish (*D. rerio*) larvae and adults, from outbred (originate from local fish shop) and AB strains maintained in the laboratory for several generations, of both sexes were used for the study. The fish were kept at 28.5 °C with a light/dark cycle of 14 h/10 h and fed twice daily. Breeding and raising was done according to Westerfield [51]. The permit to carry out these studies was obtained from the Committee for Animal Experiments of Abo Akademi University and the Office of the Regional Government of Western Finland.

2.2. Cloning of zebrafish H₁, H₂ and H₃ receptor

Sequences showing high similarity to the human H₁, H₂ and H₃ receptor were identified using the Sanger Institute zebrafish blast pages (http://www.ensembl.org/Danio_rerio). The predicted coding sequences for the receptors, based on homology analysis with known histamine receptors, were amplified using PCR. cDNA reverse transcribed (First-Strand Synthesis Kit, Amersham Biosciences, Uppsala, Sweden or SuperScript™ First-Strand Synthesis System for RT-PCR, Invitrogen, BV, Groningen, The Netherlands) from RNA extracted (RNA-wiz™, Ambion Inc., Austin, TX) from zebrafish brains was used as template in the reactions. For the H₁ receptor (Genbank accession no. DQ647806) the following primers were used: forward 5'-TGT CTC TCC TCA CCG TCA TC-3' and reverse 5'-CAA TCC CTT CAT TTA CCC GCT CT-3'. The H₂ receptor coding sequence was amplified using (Genbank accession no. DQ647808), the forward primer 5'-ATG CAA TTT ATA TTC AGC GAT-3' and reverse primer 5'-ACT ACC TTT TAC TCC CAT TAG C-3' were used. The corresponding primers for the H₃ receptor (Genbank accession no. DQ647807) coding sequence were 5'-ATG GAG AGA GAA AAC GCG-3' and 5'-AAT ACT TCT GGT CAA TGT TC-3'. Amplified PCR products were sequenced from both directions by automatic sequencing (PRISM 310; ABI, Norwalk, CT).

2.3. mRNA expression by RT-PCR

RNA was extracted from different developmental stages (3, 6, 11, 24, 48 and 72 hpf, 5, 7 and 10 dpf) and from fresh isolated brains, eyes, gills, intestine, heart, liver, muscle and spleen from 10 adult fish and reverse transcribed as described above. cDNA originating from 0.25 μ g total RNA was used for the PCR

reactions which were carried out with the same primer pairs as above for H₁, H₂ and H₃ receptor, respectively.

2.4. Sequence comparisons and analysis

Multisequence alignments were performed using ClustalW on the Web pages provided by the European Bioinformatics Institute (EMBL-EBI), sequence analyses were carried out with GENSCAN Web Server at MIT. Sequence comparisons were carried out with ALIGN provided by (EMBL-EBI) using the needle method.

2.5. [¹²⁵I]iodoaminopotentidine binding to histamine H₂ receptors

[¹²⁵I]iodoaminopotentidine ([¹²⁵I]APT) was synthesized as described [26] and was received as a kind gift from Professor Rob Leurs and Professor Henk Timmerman (Vrije Universiteit, Amsterdam, The Netherlands). Brain sections were incubated for 3 h at room temperature with 50 mM Na₂/K phosphate buffer, pH 7.5, containing 0.2% gelatin and 0.2 nM [¹²⁵I]APT, 2000 Ci/mmol. To determine non-specific binding, adjacent sections were incubated in the presence of [¹²⁵I]APT and 10 μM tiotidine. The sections were rapidly rinsed in ice-cold 50 mM Na₂/K phosphate buffer, pH 7.5 and then washed 3 × 10 min in cold, fresh buffer. The slides were dipped into ice-cold water and rapidly dried under cold airflow. The sections were apposed to Kodak BioMax MR film (Kodak, Rochester, NY) for 1 day.

2.6. Treatment of zebrafish with histamine receptor ligands

Ligand treatment of young fish was done in 6-well plates. Thirty-five fish were kept in 3 ml E3 (5 mM NaCl, 0.17 mM KCl, 0.4 mM CaCl₂, and 0.16 mM MgSO₄) with 1, 10 or 100 μM, pyrilamine, cimetidine, thioperamide or imipemip. The control groups were kept in E3. The treatment was started at 2–3 h post fertilization (hpf). At 1-day post fertilization (dpf) the fish were dechorionated to ease the penetration of ligand into embryo. The E3 and ligands were changed once every day to fresh solutions. The behavior was tracked at 5 dpf.

Thioperamide and imipemip were kind gifts from Professor Rob Leurs and Professor Henk Timmerman (Vrije Universiteit, Amsterdam, The Netherlands). Pyrilamine and cimetidine were purchased from Sigma (St. Louis, MO).

2.7. Video recording of behavior

Young zebrafish exposed to histaminergic ligands for 5 dpf were used for studies of behavioral changes.

In order to minimize effects on social behavior the fish were individually tracked in separate observation tanks. In the test the swimming performance of young fish were recorded simultaneously [14]. The experiments were done in a calm sealed-off area.

The swimming performance of young fish was recorded with a digital video camera connected to a standard PC computer system running the EthoVision Pro 3.1 software (Noldus Information Technology, Wageningen, The Netherlands). Due

to the size of young fish they were placed individually in 48-well plates where they were tracked for 10 min using sample rate 5 samples/s. The coordinates of the swimming performance were stored and used for further software analysis of the behavior. Analysis was performed with input filter 0.2 cm for minimum distance moved to get rid of system noise. One exclusion criterion were used; when the sample size did not exceed 90% of the maximum sample size the tracks were excluded. Statistical analysis was performed with Student's t-test (GraphPad Prism, GraphPad Software Ink, CA).

3. Results

3.1. Cloning and expression of zebrafish H₁, H₂ and H₃ receptors

The zebrafish histamine receptors were cloned to provide evidence of the molecular identity. The zebrafish putative H₁ receptor gene is located on chromosome 8. The intronless gene codes for a 534 aa long peptide. This is 57 aa longer than the human H₁ receptor and 2 aa shorter than the predicted canine H₁ receptor. The zebrafish H₁ has a 20 aa extension in the 5' end and a 12, 3 and 5 aa inserts in the intracellular loop (IC) 3 compared to human H₁. The zebrafish peptide has a second methionine 17 aa downstream from the first, based on Clustal W multiple sequence analysis this is more likely the real translation start site (Figure 1 in supplementary data). Peptide comparisons display 40–46% identity with the corresponding mammalian receptor (Table 1).

The putative zebrafish H₂ receptor is an intronless gene located on chromosome 15, which displays 43–47% identity on the peptide level with the corresponding peptides from dog, human and mouse (Figure 2 in supplementary data). The zebrafish H₂ is 51 aa longer than the corresponding receptor from human, with 27 aa more in the 5' end, a 6 aa longer IC3 and 13 aa longer 3' end.

Sequence comparisons with the corresponding mammalian receptor reveal highest identities in the transmembrane (TM) regions for both zebrafish H₁ and H₂ receptor. The aspartic acid in TM2, conserved in all studied aminergic receptors, exists also in the zebrafish H₁ and H₂ receptor (Figures 1 and 2 in supplementary data).

In the zebrafish H₂ receptor the alanine-lysine-arginine (AKR) and glutamic acid-histidine-lysine-alanine (EHKA) motifs in IC3 suggested to be involved in G_s binding are conserved.

The putative zebrafish H₃ receptor gene contains three exons and is located on chromosome 7. The first exon contains the start site and continues to the middle of TM2, the second exon runs to the beginning of IC2. The third exon contains the rest of the coding sequence. Peptide comparisons reveal 42–50% identities with the corresponding sequences from mouse, human and dog (Table 1, Figure 3 in supplementary data). The 1422 nucleotide long sequence has two possible translation start sites. The correct ATG is most likely the later (nucleotide 103–105, Genbank accession no. DQ647807). When compared to rat histamine H₃ receptor isoforms A, B, C, D and E, the zebrafish peptide shows highest identities to rat H_{3A}. As for the H₁ and H₂ receptor, the highest identities compared to other H₃

		TM1	
ratA	MERAPPDGLMNASGTLAGEAAAAGGARGFSAAWTAVLAALMALLIVATVLGNALVMLAFV	60	
ratB	MERAPPDGLMNASGTLAGEAAAAGGARGFSAAWTAVLAALMALLIVATVLGNALVMLAFV	60	
ratC	MERAPPDGLMNASGTLAGEAAAAGGARGFSAAWTAVLAALMALLIVATVLGNALVMLAFV	60	
ratD	MERAPPDGLMNASGTLAGEAAAAGGARGFSAAWTAVLAALMALLIVATVLGNALVMLAFV	60	
ratE	MERAPPDGLMNASGTLAGEAAAAGGARGFSAAWTAVLAALMALLIVATVLGNALVMLAFV	60	
zebrafish	MERENATSLADLD-TFENRRAQYG---QFSPSTSIFLTVMTLTLLVFATVLGNALVILAFV	56	
	*** * * * *		
	TM2		
ratA	ADSSLRTQNNFLLNLAI SDFLVGAFCIPLYVPYVLTGRWTFGRGLCKLWLVDYLLCAS	120	
ratB	ADSSLRTQNNFLLNLAI SDFLVGAFCIPLYVPYVLTGRWTFGRGLCKLWLVDYLLCAS	120	
ratC	ADSSLRTQNNFLLNLAI SDFLVGAFCIPLYVPYVLTGRWTFGRGLCKLWLVDYLLCAS	120	
ratD	ADSSLRTQNNFLLNLAI SDFLVGAFCIPLYVPYVLTGRWTFGRGLCKLWLVDYLLCAS	120	
ratE	ADSSLRTQNNFLLNLAI SDFLVGAFCIPLYVPYVLTGRWTFGRGLCKLWLVDYLLCAS	120	
zebrafish	VEKSLRTQGNFFFLNLAIADFLVGGFCIPVYIPYVLTGEWRLGRGLCKLWLVDYMLCTA	116	
	***** **		
	TM3	TM4	
ratA	SVFNIVLISYDRFLSVTRAVSYRAQQGDTTRRAVRKMALVWVLAFLLYGPAILSWEYLSGG	180	
ratB	SVFNIVLISYDRFLSVTRAVSYRAQQGDTTRRAVRKMALVWVLAFLLYGPAILSWEYLSGG	180	
ratC	SVFNIVLISYDRFLSVTRAVSYRAQQGDTTRRAVRKMALVWVLAFLLYGPAILSWEYLSGG	180	
ratD	SVFNIVLISYDRFLSVTRAVSYRAQQGDTTRRAVRKMALVWVLAFLLYGPAILSWEYLSGG	180	
ratE	SVFNIVLISYDRFLSVTRAVSYRAQQGDTTRRAVRKMALVWVLAFLLYGPAILSWEYLSGG	180	
zebrafish	SVFNIVLISYDRFLSVTKAVSYRCQGITKDAVLKMLCVWLAFLLYGPAISWEHITGG	176	
	***** **		
	TM5		
ratA	SSIPEGHCYAEFFYNWYFLITASTLEFFT PFLSVTFFNLSIYLNQRRTRLRLDGGREAG	240	
ratB	SSIPEGHCYAEFFYNWYFLITASTLEFFT PFLSVTFFNLSIYLNQRRTRLRLDGGREAG	240	
ratC	SSIPEGHCYAEFFYNWYFLITASTLEFFT PFLSVTFFNLSIYLNQRRTRLRLDGGREAG	240	
ratD	SSIPEGHCYAEFFYNWYFLITASTLEFFT PFLSVTFFNLSIYLNQRRTRLRLDGGREAG	240	
ratE	SSIPEGHCYAEFFYNWYFLITASTLEFFT PFLSVTFFNLSIYLNQRRTRLRLDGGREAG	240	
zebrafish	SVVPDGECYAEFFYNWYFLMTASTVEFFT PFSVTFHLSIYINIRNRCAMREEQPTYVR	236	
	* * * *****		
ratA	PEPPPDAPSPPPAPPSCWGCWPKGHGEAMPLHRYGVGEAGPGVEAGEAALGGGSGGGAA	300	
ratB	PEPPPDAPSPPPAPPSCWGCWPKGHGEAMPLH-----	273	
ratC	PEPPPDAPSPPPAPPSCWGCWPKGHGEAMPLH-----	273	
ratD	PEPPPDAPSPPPAPPSCWGCWPKGHGEAMPLHRYGVGEAGPGVEAGEAALGGGSGGGAA	300	
ratE	PEPPPDAPSPPPAPPSCWGCWPKGHGEAMPLH-----	273	
zebrafish	LR-SFKMKPLGAGDVQRVFFVRPVEESRVADLASRSRCCRLASTAKVSAAEFNGRQSKR	295	
	* * *		
	CA		
ratA	ASPTSSSGSSSRGTERPRSLKRGSKPSAS-SASLEKRMKMVSQSITQRFRLSRDKKVAKS	359	
ratB	-----SSGSSSRGTERPRSLKRGSKPSAS-SASLEKRMKMVSQSITQRFRLSRDKKVAKS	327	
ratC	-----RSGSKPSAS-SASLEKRMKMVSQSITQRFRLSRDKKVAKS	311	
ratD	ASPTSSSGSSSRGTERPRSLKRGSKPSAS-SASLEKRMKMVSQSITQRFRLSRDKKVAKS	359	
ratE	-----SSGSSSRGTERPRSLKRGSKPSAS-SASLEKRMKMVSQSITQRFRLSRDKKVAKS	327	
zebrafish	RDSTLADLPPLQVEERILAASEAQFHYVDHSAGPHRHRPDMVASLANRSCLSRDKKVAKS	355	

	TM6	TM7	
ratA	LAIIVSIFGLCWAPYTLLMIIRAACHGRCIPDYWYETS-----FWLLWANSVNPVLY	412	
ratB	LAIIVSIFGLCWAPYTLLMIIRAACHGRCIPDYWYETS-----FWLLWANSVNPVLY	380	
ratC	LAIIVSIFGLCWAPYTLLMIIRAACHGRCIPDYWYETS-----FWLLWANSVNPVLY	364	
ratD	LAIIVSIFGLCWAPYTLLMIIRAACHGRCIPDYCVERLGKLEASLLPLWMFSGRWRRRK	419	
ratE	LAIIVSIFGLCWAPYTLLMIIRAACHGRCIPDYCVERLGKLEASLLPLWMFSGRWRRRK	387	
zebrafish	LSVIVCVFGLCWAPYTLLMIIRAACHGQCQVHYLYEIS-----FWLLWINSSINPILY	408	
	* ** *****		
ratA	PLCHYSFRRRAFTK-----LLCP---QKLKVQPHGSLEQC-----WK	445	
ratB	PLCHYSFRRRAFTK-----LLCP---QKLKVQPHGSLEQC-----WK	413	
ratC	PLCHYSFRRRAFTK-----LLCP---QKLKVQPHGSLEQC-----WK	397	
ratD	HVCELDVPWMFNQERQNCRGARGWIGRCGLPRPPPSVLQLPAEPRQLLLPAPPPLGRWP	479	
ratE	HVCELDVPWMFNQERQNCRGARGWIGRCGLPRPPPSVLQLPAEPRQLLLPAPPPLGRWP	447	
zebrafish	PLCHSSFKRAFSK-----LLCP---SKTKIQPNIDQKY-----	439	
	* * *		
ratA	-----		
ratB	-----		
ratC	-----		
ratD	CPACPVTIRIWGWVVMG	497	
ratE	CPACPVTIRIWGWVVMG	465	
zebrafish	-----		

Fig. 1 – Sequence identities. Peptide sequence comparisons of the zebrafish H₃ receptor with the different isoforms of H₃ receptor in rat. Amino acids identical in all peptides are marked by an asterisk. The transmembrane regions 1–7 are marked TM1–7 and printed with gray letters. The area involved in constitutive activity of the receptor is marked CA and shaded.

Table 1 – Peptide comparison of the zebrafish H₁ (GenPept accession no. ABF71708), H₂ (GenPept accession no. ABF71710), and H₃ (GenPept accession no. ABF71709) peptide to the corresponding sequence from human (H₁ GenPept accession no. NP_000852, H₂ GenPept accession no. NP_071640, H₃ GenPept accession no. NP_009163), mouse (H₁ GenPept accession no. NP_032311, H₂ GenPept accession no. AAM47009, H₃ GenPept accession no. NP_598610), dog (H₁ GenPept accession no. AAD14167, H₂ GenPept accession no. AAD14163, H₃ GenPept accession no. AAO63755), predicted sequence from chick (H₁ GenPept accession no. XP_425153, H₂ GenPept accession no. XP_425153, H₃ GenPept accession no. XP_425117) and urchin (H₁ GenPept accession no. AAW88352, H₂ GenPept accession no. XP_794263, H₃ GenPept accession no. XP_794616)

H ₁	Human	Mouse	Dog	Chick	Urchin	Zebrafish	zf H ₂	zf H ₃
Human	100						19	25
Mouse	80	100					18	26
Dog	79	70	100				18	24
Chick	46	46	43	100			20	24
Urchin	27	28	29	28	100		15	16
Zebrafish	40	46	40	40	30	100	17	17
H ₂	Human	Mouse	Dog	Chick	Urchin	Zebrafish	zf H ₁	zf H ₃
Human	100						16	16
Mouse	85	100					17	17
Dog	87	80	100				15	16
Chick	56	58	57	100			16	16
Urchin	22	22	21.4	21	100		14	15
Zebrafish	44	47	43	45	20	100	17	20
H ₃	Human	Mouse	Dog	Chick	Urchin	Zebrafish	zf H ₁	zf H ₂
Human	100						18	17
Mouse	94	100					18	17
Dog	75	75.5	100				18	16
Chick	59	59	48	100			14	15
Urchin	24	24	26	14	100		16	16
Zebrafish	50	49	42	30	21	100	17	20

zf: zebrafish.

receptors are found in the TM regions and the aspartic acid (D) in TM2 is conserved. The 3' end of IC3, suggested to be involved in constitutive activity, is well conserved in the zebrafish H₃ receptor (Fig. 1). The rest of the IC3 shows low homologies compared to the corresponding receptor from human and rat. The aa 119 and aa 122 are known to be involved in the species differences of the H₃ pharmacology (REF). The zebrafish has threonine as aa 119 and valine as aa 122, this is similar to the canine receptor.

3.2. Histamine receptor expression

A signal in the RT-PCR reaction for the H₁ and H₂ receptor was detected in all developmental stages tested, indicating that mRNA for these receptors is expressed throughout the development (Fig. 2A). The product in the H₃ receptor RT-PCR reaction from 3 hpf was very weak, at 6 hpf the signal was more clearly visible.

RT-PCR was done on RNA from adult zebrafish brain, eye, gill, heart, intestine, liver, muscle and spleen for the histamine receptors (Fig. 2B). A product in the brain was detected for all three receptors. In addition, primers for the H₁ receptor gave clear products in the RNA from intestine, liver and spleen. The H₂ receptor product was also visible in gills, heart and spleen. The H₃ receptor expression was strong, in addition in the

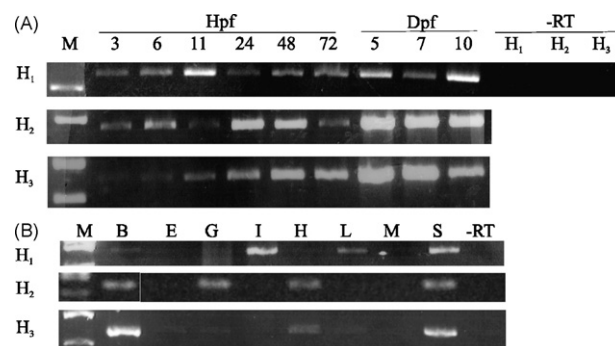


Fig. 2 – Expression of histamine receptors during development and in different adult tissues. RT-PCR the H₁, H₂ and H₃ receptor. (A) Developmental stages 3, 6, 11, 24, 48, 72 hpf and 5, 7 and 10 dpf showing PCR products for the H₁, H₂ and H₃ receptor. (B) RT-PCR products for H₁, H₂ and H₃ receptor from different adult tissues. Brain (B), eye (E), gill (G), intestine (I), heart (H), liver (L), muscle (Mu), spleen (S). RT controls are processed as the samples, contains RNA but lacks reverse transcriptase. In (B) the H₂ panel is reordered by cutting and pasting to have the samples in the same order for all three panels.

GenPept accession numbers for the rat receptors; AF237919 (H_{3A}), BAA88767 (H_{3B}), BAA88768 (H_{3C}), AAZ93637 (H_{3D}), AAZ93638 (H_{3E}).

brain, in heart and spleen, weaker bands were seen in liver, eyes and gills.

3.3. H_2 receptor binding

Binding sites for [125 I]APT were identified in most parts of the zebrafish brain. High densities of binding were observed in optic tectum, hypothalamus, locus coeruleus and the superior reticular formation (Fig. 3).

3.4. Changes in behavior after ligand exposure

The swimming behavior of young fish was analyzed 5 dpf. Parameters analyzed were total distance moved (cm/recorded time), velocity (cm/s), turn angle ($^{\circ}$), angular velocity ($^{\circ}$ /s) and meander ($^{\circ}$ /cm). The H_2 receptor antagonist cimetidine

decreased the total distance moved at all three concentrations used. The distance moved compared to the control group was decreased to 75% with 1 μ M, 75% with 10 μ M and 62% with 100 μ M cimetidine. For pyrilamine, thioperamide and imipip a significant decrease in swimming behavior was detected only with the highest, 100 μ M, dose at 5 dpf (Fig. 4A). Pyrilamine was the ligand giving the biggest effect, at 100 μ M concentration the fish swam only 14% of the control distance, the 1 and 10 μ M concentration decreased the distance moved to 75% compared to the control. Thioperamide, at the concentrations 1, 10 and 100 μ M decreased the swimming distance to 89%, 95% and 58%, correspondingly. Fish treated with imipip swam 88% (1 μ M), 89% (10 μ M) and 73% (100 μ M) of the control distance. The turning behavior of the fish was affected only by the 1 μ M concentration of pyrilamine, which gave a slightly decreased turning behavior

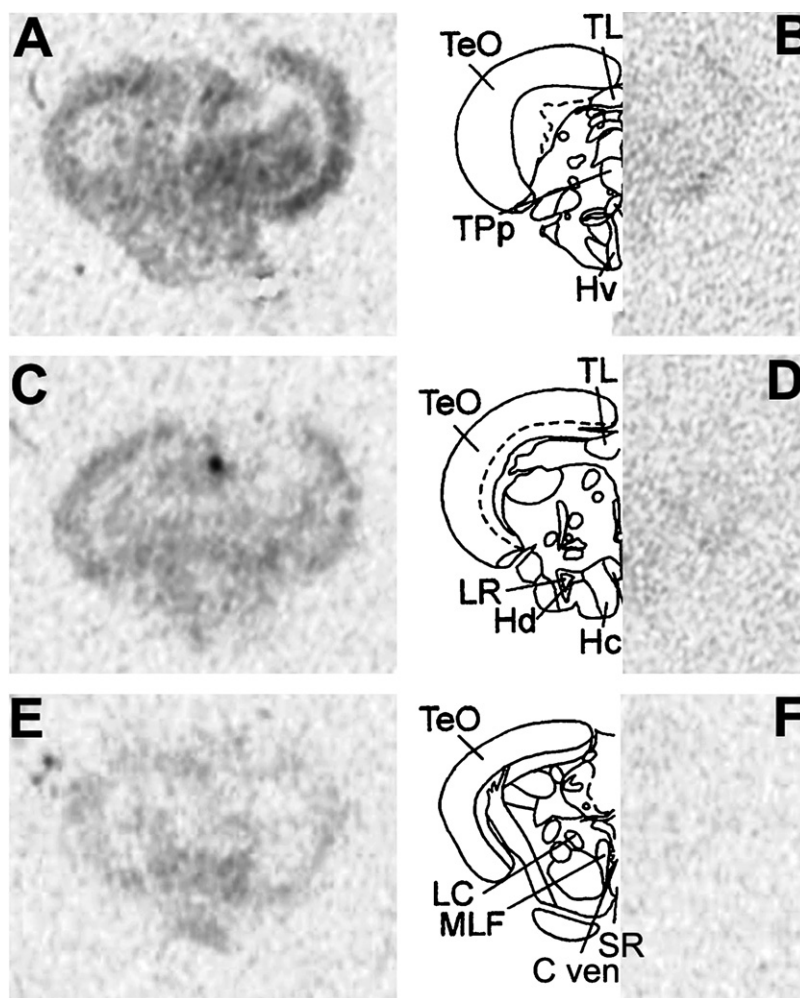


Fig. 3 – H_2 receptor binding sites in zebrafish brain revealed with [125 I]iodoaminopotentidine. (A) Binding sites in optic tectum (TeO) and the central of diencephalon are shown. (B) The corresponding section blocked with 10 mM tiotidine and a schematic drawing of the corresponding level. (C) A more caudal section compared to (A) shows the high binding in TeO and hypothalamic regions. (D) The signal is almost completely blocked in sections treated with 10 mM tiotidine. (E) Caudal part of the hindbrain revealing binding in locus coeruleus (LC) and the superior reticular formation (SR). (F) A corresponding section with the signal blocked with tiotidine is displayed together with a schematic drawing of the section level. C ven, commissura ventralis rhombencephali; Hc, caudal zone of periventricular hypothalamus; Hd, dorsal zone of periventricular hypothalamus; Hv, ventral zone of periventricular hypothalamus; LR, lateral recess of diencephalic ventricle; MLF, medial longitudinal fascicle; TL, torus longitudinalis; TPp, periventricular nucleus of posterior tuberculum.

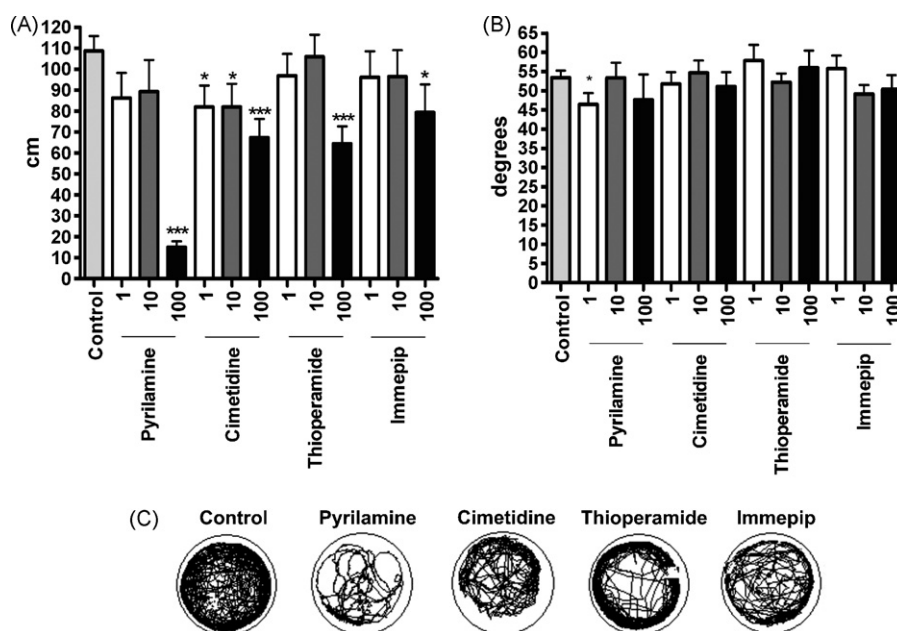


Fig. 4 – Swimming patterns of young zebrafish after treatment with histamine receptor ligands. (A) Total distance moved of young fish at 5 dpf. Bars indicate means and standard deviation. Statistical analysis preformed with Student's t-test, *** $P < 0.001$, * $P < 0.05$. N varies in the different groups; at 5 dpf control = 81, pyrillamine 1 μM = 34, 10 μM = 35, 100 μM = 36, cimetidine 1 μM = 38, 10 μM = 37, 100 μM = 37, thioperamide 1 μM = 38, 10 μM = 42, 100 μM = 45, immeipip 1 μM = 38, 10 μM = 41, 100 μM = 36. **(B)** Turning angel, the data represents the turning behavior of the same group of fish as in (A). **(C)** Typical swimming patterns for the different treatments at 100 μM ligand concentration at 5 dpf.

(turn angle) in the 5-day-old fish (Fig. 4B). Typical swimming tracks for the 100 μM treated groups at 5 dpf are shown in Fig. 4C. The pattern of the swimming did not change significantly after larvae were treated with histamine ligands. Thioperamide-treated fish tend to swim along the walls of the well, while other histamine ligand treated larvae exhibited similar patterns of middle crossing/wall circling as the control group.

4. Discussion

Cloning of the sequences for the putative H_1 , H_2 and H_3 receptors in zebrafish based on homology searches of the genbanks suggests that behavioral effects observed in fish after treatment with histamine receptor ligands are potentially mediated through specific receptors. No putative zebrafish H_4 sequences were found at the time of the searches, using the human and rat H_4 sequence information. No other closely related sequences were identified in the databases.

Amino acids known to be of importance for ligand binding to the H_2 receptor are an aspartic acid in TM3 and aspartic acid and threonine in TM5 [52]. Aspartic acid-arginine-tyrosine (DRY) residues in the IC2 have been reported to be of importance for the receptor stability [53], and this motif is conserved in zebrafish. The AKR and EHKA motifs in IC3 are suggested to be involved in G_s G-protein binding [54]. Both the residues are crucial for ligand binding and G_s binding are conserved in the putative zebrafish H_2 receptor. The results in this study on zebrafish H_2 receptor indicate that the

pharmacology and signaling mechanisms resemble those of higher vertebrates, although agonist/inverse agonist properties of ligands are difficult to predict.

A single form of putative zebrafish H_3 receptor was identified in this study. This resembles the isoform H_{3A} from rat and isoform H_3 1 from human, the so called full-length receptor. Further studies are needed to clarify if there are isoforms of zebrafish H_3 receptor.

The constitutive activity of the H_3 receptor is ensured by the end of the IC3 and the two first amino acids of TM6 [55]. This area is almost completely conserved in zebrafish, indicating the possibility of a similar constitutive activity, as in rodents, of histamine H_3 receptor. This would further strengthen the proposed involvement of zebrafish histamine H_3 receptor in modulation of higher behaviors. Furthermore, the aspartic acid residue in TM3 known to be conserved in aminergic receptors [56], is conserved in the three zebrafish histamine receptors H_1 , H_2 and H_3 receptor.

Using autoradiography and cloning of the putative peptide we indicate here that the zebrafish has a histamine H_2 receptor, which has not been characterized earlier in fish. Earlier pharmacological studies suggest the presence of H_2 receptors in peripheral tissues of cod [57]. A putative H_3 receptor, which may correspond to the receptor ligand binding published earlier [11], was cloned and found to display 42–50% identity with the corresponding mammalian receptor and 30% to the predicted chick H_3 receptor. All three zebrafish histamine receptors were expressed in the brain as revealed with RT-PCR, suggesting involvement in CNS regulation. In addition to this the H_1 receptor is expressed in intestine, liver

and spleen. Expression of H₂ receptor was found in the periphery in gills, heart and spleen and the H₃ receptor clearly in heart and spleen. Interestingly, all three receptors were also expressed very early during development, which may indicate developmental regulatory roles.

Binding sites for the H₂- and H₃-ligands are identified in the zebrafish brain in corresponding areas as in mammals. The aspartic acid residue in TM3 known to be conserved in aminergic receptors [56], is conserved in the three zebrafish histamine receptors H₁, H₂ and H₃ receptor. This data indicates that the H₁, H₂ and H₃ receptors have remained highly conserved between zebrafish and mammals.

The binding pattern for the H₃-like receptor was described earlier [11]. Attempts to map the binding of the H₁-ligand [³H]mepyramine in adult zebrafish brain sections failed due to the low intensity of the signal/small size of the tissue sections and long exposure times, making it difficult to distinguish the specific signal from background. H₂ receptor binding with [¹²⁵I]APT revealed autoradiography signals throughout the brain with high densities in optic tectum, hypothalamic areas, locus coeruleus and the superior reticular formation in a pattern, comparable with the widespread, distribution of this receptor in mammals [33]. H₂ receptor ligand binding signals were abolished with tiotidine, indicating that the detected signal is specific for the H₂ receptor. We have earlier identified the binding sites for an H₃-like receptor in zebrafish brain [11]. Binding data for both H₂ and H₃ receptors indicated that these receptors regulate integrated functions in several areas of zebrafish brain, e.g. optic tectum.

To assay possible behavioral involvement of these receptors young zebrafish were exposed to ligands for the H₃ receptor through water. The drug concentration in the water was 1, 10 or 100 μM. The exact amounts that the fish take up from water are difficult to determine, which may affect the behavioral results obtained. The young fish in this study showed a declined distance moved after exposure to 100 μM of all histamine receptor ligands studied. The H₁ antagonist pyrilamine decreased the distance moved to 14% compared to the control group. This is in agreement with the data from H₁ knock-out mice, which display a reduced exploratory behavior and decreased locomotor behavior [18]. Classical H₁ antagonists to which pyrilamine belongs, or antihistamines, are well known to cause a sedative effect, leading to a decreased movement. Interestingly, since the H₂ knock-out mice seem to lack a CNS phenotype [58], the H₂ receptor antagonist cimetidine gave rise to a significant decline in movement already at the concentration 1 μM. The H₃ receptor agonist immpip and inverse agonist thioperamide induced similar changes in fish behavior with the highest dose used. The decrease seen with thioperamide was, however, to 60% of control, whereas that of immpip showed a smaller decline to 74% of the control group value. This similar type of behavior may be due to agonist/inverse agonist properties of H₃ receptor ligands that differ from those at mammalian receptors, or regulation of unrelated phenomena that give similar results.

The behavioral changes in zebrafish after exposure to agonist/antagonist for the receptors have, to some extent been identified in mammals. Our data indicate that histamine receptors are functional in zebrafish brain and we provide here

the first, to our knowledge, molecular evidence of zebrafish histamine receptor genes. These results on the zebrafish histaminergic system is involved in measurable behaviors and further examination of its genomic organization will help reveal the evolutionary functions of histamine. The results also suggest that zebrafish larvae could be a feasible model to test pharmacological compounds in vivo, affecting the locomotor activity. However, the functional properties of the zebrafish H₁, H₂ and H₃ receptor need to be elucidated, with e.g. in vitro cell system with zebrafish signaling properties.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at [doi:10.1016/j.bcp.2007.01.014](https://doi.org/10.1016/j.bcp.2007.01.014).

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