**THE CAUDAL BURSA IN THE HELIGMONELLIDAE (NEMATODA: TRICHOSTRONGYLLINA). CHARACTERIZATION AND HYPOTHESIS ON ITS EVOLUTION**

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**Summary:**

The different patterns of the caudal bursa of the Heligmonellidae (Nematoda) are redefined, taking into account the grouping of rays 2-6 and the sequence of origin of these rays from their common trunk. The type of symmetry of the caudal bursa is also redefined. The following patterns were observed and characterized: the basic patterns: types 2-3, 2-2-1, 1-3-1 and 1-4 and the intermediary patterns: type 2-3 trending to type 2-2-1, type 2-2-1 trending to type 1-3-1, type 1-3-1 trending to type 1-4 and type 2-2-1 trending to type 1-4. An evolutionary interpretation of the patterns is attempted and seems to follow the direction: 2-3 to 2-2-1 to 1-3-1 to 1-4. Seven atypical patterns are described. The caudal bursae were classified based on their symmetry: subsymmetrical, dissymmetrical and asymmetrical. Independently of the type of symmetry, the two latero-ventral lobes may have the same or different patterns. The type of symmetry, the ratio between the two latero-ventral lobes and a characteristic pattern were utilized to characterize the caudal bursae at the level of the genus and the subfamily. The combination of the right/left ratio and the type of symmetry gives heterogeneous results, with no real association between these characters. The most conspicuous asymmetries and dissymmetries were found among the Nippostrongylinae. The most frequent pattern in the Heligmonellidae is the basic type 2-2-1; types 1-3-1 and 1-4 are less frequent but are characteristic of several genera; type 1-4 is absent from the Heligmonellidae. Whatever the pattern, in the Heligmonellidae rays 4 and 5 are the last to diverge from the common trunk of rays 2-6.

**KEY WORDS:** Nematoda, Heligmonellidae, bursal pattern, bursal symmetry, evolution.

**INTRODUCTION**

The Trichostrongylina have a common origin with the other suborders of the order Strongylida, arising from an ancestor close to the Rhabditida (Durette-Desset et al., 1994; Blaxter et al., 1998, 2001). In the strongylid nematodes, unlike the Rhabditida, the tail of the male widens in order to form a caudal bursa made up of two latero-ventral lobes and one dorsal lobe. Therefore the caudal bursa is a derived character regarded as a synapomorphy for the Strongylida. Durette-Desset & Chabaud (1981) and Durette-Desset (1985), proposed a classification of the Trichostrongyloidea which they divided into three supra familiar groups: the “Trichostrongylids”, the “Molineids” and the “Heligmosomids”. In these classifications, they highlighted various types of caudal bursae, based mainly on the grouping of rays 2 to 6 (i.e. the rays supporting the latero-ventral lobes) (Fig. 1). In 1981, four patterns were defined: 1-3-1, characteristic of the “Trichostrongylids”, 2-1-2, characteristic of the “Molineids” and 2-2-1 and 3-2, characteristic of the “Heligmosomids”. Durette-Desset (1985) added type...
2-3 and specified that each grouping is characteristic of a given evolutionary line or a given family. Then, Durette-Desset & Chabaud (1993) raised the Trichostrongyloidea to a suborder: the Trichostrongyloidea and each supra familiar group became a superfamily: the Trichostrongyloidea, the Molineoidea and the Heligmosomoidea.

This descriptive system however seems to us to be insufficient to properly describe the caudal bursae of some Nippostrongylinae, especially the genera such as Heligmonina Baylis, 1928 and Stilestrongylus Freitas, Lent & Almeida, 1937, which do not show the same pattern in each lateral lobe. Within the framework of a revision of the Heligmonellidae, the aim of this work is to redefine the different patterns of the caudal bursa found within this family, taking into account not only the grouping of rays 2-6 but also the sequence of origin of these rays from their common trunk. This enables us not only to make the descriptions of the caudal bursae more accurate, but also to highlight the characteristic type(s) of pattern within each genus and to attempt an evolutionary interpretation of the patterns.

MATERIALS AND METHODS

The elements usually considered in the description of a caudal bursa (CB) are: symmetry, pattern of latero-ventral lobes, development of the dorsal lobe and characters of the dorsal ray and rays 8. The characters of the dorsal ray and rays 8 are not treated in this work. Some terms concerning the symmetry and the pattern of the latero-ventral lobes are redefined as follows.

TYPES OF SYMMETRY

There are three types, which concern the degree of development of the latero-ventral lobes in relation to the sagittal axis of the worm:
- Subsymmetrical bursae (CB SS): both lobes are of similar size and shape in relation to the axis of the dorsal ray, which passes through the sagittal axis of the worm (Fig. 2a).
- Dissymmetrical bursae (CB DS): One lobe is larger than the other in relation to the axis of the dorsal ray, which passes through the sagittal axis of the worm. When the right lobe (RL) is larger it is cited as CB DS RL+. When the left lobe (LL) is larger it is cited as CB DS LL+ (Fig. 2b).
- Asymmetrical bursae (CB AS): The axis of the dorsal ray does not pass through the sagittal axis but is displaced to the right or the left side of the worm. One lobe may or may not be larger than the other in relation to this axis. When the right lobe (RL) is larger it is cited as CB AS RL+. When the left lobe (LL) is larger it is cited as CB AS LL+ (Fig. 2c).

TYPES OF PATTERN

The main features considered in the definition of the patterns are the grouping of the rays and the sequence of origin (or divergence) of these rays from their common trunk. The level of divergence of a ray from a common trunk (rays 2 to 6, 3 to 5 or 3 to 6) is calculated on a straight line represented by ray 4; at the level of divergence another line is traced perpendicularly to the first one, as shown in Fig. 2d.
- Basic patterns: rays 2 to 6 show a characteristic arrangement of the following types: type 2-3 (Fig. 3a); type 2-2-1 (Fig. 3b); type 1-3-1 (Fig. 3c); type 1-4 (Fig. 3d).
- Intermediary patterns (or transitional types from a basic type to another): rays 2 to 6 show an intermediary arrangement with features of two basic types: type 2-3 tending to type 2-2-1 (2-3 t 2-2-1) (Fig. 3e); type 2-2-1 tending to type 1-3-1 (2-2-1 t 1-3-1) (Fig. 3f); type 1-3-1 tending to type 1-4 (1-3-1 t 1-4) (Fig. 3g); type 2-2-1 tending to type 1-4 (2-2-1 t 1-4) (Fig. 3h).

LITERATURE DATA

The data were compiled from descriptions published in the literature. We took into account only the caudal bursae where both latero-ventral lobes, or at least one, was spread out. In the latter case, the lobe was treated separately and the pattern was not extrapolated to the entire bursa.
The systematic position of some species remains uncertain since, even if the caudal bursa is completely described, this is not the case with the synlophe. These species have therefore not been considered in this work. They are as follows: *Duretstrongylus travassosi* (Lent & Freitas, 1938), *Heligmonella vladimiri* Sadovskaja, 1952, *Heligmonina vogeli* Khalil, 1931, *Heligmonoides crassidorsualis* Franco, 1967, *Heligmonoides mirzai* Smales, 2008, *Heligmosomum delta* Travassos, 1921, *Longistiata degusi* Babero & Cattan, 1975, *Longistiata castrosilvai* Almeida, 1934, *Longistiata fortuita* Freitas, Lent & Almeida, 1937, *Longistiata perfida* Travassos, 1943, *Gobindonema boodugi* Sood & Parshad 1974, *Moganiella cricetuli* Yin & Zhang, 1981. They are here considered as Heligmonellidae incertae sedis since their generic allocation is not possible. In the particular case of *H. crassidorsualis* and *H. mirzai*, both were considered as Nippostrongylinae incertae sedis by Durette-Desset & Digiani (2010). Another species not treated is *Nesomystrongylus fissicauda* Durette-Desset, Lehtonen & Haukisalmi, 2002. This species shares more characters with the Heligmosomidae than with the Heligmonellidae, *i.e.* presence of caudal spine in the female tail, caudal bursa with rays 2 and 3 well developed, small dorsal lobe, and axis of orientation of synlophe subfrontal. It is likely that the monospecific genus *Nesomystrongylus* Durette-Desset, Lehtonen & Haukisalmi, 2002 should be transferred to the Heligmosomidae.

Figs. 2a-c. – Types of symmetry of the caudal bursa.

a. Subsymmetrical: both lobes are of similar size and shape in relation to the axis of the dorsal ray, which passes through the sagittal axis of the worm. Ex: *Pudica gamma* (Travassos, 1918). After Travassos (1921), modified.

b. Dissymmetrical: one lobe is better developed in relation to the axis of the dorsal ray, which passes through the sagittal axis of the worm. Ex: *Malvinema scapteromys* (Suriano & Navone, 1996). In this case, it involves the right lobe. After Digiani et al. (2003), modified.

c. Asymmetrical: the axis of the dorsal ray does not pass through the sagittal axis but is displaced to the right or left side of the worm. Both lobes may or may not have the same degree of development with respect to the axis of the dorsal ray. Ex: *Malvinema victoriae* Digiani, Sutton & Durette-Desset (2003). In this case, the right lobe is slightly better developed. After Digiani et al. (2003), modified.

Fig. 2d. – Determination of the point of divergence of rays 2 to 6 from their common trunk.

The levels of divergence of rays 2 to 6 from the common trunk are indicated by the dotted lines, which are perpendicular to a main axis represented by a straight line passing through ray 4. Ex: *Malvinema carolinae* Digiani, Sutton & Durette-Desset (2003). In the right lobe, ray 2 is the first (the most proximal ray) to diverge from the common trunk; the point of divergence of ray 3 is distal to that of ray 6. Rays 4 and 5 are the last to diverge. After Digiani et al. (2003), modified.

Scale-bars Figs. 2a-d: 50 μm.
RESULTS

DESCRIPTION OF THE PATTERNS

• Basic and intermediary types

  a. Type 2-3 (Fig. 3a)
  - Rays 2 and 3 grouped, arising first and together from common trunk of rays 2 to 6.
  - Rays 4 to 6 having common trunk.
  - Divergence of rays 4, 5 and 6 at same level.

  b. Type 2-2-2 (Fig. 3b)
  - Rays 4 to 6 having common trunk.
  - Rays 2 and 3 grouped, arising first and together from common trunk of rays 2 to 6.
  - Ray 6 arising at the same level as rays 3.

  c. Type 1-3-1 (Fig. 3c)
  - Rays 2 and 6 arising first and at same level from common trunk of rays 2 to 6.
  - Rays 3 to 5 having short common trunk.
  - Distal divergence of rays 4 and 5.

  d. Type 2-2-1 (Fig. 3d)
  - Rays 2 and 3 grouped, arising first and together from common trunk of rays 2 to 6.
  - Ray 6 arising at the same level as rays 3.

  e. Type 2-3 (Fig. 3e)
  - Rays 2 and 3 grouped, arising first and together from common trunk of rays 2 to 6; rays 4 to 6 have a common trunk and diverge at the same level. Ex: Paraheligmonella interrogaass (Lent & Freitas, 1938). After Durette-Desset (1968b), modified.

  f. Type 2-2-1 (Fig. 3f)
  - Rays 4 to 6 having common trunk.
  - Rays 2 and 3 grouped, arising first and together from the common trunk of rays 2 to 6; ray 6 arises at the same level as ray 3; rays 4 and 5 are the last to diverge. Ex: Heligmostrongylus crucifer (Travassos, 1943). After Travassos (1943), modified.

  g. Type 1-3-1 (Fig. 3g)
  - Rays 2 and 6 arising first and at same level from common trunk of rays 2 to 6; rays 3 to 5 have a common trunk.
  - Ray 3 arising just distally to level of divergence of rays 2 and 6.
  - Distal divergence of rays 4 and 5.

  h. Type 1-4 (Fig. 3h)
  - Rays 2 arising first from common trunk of rays 2 to 6.
  - Rays 3 to 6 having long common trunk.

Figures 3a–d. – Basic patterns. Scale-bars: 50 µm.

Figs. 3e–h. – Intermediary patterns. Scale-bars: 50 µm.

a. Type 2-3. Rays 2 and 3 are grouped, arising first and together from the common trunk of rays 2 to 6; rays 4 to 6 have a common trunk and diverge at the same level. Ex: Paraheligmonella interrogaass (Lent & Freitas, 1938). After Durette-Desset (1968b), modified.

b. Type 2-2-2. Rays 2 and 3 are grouped, arising first and together from the common trunk of rays 2 to 6; ray 6 arises at the same level as ray 3; rays 4 and 5 are the last to diverge. Ex: Heligmostrongylus crucifer (Travassos, 1943). After Travassos (1943), modified.

c. Type 1-3-1. Rays 2 and 6 arise first and at the same level from the common trunk of rays 2 to 6; rays 3 to 5 have a common trunk. Ex: Hypocristata tercera Durette-Desset & Guerrero, 2006. In this case ray 3 diverges first from the common trunk of rays 3 to 5 in the right lobe and at the same level as ray 5 in the left lobe. After Durette-Desset & Guerrero (2006), modified.

d. Type 1-4. Ray 2 arises first from the common trunk of rays 2 to 6; rays 3 to 6 have a long common trunk. Ex: Fuellebornema bocqueti (Durette-Desset, 1970a). In this case ray 3 diverges at the same level as ray 6 on the common trunk of rays 3 to 6. After Durette-Desset (1970a), modified.

e. Type 2-3. Rays 2 and 3 are grouped, arising first and together from the common trunk of rays 2 to 6; rays 4 to 6 have a common trunk; ray 6 diverges close to the level of the divergence of ray 3 and proximally to that of rays 4 and 5. Ex: Nesbeligononella acomynti Durette-Desset & Gibson, 1990. In this case rays 2 and 3 are apposed for much of their length. After Durette-Desset & Gibson (1990), modified.

f. Type 2-2-1 t 1-3-1. Rays 2 and 6 arise first and at the same level from the common trunk of rays 2 to 6; rays 3 arise just distally to the level of divergence of rays 2 and 6; rays 4 and 5 are the last to diverge. Ex: Pudica pudica (Travassos, 1921). After Cassone & Durette-Desset (1991), modified.

g. Type 1-3-1 t 1-4. Rays 2 and 6 arise first from the common trunk of rays 2 to 6; rays 3-6 have a short common trunk; rays 6 arise at the same level as rays 3; rays 4 and 5 are the last to diverge. Ex: Hypocristata anguillula (Durette-Desset, 1971). Ray 3 is still grouped with ray 2 since its extremity supports the ventral lobe. After Durette-Desset (1971), modified.

- Divergence of ray 6 proximal to divergence of rays 4 and 5 and approximately at same level of divergence of ray 3.

• Atypical patterns

Seven atypical patterns were found among the Heligmonellidae:
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. Type 1-1-2-1 (Fig. 4e, LL)
- Ray 2 arising first from common trunk of rays 2 to 6.
- Ray 3 completely separated and parallel to ray 2, arising at the same level as ray 6.
- Distal divergence of rays 4 and 5.

. Type 1-3-1 t 4-1 (Fig. 4f, RL)
- Ray 6 arising first from common trunk of rays 2 to 6.
- Ray 2 arising just distally to level of divergence of ray 6.
- Rays 2 to 5 having very short common trunk.

. Type 4-1 (Fig. 4f, LL)
- Ray 6 arising first from common trunk of rays 2 to 6.

EVOLUTIONARY INTERPRETATION OF THE PATTERNS

Durette-Desset (1985) highlighted two evolutionary trends treating the Strongylida as a set: (1) reduction of the dorsal lobe (in the Rhabditida the tail of the male is vertically elongated) and (2) lengthening of rays 4, as papillae 4 of the Rhabditida are close to the anus (see Osche 1958; Chabaud et al., 1970 for the homology between the papillae of the Rhabditida and the bursal rays of the Strongylida) (Figs. 5a, 5b). Therefore, Durette-Desset (1985) suggested that a caudal bursa with a short dorsal lobe and long rays 4 should be considered as highly evolved. Though these trends occur in each type of bursa, in general, types 2-1-2 and 2-3 (among the five types of pattern recognized) were considered as basal types for two reasons: they are the closest to the pattern of the Rhabditida with a well developed dorsal lobe and short rays 4; and they are also present in all four suborders of the Strongylida. However, no further attempt was made to explain the other patterns from an evolutionary point of view. We propose the following interpretation (Fig. 5c).

In the Heligmonellidae the most basal pattern found is of type 2-3 (Fig. 3a). In this type, the dorsal lobe is relatively long as is ray 6, which diverges distally from the common trunk of rays 4 to 6. The transition from type 2-3 to type 2-2-1 involves the migration of ray 6 towards the base of the trunk of rays 2 to 6, as seen in the intermediary type 2-3 t 2-2-1 (Fig. 3e). In type 2-2-1 (Fig. 3b), ray 6 has achieved this migration and diverges from the common trunk at the same level as rays 3.
The transition from type 2-2-1 to type 1-3-1 involves a progressive migration of ray 3 towards ray 4 along with a progressive shortening of the dorsal lobe and ray 6, as observed in the intermediary type 2-2-1 t 1-3-1 (Fig. 3f). In type 1-3-1 (Fig. 3c), the migration of ray 3 towards rays 4 is achieved, with the appearance of a common trunk of rays 3-5, and ray 6 diverges from this common trunk at same level as ray 2.

The transition from type 1-3-1 to type 1-4 involves a distal migration of ray 6, with the appearance of a common trunk of rays 3 to 6, and a progressive shortening of this trunk, as observed in the intermediary type 1-3-1 t 1-4 (Fig. 3g). In type 1-4 (Fig. 3d), only ray 2 arises first from the common trunk. This type of caudal bursa widens laterally and its width is greater than its length.

The evolutionary steps then seem to follow the direction: 2-3 to 2-2-1 to 1-3-1 to 1-4, which is consistent with the early interpretation by Durette-Desset (1985) with respect to the basal patterns and the caudal bursae in which the width is greater than the length.

Several descriptions of caudal bursae, especially in species of *Heligmonina* where the pattern was described as being of type 1-4 tending to type 1-3-1 (see for example Diouf et al., 2005) are therefore inaccurate from an evolutionary point of view.

Some lobes may show a pattern of intermediary type 2-2-1 t 1-4 (Fig. 3h). This transition from type 2-2-1 involves the shortening from the base of the set of rays 3-6 just after the divergence of ray 2, with the appearance of a short common trunk of rays 3 to 6 (intermediary type 2-2-1 t 1-4).

A particular 2-3 type was observed in some bursae or isolated lobes of the genera *Srivastavanema* (Singh, 1962) (Brevistriatinae) and *Maleinema* Digiani, Sutton & Durette-Desset, 2003 (Nippostrongylinae). In these cases one or both lateral lobes show an arrangement.

Figs. 5a-h. – Homology of the bursal rays of the Rhabditida and the Strongylida. Scale-bars: 50 μm.

a. Bursal rays of the Rhabditida sensu Osche (1958). Papillae 7 are fused with papillae 8. After Osche (1958), modified.

b. Bursal rays of the Strongylida sensu Chabaud et al. (1970). Papillae 7 are situated on the genital cone. After Chabaud et al. (1970), modified.

Fig. 5c. – Schematic representation of the bursal patterns of rays 2 to 6 showing the hypothetical evolutionary steps.

From the basal pattern of type 2-3, the evolutionary steps follow the direction: 2-3 tending to 2-2-1 t 1-3-1 to 1-4 with the intermediary patterns: 2-3 tending to 2-2-1, 2-2-1 tending to 1-3-1, 1-3-1 tending to 1-4. Some lobes have a pattern of type 2-2-1 t 1-4.

A. Type 2-3. B. Type 2-3 t 2-2-1. C. Type 2-2-1. D. Type 2-2-1 t 1-3-1. E. Type 1-3-1. F. Type 1-3-1 t 1-4. G. Type 1-4. The small arrows indicate the transition from a type to another. The dotted arrow indicates the hypothetic evolutionary progress of the pattern of the caudal bursae.
of rays 2 to 6 corresponding to a type 2-3, however the dorsal lobe is greatly reduced, the caudal bursa is laterally elongated and they are typically complementary with type 1-4. This means that in two species of Maltevinea, M. caroliniae Digiani, Sutton & Durette-Desset, 2003 and M. victoriae Digiani, Sutton & Durette-Desset, 2003, whereas one lobe is of type 2-3, the other lobe shows a pattern of type 1-4 (Figs. 2c, 2d); the remaining two species, M. scapteromysis (Suriano & Navone, 1996) and M. yagoi Digiani & Durette-Desset, 2003, show the type 1-4; in two species of Stivastavanema, S. cynocephali Durette-Desset & Purwaningsih, 1999 and S. yapi Durette-Desset & Lim Boo Liat, 1975, both lobes are of type 2-3 whereas a third species S. bhaguanstingbi Durette-Desset & Lim Boo Liat, 1975 shows a type 1-4 in both lobes. We thus consider this type 2-3 as different from that of the most basal species. It may be derived from the type 1-4 by the migration of ray 3 towards ray 2, whereas rays 4 to 6 still have a long common trunk.

- Atypical patterns

Type 1-1-2-1 (Fig. 4e) is present in the left lobe of Cordicauda cordicauda (Durette-Desset, 1966) (Bre-vistriatinae). This type seems to have been derived from type 2-2-1 by the isolation of ray 2 and short migration of ray 3 on the common trunk of rays 2 to 6.

Type 1-3-1 t 4-1 (Fig. 4f, RL) is present in the right lobes of Nippostrongylus magnus (Mawson, 1961) and N. marbaeniae Hasegawa & Syafuddin, 1995 (Nippostrongylinae). The transition from type 1-3-1 to type 4-1 involves a distal migration of ray 2 towards ray 3, forming a short common trunk of rays 2 to 5. Ray 6 is the most proximal ray to diverge from the common trunk of rays 2-6 and ray 2 arises just slightly distally to ray 6.

Type 4-1 (Fig. 4f, LL) is present in both lobes of Nippostrongylus sembeli Hasegawa & Tarore, 1995; and in the left lobes of N. magnus, N. rauschi Chabaud & Desset, 1966 and N. typicus (Mawson, 1961). This type seems directly derived from type 3-1-1, involving a marked distal migration of ray 2 towards ray 3. Ray 6 is the most proximal ray to diverge from the common trunk of rays 2-6, and rays 2 to 5 form a long common trunk on which ray 2 arises distinctly distally.

Type 3-1-1 (Fig. 4g, LL) is present in the left lobes of four species of Nippostrongylus: N. djumachani (Tenora, 1969), N. marbaeniae, N. rysswyi (Erhardova, 1959) and N. utterbergi Greenberg, 1972. It seems directly derived from type 4-1 by the migration of ray 5 from its base towards ray 6. Ray 5 diverges from the common trunk at the same level as ray 6 and there is persistence of the common trunk of rays 2 to 4.

Type 1-2-1-1 (Fig. 4h, LL) is present in the left lobes of Sciuricola dremomys (Yen, 1973) and S. moreli (Gibbons, Durette-Desset & Daynes, 1977) (Heligmonellae). It seems directly derived from type 1-3-1 by the migration (but not apposition) of ray 5 towards ray 6, with persistence of the group formed by rays 3 and 4.

Type 1-2-2 (Fig. 4i, RL) is present in the right lobe of Tricholinstowia maseri Durette-Desset & Vaucher, 1974 (Heligmonellae). It seems to be derived from type 1-3-1 (present in the other species of Tricholinstowia) by the migration of ray 5 towards ray 6. Type 2-1-2 (Fig. 4i, LL) is present in the left lobe of the same species. It could be derived from type 1-2-2 by the migration of ray 3 towards ray 2, along with the absence of common trunk between rays 3 and 4.

A pattern of type 2-1-2 is characteristic of the Molineoidae. However, type 2-1-2 of the Molineoidae is interpreted as a basal pattern, also characterized by short rays 4, whereas the pattern found in T. maseri, with long rays 4, is considered as derived from a 1-3-1 type. In this case, we consider the presence of this type in one lobe of Heligmonellidae as a convergence.

CRITERIA UTILIZED FOR THE CHARACTERIZATION OF THE CAUDAL BURSAE OF THE HELIGMONELLIDAE

- Type of symmetry (definition and examples above)

At the specific level, a caudal bursa may be subsymmetrical (CB SS); dissymmetrical with right lobe larger (CB DS RL+); dissymmetrical with left lobe larger (CB DS LL+); asymmetrical (CB AS); asymmetrical with right lobe larger (CB AS RL+); asymmetrical with left lobe larger (CB AS LL+). At the generic level, we consider that a given genus may be characterized by the type of symmetry most frequently found among the species belonging to this genus.

- Evolutionary comparison between lobes

As mentioned above, a species can have the same or a different pattern in both latero-ventral lobes. On the other hand, as seen above, the different patterns may be interpreted from an evolutionary point of view. This determines three types of “evolutionary ratio” between both lateral lobes (“right/left ratio”) in a caudal bursa: (1) the same pattern in both lobes; (2) the pattern of the right lobe is derived with respect to that of the left lobe (“RL derived”); or (3) the pattern of the left lobe is derived with respect to that of the right lobe (“LL derived”).

- Characteristic pattern

Within a given genus the bursal pattern usually varies among the species. Moreover, as seen immediately above, most species have a different pattern in each lobe. This means that in some genera we can find several different patterns. However it seems possible to choose one or two characteristic patterns for each
genus. The characteristic pattern is determined by the type most frequently found, which is here interpreted as the most frequent basic type plus the contiguous intermediary types. For example, for the genus *Sciurodenendrium* Durette-Desset, 1971 (Pudicinae, seven species) we propose that the characteristic pattern is of type 2-2-1 because amongst the 11 treatable lobes, four are of type 2-2-1, three of type 2-3 t 2-2-1, and two of type 2-2-1 t 1-3-1. Some genera may have two characteristic patterns. For example in *Heligmonina* (Nippostrongylinae, 27 species) two characteristic patterns are proposed: 1-3-1 and 1-4; type 1-3-1 being the most frequent in the right lobe and type 1-4 in the left lobe (21 and 19 lobes respectively).

**Characterization of the Caudal Bursae of the Genera of the Heligmonellidae**

- *Heligmonellidae* (Skrjabin & Schikhobalova, 1952, tribe) Durette-Desset, 1971 (four subfamilies, 56 genera, 329 species; for each genus, in parentheses, number of species examined / number of species described)
  - *Heligmonellinae* (Skrjabin & Schikhobalova, 1952, tribe) Durette-Desset, 1971 (four subfamilies, 56 genera, 329 species; for each genus, in parentheses, number of species examined / number of species described)
  - *Heligmonella* Mönnig, 1927 (six genera, 22 species)
    - *Heligmonella* Mönnig, 1927 (3/4)
      - CB SS (2 spp.) or CB DS RL+ (1 sp.). CB with the same pattern in both lobes (2 spp.) or with LL derived (1 sp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-4. Characteristic pattern: 2-2-1.
    - *Parabeligmonella* Durette-Desset, 1971 (4/5)
      - CB SS. CB with same pattern in both lobes (3 spp.) or with RL derived (1 sp.). Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1, atypical 2-2-1 t 4-1. Characteristic pattern: 2-2-1.
  - *Parabeligmonella* Durette-Desset, 1971 (4/5)
    - CB SS. CB with same pattern in both lobes (3 spp.) or with RL derived (1 sp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-4. Characteristic pattern: 2-2-1.
    - *Parabeligmonella* Durette-Desset, 1971 (4/5)
      - CB SS. CB with different patterns in both lobes (1 sp.). Patterns observed: 2-3 t 2-2-1, atypical 1-2-1-1. Characteristic pattern: the data are insufficient to choose a characteristic pattern but two characters of the CB shared by both species are rays 4 and 5 separated from base and dorsal lobe distinct.
  - *Tricholinstowia* Travassos, 1937 (8/8)
    - CB SS (6 spp.) or CB DS LL+ (1 sp.), type of symmetry unknown in 1 sp. CB with the same pattern in both lobes (4 spp.) or with RL derived (1 sp.) or with LL derived (1 sp.), right/left ratio unknown in 2 spp. Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1, atypical 1-2-2 and 2-1-2. Characteristic pattern: 1-3-1; rays 5 and 6 pincer-shaped.
  - *Tricholinstowia* Travassos, 1937 (8/8)

Remark: only the right lobe is treatable in *T. mogera* (Sadovskaja, 1952) and *T. morenisi* (Cameron & Parnell, 1933).

- *Trichoposthia* C. P. Schmidt, Myers & Kuntz, 1967 (7/7)
  - CB SS (5 spp.) or CB DS RL+ (2 spp.). CB with the same pattern in both lobes (5 spp.) or with LL derived (2 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-4. Characteristic pattern: 2-2-1.
  - *Cordicauda* Durette-Desset, 1971 (5/5)
    - CB DS LL+ (3 spp.) or CB SS (2 spp.). CB with the same pattern in both lobes (3 spp.) or LL derived (2 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-4. Characteristic pattern: 2-2-1.
  - *Cordicauda* Durette-Desset, 1971 (5/5)
    - CB SS (5 spp.) or CB DS RL+ (2 spp.). CB with the same pattern in both lobes (5 spp.) or with LL derived (2 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-4. Characteristic pattern: 2-2-1.
  - *Cordicauda* Durette-Desset & Krishnansamy, 1976 (6/6)
    - CB SS (3 spp.) or CB DS LL+ (2 spp.) or CB DS RL+ (1 sp.). CB with the same pattern in both lobes (3 spp.) or with RL derived (3 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1. Characteristic pattern: 2-2-1.
  - *Kuala* Durette-Desset & Krishnansamy, 1976 (2/2)
    - CB SS. CB with the same pattern in both lobes. Pattern observed: atypical type 1-2-2.
  - *Lagostrongylus* Fukumoto, Kamiya & Ohbayashi, 1986 (3/3)
    - CB SS. CB with the same pattern in both lobes. Patterns observed: 2-3 t 2-2-1. Characteristic pattern: 2-3 t 2-2-1.
  - *Lagostrongylus* O. Y. Ow-Yang, Durette-Desset & Ohbayashi, 1983 (2/2)
    - CB SS. CB with the same pattern in both lobes. Patterns observed: 2-3 t 2-2-1. Characteristic pattern: 2-2-1.
. _Metheligmonella_ Durette-Desset, 1971 (2/2)
CB SS. CB with the same pattern in both lobes. Patterns observed: 2-3 t 2-2-1. Characteristic pattern: 2-3 t 2-2-1.

. _Parabeligmonina_ (Ortlepp, 1939) (27/28)
CB SS (12 spp.) or with a slight left (8 spp.) or right (7 spp.) dissymmetry. CB with the same pattern in both lobes (18 spp.) or with LL derived (7 spp.) or with RL derived (2 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-3-1, 1-4, atypical 2-2-1 t 4-1 and 4-1. Characteristic patterns: 2-2-1, 1-3-1.

Remark: the caudal bursa of _P. trifurcata_ (Baylis, 1928) is not included as the published illustration is not of a fully opened bursa.

. _Quentinstrongylus_ Durette-Desset, 1969 (1/1)
CB SS. CB with the same pattern in both lobes. Patterns observed: 1-3-1. Characteristic pattern: 1-3-1.

. _Srivastavanema_ (Singh, 1962) (3/5)
CB SS. CB with the same pattern in both lobes. Patterns observed: secondary 2-3, and 1-4. Characteristic pattern: 1-4.

- _Pudicinae_ (Skrjabin & Schikhobalova, 1952, tribe) Durette-Desset, 1971 (nine genera, 44 species)

. _Acanthostrongylus_ Travassos, 1937 (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 1-3-1. Characteristic pattern: 1-3-1.

. _Durettestrongylus_ Guerrero, 1982 (2/3)
CB SS. CB with the same pattern in both lobes. Patterns observed: 2-2-1. Characteristic pattern: 2-2-1.

. _Freitastrongylus_ Gonçalves, Pinto & Durette-Desset (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 1-4. Characteristic pattern: 1-4.

. _Methellobornema_ Travassos & Darriba, 1929 (6/7)
CB SS. CB with the same pattern in both lobes. Pattern observed: 1-4. Characteristic pattern: 1-4.

Remark: the caudal bursa of _F. almeidai_ Travassos, 1937 is not included as the published illustration is not of a fully opened bursa.

. _Heligmostrongylus_ Travassos, 1917 (9/9)
CB SS. CB with the same pattern in both lobes (8 spp.) or with LL derived (1 sp.). Patterns observed: 2-2-1, 2-2-1 t 1-3-1, 1-3-1. Characteristic pattern: 2-2-1.

. _Justinema_ R’Kha & Durette-Desset, 1991 (3/3)
CB SS. CB with RL derived (2 spp.) or with the same pattern in both lobes (1 sp.). Patterns observed: 2-2-1, 2-2-1 t 1-3-1. Characteristic pattern: 2-2-1.

. _Pseudobeligmosomum_ Travassos, 1937 (1/1)
CB AS LL+. CB with the same pattern in both lobes. Pattern observed: 2-3. Characteristic pattern: 2-3.

. _Pudica_ Travassos & Darriba, 1929 (12/12)
CB SS. CB with the same pattern in both lobes (11 spp.) or with RL derived (1 sp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1, 1-4. Characteristic patterns: 2-2-1, 1-3-1.

. _Sciuromordrium_ Durette-Desset, 1971 (6/7)
CB SS. CB with LL derived (3 spp.) or with the same pattern in both lobes (2 spp.), right/left ratio unknown in 1 sp. Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1. Characteristic pattern: 2-2-1.

Remarks: the caudal bursa of _S. arripense_ (Baylis, 1947) is not included as the published illustration is not of a fully opened bursa. Only the right lobe is treatable in _S. bravobollisae_ Falcón-Ordaz & Lamothe-Argumedo, 2006.

- _Nippostrongylinae_ Durette-Desset, 1971 (29 genera, 197 species)

. _Bunomystrongylus_ Hasegawa & Mangali, 1996 (2/2)
CB DS RL+ (1 sp.) and CB DS LL+ (1 sp.). CB with RL derived. Patterns observed: 2-3 t 2-2-1, 2-2-1, 1-3-1, 1-4. No characteristic pattern.

. _Carolinensis_ (Travassos, 1937) (12/12)
CB SS (7 spp.) or CB DS LL+ (4 spp.) or CB DS LL+ (1 sp.). CB with the same pattern in both lobes (9 spp.) or with RL derived (1 sp.) or with LL derived (1 sp.), right/left ratio unknown in one species. Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1, 1-3-1 t 1-4, 1-4. Characteristic pattern: 2-2-1.

Remark: only the right lobe is treatable in _C. eophonomys_ Asakawa, Kamiya & Ohbayashi, 1986 and _C. huehuetlana_ Falcón-Ordaz & Sanabria Espinosa, 1996, the left lobe being not spread out.

. _Entzetoda_ Elias & Durette-Desset, 2003 (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 2-3 t 2-2-1. Characteristic pattern: 2-3 t 2-2-1.

. _Guerrerostrongylus_ Sutton & Durette-Desset, 1991 (2/2)
CB SS. CB with the same pattern in both lobes. Patterns observed: 1-3-1. Characteristic pattern: 1-3-1.

. _Hasamuddinia_ Hasegawa & Syafruddin, 1994 (1/1)
CB SS. CB with RL derived. Patterns observed: 2-2-1 t 1-3-1, 2-2-1 t 1-4. Characteristic pattern: 2-2-1.

. _Hassalstrongylus_ Durette-Desset, 1971 (13/14)
CB SS (6 spp.) or DS RL+ (6 spp.) or DS LL+ (1 sp.). CB with the same pattern in both lobes (9 spp.) or RL derived (2 spp.) or LL derived (2 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1, 1-4. Characteristic patterns: 1-4 (type species), 2-2-1, 1-3-1. Remark: the caudal bursa of _H. mazzai_ Freitas, Lent & Almeida, 1937 is not included as the published illustration is not of a fully opened bursa.

. _Heligmonina_ Baylis, 1928 (24/26)
CB DS LL+. CB with RL derived (21 spp.) or with the same pattern in both lobes (3 spp.). Patterns observed: 1-3-1, 1-3-1 t 1-4, 1-4. Characteristic patterns: 1-3-1 on right lobe, 1-4 on left lobe.
Remarks: the caudal bursa of *H. oenomyos* Baylis, 1928 is not illustrated and therefore untreatable. The left lobe of *H. cricetomyos* Baylis, 1928 is typical of the genus *Heligmonina* i.e. of type 1-4, but the right lobe is atypical with a type 2-2-1 not retained in the patterns observed.

. *Heligmonoides* Baylis, 1928 (12/12)
CB DS LL+. CB with the same pattern in both lobes (6 spp.) or with RL derived (3 spp.) or LL derived (3 spp.). Patterns observed: 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 1-3-1, 1-4. Characteristic pattern: 2-2-1.

. *Hypocristata* Durette-Desset, 1971 (3/3)
CB SS (2 spp.) or DS RL+ (1 sp.). CB with the same pattern in both lobes (1 sp.) or with RL derived (1 sp.) or with LL derived (1 sp.). Patterns observed: 2-2-1, 2-2-1 t 1-4, 2-2-1 t 1-3-1, 1-3-1. Characteristic pattern: 1-3-1.

. *Malaistrongylus* Ow-Yang, Durette-Desset & Ohbayashi, 1983 (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 2-2-1. Characteristic pattern: 2-2-1.

. *Malvinema* Digiani, Sutton & Durette-Desset, 2003 (4/4)
CB DS RL+ (3 spp.) or CB AS RL+ (1 sp.). CB with the same pattern in both lobes (2 spp.) or with RL derived (1 sp.) or with LL derived (1 sp.). Patterns observed: 1-4, secondary 2-3. Characteristic pattern: 1-4.

. *Mammanidula* Sadovskaja, 1952 (3/5)
CB AS RL+ (2 spp.) or CB SS (1 sp.). CB with LL derived (2 spp.) or RL derived (1 sp.). Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-2-1.
Remark: the caudal bursae of *M. melomyos* (Mawson, 1961) and *M. siamensis* (Ohbayashi & Vajrasthiva, 1983) are not included as the published illustrations are not of fully opened bursae.

. *Mausonema* Smales & Heinrich, 2010 (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 2-3. Characteristic pattern: 2-3.

. *Maxomystrongylus* Hasegawa & Syafruddin, 1997 (2/2)
CB DS LL+. CB with the same pattern in both lobes (1 sp.) or with RL derived (1 sp.). Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-2-1.

. *Melomystrongylus* Smales, 2009 (2/2)
CB DS LL+ or CB DS RL+. CB with RL derived (1 sp.), right/left ratio unknown in one species. Patterns observed: 2-3, 1-3-1. Characteristic pattern: 2-3. Remark: only the left lobe of *M. sepikensis* Smales, 2009 is illustrated and treatable.

. *Montistrongylus* Smales & Heinrich, 2010 (1/1)
CB DS RL+. CB with RL derived. Patterns observed: 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-2-1.

. *Neobeligmonella* Durette-Desset, 1971 (21/23)
CB SS. CB with the same pattern in both lobes (for at least 12 spp.) or with RL derived (for at least 7 spp.), right/left ratio unknown in 2 spp. Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-3 t 2-2-1. Remarks: the caudal bursa of *N. affinis* (Baylis, 1928) is not treatable since it is not illustrated in the original description. Only the right lobe is treatable in *N. impudica* (Baylis, 1928) and *N. moennigi* (Baylis, 1928). *N. lenniscomyosi* (Durette-Desset, 1970) is the only species having a caudal bursa with a right lobe larger and with the same pattern (1-3-1) in both lobes. On the other hand, its synlophe is also very different from that of the remaining species in the genus. It is likely that this species belongs to a different genus and it is not treated herein.

. *Nippostrongylus* Lane, 1923 (9/9)
CB AS RL+ (6 spp.) or CB DS RL+ (3 spp.). CB with LL derived. Patterns observed: 1-3-1, 1-3-1 t 1-4, 1-3-1 t 4-1, 4-1, 3-1-1. Characteristic patterns: 1-3-1 and derived atypical 4-1 on right lobe; atypical 4-1 and derived atypical 3-1-1 on left lobe.

. *Odilia* Durette-Desset, 1973 (18/19)
CB SS (8 spp.) or DS RL+ (3 spp.) or DS LL+ (2 spp.), symmetry unknown in 5 spp. CB with same pattern in both lobes (at least 8 spp.) or with RL derived (at least 3 spp.), right/left ratio unknown in 5 spp. Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1, 2-2-1 t 1-3-1, 2-2-1 t 1-4, 1-3-1, 1-3-1 t 1-4, 1-4. Characteristic patterns: 2-2-1, 1-4. Remarks: the caudal bursa of *O. polyrhabdote* (Mawson, 1961) is not included as the published illustration is not of a fully opened bursa. Only one lobe is treatable in *O. brachybursa* (Mawson, 1961), *O. implexa* (Smales, 2008), *O. moeatensis* (Hasegawa, Miyata & Syafruddin, 1999), *O. similis* (Smales, 2009) and *O. uromyos* (Mawson, 1961).

. *Orientostrostrongylus* Durette-Desset, 1970 (7/8)
CB SS. CB with the same pattern in both lobes (6 spp.) or with LL derived (1 sp.). Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-2-1. Remark: the caudal bursa of *O. siamensis* (Ohbayashi & Kamiya, 1980) is not included as the published illustration is not of a fully opened bursa.

. *Parabeligmonelloides* Fukumoto, Kamiya & Suzuki, 1980 (9/9)
CB DS LL+ (4 spp.) or CB DS RL+ (3 spp) or CB SS (2 spp.). CB with the same pattern in both lobes. Patterns observed: 2-3, 2-3 t 2-2-1, 2-2-1, 1-4. Characteristic patterns: 2-2-1, 1-4. Remark: the caudal bursa of *P. singauwaensis* Smales, 2009 is treated from its redescription (Smales & Heinrich, 2010).

. *Parasabanema* Smales & Heinrich, 2010 (1/1)
CB SS. CB with the same pattern in both lobes. Pattern observed: 2-2-1. Characteristic pattern: 2-2-1.
. *Rattustrongylus* Ow-Yang, Durette-Desset & Ohbayashi, 1983 (2/2)

CB SS (1 sp.) or CB DS LL+ (1 sp.). CB with the same pattern in both lobes. Patterns observed: 2-2-1, 2-2-1 and 1-3-1. Characteristic pattern: 2-2-1.

. *Sabanema* Ow-Yang, Durette-Desset & Ohbayashi, 1985 (4/5)

CB DS RL+ (3 spp.) or CB SS (1 sp.). CB with the same pattern in both lobes (2 spp.) or with RL derived (1 sp.) or with LL derived (1 sp.). Patterns observed: 2-3 t 2-2-1, 2-2-1. Characteristic pattern: 2-2-1.

Remark: the caudal bursa of *S. kepongi* Ow Yang, Durette-Desset & Ohbayashi, 1983 is not included as the published illustration is not of a fully opened bursa.

. *Spalacina* Biserkov, Durette-Desset & Genov, 1995 (3/3)

CB SS (2 spp.) or DS LL+ (1 sp.). CB with the same pattern in both lobes (at least 2 spp.), right/left ratio unknown in one species. Patterns observed: 1-3-1 t 1-4, 1-4. Characteristic pattern: 1-4.

Remark: only the right lobe is treatable in *S. spalacis* (Sharpilo, 1973) the left lobe being not spread out.

. *Stilestrongylus* Freitas, Lent & Almeida, 1937 (24/25)

CB DS RL+ (21 spp.) or CB SS (3 spp.). CB with RL derived (1 sp.), right/left ratio unknown in one species. Patterns observed: 2-3 t 2-2-1. Characteristic pattern: 2-2-1.

Table I. – Synopsis of the caudal bursa in the Heligmonellinae.

| Genus            | Characteristic symmetry | Right/left ratio | Characteristic pattern |
|------------------|-------------------------|-----------------|------------------------|
| *Heligmonella*   | CB SS                   | Same pattern    | 2-2-1                  |
| *Parabelgmonella*| CB SS                   | Same pattern    | 2-2-1                  |
| *Sciuricola*     | CB SS                   | LL atypical     | -                      |
| *Tricholinostia* | CB SS                   | Variable        | 1-3-1                  |
| *Trichotavassosia* | CB SS                | Same pattern    | 2-2-1                  |
| *Xericola*       | CB SS                   | Same pattern    | 2-2-1                  |

CB SS: caudal bursa subsymmetrical. LL: left lobe of caudal bursa.

. *Acanthostrongylus* Ow-Yang, Durette-Desset & Ohbayashi, 1986 (15/15)

CB SS (1 sp.). CB with the same pattern in both lobes, a right/left ratio which is variable but with predominantly the same pattern in both lobes, and several types of pattern with a predominance of the characteristic type 2-2-1 (Table II); the Pudicinae by a subsymmetrical caudal bursa, a right/left ratio which is variable but with predominantly the same pattern in both lobes, and several types of patterns with a predominance of characteristic types 2-2-1 and 1-4 (Table III); the Nippostrongylinae by a caudal bursa of variable symmetry, a right/left ratio which

Based on the elements considered above, the four subfamilies may be synthetically characterized as follows: the Heligmonellinae by a subsymmetrical caudal bursa, with the same pattern in both lobes, and a characteristic pattern of type 2-2-1 (Table I); the Brevistriatinae by a predominantly subsymmetrical caudal bursa, a right/left ratio which is variable but with predominantly the same pattern in both lobes, and several types of pattern with a predominance of the characteristic type 2-2-1 (Table II); the Pudicinae by a subsymmetrical caudal bursa, a right/left ratio which is variable but with predominantly the same pattern in both lobes, and several types of patterns with a predominance of characteristic types 2-2-1 and 1-4 (Table III); the Nippostrongylinae by a caudal bursa of variable symmetry, a right/left ratio which

Table II. – Synopsis of the caudal bursa in the Brevistriatinae.

| Genus            | Characteristic symmetry | Right/left ratio | Characteristic pattern |
|------------------|-------------------------|-----------------|------------------------|
| *Brevistriata*   | CB SS                   | Same pattern    | 2-2-1                  |
| *Calypostrongylus* | CB SS                | Same pattern    | 2-2-1                  |
| *Cordicuda*      | Variable                | Variable        | 2-2-1                  |
| *Fissicuda*      | Variable                | Variable        | 2-2-1                  |
| *Kuala*          | CB SS                   | Same pattern    | 1-2-2                  |
| *Lagostrongylus* | CB SS                   | Same pattern    | 2-3 t 2-2-1            |
| *Macrostrongylus* | CB SS                   | Same pattern    | 2-2-1                  |
| *Metheligmonella* | CB SS                   | Same pattern    | 2-3 t 2-2-1            |
| *Parabelgmonina* | Variable                | Variable        | 2-2-1, 1-3-1           |
| *Quentinstrongylus* | CB SS                | Same pattern    | 1-3-1                  |
| *Srivastavanema* | CB SS                   | Same pattern    | 1-4                    |

CB SS: caudal bursa subsymmetrical.

CB SS: caudal bursa subsymmetrical. CB DS LL+: caudal bursa disymmetrical with right left lobe larger.

Table III. – Synopsis of the caudal bursa in the Pudicinae.
is also variable and several types of pattern with a predominance of characteristic types 2-2-1 and 1-4 (Table IV).

**DISCUSSION**

Of the five main patterns recognized among the Trichostrongylina treated by Durette-Desset (1985), only three were observed in the Heligmonellidae: 2-3, 2-2-1 and 1-3-1. In this article, the main feature considered was the grouping of the rays and, in the interpretation of its evolution, only two main tendencies were considered: the shortening of the dorsal ray and the lengthening of rays 4.

| Genus             | Characteristic symmetry | Right/left ratio | Characteristic pattern |
|-------------------|-------------------------|------------------|------------------------|
| Bunomysstrongylus | Variable                | RL derived       | -                      |
| Carolinea         | Variable                | Same pattern     | 2-2-1                  |
| Enzotheria        | CB SS                   | Same pattern     | 2-3 t 2-2-1            |
| Guerrerostrongylus| CB SS                   | Same pattern     | 1-3-1                  |
| Hasamuldiina      | CB SS                   | RL derived       | 2-2-1                  |
| Hasalastrongylus  | Variable                | Same pattern 1-4 | 2-2-1, 1-3-1           |
| Heligmonina       | CB DS LL+               | LL derived       | 1-3-1 RL, 1-4 LL       |
| Heligmonoides     | CB DS LL+               | Variable         | 2-2-1                  |
| Hypocrisata       | CB SS                   | Variable         | 1-3-1                  |
| Malaistrongylus   | CB SS                   | Same pattern     | 2-2-1                  |
| Malacinema        | CB DS RL+               | Variable         | 1-4                    |
| Mammamandula      | CB AS RL+               | Variable         | 2-2-1                  |
| Mauesonema        | CB SS                   | Same pattern     | 2-3                    |
| Malvastrongylus   | BC DS LL+               | Variable         | 2-2-1                  |
| Melomystrongylus  | variable                | Unknown          | 2-3                    |
| Montistromyglus   | CB DS RL+               | RL derived       | 2-2-1                  |
| Neobelmonella     | CB SS                   | Variable         | 2-3 t 2-2-1            |
| Nippostrongylus   | CB AS RL+               | LL derived       | 1-3-1, 4-1 RL          |
|                    | CB DS RL+               | 4-1, 3-1-1 LL    |
| Odilia            | Variable                | Variable         | 2-2-1, 1-4             |
| Orientostromyglus | CB SS                   | Same pattern     | 2-2-1                  |
| Parabellmonelloides| Variable               | Same pattern     | 2-2-1, 1-4             |
| Parasabanema      | CB SS                   | Same pattern     | 2-2-1                  |
| Rattistromyglus   | Variable                | Same pattern     | 2-2-1                  |
| Sahabemina        | CB DS RL+               | Variable         | 2-2-1                  |
| Spalacina         | CB SS                   | Same pattern     | 1-4                    |
| Stilestrongylus   | CB DS RL+               | Variable         | 2-2-1, 1-4             |
| Suttonemina       | CB DS RL+               | Same pattern     | 1-4                    |
| Trichofreitasia   | CB SS                   | Same pattern     | 2-2-1                  |
| Yatinema          | Variable                | RL derived       | 2-2-1                  |

CB SS: caudal bursa subsymmetrical. LL: left lobe of caudal bursa. RL: right lobe of caudal bursa. CB DS LL+: caudal bursa dissymmetrical with left lobe larger. CB DS RL+: caudal bursa dissymmetrical with right lobe larger. LL derived: left lobe of the caudal bursa having a pattern derived with respect to that of the right lobe. RL derived: right lobe of the caudal bursa having a pattern derived with respect to that of the left lobe.

Table IV. – Synopsis of the caudal bursa in the Nippostrongylinae.

In the present article, we provide an additional feature at the descriptive level, which is the sequence of the origin of the rays from the common trunk. Consequently, the presence of a new basic pattern (type 1-4) is highlighted, plus the presence of intermediary types: type 2-3 tending to 2-2-1, type 2-2-1 tending to 1-3-1, type 1-3-1 tending to 1-4 and type 2-2-1 tending to 1-4, which have enabled us to attempt an evolutionary interpretation of the patterns.

Type 2-3, the basal type, is somewhat infrequent, but not the following intermediary type, 2-3 t 2-2-1, which is common and present in all four subfamilies. The basic type, 2-2-1, seems to be the most consistent pattern in the family: it is the most frequent characteristic pattern, is present in all four subfamilies, and there are relatively few intermediary types from 2-2-1 to 1-3-1, to 1-4 or to atypical patterns. Types 1-3-1 and 1-4 are less frequent than type 2-2-1 but are characteristic of several genera; type 1-3-1 is observed in all four subfamilies, and is the origin of several atypical types (4-1, 3-1-1, 1-2-2 and 1-2-1-1); type 1-4, characteristic of laterally elongated lobes, is absent from the Heligmonellinae. Finally, the family Heligmonellidae is characterized by the fact that, regardless of the pattern, with very few exceptions, rays 4 and 5 are always the last to diverge from the common trunk of rays 2 to 6.

It is clear that a dissymmetrical or asymmetrical caudal bursa should be regarded as derived with respect to a subsymmetrical one. However, the type of symmetry has been actually considered a character of little value above the species level, since the asymmetrical bursae and mostly the dissymmetrical bursae have arisen several times during the course of evolution. The dissymmetry usually involves the transverse (lateral) lengthening of one of the lobes. This dissymmetry probably plays a role in holding the female during copulation (Durette-Desset, 1985). The most conspicuous asymmetries and dissymmetries are found among the Nippostrongylinae. In species with a slight dissymmetry the lengthening may or may not modify the bursal pattern, which usually remains the same in both lobes. In species with strong dissymmetry, species, which are usually tightly coiled spirally, the pattern is usually different for each lobe.

Independent of the type of symmetry, both lobes of the caudal bursa may have the same or different patterns. In the latter case the most frequent situation from an evolutionary point of view is one lobe having one pattern and the other lobe showing the “next” intermediary or basic type (e.g. Neobelmonella 2-3 t 2-2-1 and 2-2-1; Heligmonina 1-3-1 and 1-4); there are very few cases in which both lobes have a totally different pattern (e.g. Heligmonella asymetrica, some species of Stilestrongylus).
In some genera the same “right/left ratio” is observed in all or most species in the genus: either both lobes have the same pattern (Heligmostrongylus, Pudica), or it is always the same lobe, which is derived with respect to the other (Heligonina). Other genera are less homogeneous and the right/left ratio varies among the species of the genus (Neobelginonella, Parabelginonina).

The combination of the right/left ratio and type of symmetry gives disparate results: in genera with marked dissymmetry or asymmetry, the hypertrophied lobe is usually the same but it may show either the derived pattern (Heligmonina, LL) or the basal pattern (Nippostrongylus, RL). In genera with subsymmetrical or slightly dissymmetrical caudal bursae, not only may the right/left ratio vary but also the type of symmetry within each genus (Odilia, Parabelginonina).

It is interesting to note that the Heligmonellinae, the Brevistriatinae and the Pudicinae, in which most of the caudal bursae are subsymmetrical with the same pattern in both lobes, are parasitic in hosts, which are relatively ancient (mainly sciuriform and cavi- morph rodents, a few in insectivores and lagomorphs). The Nippostrongylinae, in which the symmetry and the pattern in both lobes are more variable, are parasitic in a group, which appeared more recently, the murid rodents (mainly cricetids and murids).

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