Immunolocalization of choline acetyltransferase of common type in the central brain mass of *Octopus vulgaris*

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Abstract

Acetylcholine, the first neurotransmitter to be identified in the vertebrate frog, is widely distributed among the animal kingdom. The presence of a large amount of acetylcholine in the nervous system of cephalopods is well known from several biochemical and physiological studies. However, little is known about the precise distribution of cholinergic structures due to a lack of a suitable histochemical technique for detecting acetylcholine. The most reliable method to visualize the cholinergic neurons is the immunohistochemical localization of the enzyme choline acetyltransferase, the synthetic enzyme of acetylcholine. Following our previous study on the distribution patterns of cholinergic neurons in the *Octopus vulgaris* visual system, using a novel antibody that recognizes choline acetyltransferase of the common type (cChAT), now we extend our investigation on the octopus central brain mass. When applied on sections of octopus central ganglia, immunoreactivity for cChAT was detected in cell bodies of all central brain mass lobes with the notable exception of the subfrontal and subvertical lobes. Positive varicose nerves fibers where observed in the neuropil of all central brain mass lobes.

Introduction

It is universally acknowledged that acetylcholine (ACh), the first neurotransmitter to be discovered,1 plays an important role in neurotransmission and is of great significance for the assessment of the neurochemical organization of the nervous system. The cholinergic system has been assumed to play various roles in motor function, but also in more complex functions, such as learning, in both vertebrates14 and invertebrates,15 and may modulate memory,16 responding to the visual and other sensory inputs in the cephalopod.10,11

The invertebrate central nervous system (CNS) is a good model to use to study the distribution of ACh and its associated enzymes, and the study of the cephalopod nervous system has played a crucial role in our understanding of the biochemical mechanism underlying cholinergic synaptic transmission.12-14 Cephalopods, such as octopus, squid and cuttlefish, are highly evolved invertebrates and have some of the largest and most complex nervous systems among the invertebrates, with processing and computational capabilities that in some tasks rival those of some vertebrates, including a marked ability to learn.15-19 Cephalopod CNS have been shown to contain the same major neurotransmitters that are found in the mammals’ brains19 and it may be possible to use immunohistochemistry to define those areas of the cephalopod CNS that are analogous to neural regions in mammals.20,21 Biochemical information is available on the distribution of ACh within the CNS of cephalopods.10-19 There is already substantial evidence showing that the involvement of cholinergic neurotransmission in learning and memory processes appear to be differentially modulated in specific CNS areas of cuttlefishb and that ACh performs an important physiological role in mediating neuronal signaling at synapses of the cuttlefish optic lobe and into the octopus statocyst sensory hair cells.22,23 The presence of ACh in the cephalopod nervous system, in particular the octopus optic lobe, and the activities of both acetylcholinesterase (AChE) and choline acetyltransferase (ChAT) has been shown,10,24,25 but very little is known about the distribution of cholinergic structures.

Bacq and Mazza13 first reported the occurrence of ACh in the octopus and the activities of both ChAT and AChE, two ACh metabolizing enzymes. The wide distribution of ACh and AChE in the octopus nervous system indicates that, presumably, the central ganglia of this animal should also contain such large amounts of ACh, as these ganglia predominantly consist of cholinergic neurons.20 ChAT, the synthetic enzyme of ACh, has been widely used as a specific marker for cholinergic neurons. ChAT has been isolated from neuronal structures of both vertebrates and invertebrates. Successful purification of ChAT proteins from vertebrates led to the production of a number of specific ChAT antibodies that has made it possible to confirm by immunohistochemistry a detailed map of the cholinergic systems of mammals20-24 and non-mammalian vertebrates.35-39 A similar approach has been used to produce antibodies against purified ChAT from invertebrates, though this has been limited to animals such as flies,40-42 moths,43,44 and worms.44 Conversely, few data are available on ChAT immunoreactive (IR) structures in octopus because none of the available ChAT antibodies cross-react sufficiently with octopus ChAT, whose chemical structure at protein and mRNA levels remains unknown.

In our recent immunohistochemical study,28 we used a new antiserum45 to label ChAT of the common type (cChAT) in the optic lobe and peduncle complex of cephalopod mollusk *Octopus vulgaris* (Cuvier 1797). Western blot and immunoprecipitation analyses indicate that cChAT antisem recognizes an octopus ChAT-like protein which is capable of producing ACh. Using this antisem, we found cChAT immunoreactivity in the retina, optic lobe, and its neighboring peduncle complex, and IR cell bodies in the cell islands of the optic lobe medulla and the cortical layer of the posterior olfactory lobe. IR fibers and nerve terminals were also found in the plexiform layer of the deep retina, within the stroma of the optic gland, and the neuropils of the optic, peduncle, and olfactory lobe. In the present study, we used the same cChAT antiserum to extend our investigation to delineate the distribution pattern of cChAT-IR neuronal cells.
and fibers in the other parts of octopus central nervous system (CNS).

Organization of the central nervous system

The octopus CNS is made up of central and peripheral ganglia. The CNS includes central brain mass and optic lobes. The central brain mass is encased in a cartilaginous capsule which is penetrated by the esophagus. It is partitioned by the esophagus into a supra- and a sub-esophageal mass, which are bilaterally interconnected. Laterally and tightly connected to the distal boundaries of the central brain mass, the paired optic lobes are situated adjacent to the eyes. Both supra-esophageal and sub-esophageal masses are further subdivided into numerous, mostly bilateral, symmetrical brain lobes that are functionally differentiated. Each lobe is generally described as being composed of a peripheral region of cell layers (cortex) surrounding a neuropil consisting of fiber network.

The magnocellular lobes lie laterally to the esophagus, linking the supra- and sub-esophageal regions, and the chromatophore lobes lie on the latero-dorsal side of the sub-esophageal mass. All lobes of the CNS lie inside a cartilaginous brain capsule except the optic lobe that lies adjacent to each eye outside the cartilage.

Materials and Methods

Tissue preparation

Specimens of octopus (Octopus vulgaris), each weighing approximately 1.3 kg, collected either from the Island Sea of Japan near Akashi or from the sea near the Straits of Messina (Sicily, Italy) were kept alive in cold seawater. Animals were killed under anesthesia by adding 2% tricaine (ethyl 3-aminobenzoate methanesulfonate salt; MS-222, Sigma, St. Louis, MO, USA) to seawater at 15°C and used for immunohistochemical examination, as described elsewhere. Central brain mass was quickly dissected out. For Western blot analysis, the central nervous system was stored at -80°C until use. For immunohistochemistry, the tissues were fixed for 24 h in ice-cold 0.1 M phosphate buffer (PB, pH 7.4) containing 4% paraformaldehyde and 0.2% picric acid. After washing with PB, the tissues were immersed for at least 24 h in PB containing 15% sucrose. Before immunohistochemical staining, the sections were incubated for at least 3 days in PB containing 0.9% NaCl (PBS) plus 0.3% Triton X-100 at 4°C to improve tissue permeability.

cChAT antibody

The production of cChAT antiserum has been described previously. The specificity of the antibody used to octopus cChAT-like molecule has been previously described. For Western blotting, aliquots of either cChAT primary antisera or pre-immune serum of the rabbit which produced cChAT antiserum, each containing 100 μg of protein, were labeled using the Peroxidase Labeling Kit-NH2 (Dojindo Molecular Technologies, Inc., Kumamoto, Japan) according to the manufacturer’s instructions.

Western-blot

Octopus central brain mass was homogenized with 10% (w/v) cold Tris-HCl (50 mM, pH 7.4) containing protease-inhibitor cocktail (P-2714, Sigma) using a Polytron device. After centrifugation at 12,000g for 20 min, the supernatant was collected as a crude extract of octopus central brain mass. Protein concentration was measured using a protein assay kit adapted from the method of Bradford (Bio-Rad Laboratories, Tokyo, Japan) with bovine serum albumin as a standard. Aliquots of the supernatant containing 25 μg of protein were electrophoresed on a 5-20% gradient dodecyl sulfate polyacrylamide gel under reductive conditions and then electro-transferred on polyvinylidene difluoride membrane as described previously. After transfer, membrane was fixed in 4% paraformaldehyde in 0.1 M Tris-HCl buffered saline (pH 7.4) (TBS) for 20 min at room temperature. After extensive washing in TBS, strips of the membrane were incubated for 1 h with 10% skim milk in TBS, then incubated overnight at 4°C either with the peroxidase labeled primary antiserum against cChAT or the peroxidase labeled pre-immune serum in an immunoreaction enhancer solution (Can Get Signal, Toyobo, Osaka, Japan). After extensive washing in TBS containing 0.5% tween 20, peroxidase labeled bound antibodies on membrane were directly visualized by reaction with enhanced chemiluminescence reagent (Chemi-Lumi One Super, Nacalai Tesque Inc., Kyoto, Japan) and the signal

Table 1. Semi-quantitative analysis of cChAT immunoreactivity in transverse sections of the octopus central brain mass. Localization and relative abundance of cChAT-IR structures. Lobes are listed along the anterior-posterior axis. Frontal, subfrontal, buccal, vertical, subvertical and basal are supraesophageal lobes; brachial and pedal are subsesophageal lobes; magnocellular and chromatophore lobes lie laterally to the esophagus.

<table>
<thead>
<tr>
<th>Lobes</th>
<th>Immunoreactive cell bodies</th>
<th>Immunoreactive fibers</th>
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<tbody>
<tr>
<td><strong>Buccal</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Superior</td>
<td>++ (m)</td>
<td>+++</td>
</tr>
<tr>
<td>Posterior</td>
<td>++ (m)</td>
<td>+</td>
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<tr>
<td><strong>Frontal</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Superior</td>
<td>+ (s)</td>
<td>++</td>
</tr>
<tr>
<td>Inferior</td>
<td>++ (s)</td>
<td>+++</td>
</tr>
<tr>
<td><strong>Subfrontal</strong></td>
<td>-</td>
<td>+++</td>
</tr>
<tr>
<td><strong>Magnocellular</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Dorsal</td>
<td>++ (s)</td>
<td>++</td>
</tr>
<tr>
<td>Ventral</td>
<td>++ (s) ++ (l)</td>
<td>+</td>
</tr>
<tr>
<td><strong>Brachial</strong></td>
<td></td>
<td></td>
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<tr>
<td>Anterior</td>
<td>+ (s) +++ (m)</td>
<td>+</td>
</tr>
<tr>
<td>Posterior</td>
<td>+ (s) +++ (m)</td>
<td>+</td>
</tr>
<tr>
<td><strong>Vertical</strong></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Subvertical</td>
<td>-</td>
<td>+++</td>
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<tr>
<td><strong>Basal</strong></td>
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<tr>
<td>Dorsal</td>
<td>-</td>
<td>+++</td>
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<tr>
<td>Medial</td>
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<tr>
<td>Lateral</td>
<td>+/- (s)</td>
<td>+</td>
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<tr>
<td><strong>Chromatophore</strong></td>
<td>+++ (s) +++ (l)</td>
<td>+</td>
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</tbody>
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+++, Countless positive cells and/or fibers; ++, numerous positive cells and/or fibers; +, discrete number of positive cells and/or fibers; +/-, scattered positive cells and/or fibers; -, no positive cells and/or fibers. Cell sizes: s, small (5-15μm-15 μm); m, medium (20-30μm-30 μm); l, large (30-50μm-50 μm).
emitted was recorded for 10 min using the Lumino-Image Analyzer LAS-4000 (Fujifilm, Tokyo, Japan).

**Immunohistochemistry**

To inactivate the endogenous peroxidase activity, the sections were pre-treated for 30 min at room temperature with PBS containing 0.3% Triton X-100, 0.1% sodium azide and 0.5% H₂O₂, and, to avoid non-specific binding of serum proteins, incubated for 30 min at room temperature with normal donkey serum 1:50 in PBS containing 0.3% Triton X-100 and 0.5% bovine serum albumin (BSA, Sigma). Serial sections were then incubated for 72 h at 4°C in the primary antibody solution rabbit anti cChAT (diluted 1:50,000). The sections were then incubated with a biotinylated donkey anti-rabbit IgG (Jackson Immunoresearch Laboratories, West Grove, USA; diluted 1:1000) for 2 h at room temperature and then for 1 h at room temperature with avidin-biotin-peroxidase complex (ABC Elite, Vector Laboratories, Burlingame, USA; diluted 1:2000). PBS containing 0.3% Triton X-100 was used for diluting all the reagents and washing sections after each step. The localization of peroxidase activity was visualized by reacting the sections for 3 min at room temperature with a solution containing 0.04% 3,3’-diaminobenzidine tetrahydrochloride (DAB; Fluka, Buchs, Switzerland), 0.4% nickel ammonium sulfate, and 0.003% H₂O₂ in 0.05M Tris-HCl buffer, pH 7.6 giving a dark blue granular precipitate. The stained sections were mounted on glass slides and air-dried. After staining, the sections were then dehydrated, cleared and coverslipped with Permount (Fisher Scientific, Pittsburg, PA, USA) for microscope observation (AX70 Provis, Olympus Optical, Tokyo, Japan). For control experiments, the primary antiserum was substituted with buffer or normal rabbit serum or primary antiserum was pre-absorbed overnight at +4°C with the antigen used for its production (2 µg/mL antiserum at working dilution). None of the control sections showed positive immuno-staining (Figure 1A).

**Image analysis for cells**

We studied cChAT-positive neuronal cells on well-stained central brain mass sections randomly selected from 3 octopuses. We used a computer-assisted image analyzer equipped with a software tool (Hyperfocus, IAS 2000, Deltasistemi, Roma, Italy) to obtain high-resolution microphotographs and also to estimate the cell size of cChAT-positive somata. For pear-shaped somata we measured both their major and minor axes, and for round or ovoid somata we assessed only their mean axes. Cell size values are expressed as the mean ±SD.

**Results**

**Western blot**

We previously reported that a number of octopus tissue molecules reacted with secondary antibodies against anti-rabbit IgG used for detecting cChAT antibody on our Western blot after SDS-PAGE, resulting in non-specific signals. Then, in order to avoid the use of peroxidase labeled secondary antibodies, peroxidase labeled cChAT antiserum and peroxidase labeled pre-immune serum were used. In these conditions, a clearly stained band was observed on immunoblot of SDS-PAGE of crude extract from octopus central brain mass, while the pre-immune serum display no signals (Figure 1B). The molecular weight of the band is approximately 81-kDa which is similar to the size of octopus ChAT enzyme previously reported in optic lobe.29 It should be noted that a number of faint bands were observed but that these were considerably less intense than the major 81kDa. Next, we used sections of pedal lobe, which are normally highly immunoreactive to the cChAT antiserum (Figure 1A), and incubated them with the cChAT antiserum pre-absorbed with antigenic peptide (Figure 1B).

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**Figure 1.** Western blot of crude protein extract (left panel) from the central brain mass of Octopus vulgaris revealed using peroxidase labeled pre-immune serum (left column) and peroxidase labeled cChAT (right column). A clear band at approximately 81-kDa (arrowhead) is recognized by cChAT antiserum. Molecular weight marker is Bench Mark Prestained Protein Ladder (Invitrogen, Carlsbad, CA, USA). cChAT immunoreactivity in the octopus pedal lobe (right panel) using cChAT antiserum (a) or cChAT antiserum pre-absorbed with antigenic peptide (b). Scale bar: 30 µm.
A dramatic decrease in signal intensity was observed on the section incubated with cChAT antiserum pre-absorbed by the antigenic peptide, providing further confirmation of the cross-reaction of the rat cChAT antibody with the octopus cChAT in the tissue used for immunohistochemistry.

**cChAT-immunoreactivity**

Transverse sections of the octopus central brain mass has shown cChAT immunoreactivity in several lobes belonging to both the supraesophageal (frontal, subfrontal, buccal, vertical, subvertical and basal lobes) and subesophageal (brachial and pedal lobes) masses (Table 1). Geographically distinct specimens of octopus were examined but no difference in distribution of immunoreactivity in the central brain mass was found.

**Frontal lobe**

The frontal lobe can be divided into two main parts: superior and inferior frontal lobe. The superior frontal lobe can be further divided into median (the median superior frontal lobe, MSF) and lateral part. We found a discrete number of cChAT-IR small neurons (12.5±1.1 µm) in the MSF (Figure 2A). These IR cells come from the small cell layers surrounding the surface of the lobe. A network of IR fibers in the MSF, weaker than in the lateral superior frontal lobe (Figure 2A and B) was found. The inferior frontal lobe showed small cChAT-IR cells (12.5±1.7 µm) and an intensely IR fiber network in the neuropil. The neuropil of the frontal lobe showed two distinct zones: the outer and the inner zone. In both zones, irregularly tangled cChAT-IR fibers were found (Figure 2B).

Below MSF, in the neuropil of two subfrontal lobes, we observed an intensely cChAT-IR fiber network.

**Buccal lobes**

Buccal lobes include a superior buccal and two posterior buccal lobes. All buccal lobes appeared strongly cChAT-IR. The neuropil of the superior buccal lobe showed a dense network of intensely IR fibers surrounded by numerous medium sized (25±1.9 µm) cChAT-IR neurons. In the posterior buccal lobes, medium sized (25±2.2 µm) cChAT-IR neurons surrounded the neuropil (Figure 2C) where a ring of gathered cChAT-IR fibers surrounded an irregular tangle of small and large IR fibers. In the posterior buccal commissure, strongly cChAT-IR fibers were observed (Figure 2D).

**Magnocellular lobes**

In the cellular layer of both dorsal and ventral magnocellular lobes, several small cChAT-IR cells (7.5±1.1 µm) were found. In the ventral part, large pear shaped cChAT-IR cells (60±2.1×35±0.9 µm) were also found. cChAT-IR fibers were irregularly distributed throughout the neuropil (Figure 2E).

**Brachial lobe**

In the cellular layer of both anterior and posterior brachial lobes, numerous medium sized cChAT-IR pear shaped neurons (40±2.2×25±1.3 µm) were found together with small cells (12.5±0.8 µm). Irregularly tangled cChAT-IR fibers were observed in the neuropil (Figure 2F).

**Vertical lobe**

The vertical lobe, consisting of five gyri, occupies the top of the supraesophageal mass. The gyri communicate between each other by numerous tracts of fibers. The amacrine...
cells, whose fibers do not extend beyond the vertical lobe, were the most numerous in this lobe and had no cChAT-IR. Ventrally the gyri are connected with the subvertical lobe. Faint, large cChAT-IR neurons (52.5±2.4 µm) were observed between the amacrine cell layer and the neuropil. Intensely stained cChAT-IR fibers were found throughout the neuropil and in the connection with the subvertical lobe (Figure 3A). cChAT-IR fibers form a ring of surrounding cChAT-IR varicose fibers in the centre of the neuropil.

In the subvertical lobe, numerous cChAT-IR fibers were observed but no cChAT-IR cells were found (Figure 3B).

**Basal lobes**

In the basal lobes, we can distinguish dorsal, median and lateral lobes. Throughout the basal lobes, a dense IR nerve fiber network was found, more intense than in the dorsal basal lobe (Figure 3C). Scattered small cChAT-IR cells (12.5±0.8 µm) were found in the cellular layer of the median and the lateral basal lobe.

**Chromatophore lobes**

Two numerous groups of cChAT-IR neurons were shown: large pear shaped (65±2.5×40±2.0 µm) and small (15±1.8 µm) neurons. The large neurons were regularly arranged at the periphery, while the smaller neurons were found in the inner side of the cell layer. A dense cChAT-IR rather regular fiber network was shown in the neuropil (Figure 3D).

**Pedal lobes**

Numerous large cChAT-IR pear shaped neurons (60±2.3×45±1.6 µm) were detected in the cellular layer together with small cells (20±1.8 µm) and the neuropil showed a dense cChAT-IR fiber network (Figure 3E).

**Discussion**

A great deal of information is available on the neuroanatomy and the functions of octopus CNS. Five functionally different areas can be distinguished: auxiliary memory centers, receptor analyzers, higher motor centers, lower motor centers and intermediate motor centers. We used histochemical procedures to describe and map the distribution of structures in the octopus central brain mass where cholinergic synaptic transmission is prevalent; this has not been described elsewhere. In agreement with previous data, our results demonstrate the presence of cholinergic structures in lobes which are involved in learning, memory and movement.

Examination of central brain mass sections gave us the opportunity to study cChAT -immunoreactive cell bodies and fibers and their distribution and intensity among the different lobes. Widespread cChAT-IR fibers were observed in the superior frontal, vertical and sub-vertical lobes that belong to auxiliary memory centers; these are areas important for establishing and storing visual or tactile memories. The MSF/vertical lobe system (MSF-V) in octopus is a brain region that has been known for over 50 years to be closely involved in learning and memory. The vertical lobe system also plays a part in tactile learning and is, together with the optic lobes, the main component of the visual learning system. In experiments in which the vertical lobe of *Octopus vulgaris* was lesioned, the ability to learn visual discriminations was drastically impaired and the disruption of the cholinergic systems in the higher motor centers interferes with both learning behaviors and memory recall.

The vertical lobe contains two types of morphologically identified neurons: approximately 25 million small amacrine interneurons that synapse onto only approximately 65,000 large efferent cells. These neurons and their connections together with their afferent fibers from the MSF form an association matrix analogous to the vertebrate hippocampus. So the high numbers of interneurons suggest the importance of a large number of units to form a high redundancy of connections. As these features are found in both octopus MSF-V and in vertebrate hippocampus, it would appear that they are needed to create a large capacity for memory associations. Our immunohistochemical results support the hypothesis that cholinergic inputs from other brain structures convey modulator input signals into the vertical lobe, as do the acetylcholine fibers innervating the vertebrate hippocampus. Therefore,
the organization of areas with comparable function in octopus and in mammals indicates that similar cellular organization has been retained between phylogenetically distinct animals with complex forms of learning. Furthermore, the vertical lobe of the octopus brain is also involved in complex forms of long-term learning and memory.\textsuperscript{39,67} In fact, Hochner \textit{et al.}\textsuperscript{44} found that the octopus vertical lobe shows two different types of mechanisms for induction of long-term potentiation as in different parts of the vertebrate hippocampus. One type appears to depend on the postsynaptic response, while the other does not require a strong postsynaptic response for induction.\textsuperscript{43}

Our results could suggest that, like serotonin,\textsuperscript{41} ACh may act as a neuromodulator in a reinforcing/reward signaling system. These findings reinforce the conclusion that the octopus vertical lobe and the vertebrate hippocampus seem to show a striking example of evolutionary convergence, so the CNS of cephalopods will improve our understanding of the evolution of such associative systems.\textsuperscript{35}

Our observations of ChAT immunoreactivity in the subfrontal lobe are in agreement with its function as a memory store, confirming the putative involvement of cholinergic neurotransmission in central brain learning processes.\textsuperscript{66} It is possible that ChAT-IR cell bodies and fibers may have an effect on memory storage. Indeed, available evidence suggests that the subfrontal lobe seems to be necessary for the performance of tactile discrimination and tactile memory.\textsuperscript{43,66,67}

In the subfrontal lobe, ChAT-IR cell bodies were not detected. This suggests that the numerous ChAT-IR fibers observed in the neuropil have an extrinsic origin. It is possible that this bundle is composed of fibers originating from the inferior frontal lobe and from the posterior buccal lobes, where we clearly see ChAT-IR cell bodies. Some of them end here; others run on with the cerebral tracts to the subvertical lobes. It is presumed that the buccal tracts carry signals of taste from the mouth to the learning centers of the inferior frontal and superior frontal/vertical systems.\textsuperscript{44}

A dense network of intensely stained fibers surrounded by numerous IR cell bodies was found in the superior buccal lobe. Together with the subfrontal lobe, the superior buccal lobe is a system concerned with tactile learning.\textsuperscript{44} A great number of ChAT-IR fibers and large IR neurons were found in the posterior buccal lobes, the center of learning chemotactile discrimination. They receive fibers from the arms and buccal mass, and they send fibers downwards to the arm nervous centers and backwards to the optic and superior frontal/vertical lobe systems.\textsuperscript{44} In octopus, it has been shown that nitric oxide neurotransmission is needed for both visual and chemo-
tactile learning.\textsuperscript{66,67} On the other hand, ACh is known to regulate nitric oxide synthase expression in mammals as well as in snail, another mollusk.\textsuperscript{76,11} The observation here may suggest that regulation of nitric oxide by ACh may also occur in the chemotactile center of octopus.

The presence of numerous ChAT-IR fibers in the dorsal and median basal lobes, higher motor centers, agrees with the involvement of cholinergic system also in movement.\textsuperscript{49} In fact, these lobes receive inputs from both the visual and gravitational (statocysts) systems, and have cerebellar-type effects on motor function.\textsuperscript{39} They form important cell stations on some of the main channels between the optic lobes and other receptor centers, and the intermediate and final motor cells of the subophageal lobes. They also regulate movement, including attack. The setting-up of a memory representing association of a given situation with a shock is a property of the optic lobes shared with the basal lobe, but the persistence of the representation depends upon the presence of the vertical lobe.\textsuperscript{43,63} In the intermediate and in the lower motor centers our findings agree with the involvement of cholinergic system in movement.

Numerous neurons and intensely ChAT-IR fibers were seen in the chromatophore lobes. At a higher level, the chromatophores are controlled by the optic lobes that act on visual information and select specific motor programs. At a lower level, the chromatophores are controlled by the motoneurons in the chromatophore lobes that carry out the programs, so their activity or inactivity produces the patterning seen in the skin.\textsuperscript{22,71} Two different color classes of chromatophore motoneurons, that excite black chromatophores or yellow, orange and red chromatophores, are inhibited by descending cholinergic pathways derived from optic lobes.\textsuperscript{22,25,74,75} Our findings agree with previous investigation on AChE presence in the neuropil of all the chromatophore lobes and the optic lobes,\textsuperscript{22} and with ChAT immunoreactivity in the optic lobes.\textsuperscript{22} In the magnocellular lobes, we found many ChAT-IR cell bodies and fibers. The magnocellular lobes are involved in the escape behavior and receive fibers from the optic lobe; therefore, they are responsible for visual evoked activity.\textsuperscript{52} It has been postulated that visual inputs related to a potential predator, and thus likely to evoke an escape response, are communicated via identified pathways to the magnocellular lobes, so they are involved in direct motor responses.\textsuperscript{66}

A great number of ChAT-IR neurons and intensely stained fibers were found in the pedal lobes. The pedal lobes, intermediate motor centers, integrate signals that result in various movements of the funnel, tentacles, head including movements of the eyes (ocular motor centre).\textsuperscript{67} In this context, it is interesting that Uemura \textit{et al.}\textsuperscript{74} have reported the presence of serotonin-IR cells localized in some of the ganglia we had examined in our study. In the buccal lobes, they found large serotonin-IR cells,\textsuperscript{74} while we detected small, scattered ChAT-IR cells. In the chromatophore lobes, they have reported the presence both of some large and small serotonin-IR cells\textsuperscript{74} and we also found large and small ChAT-IR cells. In the pedal lobes, they have found some large serotonin-IR cells\textsuperscript{74} and we visualized numerous large ChAT-IR cells together with some small ones. Serotonin is a chemical mediator for the control of the change in color, in addition to the cholinergic innervation of the chromatophore system.\textsuperscript{55,74} Investigations still need to be carried out to determine whether cholinergic and serotoninergic systems found in chromatophore and pedal lobes interact.

Cephalopods, a fascinating group of animals, are capable of producing a wide repertoire of behavior and provide an excellent invertebrate model for understanding the neural substrate of learning and memory, where the cholinergic system plays an important part. The roles of octopus brain cholinergic neurons in memory and motor function are reminiscent of those seen in higher vertebrates, including mammals.

\textbf{References}


