Chapter 1

General Introduction

The phylogenetic position of the Bryozoa is subject to ongoing discussions. Bryozoa or Ectoprocta are a group of sessile, colonial invertebrates which occur, with probably 8,000 recent species worldwide, in marine and limnetic habitats (introductory reviews in Ryland 1970, 2005, Mukai et al. 1997, Nielsen 2001). Since Hatschek (1891), the predominant opinion was that they form, together with Brachipoda and Phoronida, the Tentaculata or Lophophorata (Hyman 1959). These were assigned either to protostomes or to deuterostomes or were placed "intermediate". Despite varying views about the monophyly of lophophorates, most later approaches united them with Deuterostomia as Radialia (Jefferies 1986, Eernisse et al. 1992, Backeljau et al. 1993, Ax 1989, Brusca and Brusca 2003). This was based chiefly on the coelomic organization and the apparent homology of the tentacular apparatus to that of hemichordates and crinoid echinoderms. An alternative hypothesis, based mainly on larval morphology and similarities in metamorphosis, has been put forward by (Nielsen 1971, 1977, 2001, 2002), who agreed to the deuterostome affinity of Brachiopoda and Phoronida, but suggested a sister-group relationship between Bryozoa and Kamptozoa, thus a protostome affinity of bryozoans.

In a phylogenetic study using 18S rDNA data, Halanych et al. (1995) revealed evidence for a common ancestry of the lophophorate taxa, annelids and molluscs united in a clade named Lophotrochozoa. This new evidence challenged the long-held view of a deuterostome relationship of all or some lophophorate groups and many of the ideas that supported this. Subsequent studies of more extensive datasets and further molecular markers almost consistenly corroborated Lophotrochozoa (see Halanych 2004 for review). Since at the same time many cladistic analyses of broad morphological datasets still supported deuterostome affinities (Zrzavy et al. 1998, 2003, Sorensen et al. 2000, Peterson and Eernisse 2001, Nielsen 2001, Glenner et al. 2004), a serious conflict appeared. However, until presently molecular approaches either fail to resolve the exact position of Bryozoa (as well as that of several other lophotrochozoan taxa) or provide conflicting evidences. Bryozoa have, for example, been suggested to be basal Lophotrochozoans (Halanych et al. 1995, Passamaneck and Halanych 2006), basal prostomes (Giribet et al. 2000), polyphyletic (Mackey et al. 1996, Passamaneck and Halanych 2006), polyphyletic with phoronid affinities (Helmkampf et al. 2008), sister to Brachiopoda (Waeschenbach et al. 2006), sister to Kamptozoa (Hausdorf et al. 2007).

Concerning the molecular data, several sources of error have been suggested that could cause the current problems (Telford et al. 2005, e.g., Philippe and Telford 2006, Baurain et al. 2007). Although some authors (Scotland et al. 2003) question in principle the value of morphology for phylogenetic inference, most researchers agree that morphology represents a complex data set whose careful analysis can serve as an independent test for molecular based phylogenies (Wiens 2004, Jenner 2004). Morphology-based phylogenies depend heavily on the characters chosen and the accuracy of the used details (Jenner and Schram 1999). Morphological data sets suffer chiefly from two problems. One is that the number of phylogenetically informative characters is relatively low in comparison to DNA sequence data. Especially in "phylum-level" phylogenies of the metazoa only few characters are comparable across the different groups. Thus, a single character has potentially a relatively high impact on the result of the analysis. This poses the second problem: Many characters are only insufficiently investigated in certain groups, leading to a high degree of uncertainty, when coding character states in a data matrix. This also applies for Bryozoa: looking at the most recent morphological data matrices that have been used for analyzing metazoan phylogeny (Zrzavy et al. 1998, 2003, Sorensen et al. 2000, Peterson and Eernisse 2001, Nielsen 2001, Glenner et al. 2004), it turns out that a majority of those characters that can be suspected to contain essential information, are scored "unknown" for Bryozoa. This is especially the case for characters of embryogenesis, neural organization and mesodermal structures. In contrast to this situation, detailed comparable data from many other metazoa taxa have become available by a growing field of comparative developmental biology and neuroanatomy during the last years (see e.g., Nielsen 2004, 2005, Minelli and Fusco 2008, Lichtneckert and Reichert 2007). Hence, in this thesis, a selection of these characters have been investigated with the central aim to provide new phylogenetically significant data.

Commonly the three bryozoan subtaxa Phylactolaemata, Stenolaemata, and Gymnolaemata, which differ in terms of both morphology and life-style, are distinguished. Gymnolaemata comprise Cheilostomata and Ctenostomata. In the traditional phylogenetic hypotheses, the morphology of Phylactolaemata has been regarded as especially significant, because several similiarities to Phoronida had been recognized. Indeed, Phylactolaemata differ considerably from Stenolaemata and Gymnolaemata, hindering assumptions about ancestral character states for Bryozoa. Chapter 2 *Ganglion ultrastructure in phylactolaemate Bryozoa: Evidence for a neuroepithelium*, addresses the question whether the cerebral ganglion in Phylactolaemata is, in contrast to that of Gymnolaemata, hollow and formed by an invagination process comparable to the vertebrate neurulation. Chapter 3: *Ultrastructure of the body cavities in phylactolaemate Bryozoa* aims to answer the question, whether the epistome, an upper-lip like organ contains an independent secondary body cavity, as implied by earlier morphologists.

The evolution of larval types within Bryozoa is still elusive. Stenolaemata and Phylactolaemata exhibit larvae, which are hardly comparable to those of gymnolaemates as well as other metazoan groups (Reed 1991). In gymnolaemates, the planktotrophic cyphonautes, as well as several types of lecithotrophic larvae occur (Zimmer and Woollacott 1977). Chapter 4 *Muscular systems in gymnolaemate bryozoan larvae* focuses on the phylogenetic significance of larval musclature for the evolution of bryozoan larval types. During the last ten years, larval neural anatomy has regained much attention as complex character, of possibly high phylogenetic significance. Only few results on bryozoan larvae exist, which do not provide a consistent picture by now. Thus, in chapter 5 *Serotonergic and FMRFamidergic nervous systems in gymnolaemate bryozoan larvae*, data on nervous system characters of different larvae are collected and a putative ancestral pattern is proposed.

Early embryology in Bryozoa has been studied by several earlier workers, but the origin of mesoderm is still essentially unknown. As data on mesoderm formation have been acquired in many metazoan taxa, namely candidate sister groups like Phoronida and Brachiopoda (Freeman 2000, 2003, Lüter 2000, Freeman and Martindale 2002), knowledge on this character in Bryozoa has the potential to contribute arguments for the phylogenetic position of Bryozoa. In chapter 6 *Mesoderm origin in Membranipora membranacea*, developmental stages of a species exhibiting a cyphonautes larva are examined on the ultrastructural level.

Chapter 7 summarizes the most important results of this thesis and discusses their impact on the conceptions about phylogeny and position of the Bryozoa .

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