

A morphometric study of the amygdala in the common shrew

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The characteristic features of the common shrew amygdala (CA), as shown by volumetric comparisons of the individual nuclei, are the poor development of the lateral (LA) and basomedial (BM) nuclei as well as the particularly strong formation of the basolateral (BL) and lateral olfactory tract (NLOT) nuclei. The central (CE), cortical (CO) and medial (ME) nuclei are also well organised in this species. All these features are even more distinctly visible when the total number of neurons in the nuclei referred to are compared. A comparison of the densities of neurons in the individual nuclei with the mean numerical density of cells in the CA indicates that there are the 3 different regions within the common shrew's CA. The densities of neurons in the LA, BL, and BM are significantly lower than the mean density of cells in the CA ($p < 0.05$). In the CE this value does not differ from the mean ($p > 0.05$). In the CO, ME and NLOT the density values are significantly higher than the mean ($p < 0.05$). Furthermore, a similar division of the shrew's CA can, to some extent, be performed using the size parameters of the amygdaloid neurons as a marker. Interestingly, the large neurons populate less densely organised CA areas like the LA, BL and BM, whereas the small cells populate the ME and NLOT, where the neurons are densely arranged. The CE and CO occupy intermediate positions, with the neurons similar in size to the mean for the shrew's CA.

Key words: amygdala, common shrew, morphometric analysis, 3-D reconstructions

INTRODUCTION

The amygdala (CA) is an extremely complicated and highly organised collection of nuclei and cortical areas, varying in relation to cellular structure [11, 18, 22, 25], connections [56, 68] and neurotransmitters expression patterns [2, 41, 43, 45, 46, 52, 55, 68]. It is currently suggested that there are more than 10 separate structures within the CA [56]. The cellular organisation of these areas is often so diversified that they require additional divisions [18, 25, 56, 68].

In spite of the large amount of data dedicated to the CA, quantitative aspects of its organisation have

remained obscure. For instance, almost nothing is known about the size of the individual nuclei in the CA of species most studied. To our knowledge, only a few volumetric reports are available. The size parameters of the basolateral amygdala were described by Breathnach and Goldby [5] in the porpoise as well as by Berdel et al. [1] in the developing brain of the rat. More detailed volumetric relationships between various amygdaloid nuclei were reported by Maksymowicz [47] in the dog and by Śmiałowski [69] in the macaque. The most extensive volumetric studies on the CA were put forward by Stephan et al. [65–67], although the data presented there are restrict-

ed to insectivora and primates only and, moreover, concern exclusively larger areas of the CA than the single nuclei. Of the individual structures only the NLOT and the large-celled portion of the BL are taken into account. Quantitative aspects of the cellular populations in the mammalian CA have so far received even less attention. For instance, to date no detailed data concerning the density of neurons in the individual CA nuclei are, to our knowledge, available. Where reports on this subject do exist, they are dedicated only to the selectively stained and highly specialised subpopulations of amygdaloid interneurons [2, 41, 46, 55]. The total number of neurons in the CA has only been studied in the selected amygdaloid nuclei of the rat [62]. The size of amygdaloid neurons is usually reported by Golgi studies [22, 38, 49]. However, this technique stains only 3–5% of neurons from the total cellular population in the area studied and the stained sample is not necessarily representative [64]. A detailed morphometric analysis of the amygdaloid neurons has only been carried out by Dzięwiątkowski et al. on the rabbit [11].

The aim of our study, therefore, was to give the first detailed morphometric characteristics of the CA in the common shrew. Since most mammalian species are thought to have had their phylogenetic origin in insectivore-like ancestors, the data on the common shrew can represent a good reference base for evaluating the different trends in brain structure evolution during mammalian phylogeny.

MATERIAL AND METHODS

A total of 6 brains were used to study the morphometric parameters of the CA in the common shrew. All efforts were made to minimise animal suffering and to minimise the number of animals used. The brains were removed from the skulls, embedded in paraffin and sectioned in 50 μm thick slices in the coronal plane.

All sections were analysed cytoarchitectonically and morphometrically with a calibrated image analysis system consisting of a computer equipped with morphometric software (Multi-Scan 8.2, Computer Scanning Systems, Poland) and a light microscope coupled with a digital camera (CM40P, VideoTronic, Germany). The sequence of the successive operations in the single animal is described in detail below.

The three-dimensional reconstructions of the CA. The microscopic fields were viewed at low magnification through the 5-fold objective. The 512 \times 512 pixel partial microscopic images of a single section were digitally recorded by means of a camera cou-

pled with a microscope and computer. The scans were subsequently joined with those of neighbouring sections to form larger areas comprising the whole CA and adjoining structures (so-called "digital slices"). All the available sections were taken into account.

The digital slices were registered by means of TPS (USA) and Morpheus (USA) morphometric software according to the Generalised Procrustes Analysis (GPA) that superimposes the landmark configurations using the least-squares estimates for the translation and rotation parameters [3, 13, 57, 58]. To achieve maximum precision, the slices were registered in pairs. The first image in the pair was the reference slice for the second slice. The landmark configurations were generated on both slices in the pair. The data consisted then of the point's co-ordinates in the reference slice and the corresponding landmarks in the unregistered target slice. The second slab in the pair was translated and rotated into alignment based on these landmarks with the help of the built-in Morpheus GPA procedure. After registration the second slice in the pair was the reference slice in the next pair. The sequence was then as follows: 1–2, 2–3, 3–4 ... 75–76.

All the registered digital slices of the CA in the single animal were incorporated into one multi-sliced 3-dimensional image (a so-called "stack"). In this kind of 3D data set the digital slices were the consecutive serial sections oriented in perpendicular to the long axis of the CA and covering its whole rostrocaudal extent. On each digital slice the boundaries of the individual amygdaloid nuclei were outlined with different colours by a mouse-driven cursor. All the outlines were drawn under the control of the light microscope. The definite stack was then composed of the serially aligned raw images and the incorporated outlined structures of the CA. With the help of the IGL Trace software the 2-dimensional outlines were transformed to the 3-dimensional slabs. The 3-dimensional reconstruction of the CA for the single animal was simply a sum of all the serially aligned slabs.

The volumetric analysis. To evaluate the volume of the individual amygdaloid nuclei the volumes of slabs traced with the same colour were totalled according to the formula proposed by Cavaliero [14, 32]. The total CA volume presented in this study was the sum of the volumes of the 7 amygdaloid nuclei studied, namely the lateral, basolateral, basomedial, central, cortical, medial and lateral olfactory tract nuclei.

The numerical density counts. To evaluate the numerical density of cells in each of the amygdaloid

nuclei, the optical dissector method was implemented with the guidelines described by West et al. [71].

Evaluation of the total numbers of neurons.

The total numbers of neurons in each of the nuclei was calculated by multiplying the volume of the given nucleus by the numerical density of cells in it [71].

The morphometric parameters of neurons. In each of the individuals studied the morphometric analysis of CA neurons was performed using Multi-Scan 8.2 morphometric software. Each neuron was characterised by a set of morphometric parameters: the length (the long axis of the soma), the width (the short axis of the soma), the size (the sum of both axes) and the shape factor (the ratio of both axes). The test frames were arranged this way to cover the total cross-sectional area of the nucleus studied. In the common shrew 5 cross-sections per nucleus in the single animal were taken into account.

The statistical analysis. The statistical analysis was performed using CSS: Statistica v.5.0 (Statsoft, USA). The analysis of variance (Anova) was performed followed by post hoc tests (Duncan's post hoc analysis) and planned comparison tests (Student's paired t-test) between the consecutive pairs of means. The level of the statistical significance was set at $p < 0.05$. To evaluate the precision of all measurements the coefficient of error was calculated. Values below 0.1 were achieved for all the parameters examined.

RESULTS

The nuclear pattern of the common shrew's CA

The position of the various amygdaloid nuclei in the common shrew's CA and the pattern of their delimitations are shown in Figure 1 and 2, respectively.

CA of the common shrew can be portioned into 10 nuclei, classified according to the similarities in the cellular structure into 3 distinct groups: the basolateral (BLC), central (CC) and corticomедial (CMC). BLC is composed of the lateral (LA), basolateral (BL) and basomedial (BM) nuclei. The anterior amygdaloid area (AAA) as well as the central (CE) and intercalated nuclei (I) were included in CC. CMC are composed of the cortical (CO) and medial nuclei (ME) as well as the nucleus of the olfactory tract (NLOT) and the amygdalo-hippocampal area (AHA).

The position, general morphology and cellular structure of these different regions in the shrew's CA as well as their parcellation into the finer parts are very similar to those described by Krettek and Price in the rat [25]. Briefly, for the more subtle differences in density and cell size, LA, BL and BM are

further subdivided into the anterior (LAa, BLa and BMa) and posterior (LAp, BLp and BMp) parts. Two subdivisions can also be distinguished within the CE of the shrew, although the boundaries between them are less sharp than within the basolateral nuclei. The medial part (CEm) is located more rostrally and is composed of the larger, darker and more tightly packed cells. The lateral part (CEl) contains the smaller, lighter and more loosely arranged neurons. Finally, CO consists of 3 distinct regions: the anterior cortical nucleus (COa), the periamygdaloid cortex (COc) and the posterior cortical nucleus (COp). The ME, NLOT, AHA and AAA are homogenous regions.

Although the borders of the various nuclei in the shrew's CA were generally easy to recognise, some CA regions were poorly delineated from the surrounding nuclei. As a result of general delimitation problems, such structures as AAA, AHA and I were excluded from the volumetric investigation in the present study. The minor difficulties which were encountered in delineating the boundary between BM and CO as well as the rostral borders of ME and CE were overcome by taking into consideration the borders in the adjacent sections. Since all the sections are serially aligned, the additional support in delineating the obscure portions in the shrew's CA provided thick digital sections (200-300 μm) composed from single 50 μm digital scraps.

The morphometric parameters of the CA of the common shrew

The morphometric data concerning individual CA nuclei are listed in the subsequent rows of Table 1. The values for CA as a whole are presented in the last row.

Volume. The characteristic features of the common shrew's CA are poor development of LA and BM, whereas BL and NLOT are particularly strongly formed. CE, CO and ME also appear well developed in this species. The smallest average volume in the common shrew's CA is that of BM, while the largest is that of BL. According to the raw volumetric measurements and percentages, almost all the nuclei studied are significantly different ($p < 0.05$), although BL does not differ statistically from CO ($p > 0.05$).

Numerical density. The density of neurons in the common shrew's CA is smallest in BL, while that in ME is the largest. Comparison of the densities of neurons in the individual amygdaloid nuclei with the average for the total CA indicates that there are 3 different regions within the common shrew's

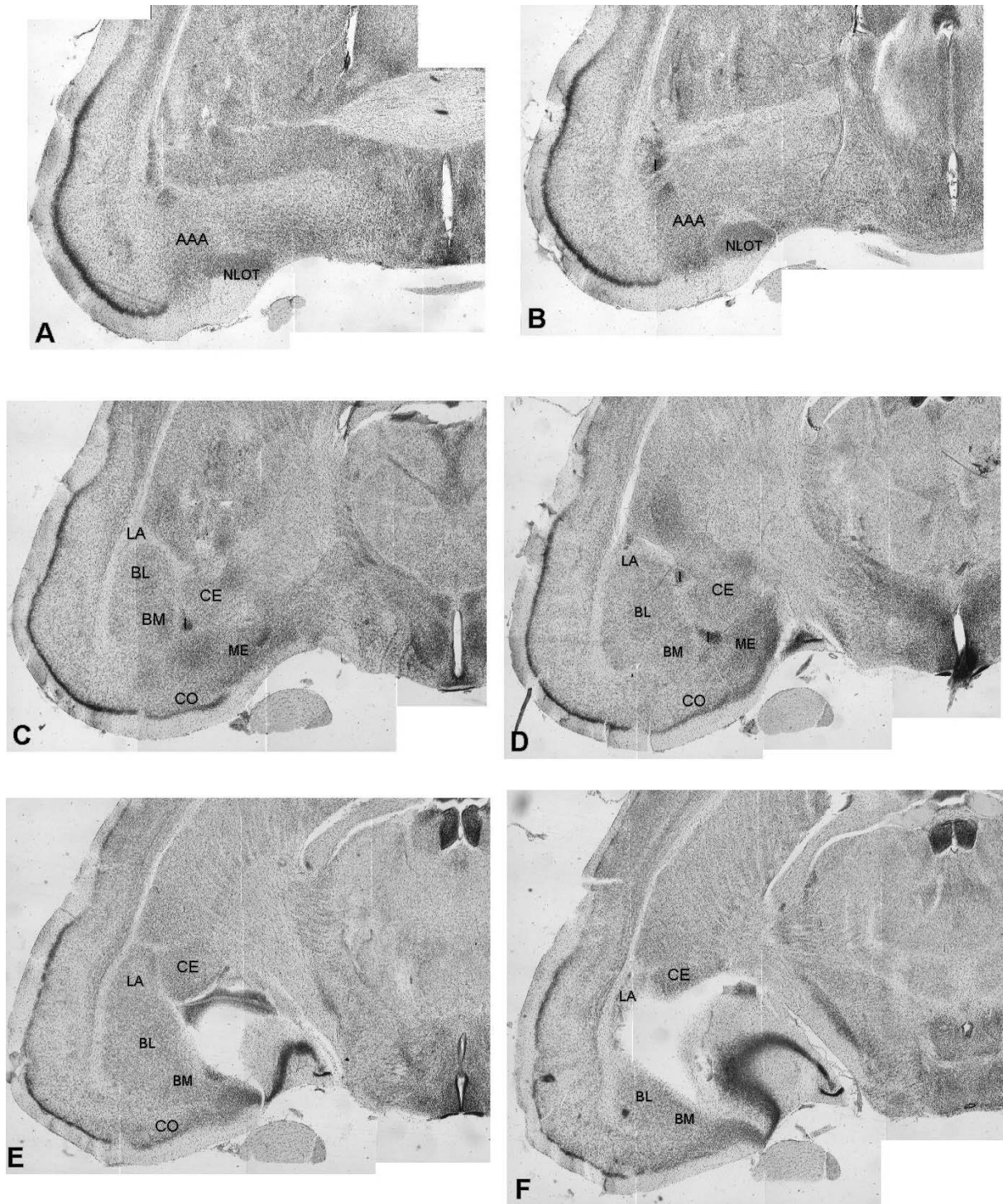


Figure 1. The position of the various amygdaloid nuclei in the common shrew's CA. **A-F** the subsequent coronal sections through the common shrew's CA in the rostro-caudal direction. Scale bar: 1 mm

CA. The densities of neurons in LA, BL and BM are significantly lower than the mean density of cells in CA ($p < 0.05$). In CE it does not differ from the mean ($p > 0.05$). In the CO, ME and NLOT these

values are significantly higher than the mean ($p < 0.05$). It should be noted, however, that the density of neurons in LA does not differ significantly from that in BL and BM ($p > 0.05$). The density of

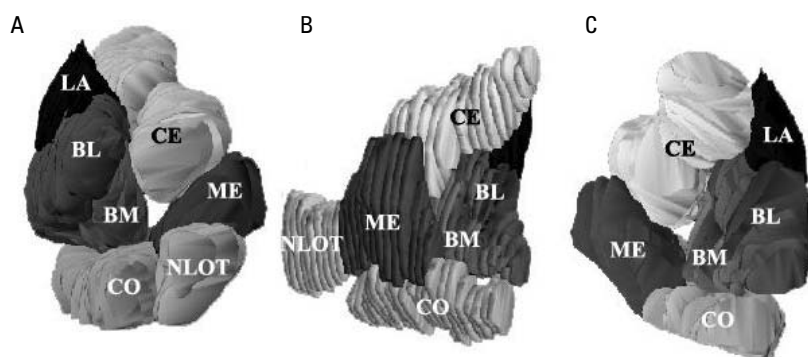


Figure 2. 3-D reconstruction of the common shrew CA: **A, B, C** — rostral, postero-medial and caudal views respectively

Table 1. The morphometric parameters of the individual amygdaloid nuclei of the common shrew CA.

Nucleus	Volume (mm ³)	Percentage of volume (%)	Numerical density (N/mm ³)	Total number of neurons (N)	Percentage of neurons (%)
LA	0.07 ± 0.01	10.78 ± 0.5	195799 ± 8888.56	14052.6 ± 2476.68	8.68 ± 0.74
BL	0.14 ± 0.02	20.64 ± 0.92	182643.3 ± 7412.85	25054.97 ± 4048.09	1531 ± 1.43
BM	0.05 ± 0.01	7.25 ± 0.66	195140.1 ± 9548.45	9412.14 ± 1710.36	5.82 ± 0.68
CE	0.12 ± 0.02	17.97 ± 0.79	253820.4 ± 141 80.9	30249.45 ± 4336.92	18.72 ± 1.12
CO	0.13 ± 0.01	19.62 ± 1.06	262600.3 ± 121 81.3	34073.61 ± 4034.12	21.18 ± 1.75
ME	0.1 ± 0.01	14.53 ± 0.29	332239.8 ± 13178.05	31889.83 ± 2904.33	19.82 ± 0.6
NLOT	0.06 ± 0.01	9.22 ± 0.2	271280.3 ± 13426.71	16515.4 ± 1431.85	10.27 ± 0.48
CA	0.66 ± 0.08	100 ± 0	241931.9 ± 5210.85	161248 ± 18111.13	100 ± 0

cells in the CO is also statistically similar to that in CE and NLOT ($p > 0.05$).

The total number of neurons. The smallest number of neurons in the common shrew's CA is in BM, whereas the largest cellular population is in CO. It should be added that according to the raw measurements and percentages almost all the nuclei studied are significantly different ($p < 0.05$), although ME does not differ statistically from CE and CO ($p > 0.05$).

The morphometric parameters of neurons in the common shrew's CA

The morphometric parameters of neurons in individual CA nuclei are listed in the subsequent rows of the Table 2. The mean values for CA are presented in the last row.

The length of neurons. The smallest average length of the soma in the common shrew's CA is displayed by the neurons located in the I. The cells in BL are the longest. The average lengths of neurons in LA, BL and BM are significantly higher than the length

of the mean CA neuron ($p < 0.05$). These values in CE ($p > 0.17$) and CO ($p > 0.73$) do not differ statistically from the mean. In the remaining nuclei the average lengths of neurons are significantly lower than the mean ($p < 0.05$). According to this parameter almost all the nuclei studied are significantly different ($p < 0.05$), although CE does not differ statistically from CO ($p > 0.05$).

The width of neurons. The smallest average width of the soma in the common shrew's CA is displayed by the cells in the I, whereas the largest perikarya, according to this parameter, are present in BL. The average width of neurons in LA, BL, BM and CO is significantly higher than the width of the mean CA neuron ($p < 0.05$). In the remaining nuclei these values are significantly lower than the mean ($p < 0.05$). According to this parameter, all the nuclei studied are significantly different ($p < 0.05$).

The average size of neurons. The size of the neuron in the common shrew's CA can be expressed by the sum of the length and width of the soma. The smallest size values in the common shrew's CA are

Table 2. The morphometric parameters of the neurons in the individual amygdaloid nuclei of the common shrew CA

Nucleus	Length of neurons [μm]	Width of neurons [μm]	Size of neurons [μm]	Shape of neurons
LA	13,3 \pm 0.78	8.82 \pm 0.55	22.19 \pm 1.65	1.51 \pm 0.09
BL	15.44 \pm 0.44	9.1 \pm 0.48	24.54 \pm 0.92	1.70 \pm 0.05
BM	13.78 \pm 0.48	7.98 \pm 0.69	21.76 \pm 1.13	1.73 \pm 0.09
CE	11.45 \pm 0.51	6.24 \pm 0.28	17.78 \pm 0.94	1.84 \pm 0.09
CO	1134 \pm 0.53	7.18 \pm 0.46	18.52 \pm 0.99	1.58 \pm 0.02
I	6.24 \pm 0.55	4.55 \pm 0.39	10.79 \pm 0.78	1.37 \pm 0.09
ME	8.71 \pm 0.32	5.71 \pm 0.21	14.42 \pm 0.34	1.53 \pm 0.09
NLOT	10.53 \pm 0.79	6.18 \pm 0.53	16.71 \pm 0.64	1.70 \pm 0.09
CA	11.35 \pm 2.93	6.97 \pm 1.59	18.34 \pm 4.48	1.62 \pm 0.15

found in the cells in I. The neurons in BL are the largest. The average sizes of neurons in LA, BL and BM are significantly higher than the size of the mean CA neuron ($p < 0.05$). This value in CO does not differ statistically from the mean ($p > 0.06$). In the remaining nuclei the average sizes of neurons are significantly lower than that of the mean CA neuron ($p < 0.05$). According to this parameter almost all the nuclei studied are significantly different ($p < 0.05$). LA alone does not differ statistically from BM ($p > 0.05$).

The shape of neurons. The shape of the neuron in the common shrew's CA can be expressed by the ratio of the length and width of the soma (the shape factor). The smallest average value of this coefficient was noted in the I, whereas the largest mean was observed in CE. The shape factor values in BL, BM, CE and NLOT are significantly higher than the mean value in the shrew's CA ($p < 0.05$). In the remaining nuclei these values are significantly lower than the mean ($p < 0.05$). According to this parameter almost all the nuclei studied show significant differences ($p < 0.05$). BM, however, does

not differ statistically from BL and NLOT ($p > 0.05$), nor does LA differ statistically from ME ($p > 0.05$).

The following represents the common shrew's CA nuclei series according to the increasing values of the morphometric parameters (Table 3).

DISCUSSION

Comparison of the present results with those obtained in previous studies. This is the first investigation that provides a detailed morphometric analysis of the 8 nuclei in the common shrew's CA. The only previous quantitative studies of the shrew's CA were proposed by Stephan et al. [65–67] in a series of comparative reports. However, these reports provided only selective volumetric measurements on the total size of CA and some of its larger components. Of the finer parts of CA only NLOT and the large-celled portion of BL were analysed. To our knowledge, no morphometric data concerning the number, density and size of the neurons in the common shrew's CA have been available to date.

Table 3. The arrangement of the common shrew's CA nuclei according to the increasing values of the morphometric parameters

Volume of CA nucleus		BM	NLOT	LA	ME	CE	CO	BL
Numerical density of neurons		BL	BM	LA	CE	CO	NLOT	ME
Total number of neurons		BM	LA	NLOT	BL	CE	ME	CO
Length of neurons	I	ME	NLOT	CO	CE	LA	BM	BL
Width of neurons	I	ME	NLOT	CE	CO	BM	LA	BL
Size of neurons	I	ME	NLOT	CE	CO	BM	LA	BL
Shape of neurons	I	LA	ME	CO	NLOT	BL	BM	CE
		Min.	—————>					Max.

Stephan et al. [67] put forward volumetric comparisons of the two larger but clearly delineated components of CA. The first, known as the lateral amygdala (LAM), combines LA, BL, BM and CO. The second, defined as the medial amygdala (MAM), is composed of CE and ME as well as AAA and NLOT. LAM percentage in relation to the size of the total CA in the common shrew was reported by Stephan et al. [67] to be 51.4%. LAM ratio calculated in the same species as our results yields a value of 58.7%. The figure of Stephan et al. [67] seems to be relatively low in comparison to our finding. However, LAM percentages in the work of Stephan et al. [67] were taken from CA in which AAA was included. Since this structure is comparatively large, it can be expected to influence the total volume of CA in the species. After exclusion of AAA from CA the percentage of LAM group presented by Stephan et al. [67] should be considerably larger and thus in good agreement with our results. Direct comparison of MAM group with our results is not possible since AAA was not taken into account in the present investigations.

The recent studies of Dziewiątkowski et al. [11] in the rabbit have shown that the individual nuclei in CA may be precisely labelled using the morphometric parameters of their neurons as a marker. The largest neurons in the rabbit CA were found in BL, while the smallest cells were localised in ME. The neurons populating CE and CO occupy an intermediate position and are very similar to each other in terms of size parameters. The neuronal size distribution depicted by Dziewiątkowski et al. [11] in the rabbit corresponds very well with our results in the shrew (Table 2). The only difference is that the neurons in the rabbit are considerably larger than those in the shrew. The positive correlation between the size of the cells and the size of the brain was described by Morgane et al. [51] for the limbic cortices of several Cetacean species and was also confirmed for CA by Równiak in a comparative study of the mammalian amygdala [61].

The model of the common shrew's CA organisation. The fundamental nuclear pattern and relationships between the various components in the shrew's CA conform to those found in the other species.

1. All the nuclei observed in the shrew have also been described in the mole [20], rat [25], gerbil [63], guinea pig [19], rabbit [18], cat [25, 54], dog [47], porpoise [5], bat [16], tree shrew [12], monkey [55, 69] and in humans [4]. Furthermore, the position and general morphology of these different CA regions

in the shrew are most similar to those found by Salter in the Mongolian gerbil [63]. However, the position of both parts of BM in the common shrew's CA when compared with the gerbil's BM seems to be reversed. The parcellation of the shrew's CA into finer parts was easily performed with the use of the terminology proposed by Krettek and Price in the rat [25].

2. Comparison of the density values in the common shrew's CA indicates that it can be subdivided into 3 compartments. LA, BL and BM form the amygdaloid region, characterised by a low packing density of neurons. The density values noted in all these nuclei are significantly lower than the mean density of cells in the shrew's CA. In contrast, CO, ME and NLOT form densely organised amygdaloid areas where the density values are significantly higher than the mean. According to this parameter, CE occupies an intermediate position with a numerical density value similar to the mean for the total of the shrew CA. Furthermore, a similar division of the shrew CA can, to some extent, be performed using size parameters of the amygdaloid neurons as a marker. Interestingly, the large neurons populate less densely organised CA areas, whereas the small cells occupy regions where the neurons are densely arranged. The inverse inter-relationship between the size of the neurons and the packing density is obvious when respective values in BL and ME are compared. The impression is even stronger when BL and I are analysed. Interestingly, the morphometric parcellation of CA presented in our studies corresponds fairly well with cytoarchitectonic studies performed with the Golgi methods in various mammalian species [4, 7, 22, 40, 48]. The basic cellular populations in LA, BL and BM form the pyramidal and semi-pyramidal neurons with large or medium-sized somata and spine-laden dendrites [4, 7, 22, 40, 48]. These clearly resemble neurons in the claustrum [68] and cerebral cortex [4, 7, 68]. In CE and I small and medium-sized cells prevail, which share the similar morphological features with the striatal neurons [22, 37, 50, 60, 68]. In contrast to the remaining amygdaloid nuclei, CO, ME and NLOT resemble the piriform cortex in the cellular structure [25, 68]. The basic cellular type in these nuclei determines small or medium size pyramidal cells [22, 39, 49]. It is noteworthy that the tripartite morphometric division of CA described above seems to be similar to those proposed by Swanson et al. [68], Crosby et al. [9] and Pitkänen et al. [56]. It differs from Johnston's bipartite suggestion [21] only in that Johnston's primitive

corticomedial group was subdivided into 2 separate regions (the corticomедial and central groups).

Although the general organisation of CA of the common shrew is similar to that found in other mammals, there are some peculiar features in the size and degree of differentiation of the various nuclei (Fig. 1, 2). The following are some of the peculiarities we have picked out:

1. The characteristic feature of the common shrew's CA is poor development of LA and BM. Similar conclusions were reached by Crosby et al. [10] and Stephan et al. [65, 67], although neither of these authors performed quantitative corroborations. Interestingly, hodological and cytoarchitectonic studies have shown that LA and BM are closely related anatomically [17, 44, 53, 61]. The projections of the ventral prefrontal cortices to the basolateral region of CA mainly targeted LA and BM, whereas the input to the BL, which lies between LA and BM, was very weak [17, 44, 53]. Furthermore, LA and BM, but not BL, have strong interconnections with each other and the superficial nuclei of CA [25, 31, 53, 56]. Finally, in contrast to BL, both nuclei receive afferents from the posterior thalamic region that processes auditory and somatosensory stimuli [27] as well as from the perirhinal cortex that probably processes visual and auditory signals [42]. It should be noted that the Golgi studies in various mammals suggest that LA and BM contain similar cell types [4, 7, 22]. Furthermore, the present morphometric results demonstrate that the sizes and densities of neurons in both these regions are also statistically comparable.
2. According to our data, CO, ME and, especially, NLOT are, in contrast to LA and BM, well developed in the shrew's CA (Fig. 1, 2, Table 1). Interestingly, all these superficial nuclei, in contrast to the rest of CA, receive direct input from the olfactory bulbs [23, 24, 68] and most of them also send information back to the bulbs [23, 24, 68]. COp and ME are the targets of the projection from the accessory olfactory bulb [23, 68], whereas the main olfactory bulb projects to COa and COc [24, 68]. Rostrally the main olfactory projection also terminates in NLOT [56, 68]. In addition, all these nuclei have 3 more or less distinct layers that are very characteristic for the piriform lobe areas (present results and [11, 18, 25, 68]). Moreover, the cells found in these layers clearly resemble neurons in the respective layers of the piriform cortex (present results and [22, 25, 47, 69]).

3. Since BL is, according to many authors [1, 4, 5, 7, 11, 18, 22, 29, 52, 56, 68], an integral part of BLC, one would expect it to be poorly developed in the shrew's CA. However, this seems not to be the case as BL is especially well represented in this species (Fig. 1, 2, Table 1). Surprisingly, the strong development of the BL in the macrosmatic animals has also been reported by several authors. Salter [63] found that this nucleus is particularly well formed in the gerbil's CA. Similar impressions regarding the tree shrew were reported by Flügge et al. [12]. Interestingly, LA in both these species was, as in the shrew, considerably smaller than the BL both in the coronal and sagittal sections [12, 63]. Stephan et al. [65, 67] using volumetric methods, pointed out that the magnocellular part of BL occupies much a larger area in CA of insectivora than in primates (Prosimians and Simians). It should be noted that BL shows some common features with NLOT [35]. Both these nuclei receive a relatively dense dopaminergic innervation from the lateral ventral tegmental area and the medial part of the *substantia nigra* [30] and are strongly interconnected with each other [56]. In addition, both of them are innervated by the cholinergic cells in the basal forebrain [8, 15].

Functional remarks. Since according to Stephan et al. [67] "the comparative neuro-anatomical investigations have demonstrated that the size of a given brain structure is highly inter-related with the functional requirements of its habits" the question arises as to what the functional consequences are of the type of CA organisation found in the common shrew?

LA is a major recipient for the sensory-related thalamic and cortical inputs directed to CA and LA projects back to these areas as well [25, 44, 56, 68]. This appears to be a major site of convergence of all non-olfactory sensory modalities, namely the visual [29, 42], auditory [27–29, 36, 59] and somatosensory [26, 29, 70]. BM, which has also strong inter-relationships with the sensory-related thalamic and cortical areas [6, 26, 56], seems to be linked more to contextual information [29, 33, 34]. Since both these nuclei are poorly developed in the common shrew, it is reasonable to state that non-olfactory information processed by the cortex is poorly represented inside CA of this species. Consequently, CA of the common shrew is probably more influenced by olfaction. Interestingly, the superficial nuclei of CA, like that of CO, ME and NLOT that include the secondary olfactory and vomeronasal areas [23, 24, 68], are, in contrast to LA and BM, especially well represented in CA of the shrew.

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