

NIH Public Access

Author Manuscript

Gene Expr Patterns. Author manuscript; available in PMC 2014 December 01.

Published in final edited form as:

Gene Expr Patterns. 2013 December ; 13(8): . doi:10.1016/j.gep.2013.06.005.

Expression of Arginine Vasotocin Receptors in the Developing Zebrafish CNS

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Abstract

Vasotocin/vasopressin is a neuropeptide that regulates social and reproductive behaviors in a variety of animals including fish. Arginine vasotocin (AVT) is expressed by cells in the ventral hypothalamic and preoptic areas in the diencephalon during embryogenesis in zebrafish suggesting that vasotocin might mediate other functions within the CNS prior to the development of social and reproductive behaviors. In order to examine potential early roles for vasotocin we cloned two zebrafish vasotocin receptors homologous to $AVPR_{1a}$. The receptors are expressed primarily in the CNS in similar but generally non-overlapping patterns. Both receptors are expressed in the forebrain, midbrain and hindbrain by larval stage. Of note, $AVTR_{1a}$ -expressing neurons in the hindbrain appear to be contacted by the axons of preoptic neurons in the forebrain that include $avt+$ neurons and from sensory axons in the lateral longitudinal fasciculus (LLF). Furthermore, $AVTR_{1a}$ -expressing hindbrain neurons extend axons into the medial longitudinal fasciculus (MLF) that contains axons of many neurons thought to be involved in locomotor responses to sensory stimulation. One hypothesis consistent with this anatomy is that AVT signaling mediates or gates sensory input to motor circuits in the hindbrain and spinal cord.

1. Introduction

Signaling within the CNS via the nonapeptide, arginine vasopressin (AVP) in mammals and its homolog arginine vasotocin (AVT) in nonmammalian vertebrates, regulates social and reproductive behaviors in a wide variety of species (reviewed in Donaldson and Young, 2008). Dysfunction of signaling by AVP and oxytocin, another nonapeptide implicated in social behaviors, is thought to contribute to psychiatric disorders such as autism, affective disorders, obsessive-compulsive disorder, posttraumatic stress disorder and schizophrenia (reviewed in Heinrichs et al., 2009). Furthermore, AVP/AVT released into the circulation by the posterior pituitary in response to sexual stimulation, stress and dehydration mediates a variety of peripheral effects including antidiuretic activity by the kidney (Leng and Bicknell, 1986; Nishimura and Fan, 2003).

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AVP is expressed by neurons of the supraoptic, paraventricular and suprachiasmatic nuclei of the hypothalamus in mammals (Brownstein et al., 1980; Young and Gainer, 2003) and AVT primarily by neurons of the preoptic area in fish (Venkatesh and Brenner, 1995; Acher et al., 1997). Additionally AVP is expressed by the bed nucleus of the stria terminalis and amygdala in the mammalian brain (DeVries and Buijs, 1983; DeVries et al., 1985). There are 3 AVP receptors in mammals with the AVPR1a (V1a) and AVPR1b (V1b) receptors expressed primarily in the CNS and the AVPR₂ (V2) receptor in the periphery (Caldwell et al., 2008). AVT receptors have been identified in a number of teleosts as well (Mahlmann et al., 1994; Conklin et al., 1999; Warne, 2001; An et al., 2008). In the pupfish and perhaps other teleosts there appear to be two V1a receptors and a V2 receptor for AVT (Lema, 2010). Like the mammalian V1a receptor, RT-PCR found that the pupfish V1a receptors are widely expressed throughout the CNS.

AVT is expressed by cells in the ventral hypothalamus and the preoptic area of the diencephalon, and isotocin in the preoptic area (Tessmar-Raible et al., 2007; Eaton et al., 2008; Blechman et al., 2011) during embryogenesis in zebrafish prior to the development of social or reproductive behaviors. During embryogenesis behaviors exhibited by zebrafish embryos are restricted to simple motor responses such as escape swimming evoked by sensory stimulation (Saint-Amant and Drapeau, 1998). The embryonic expression of AVT suggests that it may participate in the development of sensory and/or motor circuits early in development. As a first step in examining the function of AVT signaling in the embryonic CNS, we cloned two V1a type receptors in zebrafish and determined their expression patterns during early stages of development. The expression pattern of the V1a receptors is concordant with the hypothesis that AVT signaling may play a role in early sensory/motor function.

2. Results

2.1. Zebrafish contain two AVT receptors homologous to AVPR1a

We cloned two AVT receptors by RT-PCR from adult zebrafish brain tissue, $AVTR_{1a1}$ and AVTR_{1a2} , that were 60% and 62% identical at the amino acid level with human V1a receptor and 54% and 52% identical with human V1b receptor, respectively (blastp, www.ncbi.nlm.nih.gov) (Fig. 1A). A phylogenetic analysis of zebrafish $AVTR_{1a1}$ and $AVTR_{1a2}$ was consistent with the assignment of the zebrafish receptors as V1a type (Fig. 1B). These findings are consistent with previous findings of a duplicated V1a receptor in teleosts (Lema, 2010).

2.2. avtr1a1 and avtr1a2 are expressed primarily in the CNS in early stage zebrafish

By 25 hours postfertilization (hpf) avtr1a1 and avtr1a2 are expressed by a cluster of cells in the forebrain and by a small number of discrete cells in the hindbrain (Fig. 2A). The earliest expression seen via in situ hybridization was *avtr1a1* in the forebrain and hindbrain at 22 hpf (not shown). Examination of avtr1a1 expressing cells in embryos in which all axons are labeled with anti-acetylated -tubulin found that the forebrain cells are in apparent contact with the postoptic commissure (POC) and/or tract of the postoptic commissure (TPOC) (Fig. 2B; Chitnis and Kuwada, 1990), which is consistent with these cells projecting axons into the POC/TPOC. These *avtr1a1*+ neurons as the putative *avtr1a1*+ epiphyseal and nucleus of the posterior commissure neurons (see below) may also express HNK-1 since the pattern of early neurons/axons labeled with anti-HNK-1 and anti-acetylated -tubulin are similar (Wilson et al., 1990). Additionally there are irregularly spaced, occasional dorsal cells in the spinal cord of unknown identity that express *avtr1a1* (Fig. 2C).

By 48 hpf *avtr1a1* is expressed by a cluster of ventral forebrain cells (I), forebrain cells near the dorsal midline (II), dorsal cells located at the forebrain/tectum boundary (III) ventral cells near the forebrain/tegmentum border (IV), and two longitudinal stripes of cells in the midbrain/anterior hindbrain region (V and VI) when viewed from a dorsal perspective as well as the posterior hindbrain cells (Fig. 3A-C). Lateral views show that in the ventral forebrain cluster I is located anterior and ventral to cluster IV (Fig. 3D). Examination of cluster I avtr1a1+ cells in 48 hpf embryos with all axons labeled with antiacetylated tubulin showed that these neurons appear to project axons into the TPOC (Fig. 4A) suggesting that these are the POC/TPOC forebrain neurons observed earlier (Fig. 2B). Examination of forebrain *avtr1a1*+ cells in *otpb.A:GAL4; UAS:gfp* embryos that express GFP in preoptic neurons (Fujimoto et al., 2011) showed that cluster I avtr1a1+ cells are adjacent to the preoptic neurons at 48 hpf (Fig. 4B). Furthermore at 72 hpf avtr1a1 is expressed in the region immediately posterior to the preoptic region, either ventral or dorsal to the preoptic neurons (not shown). Thus preoptic neurons do not express *avtr1a1* during the embryonic stages examined.

The dorsal forebrain cells (II) are found at the lateral base of the epiphysis, which can be seen with DIC optics in 48 hpf embryos (Fig. 3A inset). These cells are likely neurons since they appear to extend axons in the dorsal-ventral diencephalic tract (DVDT) (Fig. 5A; Chitnis and Kuwada, 1990; Wilson et al., 1990). The cluster of neurons on the left side of the brain appears to be larger than that on the right side (Fig. 3A inset) in accord with the left-right asymmetry exhibited by the pineal complex in embryonic zebrafish (Concha et. al., 2000). The forebrain/tectum boundary cells (III) could be neurons of the nucleus of the posterior commissure (nucPC) since they appear to extend axons in the PC (Fig. 5B; Chitnis and Kuwada, 1990). The *avtr1a1* expressing cells near the midbrain/hindbrain boundary (V) at approximately 48 hpf could be neurons of the nucleus of the Medial Longitudinal Fasciculus (nucMLF) since these neurons reside in this region. nucMLF neurons express $pitx2c$ and are labeled by EGFP in $pitx2c:egfp$ zebrafish (Wolman et al., 2008). However, in situ hybridization for $avtr1a1$ in these transgenic embryos showed that the $avtr1a1$ expressing cells were in the same region as the nucMLF neurons but did not express $pitx2c:egfp$ suggesting they were not nucMLF neurons (Fig. 5C). It is noteworthy that the medial longitudinal stripe of cells (V) are also located in a position similar to that of the raphe serotonergic neurons (Lillesaar et. al., 2007).

Given that many neurons in the hindbrain are characterized by their stereotyped locations, morphology and molecular properties, we also examined the avtr1a1+ hindbrain cells in order to determine their identities. The caudal most $avtr+$ cells are located in the region of the T reticular neurons that project into the contralateral MLF and can be labeled with MAb RMO44 (Skromne et al., 2007), which recognizes a specific neurofilament expressed by these neurons. Double-labeling with RMO44 and *in situ* hybridization for *avtr1a1* showed that the caudal *avtr1a1* positive neurons are not T reticular neurons (Fig. 5D). Additionally, the double-labeled embryos demonstrated that the M cell which projects posteriorly into the contralateral MLF and are also labeled by RMO44 do not express avtr1a1.

At 48 hpf *avtr1a2* appears to be expressed by fewer cells in the midbrain/anterior hindbrain region compared with *avtr1a1* while the forebrain expression of the receptors is similar to that for avtr1a1 (Fig. 6A,D). Since *avtr1a1* and *avtr1a2* are expressed in similar patterns in the brain, it is possible that they may be co-expressed in neurons. However, double in situ hybridization of 48 hpf embryos for both receptor genes found that the two receptors appeared generally not to be co-expressed by neurons (Fig. 6B,C,E,F).

avtr1a1 is expressed by cells outside of the CNS with expression seen in the endothelial cells of the developing vasculature in the trunk at 24 hpf (Fig. 2C); the pharyngeal arches; a

midline structure ventral to the brain between the eyes (Fig. 3D); and cells in the space between the lens and the retina by 48 hpf (Fig. 3E).

2.3 The morphology and location of hindbrain avtr1a+ neurons are consistent with a role in sensory/motor responses

We determined the axonal trajectories of $avtr1a1+$ neurons by combined in situ hybridization and immunolabeling with anti-acetylated tubulin. We found that the hindbrain avtr1a1+ neurons project axons towards and into the ipsilateral MLF (Fig. 7A). The fact that these *avtr1a1*+ neurons project into the MLF indicates that these neurons may participate in regulating motor responses since many neurons with axons within the MLF are known to have a motor function in teleosts including zebrafish (Zottoli, 1977; Nissanov et al., 1990; Gahtan et al., 2002). A group of cells known as IC neurons are located in the caudal hindbrain and project axons in the ipsilateral MLF (Mendelson, 1986; Metcalfe et al., 1986). IC neurons are rhythmically active during spontaneous coiling of the body (Saint-Amant and Drapeau, 2001). At present there are no markers for these neurons so it is not possible to readily identify whether IC neurons are avtr+.

Many neurons that participate in motor responses in zebrafish receive sensory input (Gahtan et al., 2002; Kohashi and Oda, 2008). In the zebrafish, hindbrain sensory axons extend along the lateral longitudinal fasciculus (LLF). Axons in the LLF appear to course over the cell bodies of the avtr1a1-expressing neurons as seen in confocal stack images and in single focal planes (Fig. 7B), so potentially could make synapses with the $avtr1a1+$ neurons.

The finding that the axons of *avtr1a1*+ neurons project into the MLF and the proximity of their cell bodies to the RB sensory axons of the LLF are consistent with a role for $avtr1a1+$ neurons in motor responses of embryos to sensory stimulation. Further physiological and morphological analysis of $avtr1a1+$ neurons will be required to establish this hypothesis.

2.4 Preoptic neurons project axons in the vicinity of avtr1a1-positive neurons

Since avt, avtr1a1, and avtr1a2 are expressed early in zebrafish, we wondered if the avt+ neurons innervate the receptor-expressing neurons in embryos. Unfortunately anti-AVT appears not to label AVT-expressing neurons until 5 dpf (Eaton et al., 2008), so we examined transgenic embryos in which avt+ neurons expressed GFP. Preoptic neurons express AVT and isotocin, an ortholog of mammalian oxytocin, and this requires expression of the homeobox transcription factor, orthopediab, in preoptic neurons (Blechman et al., 2007; Ryu et al., 2007; Eaton et al., 2008; Lohr et al., 2009). Preoptic neurons can be visualized in the otpb.A:egfp and otpb.A:GAL4; UAS:gfp transgenic zebrafish that express GFP in preoptic neurons (Fujimoto et al., 2011). To examine whether preoptic neurons could innervate *avtr*+ neurons, transgenic *otpb.A:egfp^{caax}* embryos that express membranetargeted EGFP in preoptic neurons were examined in combination with in situ hybridization for *avtr1a1*. In *otpb.A:egfp^{caax}* embryos EGFP-labeled axons from preoptic neurons projected posteriorly. In confocal stack images the posteriorly-projecting axons extend in the medial longitudinal catecholaminergic tract (MLCT; Kastenhuber et al., 2010) in close proximity to *avtr1a1* expressing neurons in the anterior and caudal hindbrain (Fig. 8). This could be seen in single focal planes as well (not shown). Thus preoptic neurons including those presumably expressing *avt* could be presynaptic to the $avt +$ neurons. However, since preoptic neurons include those that express isotocin as well as vasotocin, physiological and anatomical examination of confirmed *avt* expressing neurons and $avt +$ neurons will be required to establish synaptic interactions.

3. Discussion

Although AVP/AVT signaling within the brain has been widely studied as an important factor for reproductive and social behaviors (reviewed in Stoop, 2012), relatively few investigations have examined the development of AVP/AVT signaling during CNS development. Indeed AVP and oxytocin binding sites are found in the mammalian neonatal spinal cord (Liu et al., 2003; Stoop, 2012). Furthermore, it is likely that these neonatal spinal neurons receive input from AVP expressing axons: AVP is expressed by paraventricular (PVN) and supraoptic hypothalamic neurons in rat embryos (Lipari et al., 2001), AVP expressing PVN cells innervate the spinal cord (Sawchenko and Swanson, 1982; Motawei et al., 1999; Hallbeck et al., 2001), and PVN axons project to the neonatal spinal cord (Leong et al., 1984; Kudo et al., 1993; Lakke, 1997). Our study demonstrates that AVT receptors are expressed in the CNS early in zebrafish development together with AVT (Eaton et al., 2008). This suggests that AVT signaling may regulate some aspect of CNS development (e.g. Boer et al., 1982a & b; Hammock and Levitt, 2012) and/or control early behaviors.

The expression of AVT and its receptors correlate with the earliest behaviors of zebrafish embryos. These are spontaneous coiling of the body and motor responses to sensory stimulation, touch-induced fast escape response and swimming (Saint-Amant and Drapeau, 1998). Thus it is possible that AVT signaling may play a role in regulating these early motor behaviors. Several findings from our study are consistent with this hypothesis. First, AVT receptor expressing neurons in the hindbrain project axons into the MLF as do many other neurons known to regulate motor behaviors in zebrafish. Second, the morphological relationship between sensory axons of the LLF and the AVTR expressing hindbrain neurons suggests that sensory neurons could synaptically activate $avtr+$ neurons. Third, $otpb$ expressing preoptic neurons, which include AVT expressing neurons, project axons into the hindbrain with the axons in close proximity to $avtr+$ neurons. Since isotocin, the fish homolog to oxytocin, is also expressed during embryogenesis (Eaton et al., 2007), it is possible that both AVT and isotocin may regulate motor behaviors in zebrafish embryos.

Our hypothesis assumes that the interaction between the $avt+$ preoptic neurons and the hindbrain *avtr*+ neurons is a synaptic one based upon release of neuropeptide from axon terminals. AVP and oxytocin, as are other neuropeptides, are found in large dense core vesicles in neurons and can be released in a Ca^{2+} dependent fashion directly from the cell bodies and dendrites (Brownstein et al., 1980; Ludwig and Leng, 2006). Recently, however, physiologically relevant synaptic release of oxytocin from axon terminals was elegantly demonstrated in the amygdala (Knobloch et al., 2012). In the case of zebrafish embryos the cell bodies of $avt+$ preoptic neurons are in the diencephalon while the $avt+$ neurons in question are located in the hindbrain. The distance between the *avt*+ preoptic cell bodies and avtr+ hindbrain neurons and the physical proximity of preoptic axons and the hindbrain neurons is consistent with a synaptic release from axon terminals of preoptic neurons. However it is also possible that AVT could act in a humoral fashion via release into the early vascular system and/or ventricles.

We hypothesize that binding of AVT by the hindbrain $avt+$ neurons increases the excitability of these neurons and thus regulates the responsiveness of the receptor expressing neurons to sensory stimulation in zebrafish (Fig. 9) much like the sensorimotor processing role proposed for AVP/AVT (Rose and Moore, 2002). In fact AVP and oxytocin depolarize motor neurons, interneurons and preganglionic neurons in the rodent spinal cord (Suzue et al., 1981; Kolaj and Renaud, 1998; Oz et al., 2001; Liu et al., 2003). Application of AVP onto facial motor neurons and oxytocin onto vagal motor neurons leads to a TTX insensitive, voltage-dependent persistent I_{Na} (Raggenbass et al., 1991) while in neurons in the lumbar spinal cord AVP suppresses resting conductance to K^+ and enhances

conductance to cations (Ogier et al., 2006). Furthermore, AVT induces an inward current in oocytes expressing flounder avtr (Warne, 2001). Finally AVP can induce rhythmic activity in the spinal cord by itself or in combination with serotonin suggesting that AVP can activate central pattern generators (Pearson et al., 2003). Interestingly, the invertebrate homologs of AVP and oxytocin, conopressin, hirudotocin and annetocin, induced activation of a central pattern generator for reproductive behaviors in the Medicinal leech (Wagenaar et al., 2010) suggesting that regulation of motor behaviors might be an evolutionarily ancient function for these neuropeptides.

4. Materials and methods

Fish breeding and maintenance

Zebrafish (Danio rerio) were maintained in a breeding facility following the guidelines set forth by the University of Michigan Animal Care and Use protocols (Zhou et al., 2008). The fish were provided daily with either dry food (Micro pellets, Kyorin, Japan) or brine shrimp (Brine Shrimp Direct, Utah). Embryos were collected and incubated at 28.5 °C and staged according to hours post fertilization (hpf; Westerfield, 1995). For in situ hybridization of the embryos older than one day post-fertilization, the embryos were kept in 0.2 mM 1-Phenyl-2 thiourea (Sigma, P-7629) to keep them transparent until they were fixed at desired developmental stages. Transgenic lines used in this paper were *otpb.A:GAL4* $(Tg(\text{otp: GAL4-VP16}, \text{my17: GFP}^{2c57})$ and $Tg(\text{UAS: GFP})$ (Fujimoto et al., 2011); otpb.A:egfp^{caax} (Tg(otpb:1EGFP)^{zc49}) (Xing et al., 2012); and pitx2c:egfp $(Tg(\text{pitx2:EGFP}^{zy8})$ (Wolman et al., 2008).

cDNA cloning of zebrafish AVTRs

cDNA cloning for AVTRs was conducted following standard protocols (Zhou et al., 2008). Whole brains were dissected from male adult zebrafish (approximately 14 months post fertilization, N=4) and total RNA was extracted following the manufacture's protocol for TRIzol® Reagent (Invitrogen, #15596-018). The RNA was reverse transcribed with oligodT primers (Invitrogen, #58862) following the manufacture's protocol for SuperScript® II Reverse Transcriptase (Invitrogen, #18064-022). For the PCR of avtr1a1 cDNA, the following primers were used to amplify the coding sequence along with 5 -UTR and 3 - UTRs; forward primer, 5 -GCTCGCGCCTTTACGCATGA-3 , and reverse primer, 5 - ACAGGAGGGTAAATGCTTTTGACT-3. To PCR *avtr1a2*, the following primers were used; forward primer, 5 -CGCGCGTATTTCCATCAATCAAGC-3 , and reverse primer, 5 - AATCGCTGTGCTTTCAGCTGGT-3 . The PCR products were purified and cloned into pGEM T-easy vector following the manufacture s instructions (Promega, #A137A). The obtained clones were sequenced at the University of Michigan Sequencing Core.

Sequence analysis

Amino acid alignments for human V1a (NP_000697), zebrafish AVTR1a1 and AVTR1a2 (Fig. 1) were constructed with ClustalW (Biology Workbench, SCSD; [http://](http://seqtool.sdsc.edu) [seqtool.sdsc.edu\)](http://seqtool.sdsc.edu). Default settings of the textshade option (Biology Workbench) were adopted except that amino acid identity (dark gray) and similarity (light gray) were colorcoded. The amino acids were numbered corresponding to that for zebrafish $AVTR_{1a1}$. The transmembrane domains for zebrafish $AVTR_{1a1}$ (black bars) were determined using MEMSAT3 (<http://bioinf.cs.ucl.ac.uk/psipred>). The white box indicates the DUF1856 domains (unknown function) conserved in the C-termini of various AVP receptors (conserved domain search; <http://www.ncbi.nlm.nih.gov/cdd>) . The intracellular loops (ICL) and extracellular loops (ECL) are also indicated. Phylogenetic tree was constructed based on the alignment of amino acid sequences for vasopressin/vasotocin receptors across vertebrate species with the MEGA 4.0 software (neighbor-joining method with pairwise deletion and

bootstrap values from 1000 replicates) as described by Lema (2010). GenBank accession numbers are provided in the parentheses.

Whole-mount in situ hybridization & immunolabeling

To synthesize RNA probes for *in situ* hybridization for *avtr* genes, partial coding sequences (685 bp for *avtr1a1 & 658 bp for avtr1a2)* were cloned into pGEM T-Easy vectors using the following primers; forward primer (avtr1a1), 5 -CTTGGGAATGTTCGCGTCCACTTA-3, reverse primer (avtr1a1), 5 -TTAGGCTGTTTCCCATGCTGGAAC-3, forward primer $(avtr1a2)$, 5 -GGACTTTCTGTGCAGGATCGTCAA-3, and reverse primer $(avtr1a2)$, 5 -TCCGCTGAACACCATGTAGATCCA-3 . The partial clones were sequenced at the University of Michigan Sequencing Core and confirmed to be identical to the complete AVTR cDNA clones described above (cDNA cloning of zebrafish AVTRs). The partial cDNA clones were used to synthesize, in vitro, digoxigenin (DIG)-labeled antisense probes with DIG RNA labeling mix (Roche) following the manufacture's instructions for mMESSAGE mMACHINE Kit (Ambion, Life Technologies). The sense probe for avtr1a1 was used as a negative control and found to generate no significant labeling (not shown). The standard protocol was followed to conduct *in situ* hybridization (Zhou et al., 2008). Anti-DIG-AP Fab fragments (Roche) was used to carry out color reactions using NBT/BCIP as a substrate (Roche). For fluorescent double-labeling, Fast Red (Roche) was used as a substrate for *in situ* hybridization and immunolabeling was carried out with primary antibodies [antiacetylated tubulin (1:1000, Sigma-Aldrich), anti-GFP (1:1000, Torrey Pines Biolabs), anti-Neurofilament-M/RMO44 (1:500, Invitrogen, Life Technologies)] and Alexa488-anti-mouse/rabbit IgG (1:500, Invitrogen, Life Technologies). The labeled embryos were mounted in Vectashield mounting medium (Vector Laboratories) and fluorescent images were acquired with a Leica SP5 laser scanning confocal microscope.

For double in situ hybridization, dinitrophenol (DNP)-labeled anti-sense riboprobes for avtr1a1 were synthesized with DNP-11-UTP (PerkinElmer), and DIG-labeled riboprobes for avtr1a2 were synthesized using DIG RNA labeling mix (Roche). Prior to in situ hybridization, the embryos were treated with 2% hydrogen peroxide, and dextran sulfate was added to the hybridization reaction as described by Lauter et al. (2011). The alkaline phosphatase-mediated color reaction was first carried out for avtr1a2 using anti-DIG-AP Fab fragments (Roche) with NBT/BCIP (Roche) as a substrate. After anti-DIG-AP was inactivated by 0.1 M glycine-HCL (pH 2.2), the color reaction for *avtrial* was conducted using anti-DNP-AP (Vector Laboratories) with INT/BCIP (Roche) as a substrate. The labeled embryos were mounted in 70 % glycerol to obtain bright-field microscope images.

Acknowledgments

We thank M. Halloran and J. Yost for the *pitx2c:egfp* transgenic fish and A. Migda for the care of zebrafish. Research was supported by grants from NINDS (RO1NS54731) and NSF (0725976) to JYK.

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Two vasotocin receptors are expressed in the CNS of zebrafish embryos as is vasotocin.

Vasotocin receptor neurons may be innervated by sensory and vasotocin neurons.

Vasotocin receptor expressing neurons may participate in locomotor responses.

Vasotocin signaling in the embryonic CNS may gate sensory input to motor circuits.

Fig. 1.

Zebrafish $AVTR_{1a1}$ and $AVTR_{1a2}$. (A) Amino acid alignment for human V1a (NP_000697), zebrafish $AVTR_{1a1}$, and zebrafish $AVTR_{1a2}$. There is a high degree of amino acid conservation between human V1a and zebrafish AVTRs. The dark gray boxes indicate amino acid identity and light gray boxes similarity. The amino acid number corresponds to that for zebrafish $AVTR_{1a1}$. The black bars mark the transmembrane domains (TMs) for zebrafish $AVTR_{1a1}$. The white box indicates the DUF1856 domains conserved in the Ctermini of various AVP receptors. The locations of the intracellular loops (ICL) and extracellular loops (ECL) are also indicated. (B) The phylogeny of AVP/AVT receptors.

Fig. 2.

Vasotocin receptors (avtr1a1 and avtr1a2) are expressed in discrete populations of cells in the CNS of 24/28 hpf embryos. (A) Dorsal views (anterior up) of in situ hybridizations of 25 hpf embryos showing that both *avtr1a1* and *avtr1a2* are expressed in the forebrain $(A1, A3)$ and the posterior hindbrain (A2, A4). Arrows denote position of the posterior border of the otocyst. Scale:100 m. (B) Combined in situ hybridization (red) and anti-acetylated tubulin labeling of axons (green) showing *avtr1a1* expressing neurons in the forebrain appear to project axons into the postoptic commissure (white arrow) and tract of the postoptic commissure (28 hpf, lateral view with anterior right and dorsal up). Yellow arrow denotes the anterior commissure. Scale: 50 m. (C) Lateral view of trunk of 24 hpf embryo showing avtr1a1 expressing neuron in the dorsal spinal cord (arrow) and apparent endothelial cells forming blood vessels (arrowhead). Scale: 100 m.

Fig. 3.

More neurons express $avtr1a1$ at 48 hpf. (A) View of the brain focused dorsally showing $avtr1a1+$ cells in the base of the epiphysis (II) and the forebrain/tectum boundary (III). Inset shows a higher magnification view of the cluster II avtr $1a1+$ cells with asterisk denoting the epiphysis and dotted line the midline. Scale: 100 m for (A-C). (B) Same view focused more ventrally than (A) showing the *avtr1a1*+ cells at the forebrain/tegmentum boundary (IV) and two rough stripes of cells that cross the midbrain/hindbrain boundary (V and VI). (C) Same view focused even more ventrally showing the anterior forebrain *avtr1a1*+ cells (I). (D) Lateral view showing the $avtr1a1+$ cells in the anterior forebrain (I), near the dorsal midline of the forebrain (II), at the forebrain/tectum boundary (III), at the forebrain/tegmentum boundary (IV) and the lateral stripe near the midbrain/hindbrain boundary (VI). Outside the brain the pharyngeal arches (arrowhead) and a midline structure in between the eyes (arrow) also express $avtr1a1$. Scale: 100 m. (E) View of eye showing $avtr1a1$ + cells (arrowhead) between the lens and the retina. Scale: 100 m.

Fig. 4.

At 48 hpf the *avtr1a1*+ cells in the anterior forebrain (I) are the POC/TPOC neurons and not preoptic neurons. (A) Dorsal perspective (anterior up) of an embryo labeled with *avtr1a1* riboprobe (red) and anti-acetylated tubulin labeled axons (green) showing that the forebrain avtr1a1+ cells appear to extend axons in the TPOC (arrow). Scale: 50 m. (B) A ventral perspective of an otpb.A:GAL4; UAS:gfp embryo labeled with avtr1a1 riboprobe showing that the anterior forebrain $avtr1a1+$ neurons (red) are located just medial to the preoptic neurons (green, yellow arrow). Blue arrow denotes the olfactory neurons that express GFP in the transgenic embryos. Scale: 100 m.

Fig. 5.

The epiphyseal and nucPC neurons but not the nucMLF nor T reticular neurons express avtr1a1 at approximately 48 hpf. (A) Lateral view (anterior left, dorsal up) of a z-stack of confocal images of a embryo double labeled for avtr1a1 and anti-acetylated tubulin (axons) showing that the cluster II *avtr1a1*+ cells at the lateral base of the epiphysis (red, asterisk) project axons into the DVDT (green, arrowhead) and the $avtr1a1+$ cluster III cells (red, star) are adjacent to the PC (arrow). The $avtr1a1+$ cells seen ventrally are the neurons in the ventral forebrain and near the forebrain/tegmentum boundary (I and IV). Scale: 50 um. (B) A single focal plane seen in a lateral view showing that the cluster III avtr1a1+ cells (red) appear to extend axons in the PC (green, arrows). Scale: 50 m. (C) Ventral perspective of $pitx2c:egfp$ embryos labeled with a *avtr1a1* riboprobe showing that the nucMLF neurons (green, bracket) do not express avtr1a1 (red). Arrowhead denotes the MLF; asterisk denotes the forebrain *avtr1a1*+ neurons. Scale: 100 m. (D) Dorsal perspective (anterior left) of the hindbrain of embryo double labeled for *avtr1a1* riboprobe (red, bracket) and MAb RMO44 (green) showing that the posterior hindbrain *avtr1a1*+ cells are not the T reticular neurons. The arrowhead indicates *avtr1a1*+ cells in the otocyst. Scale: 100 m.

Fig. 6.

avtr1a2 is expressed in a pattern similar to avtr1a1 but they are not coexpressed by neurons. Dorsal perspective showing that $avtr1a2$ is expressed in the brain (arrow and arrowhead) (A) and hindbrain (D) at 48 hpf. Double in situ hybridizations for avtr1a1 and avtr1a2 shows that brain neurons generally do not coexpress *avtr1a1* and *avtr1a2* (B and C) nor do posterior hindbrain neurons (E and F) at 48 hpf. B and C are the same field of view from the same embryo as are E and F. Scale for A-D: 100 m and for E-F: 50 m.

Fig. 7.

avtr1a1 expressing neurons in the caudal hindbrain project axons into the ipsilateral MLF and are in close proximity with the LLF at 28 hpf. (A) Lateral view (anterior left) showing caudal hindbrain, $avtr1a1+$ neurons (red) project axons (labeled with antiacetylated tubulin) into the ipsilateral MLF (arrow). Scale: 20 m. (B) Lateral view showing that the LLF (green, bracket) courses over the avtr1a1+ neurons (red). Asterisk denotes a RB sensory neuron. Inset: single confocal plane showing LLF axons coursing over $avtr1a1+$ hindbrain neurons. Scale: 50 m.

Fig. 8.

avtr1a1+ neurons are in close proximity to the axons of otpb expressing preoptic neurons at 49-51 hpf. (A) Lateral view (anterior left) in an *otpb.A:egfp^{caax}* embryo showing that the anterior hindbrain $\frac{avtr1a1}{+}$ neurons (red) are in close proximity to the axons of preoptic neurons in the MLCT (arrow). Scale: 50 m. (B) Lateral view in an *otpb.A:egfp^{caax}* embryo showing that the $avtr1a1$ + neurons in the caudal hindbrain are in close proximity to the preoptic axons in the MLCT (arrow). Scale: 50 m.

Fig. 9.

Model for the regulation of the responsiveness of $avtr+$ neurons in the hindbrain to sensory input in zebrafish embryos. In the model AVT is constitutively released at the synapse onto $avtr+$ neurons to enhance activation of these neurons by sensory stimulation and modulate motor responses mediated by the central pattern generator (CPG). Dashed arrow denotes hypothesized connection between the hindbrain avtr+ neurons and the CPG.