

PROTIST TOOLS

A Short Guide to Common Heterotrophic Flagellates of Freshwater Habitats Based on the Morphology of Living Organisms



Alexandra Jeuck, and Hartmut Arndt¹

Department of General Ecology, Zoological Institute, Cologne Biocenter, University of Cologne, Zùlpicher Straße 47b, D-50674 Cologne, Germany

Introduction

Heterotrophic flagellates (HF) are very likely the most abundant eukaryotes on Earth, hundreds of specimens occur in each droplet of water even in groundwater and the deep sea. As the main feeders on bacteria they play an essential role in aquatic and terrestrial food webs (Arndt et al. 2000; Azam et al. 1983; Bonkowski 2004). In addition, they can act as important herbivores (Arndt and Mathes 1991; Nauwerck 1963; Sherr and Sherr 1994), detritivores (Scherwass et al. 2005) and osmotrophs (Christoffersen et al. 1997; Sanders et al. 1989; Sherr 1988) as well as mixotrophs (Bird and Kalff 1986; Sanders 1991). The relative contributions to these different modes of feeding can vary within taxonomic groups and even within one and the same organism, e.g. *Ochromonas* sp. (Jones 2000; Wilken et al. 2013). Furthermore, bacterial communities are not only grazed by protozoans but are also structured by protistan grazers (e.g. Boehme et al. 2009; Güde 1979; Jürgens and Matz 2002; Pernthaler 2005).

HF are a very heterogenous group with an enormous size range between 1–450 μm (some authors refer to the species smaller than 15 μm as heterotrophic nanoflagellates “HNF”, Arndt et al. 2000). High tolerances to changes in salinity allow several species to live both in marine as well as in freshwater habitats, though several phylogenetic studies have also indicated clearly separated marine and freshwater clades (e.g.

Kim and Archibald 2013). In contrast, the dominant taxonomic groups within different marine and freshwater pelagic communities (e.g. stramenopile taxa, dinoflagellates, choanoflagellates, kathablepharids) and benthic communities (e.g. euglenids, free-living kinetoplastids, cercozoans) seem to be surprisingly similar (Arndt et al. 2000). Their modes of movement as important taxonomic characteristics comprise gliding or free-swimming forms or they may be temporarily or permanently attached to a substrate (Fenchel 1987). Feeding modes include true filter-feeding (e.g. choanoflagellates), direct interception feeding (e.g. chrysomonads) or raptorial feeding (e.g. most benthic forms) (Boenigk and Arndt 2002; Fenchel 1991).

One important prerequisite to estimate the role of HF for the flux of matter in ecosystems is the determination of their abundance. In the early days of HF quantification, the so called “HNF” were counted exclusively in fluorescently stained fixated samples (e.g. Porter and Feig 1980). However, fixation bears the problem that many species are disrupted upon the fixation process (Choi and Stoecker 1989; Sonntag et al. 2000) or are misidentified as small naked amoebae, zoospores, yeasts, or disrupted cells from a range of eukaryotes, since flagella are not always adequately preserved (Patterson and Larsen 1991). Especially benthic HF are affected by the above explained problems because they are additionally masked by sediment particles. Further, large HF can easily be overlooked on membrane filters due to their low contribution to total HF abundance and their special sensitivity to fixatives. This is crucial since they may contribute to about half of the HF biovolume (Arndt et al. 2000). Therefore, the

¹Corresponding author; fax +49 221 470 5932
e-mail hartmut.arndt@uni-koeln.de (H. Arndt).

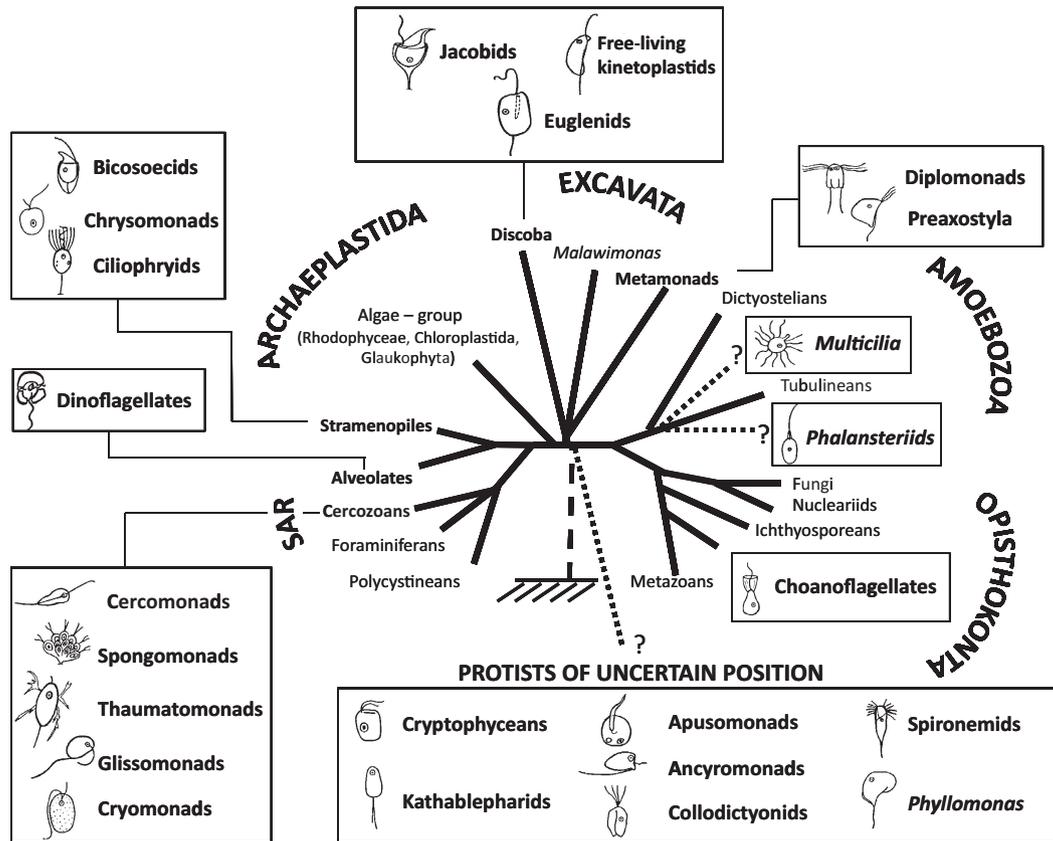


Figure 1. Selective overview of the systematic position (according to Adl et al. (2012)) of flagellate groups considered in the guide (see Charts 1–15).

taxonomic resolution of fixation processes is often very low, most quantitative studies even ignored the variety of HF.

In contrast, live counting is an essential and reliable method for determining HF abundances. It allows a combination of taxonomic and quantitative studies for understanding the role of HF in microbial food webs (Leadbeater and Green 2000; Patterson and Larsen 1991). Unfortunately, the taxonomic resolution of HF studies is hampered by the fact that the literature is often focused on only a few taxonomic groups. Therefore, we designed a (hopefully) user-friendly guide for ecologists to unravel the “HNF” black box. In contrast to the few already existing taxonomic keys/reviews (e.g. Bass et al. 2009; Cavalier-Smith and Chao 2010; Lee et al. 2000; Patterson and Hedley 1992; Patterson and Larsen 1991; Zhukov 1993), the present guide is compact and focuses on the dominant and common forms of HF occurring in pelagic and benthic freshwater habitats. Following the idea of Foissner and Berger (1996) with regard to ciliates, we designed a guide which should be useful also for biologists not specialized in the study of heterotrophic

freshwater flagellates due to its polytomous organization, the help of drawings and video sequences (Supplementary Material). According to our experience in quantitative studies of morphotypes, about 80–90 per cent of the dominant heterotrophic flagellates found in freshwater habitats should be identifiable at least to rough taxonomic categories (and functional guilds). The term morphotype is used here for taxa that can be differentiated by means of their morphology when only analysed by light microscopy. It means that e.g. genotypes may be hidden among certain morphotypes. A typical example is the species complex of *Neobodo designis* (Scheckenbach et al. 2006). The guide comprises all the main groups, so that a large number of species morphotypes can be determined at least to the level of genus. Special attention was given to the most interesting and ecologically relevant forms. The taxonomic categories were used according to the recently revised classifications by Adl et al. (2012).

Within the next decade, next generation sequencing of field samples will give more detailed insights into the structure of HF field communities

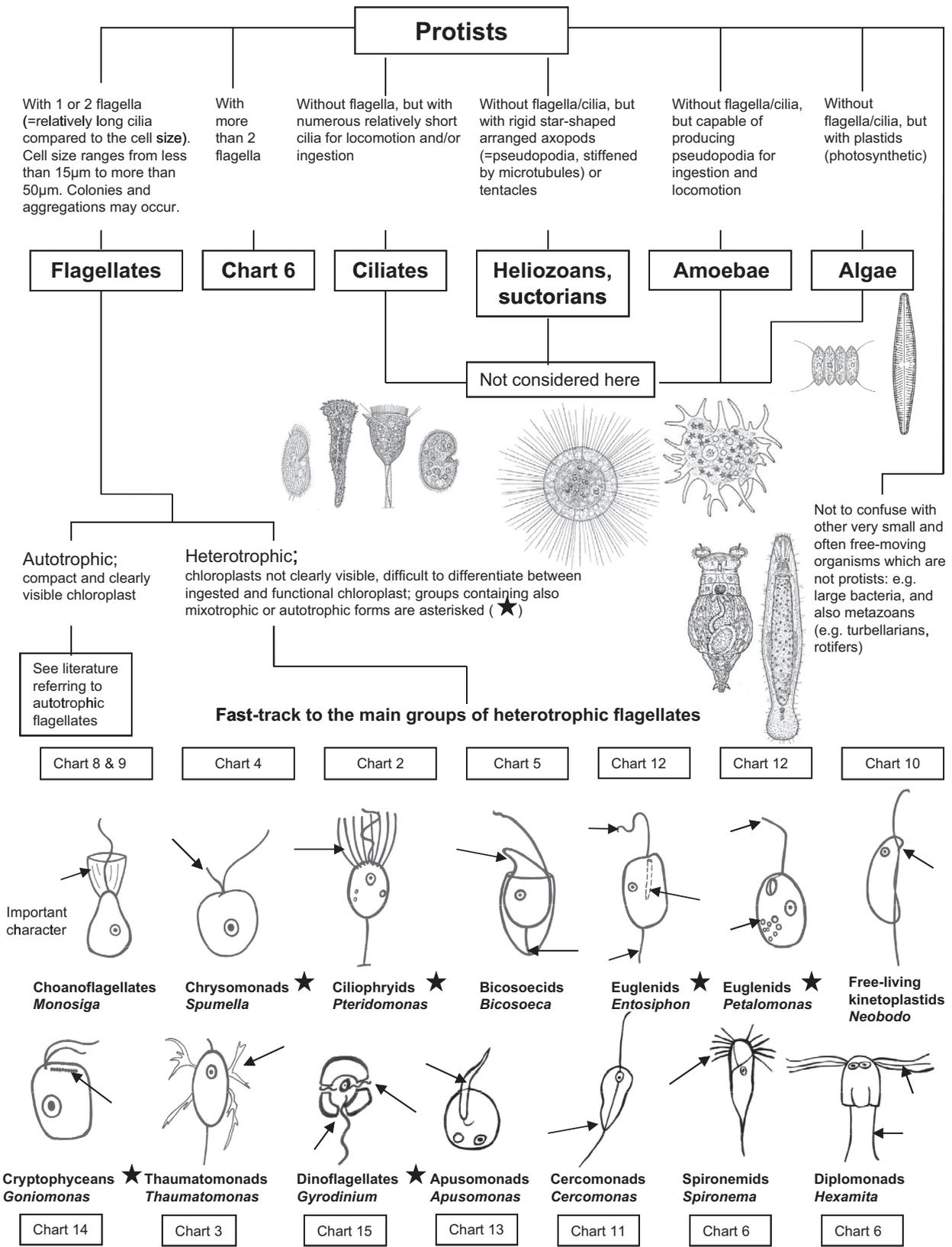


Chart 1. User-friendly guide to common heterotrophic freshwater flagellates. All scale bars in the guide indicate 10 µm.

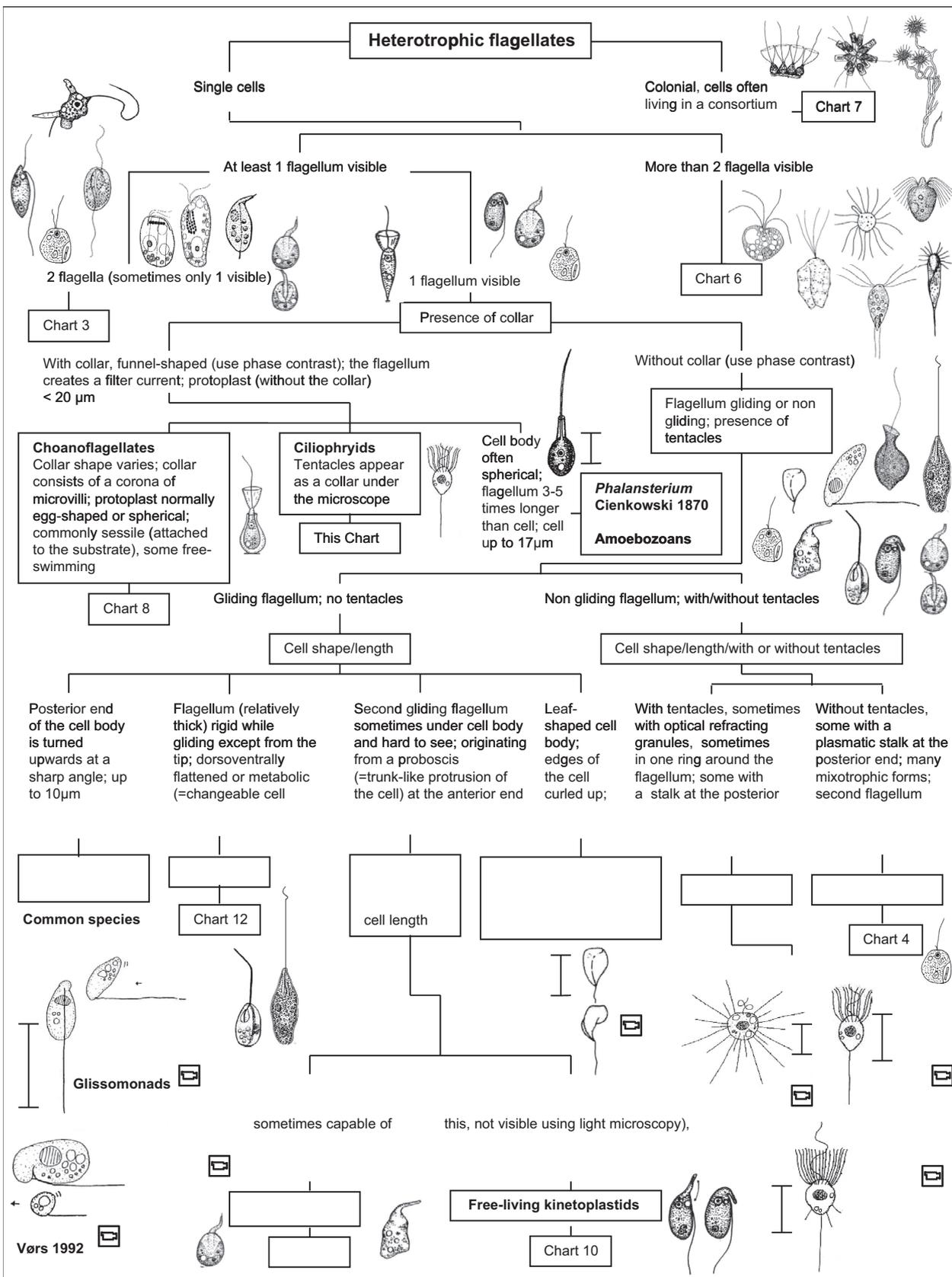


Chart 2.

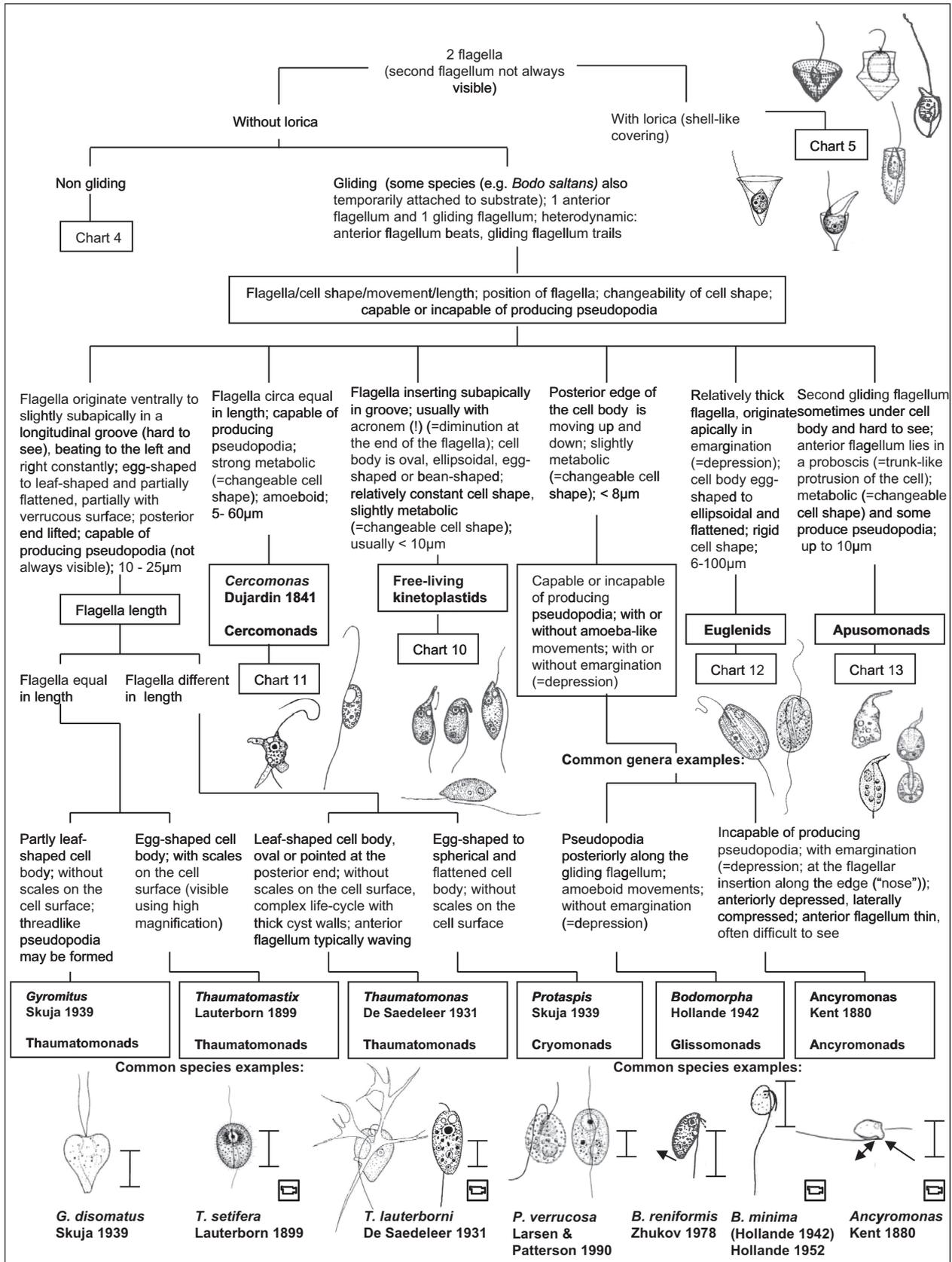


Chart 3.

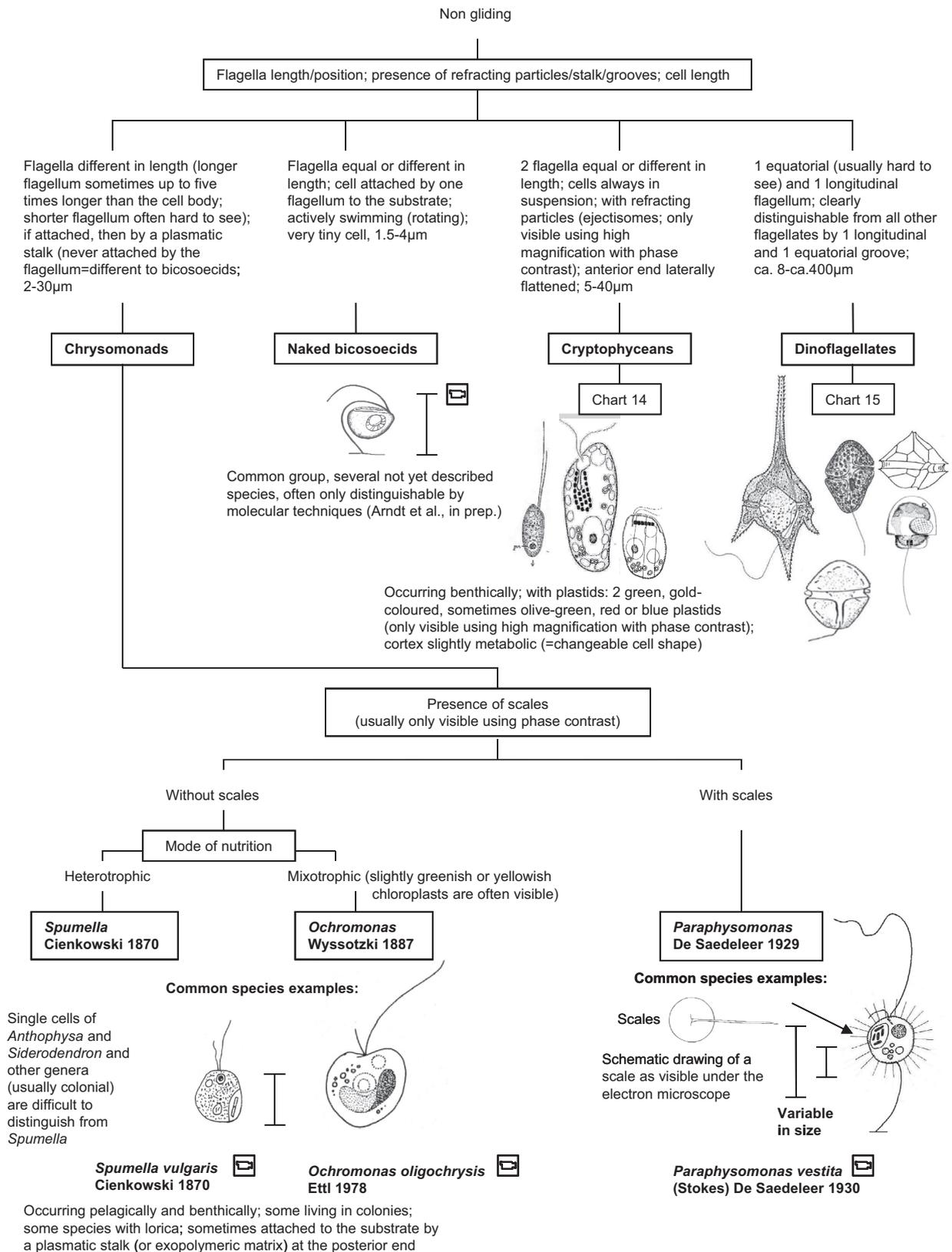
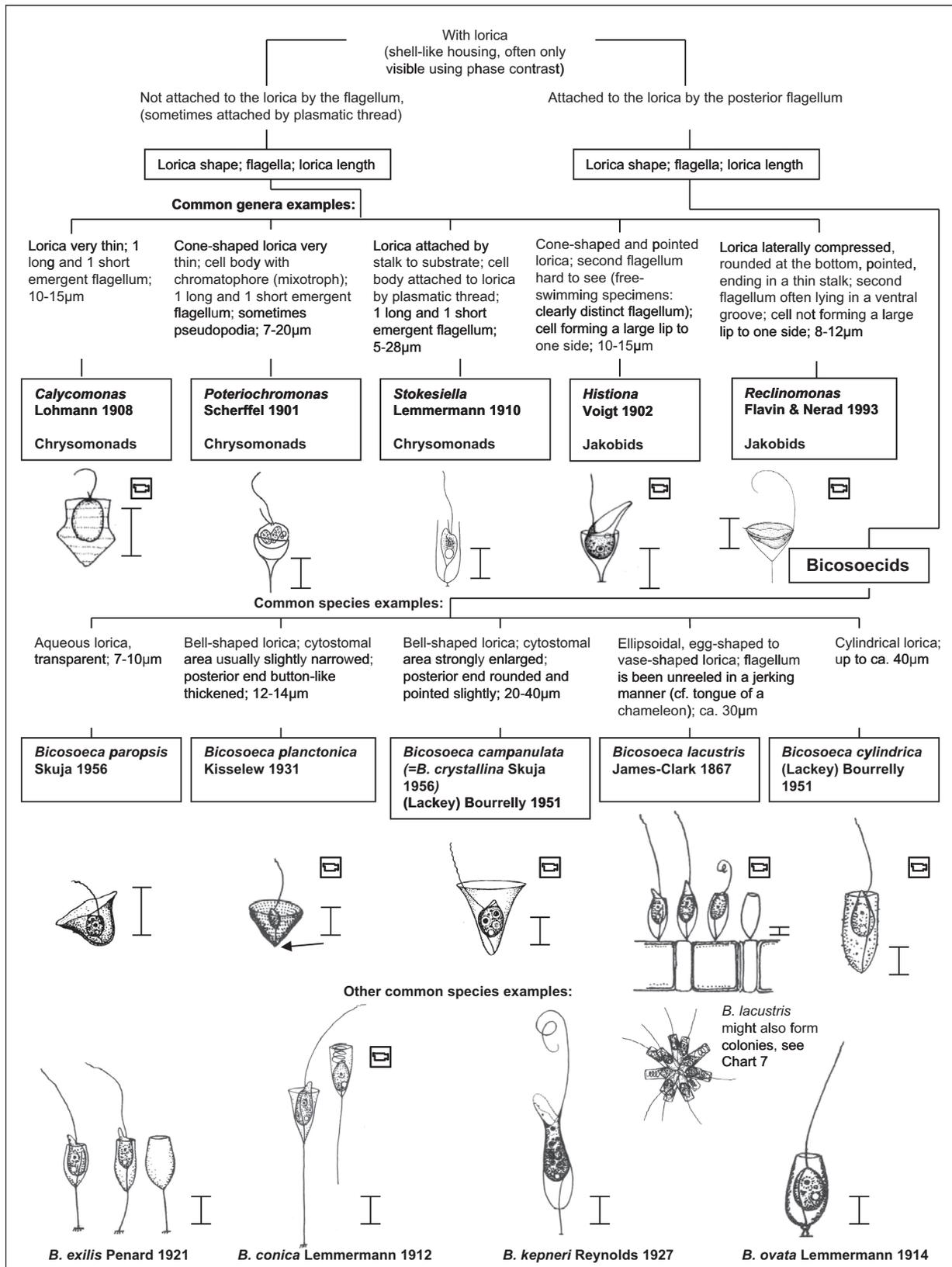


Chart 4.



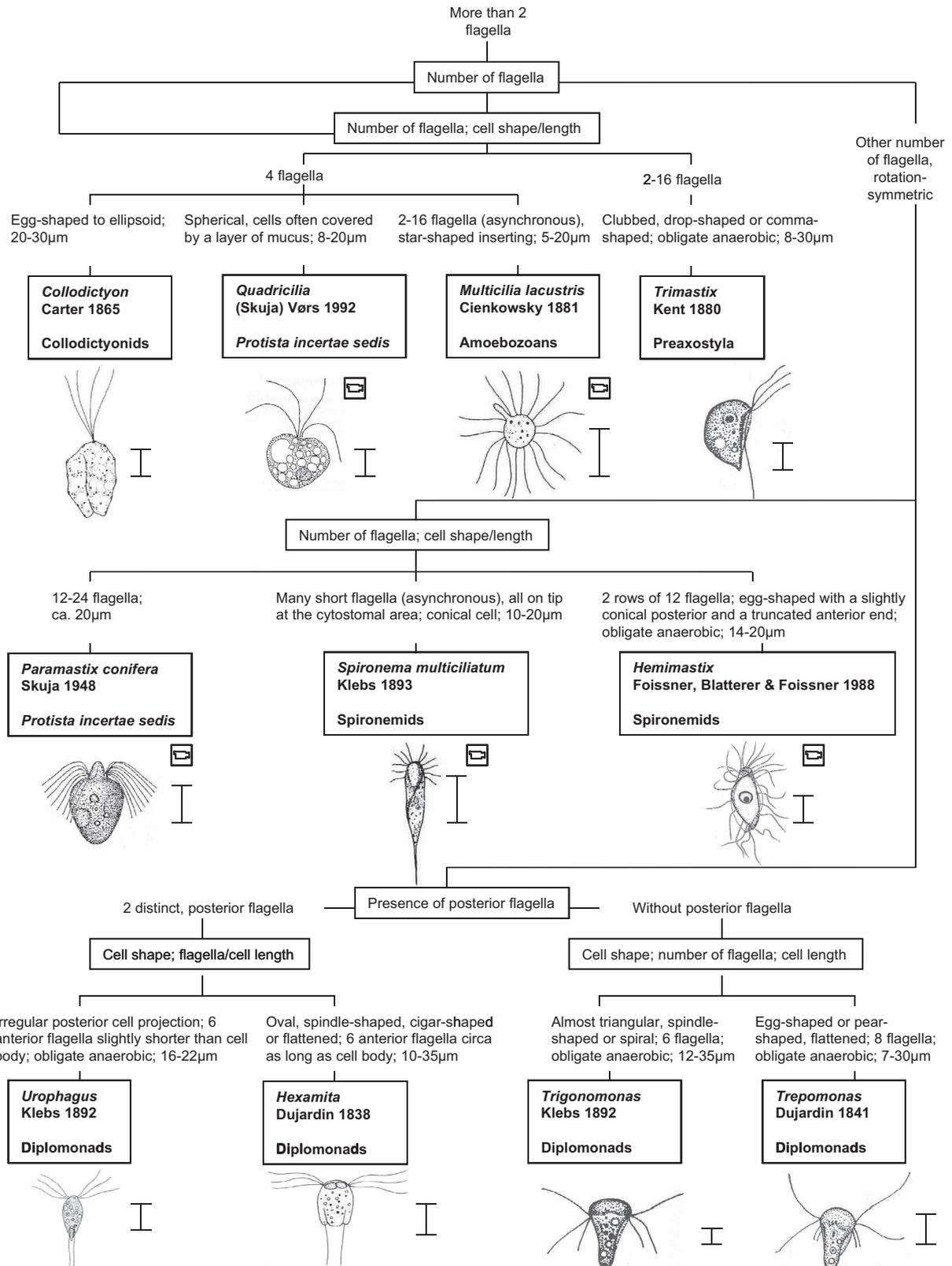


Chart 6.

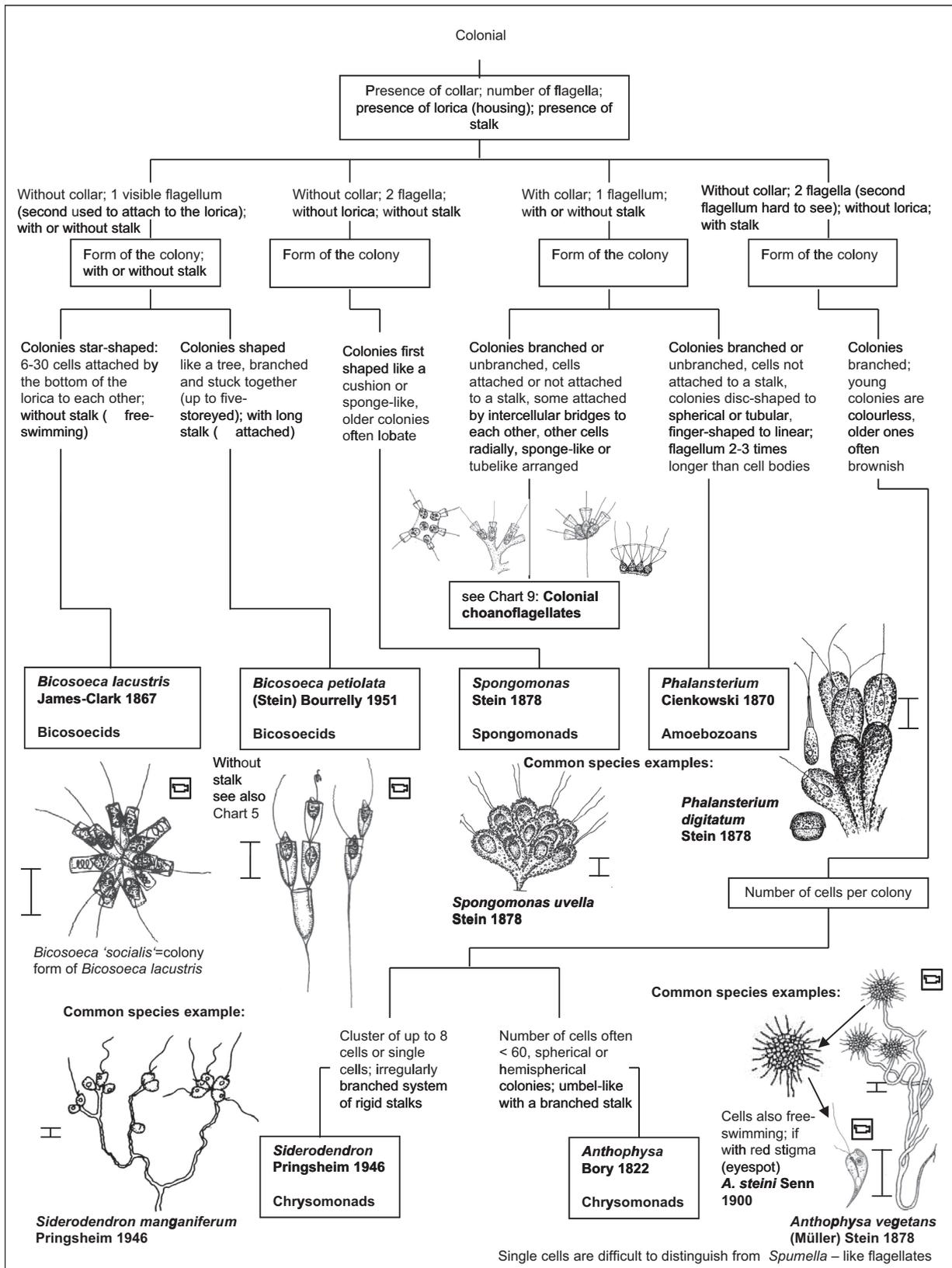


Chart 7.

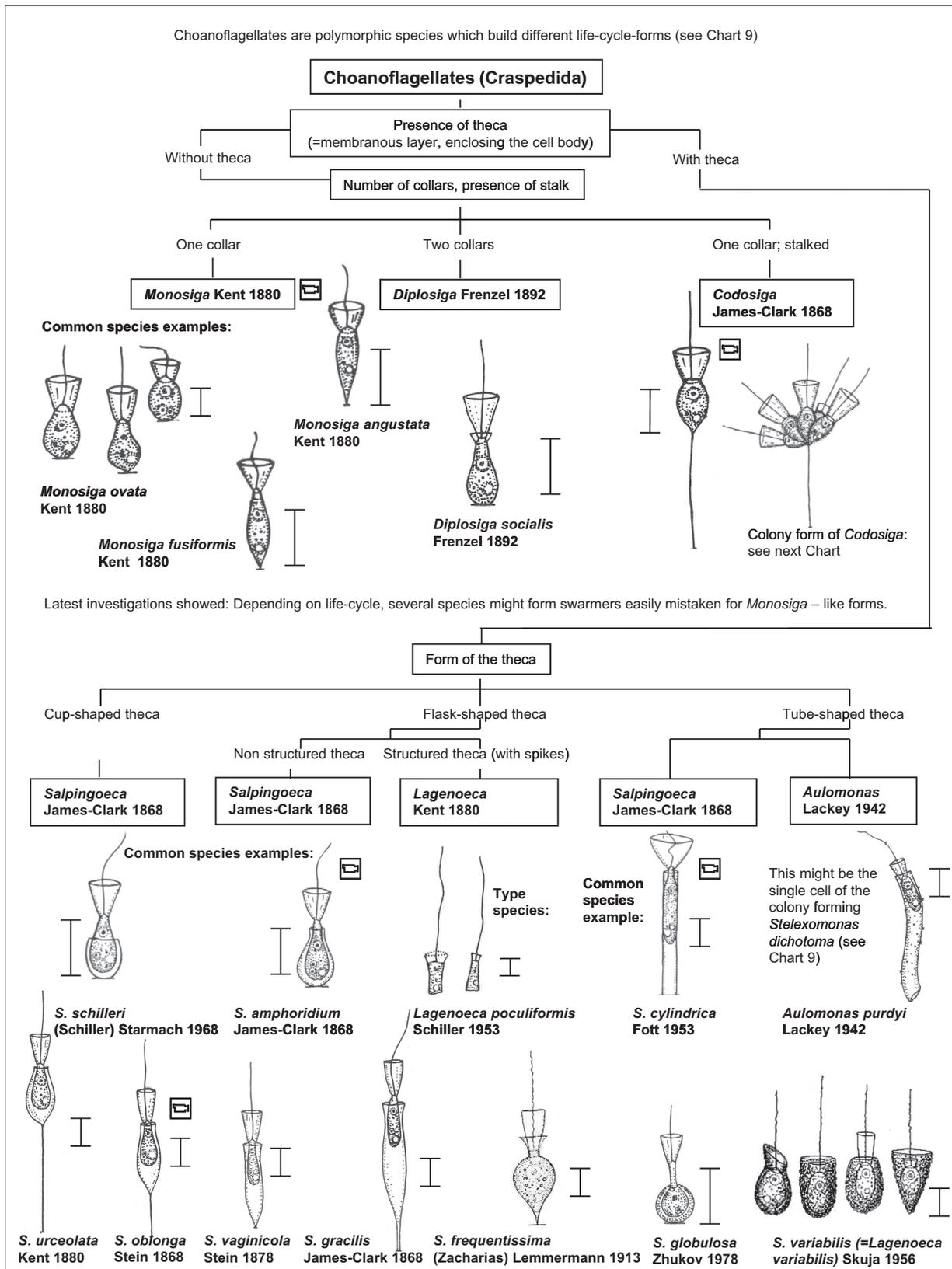


Chart 8.

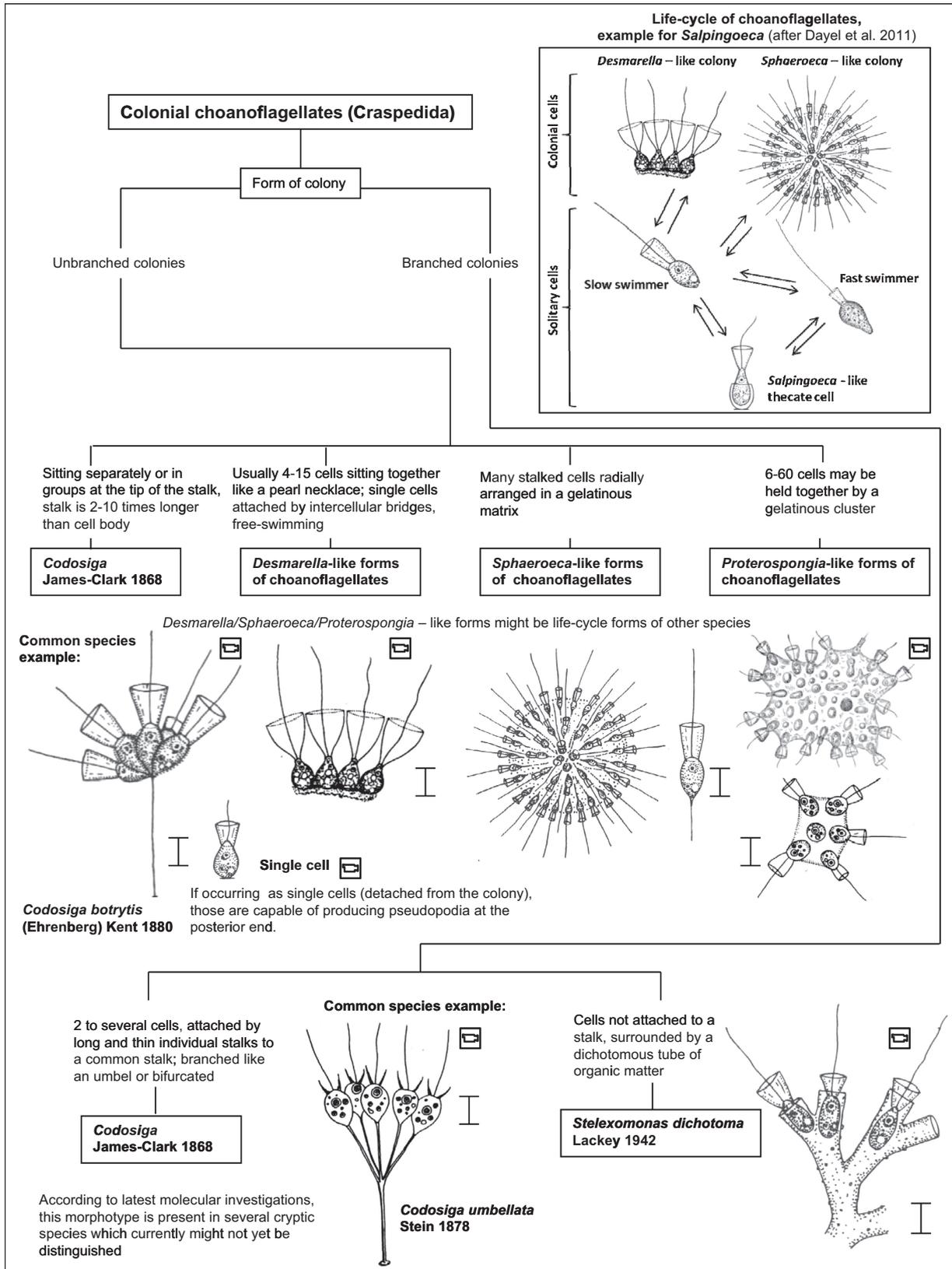


Chart 9.

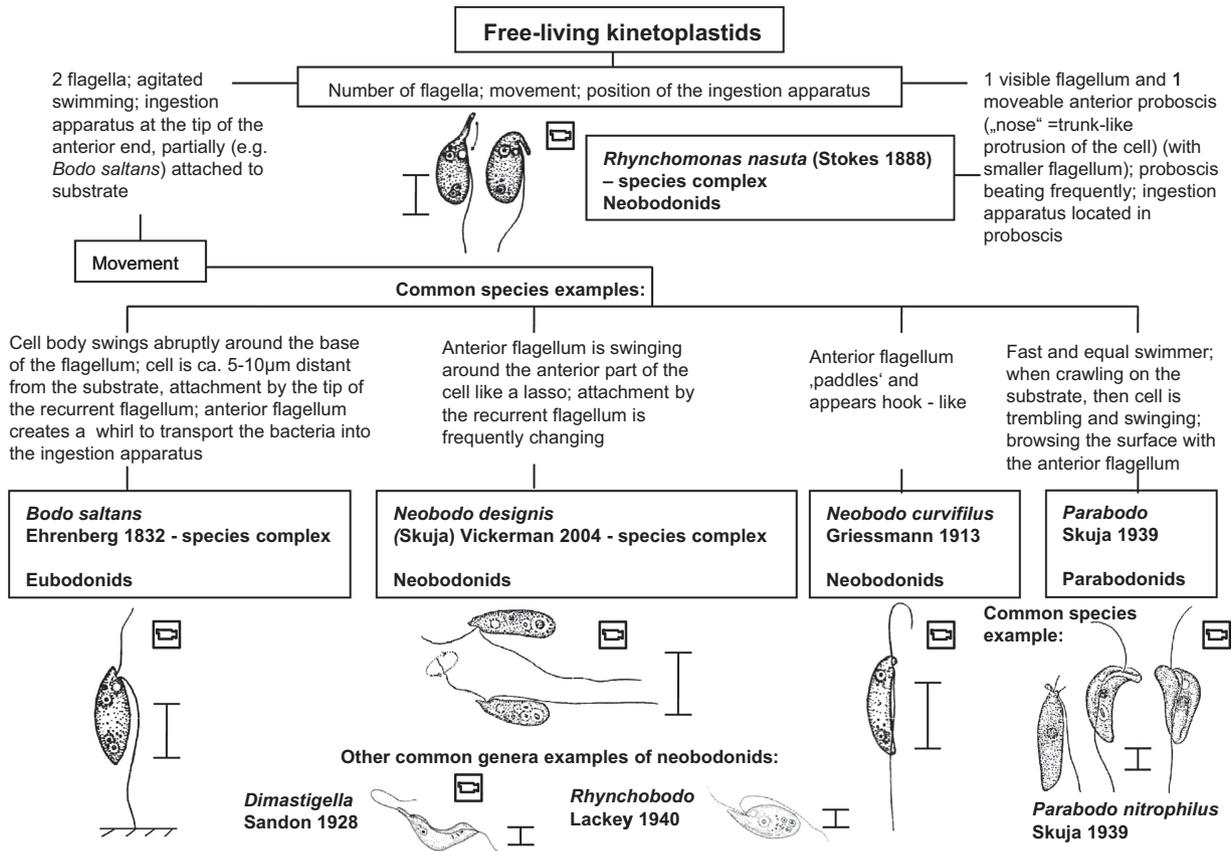


Chart 10.

(e.g. Pawlowski et al. 2011, Stoeck et al. 2010, and Domonell et al., unpubl. observ.). However, their application still bears a lot of problems concerning the following points: 1) many specific primers are necessary to cover the whole HF diversity due to the wide molecular diversity of HF (some groups will be overlooked), 2) active and inactive forms are difficult to differentiate, 3) databases are incomplete (Will and Rubinoff 2004) and contain high numbers of errors (Prosdocimi et al. 2013). Quantitative morphotype studies will provide helpful information to increase reliability of databases and offer important knowledge on the ecology and evolution of HF groups significant in ecosystems (Pawlowski et al. 2012). We would like to increase the resolution of these quantitative morphotype studies by this taxonomic guide.

How to Use the Guide

The present guide is intended to help also unexperienced researchers to identify HF morphotypes, although it is recommended to have a general

knowledge of HF (recommended literature: e.g. Hausmann et al. 2003; Leadbeater and Green 2000; Patterson and Larsen 1991). The systematic position of flagellate groups considered in the guide is given in Figure 1 following the recent revision by Adl et al. (2012). Following the idea of Foissner and Berger (1996) with regard to ciliates, we designed a polytomous guide. The first Chart contains a general overview of the most important characters of the main groups of protists. The user is led to a “fast-track” regarding the main groups of HF (Charts 1–15) by schematic drawings with arrows pointing to the important characters. If identification is already sufficient the user is led to the special Chart of the organism. If not, he should proceed to Chart 2. The user is led from obvious characters (e.g. colonial or single cells, number of flagella) to more detailed descriptions. Features of organisms which are used to discriminate between different taxa are set within boxes while the discriminator for the subsequent path of identification is shown without boxes. Figures are added to the descriptions of characters to accelerate and increase the chance of identification. All scale bars in the guide indicate

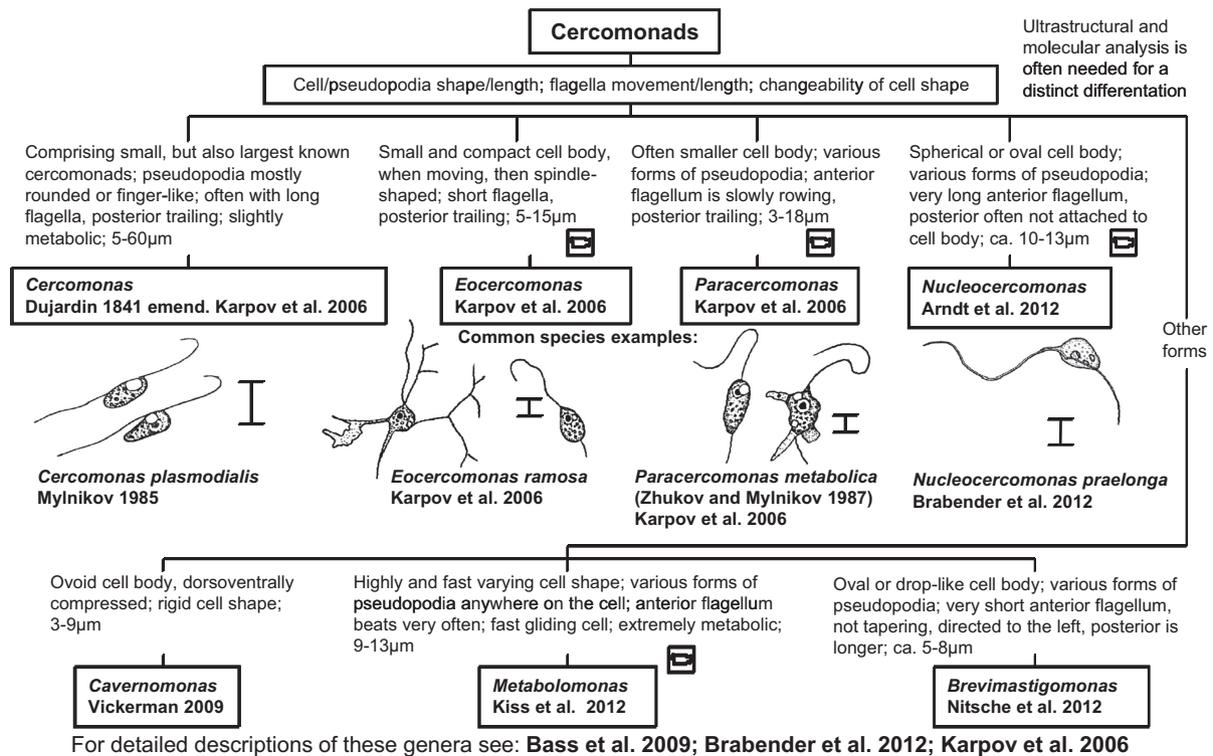


Chart 11.

10 µm. The guide leads only to the species or genus morphotype. It is widely accepted that many protistan morphotypes contain a few up to hundreds of genotypes (typical examples are the common species *Neobodo designis* (Scheckenbach et al. 2006) or *Codosiga botrytis* (Stoupin et al. 2012)), see above. If video sequences (Supplementary Material) are available for the organisms, they are marked by a video icon (📺). Some additional literature is given on the Charts (Cavalier-Smith and Chao 2010; Bass et al. 2009; Brabender et al. 2012; Karpov et al. 2006). Due to the fact that major forms of heterotrophic flagellates are ubiquitous in marine and freshwater environments, this guide may also be useful in parts for an orientation regarding flagellate groups in brackish water systems.

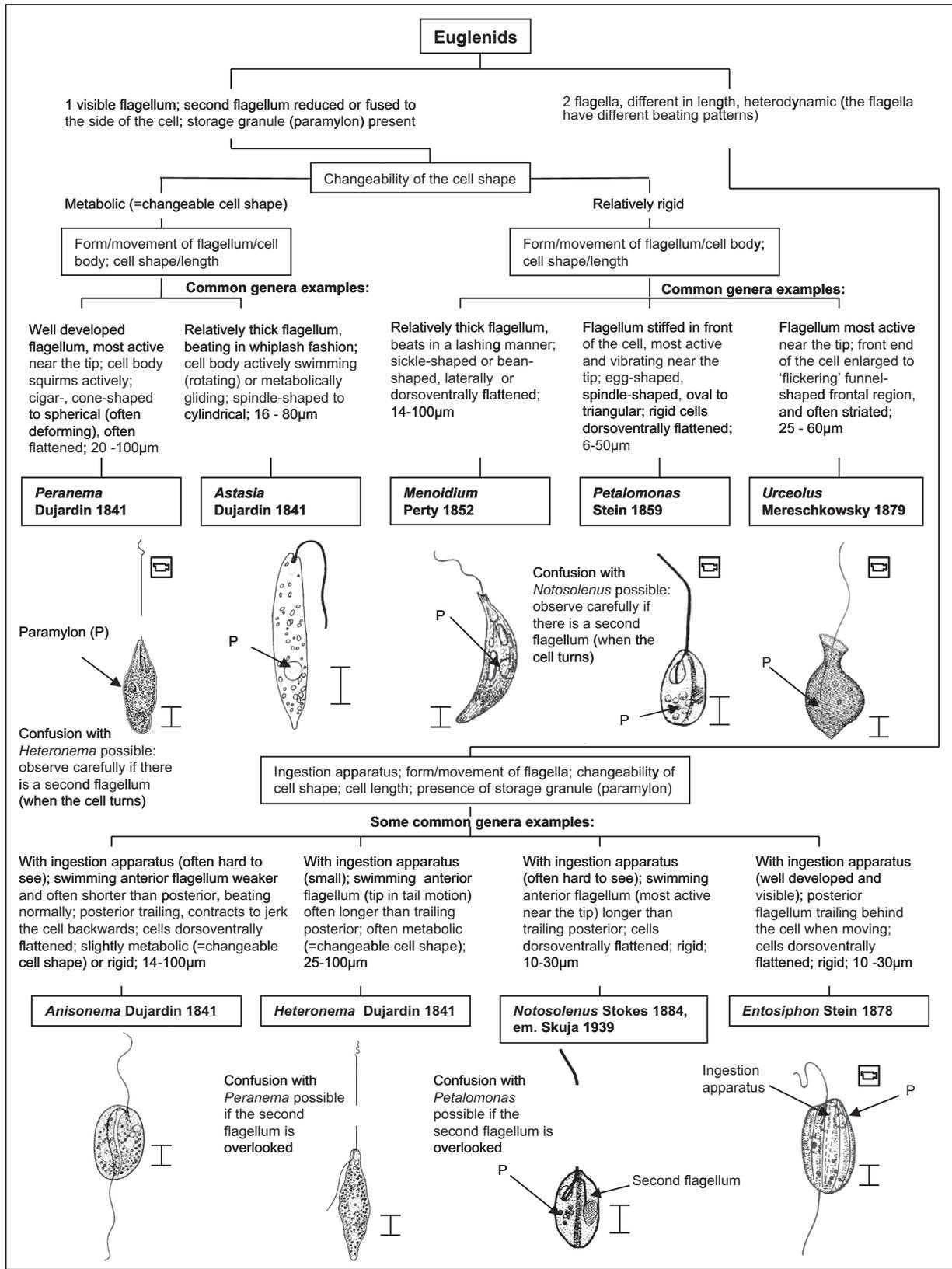
Methods

The guide is based on an analysis using the classical living-droplet method by phase contrast/DIC light microscopy (e.g. Arndt and Mathes 1991; Gasol 1993; Massana and Güde 1991). This method is reliable to detect morphological and behavioural characteristics of even very small HF regarding their main taxonomic groups – in contrast to hidden characteristics within samples treated by fixatives. Pelagic flagellates (undiluted samples) and benthic flagellates (diluted by a factor of 5- >20 with filtered water) can be studied in droplets of

5-20 µl on a prepared microscope slide or an Utermöhl chamber (Utermöhl 1958; HydroBios GmbH, Kiel, Germany; Fig. 2). The HF composition has to be analysed within one hour after sampling by means of a phase contrast microscope equipped with high resolution video-recording which is helpful for further and later identification. 20x, 40x objectives are helpful for rough morphological observations and quantitative counts. Individual samples can be analysed within a few minutes (particularly critical for pelagic samples). The use of a 63x long distance objective or water immersion objectives (63x and 100x; with a long working distance) are recommended for morphotype identifications. Phase contrast equipment is mostly indispensable to analyse position and movement of flagella, the presence of collar, lorica, or stalk of some flagellates. Differential interference contrast (DIC) equipment might additionally help to identify cell structures such as paramylon, ingestion apparatus, vacuoles or extrusomes.

A realistic compromise should be a combination of counts by the living-droplet method with counts of chemically fixed samples and analysis by epifluorescence microscopy. The following fixatives are commonly in use: 2% glutaraldehyde (e.g. Caron 1983; Choi and Stoecker 1989), 2% formaldehyde (e.g. Porter and Feig 1980; Sherr et al. 1989), buffered formaldehyde (e.g. Børshiem and Bratbak 1987; Sherr and Sherr 1983) and 0.5% acidic Lugol' solution (10 g I₂, 20 g KI, 10 g sodium acetate in 140 ml aq. dest.) plus 3% formaldehyde (e.g. Sherr et al. 1989).

The fixed samples need to be kept at 4 °C in the dark until staining. Fluorescent dyes (e.g. DAPI (= 4'6-diamidino-2-phenylindole, Sigma Lot 47H4107, Porter and Feig 1980) are suitable for staining. After staining the sample should be filtered onto a 0.2-µm membrane filter using vacuum not exceeding 150 mm Hg. For an even distribution of the particles, another membrane (0.6 µm pore size) may be placed

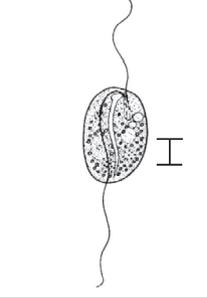


Ingestion apparatus; form/movement of flagella; changeability of cell shape; cell length; presence of storage granule (paramylon)

Some common genera examples:

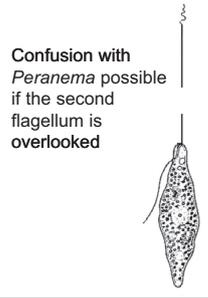
With ingestion apparatus (often hard to see); swimming anterior flagellum weaker and often shorter than posterior, beating normally; posterior trailing, contracts to jerk the cell backwards; cells dorsoventrally flattened; slightly metabolic (=changeable cell shape) or rigid; 14-100µm

***Anisonema* Dujardin 1841**



With ingestion apparatus (small); swimming anterior flagellum (tip in tail motion) often longer than trailing posterior; often metabolic (=changeable cell shape); 25-100µm

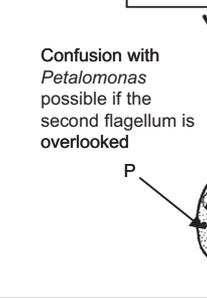
***Heteronema* Dujardin 1841**



Confusion with *Peranema* possible if the second flagellum is overlooked

With ingestion apparatus (often hard to see); swimming anterior flagellum (most active near the tip) longer than trailing posterior; cells dorsoventrally flattened; rigid; 10-30µm

***Notosolenus* Stokes 1884, em. Skuja 1939**



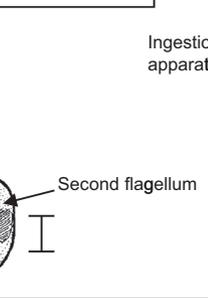
P

Confusion with *Petalomonas* possible if the second flagellum is overlooked

Second flagellum

With ingestion apparatus (well developed and visible); posterior flagellum trailing behind the cell when moving; cells dorsoventrally flattened; rigid; 10 -30µm

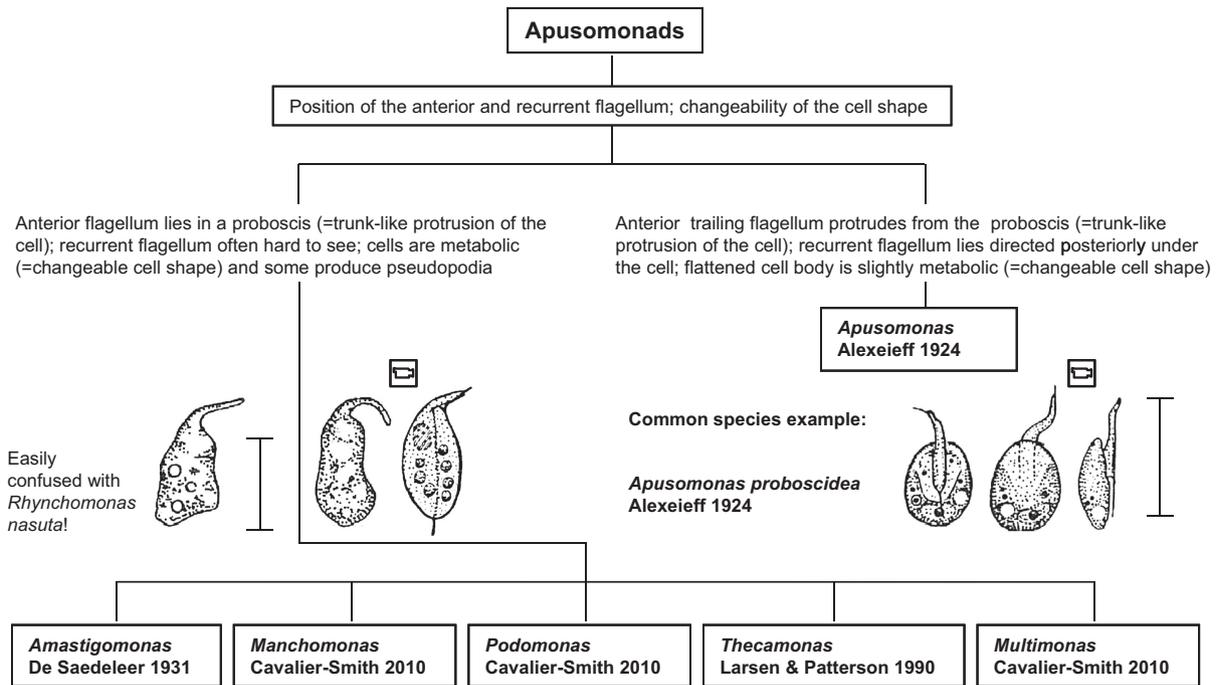
***Entosiphon* Stein 1878**



Ingestion apparatus

P

Chart 12.



For detailed descriptions of these genera see: Cavalier-Smith and Chao 2010

Chart 13.

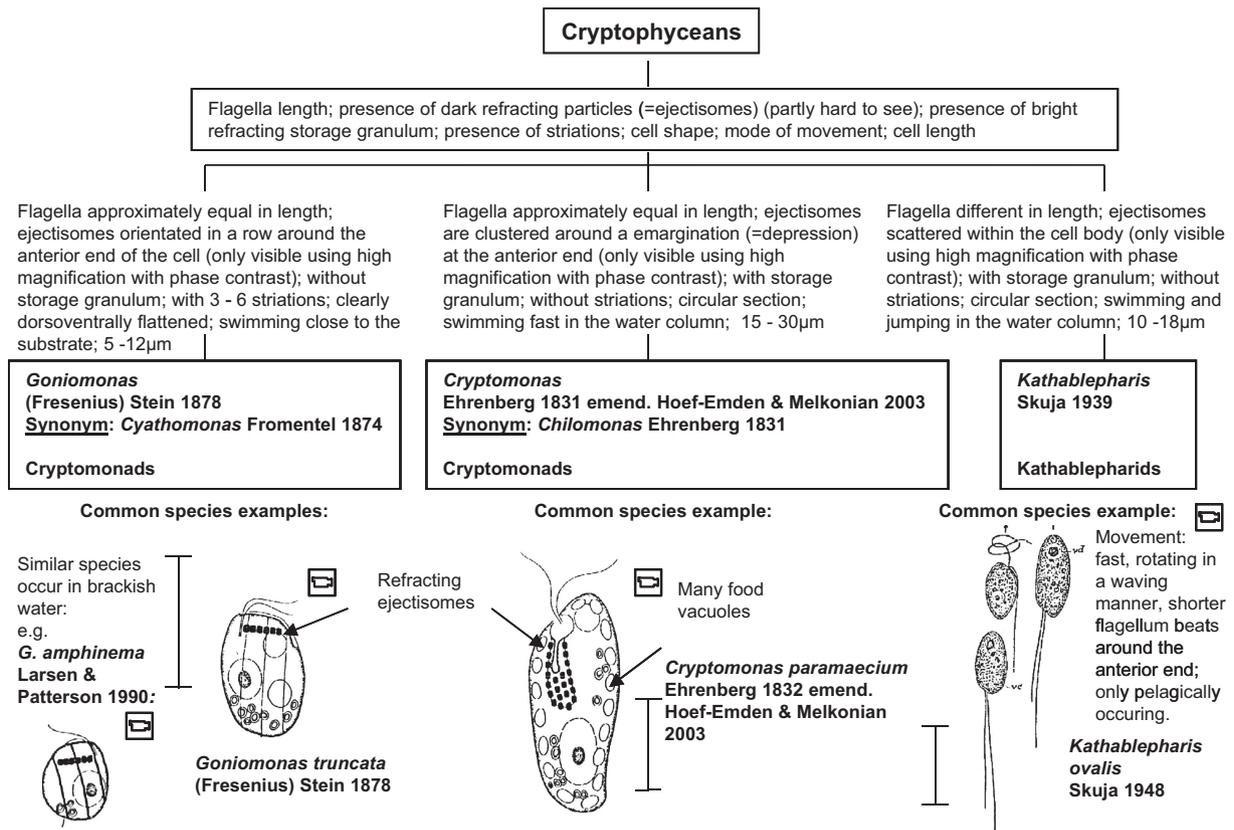


Chart 14.

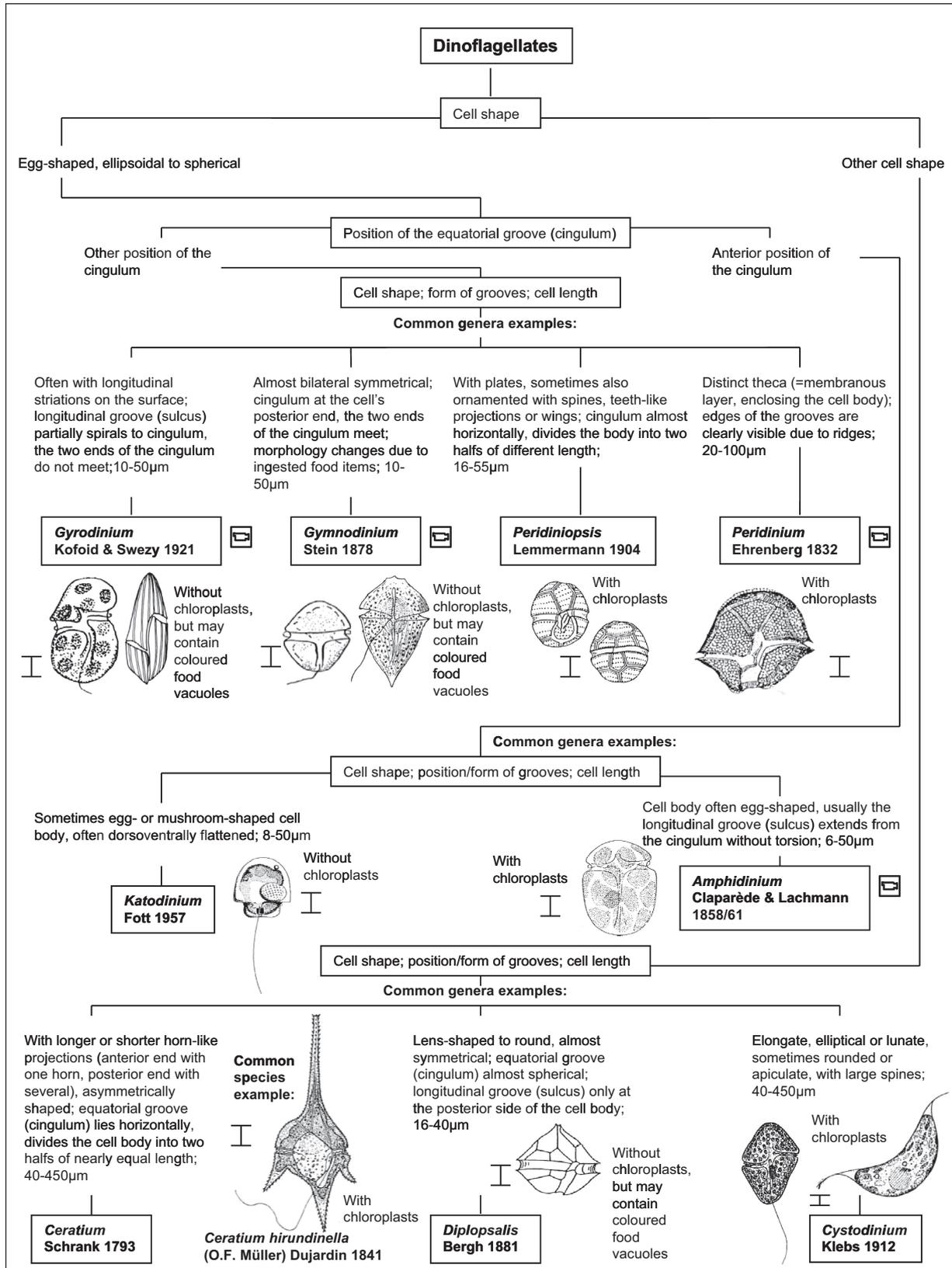


Chart 15.

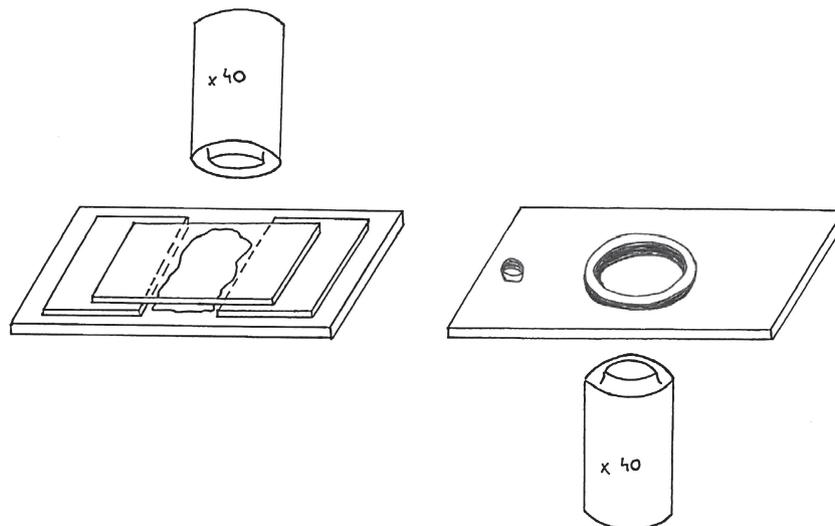


Figure 2. Schematic drawing of some helpful counting chambers. a) Simple microchamber: Two cover slips are placed on the slide (each on every end). A 10 µl droplet is placed between them and a third cover slip above the droplet and the two slips. b) Utermöhl chamber for counting living flagellates or sedimented, chemically fixed samples.

underneath the filter. The filters are generally stored deep-frozen at -20°C until examination by an epifluorescence microscope.

It is additionally recommended to establish cultures of HF isolated from the samples (e.g. simply by adding autoclaved wheat grains etc. as a carbon source for food bacteria to parts of raw samples). This should aid in morphotype identification of bacterivorous forms and to check whether the determined morphotypes are life-cycle forms (swarmer, colony etc.) of other types, as it is often the case with e.g. choanoflagellates where extrinsic factors can induce the formation of multicellular forms (Alegado et al. 2012; Dayel et al. 2011). However, it has to be considered that all cultivation and fixation methods (and also molecular methods) can be very selective. According to our own experience, the percentage of live-counted organisms appearing in cultures ranges between 1 and 30 per cent.

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Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.protis.2013.08.003>.

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