Evolution of reproductive development in the volvocine algae

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Abstract The evolution of multicellularity, the separation of germline cells from sterile somatic cells, and the generation of a male–female dichotomy are certainly among the greatest innovations of eukaryotes. Remarkably, phylogenetic analysis suggests that the shift from simple to complex, differentiated multicellularity was not a unique progression in the evolution of life, but in fact a quite frequent event. The spheroidal green alga *Volvox* and its close relatives, the volvocine algae, span the full range of organizational complexity, from unicellular and colonial genera to multicellular genera with a full germ–soma division of labor and male–female dichotomy; thus, these algae are ideal model organisms for addressing fundamental issues related to the transition to multicellularity and for discovering universal rules that characterize this transition. Of all living species, *Volvox carteri* represents the simplest version of an immortal germline producing specialized somatic cells. This cellular specialization involved the emergence of mortality and the production of the first dead ancestors in the evolution of this lineage. Volvocine algae therefore exemplify the evolution of cellular cooperation from cellular autonomy. They also serve as a prime example of the evolution of complex traits by a few successive, small steps. Thus, we learn from volvocine algae that the evolutionary transition to complex, multicellular life is probably much easier to achieve than is commonly believed.

Keywords Cell differentiation · Multicellularity · Reproduction · Green algae · *Volvox carteri* · *Chlamydomonas reinhardtii*

Introduction

Since the early work of August Weismann on the continuity of germ plasm at the end of the nineteenth century (Weismann 1893, 1892, 1889), the spheroidal green alga *Volvox* and its close relatives have been seen as suitable model organisms for addressing fundamental issues in the evolution of multicellularity and the development of a germ–soma dichotomy (Kirk 1998, 2001, 2003, 2005; Nozaki 2003; Hallmann 2003b; Schmitt 2003; Kirk and Kirk 2004; Michod et al. 2006). *Volvox carteri* illustrates the concept of a germ–soma dichotomy with diagrammatic clarity (Fig. 1): an adult *V. carteri* spheroid has only two cell types, ~2,000 small *Chlamydomonas*-like somatic cells, and ~16 large asexual reproductive cells (gonidia). The potentially immortal gonidia are nonmotile cells that are specialized for growth and reproduction; these cells constitute the germline. In contrast, the biflagellate somatic cells are specialized for extracellular matrix (ECM) biosynthesis, motility, and phototaxis. They are also incapable of dividing and are programmed to die when only a few days old. Therefore, of all living species, *Volvox* represents the simplest version of an immortal germline producing mortal soma.

Particularly with regard to questions about the evolution of multicellularity and of the germ–soma dichotomy, both *Volvox* and its close relatives, the volvocine green algae are of special interest (Kirk 1998, 2005; Prochnik et al. 2010). The volvocine green algae form a group of genera closely related to the genus *Volvox* within the order...
Volvocales (Chlorophyta). This group spans the full range of organismal complexity, from unicellular organisms, such as *Chlamydomonas reinhardtii*, to colonial organisms without a division of labor, such as *Gonium pectorale*, *Pandorina morum*, and *Eudorina unicocca*, to multicellular organisms with a partial or full germ–soma division of labor, such as *Pleodorina californica* and *V. carteri*, respectively (Fig. 1).

To facilitate detailed molecular analyses in volvocine algae, an array of important modern analytical tools has been developed. For example, stable nuclear transformation is possible (Fernández et al. 1989; Debuchy et al. 1989; Kindle et al. 1989; Mayfield and Kindle 1990; Schiedlmeier et al. 1994; Lerche and Hallmann 2009), unselectable markers can be efficiently co-transformed together with selectable markers (Kindle et al. 1991; Gruber et al. 1992; Schiedlmeier et al. 1994; Minko et al. 1999; Lerche and Hallmann 2009), suitable reporter genes and both inducible and constitutive promoters are available (Goldschmidt-Clermont 1991; Hallmann and Sumper 1994; Lerche and Hallmann 2009), expression of chimeric genes works (Blankenship and Kindle 1992; Hallmann and Sumper 1994; Lerche and Hallmann 2009), foreign genes can be expressed (Hallmann and Sumper 1996; Stevens et al. 1996; Hallmann and Rappel 1999; Lerche and Hallmann 2009), dominant selectable markers are available (Nelson et al. 1994; Hallmann and Rappel 1999), gene replacement by homologous recombination is feasible (Sodeinde and Kindle 1993; Hallmann et al. 1997), plasmids or transposons can be used for gene tagging (Tam and Lefebvre 1993; Schnell and Lefebvre 1993; Miller and Kirk 1999; Ueki and Nishii 2008), and the genomes of *C. reinhardtii* and *V. carteri* have been sequenced (Merchant et al. 2007; Prochnik et al. 2010).

In summary, the existence of a germ–soma division of labor in *Volvox*, the range of complexity from unicellular to multicellular species in volvocine algae, and the availability of an extensive molecular toolkit make volvocine green algae useful model organisms for studying the evolution of multicellularity and of sterile somatic cells.

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**Fig. 1** Rough approximation of the evolution of volvocine green algae from unicellular forms to colonial and multicellular forms with increasing complexity. Six representative species with characteristic developmental traits were arranged such that there is a progressive increase from left to right in morphologic and developmental complexity (Kirk 1998, 2000; Prochnik et al. 2010). Check marks indicate that a given trait is present in the respective species. Graded differences in a given trait are indicated by 1–5 plus signs and ± indicates ambiguity or occasional occurrence. The photomicrographs show *Chlamydomonas reinhardtii* (SAG 11-32b), *Gonium pectorale* (SAG 12.85), *Pandorina morum* (SAG 60-1d), *Eudorina unicocca* (SAG 24-1c), *Pleodorina californica* (SAG 32.94), and *Volvox carteri* (Eve). The exact phylogenetic position of these six species is indicated with filled black circles in the evolutionary tree in Fig. 3.
A phylogenetic perspective on the transition from unicellularity to multicellularity

In many textbooks, volvocine algae are used to illustrate the evolutionary transition from unicellular to multicellular organisms, and some readers might therefore mistakenly arrive at the conclusion that this transition happened only once in the evolution of all multicellular life. However, multicellularity has arisen independently again and again in the tree of life, over billions of years. In fact, these transitions were not even restricted to eukaryotes but occurred in prokaryotes as well: The first evidence of a shift to multicellularity comes from fossilized Cyanobacteria-like prokaryotes, which lived 3–3.5 billion years ago (Schopf 1993; Knoll 2003), and the first multicellular eukaryotes may have existed about 1 billion years ago (Knoll et al. 2006). Between about 530 and 570 million years ago, a burst of diversification occurred, with the eventual appearance of the lineages of almost all metazoa living today (Carroll 2001; Knoll 2003; King 2004; Maynard Smith and Szathmáry 1995; Pfeiffer and Bonhoeffer 2003; Grosberg and Strathmann 2007). Among the eukaryotes, multicellular organisms independently originated at least 25 times from unicellular ancestors (followed in some cases by secondary losses) (Buss 1987; Cavalier-Smith 1991; Maynard Smith and Szathmáry 1995; Kirk 1998; Bonner 1998, 2000; Carroll 2001; Kaiser 2001; Medina et al. 2003; Baldauf 2003; King 2004; Grosberg and Strathmann 2007; Rokas 2008; Prochnik et al. 2010).

Almost every lineage of the eukaryotic tree of life includes multicellular forms; the most well-known lineages are the Embryophyta (land plants), Chlorophyta (green algae), Rhodophyta (red algae), Phaeophyceae (brown algae), Bacillariophyta (diatoms), fungi, and animals (Fig. 2).

For most multicellular lineages, it has been quite challenging to investigate the shifts to multicellularity because the transitions occurred so long ago (Herron and Michod 2008). Therefore, a molecular understanding of the genetic changes underlying this transition might even be unattainable in these lineages. In contrast, the volvocine algae represent a unique opportunity to study the transition to multicellularity at a molecular level because the multicellular members of this group diverged relatively recently from their unicellular relatives: the last common ancestor of the unicellular alga *C. reinhardtii* and the multicellular alga *V. carteri* lived only 50–200 million years ago (Rausch et al. 1989; Herron and Michod 2008; Herron et al. 2009). Moreover, extant volvocine species display a range of intermediate grades between unicellular forms and multicellular forms with a complete separation of germ and soma (Fig. 1).

Taxonomically, the volvocine algae belong to the order Volvocales. Together with orders such as the Dunaliellales, Chlorococcales, Oedogoniales, and Chaetophorales, the Volvocales belong to the Chlorophyceae, one of the four classes of Chlorophyta (green algae) (Fig. 2). Detailed molecular phylogenetic analyses of the volvocine algae have revealed that the history of the whole group is quite complex, and not just a simple, linear progression in size and complexity (Nozaki et al. 2000; Nozaki 2003; Nakada et al. 2008; Herron and Michod 2008; Herron et al. 2010). Therefore, the few volvocine species arranged in order of increasing complexity in Fig. 1 reflect only a rough approximation of the evolution of volvocine green algae. The detailed molecular phylogenetic reconstruction of volvocine algae shown in Fig. 3 confirms that volvocine algae constitute a robust monophyletic group with two major families within this group, the Goniaceae and the Volvocaceae (Larson et al. 1992; Nozaki et al. 1995, 1999, 2000, 2002; Kirk 1998; Coleman 1999; Nozaki 2003;...
Nakada et al. 2008; Herron and Michod 2008; Herron et al. 2010). However, the evolutionary tree of volvocine algae is highly branched, and several taxa, including the genus *Volvox*, are found on more than one branch. The structure of volvocine taxonomy results from the fact that early classifications of this group were based on morphological correlations rather than phylogenetic relationships; several nominal taxa appear to represent convergent morphologies rather than monophyletic relationships. Therefore, the genus and species names of volvocine algae probably should be revised based on molecular phylogenetic reconstructions (Fig. 3).

A detailed molecular phylogenetic analysis indicates that multicellularity evolved only once in volvocine algae (Fig. 3) (Larson et al. 1992; Nozaki et al. 1995, 1999, 2000, 2002; Kirk 1998; Coleman 1999; Nozaki 2003; Nakada et al. 2008; Herron and Michod 2008; Herron et al. 2010). After this transition to multicellularity, a partial germ–soma division of labor emerged independently in three different lineages. Organisms are characterized as exhibiting a “partial division of labor” if their gonidia are derived by redifferentiation of cells that had initially differentiated as biflagellate (somatic-like) cells and contributed to the organism’s motility, whereas organisms are characterized as exhibiting a “full” or “complete division of labor” if their gonidia differentiate without first developing functional flagella and contributing to the motility of the organism (Herron et al. 2010). In volvocine algae, a full germ–soma division of labor evolved three times (within one of the three lineages that show partial germ–soma division) and led to the species *Volvox gigas*, *Volvox africanus*, *Volvox obversus*, and *V. carteri* (Fig. 3) (Nozaki et al. 1997; Herron and Michod 2008; Herron et al. 2009, 2010). The evolution of germ–soma division also reflects the evolution of a sterile soma, which involves the emergence of mortality and the production of the first dead ancestors in the evolution of this lineage. Remarkably, the innovation “division of labor” was actually lost twice in volvocine algae (Nozaki et al. 1997; Herron and Michod 2008; Herron et al. 2009, 2010).

Fig. 3 Evolutionary tree of volvocine algae based on the nucleotide sequences of five chloroplast genes. This phylogenetic analysis indicates that multicellularity evolved only once in this group. In contrast, a partial germ–soma division of labor evolved independently in three different lineages and was lost twice (Nozaki et al. 1997; Herron and Michod 2008; Herron et al. 2009). A full germ–soma division also evolved three times. The meanings of symbols and letters are given in the figure. Names of strains are given in gray behind the species names. The species shown on the photomicrographs in Fig. 1 are indicated with filled black circles. This tree was adapted from Herron and Michod (2008) and others (Sachs 2008; Herron et al. 2010); some additional information was added (Lerche and Hallmann 2009; Ueki et al. 2010).
Driving forces in the evolutionary transition toward differentiated multicellularity

Many previous studies address the question why and how multicellularity and germ–soma differentiation evolved in volvocine algae (and elsewhere) (Bonner 1965, 1974, 1988, 1993, 1998, 2000, 2003; Kirk 1988, 1994, 1995, 1997, 1998, 1999, 2000, 2001, 2003, 2005; Kirk et al. 1990, 1993; Koufopanou and Bell 1991, 1993; Desnitski 1993, 1995; Koufopanou 1994; Bell 1998; Schmitt 2001, 2003; Miller 2002, 2010; Stark and Schmitt 2002; Hallmann 2002, 2003a, b, 2006b; Michod et al. 2003, 2006; Pfeiffer and Bonhoeffer 2003; Kirk and Kirk 2004; Nedelcu and Michod 2004; Cheng et al. 2005; Solari et al. 2006a, b, 2007; Grosberg and Strathmann 2007; Willensdorfer 2008, 2009; Sachs 2008; Rokas 2008; Herron et al. 2009; Gavrilets 2010). The multicellular alga *C. reinhardtii* is dioecious (i.e., it produces separate male and female organisms), but the asexual life cycles of both sexes (mating types) are indistinguishable (Figs. 5, 6); the same is true for the unicellular alga *C. reinhardtii* (Fig. 4) (Kirk 1998; Harris et al. 2009).

Volvocine algae switch from their principal, asexual mode of reproduction to a sexual mode when life-threatening conditions approach (Figs. 4, 6) (Kirk 1998; Harris et al. 2009). Thus, sexuality is a form of emergency response program for volvocine algae. Although the sexual cycle results in the production of resistant zygospores, as well as genetic recombination and repair, it does not result in any significant amount of increase in population size.

Key evolutionary innovations in the asexual life cycle

When multicellular organisms evolved from unicellular ones, previous achievements had to be modified and developed to fulfill the requirements of multicellularity (Szathmáry and Smith 1995; Bonner 1998; Grosberg and Strathmann 2007; Prochnik et al. 2010). A comparison of the developmental and reproductive traits of the modern multicellular alga *V. carteri* with those of the extant unicellular alga *C. reinhardtii* and species with intermediate complexity provides insight into the evolutionary innovations that were required for multicellular life.

The cell division cycle of most asexually reproducing volvocine algae is different from the cycles seen in most other organisms (Kirk 2003, 1998). The asexual reproductive cells of most volvocine algae grow 2^n-fold in size and then divide rapidly n times by multiple fission to produce 2^n offspring cells. This type of cell division is known as palintomy and multiple fission (Sleigh 1989; Desnitski 1995). The n has a value from 2 to 15 depending on the volvocine species and, to some extent, the environmental conditions (Kirk 2003, 1998). In *C. reinhardtii*, the n is usually 2 or 3 (Fig. 4a, b), and in *V. carteri*, it is usually 11 or 12 (Fig. 5). However, there is an important difference between the multiple fission programs of these two species: the offspring cells in *C. reinhardtii* almost always separate from each other (complete cytokinesis) and eventually become unicellular individuals, while the offspring cells in multicellular volvocine algae stay linked to each other by cytoplasmic bridges throughout the rest of embryogenesis due to an incomplete cytokinesis (Fig. 1). Hence, incomplete cytokinesis and an increase in the number of cleavage divisions led to an evolutionary rise in adult cell number in volvocine algae, ranging from one cell in *C. reinhardtii* to many thousands of cells in various species of *Volvox* (Figs. 1, 6a, b).

At the end of cleavage, each embryo of colonial or multicellular volvocine algae contains all of the cells that will be present in an adult of the next generation. However,
The orientation of the cells of the embryo is inside out with respect to the adult configuration; that is, the cells have their flagellar ends pointing toward the interior, rather than toward the exterior where they will be needed to function in locomotion. To correct this maladaptive arrangement and to achieve the adult configuration, the embryo turns itself right-side out in a gastrulation-like morphogenetic process called embryonic inversion (Figs. 1, 5) (Halmann 2006a). The cytoplasmic bridges that link all embryonic cells play a major role in inversion (Kirk 1998; Kirk and Nishii 2001).

The shift to organisms with more than one cell also required the elaboration of a multifunctional ECM out of the simple ECM (usually called cell wall) of the Chlamydomonas-like ancestor. In all volvocine algae, the structured ECM meshwork is assembled mainly from hydroxyproline-rich glycoproteins (Miller et al. 1974; Kirk et al. 1986; Sumer and Hallmann 1998; Hallmann 2003b, 2006b). In volvocine algae with high levels of organismal complexity like Volvox, the ECM not only embeds the cells in the surface of a transparent sphere, holds them together and allows for their suitable orientations but also represents a rather dynamic and multifunctional interface between each cell and its neighboring cells and/or environment. The ECM is continuously adapted to changing intracellular, organism-specific or environmental conditions, and ECM remodeling is essential for embryonic development, morphogenesis, and reproduction (Nagase and Woessner 1999; Hallmann 2003b). As the number of cells increased during the evolution of the volvocine lineage, the size of organisms increased even more rapidly due to an exponential increase in the volume of ECM per cell (Fig. 1). In adult V. carteri spheroids in the stage shortly before the release of juveniles (Fig. 5), the ECM constitutes up to 99% of the spheroid volume, whereas C. reinhardtii has only a relatively thin ECM (Fig. 1). The evolution of large multicellular spheroids with a voluminous ECM might result in an improved protection of the next generation, which develops inside the parent spheroid (Fig. 5). Additionally, predation spares larger organisms: small green flagellates are subject to predation by invertebrate filter feeders such as rotifers and small crustaceans, but most colonial volvocine algae are too large to be consumed by these filter feeders (Bell 1985; Kirk 1998).

Traits that impact forward swimming and phototaxis also required evolutionary adaptation during the transition to an organism with more than one cell. In multicellular organisms, phototactic swimming requires a proper and stable orientation of cells with respect to one other and some kind of coordination between cells (Ueki et al. 2010).
Furthermore, unicellular volvocine algae like *Chlamydomonas* show a characteristic breaststroke-type swimming motion, which is inappropriate for propelling a multicellular spheroidal organism. In multicellular volvocine algae, the effective strokes of both flagella of each cell beat in more or less the same direction to push the organism forward (Gerisch 1959; Hoops and Floyd 1983; Hoops 1993; Ueki et al. 2010). This change in beating was accomplished by an evolutionary rotation of the basal bodies, which underlie the flagella and determine their orientations (Fig. 1) (Gerisch 1959; Greuel and Floyd 1985; Kirk 2005). In multicellular volvocine algae, flagellar beating also evolved in such a way that it results in rotation of the spheroid instead of rotation of the cell, because even in the multicellular species, the simple eyespot of each cell, a primitive visual system, is used like radar to scan the environment for light sources.

The evolutionary adjustment of flagellar beating direction in colonial and multicellular volvocine species also coincided with the invention of organismal polarity and an anterior–posterior axis, which is defined by the direction of swimming (Fig. 1). The colonial and multicellular algae swim in a posterior-to-anterior direction because their flagella beat in an anterior-to-posterior direction. In volvocine algae with increased organisal complexity, there is also a gradient in cell size along the anterior–posterior axis, with the largest cells at the anterior pole and the smallest at the posterior pole (Kirk 1998; Ueki et al. 2010). The cell-to-cell distances at the anterior pole are also larger than at the posterior pole. In volvocine species with a germ–soma division of labor, the reproductive cells are normally located within the posterior hemisphere. Another sign of organismal polarity is a gradient in eyespot size, with cells at the anterior pole having the largest and most light-sensitive eyespots; at the posterior pole, the eyespots are tiny or absent, making the corresponding cells appear to be blind (Ueki et al. 2010). The differences in eyespot size produce a photoresponsive gradient decreasing from the anterior to the posterior pole of the spheroid, with the highest responses at the anterior and no response at the posterior. The differences in cell and eyespot sizes reflect functional differences between previously identical cells and thus the evolution of specialization.

The transition to differentiated multicellularity also greatly affected reproductive traits. In *C. reinhardtii* (Figs. 1, 4) and other less-complex volvocine algae like *G. pectorale* and *P. morum* (Fig. 1), there is only one cell type. For asexual reproduction, the biflagellate cells grow,
redifferentiate as nonmotile cells, and then undergo multiple fission to form daughter colonies. Once all cells of these daughter colonies have developed flagella, the daughters hatch and swim away.

In volvocine algae with an intermediate level of organismal complexity namely, certain species and strains of *Eudorina*, a few of the cells at the anterior pole of the embryo retain their flagella and fail to divide. In other words, the first sterile somatic cells emerged in evolution of this lineage (Fig. 1). The formation of sterile somatic cells is also graded along the anterior–posterior axis, with the sterile cells at the anterior pole (Kirk 2005).

The first clear sign of cell differentiation and germ–soma dichotomy in the volvocine lineage becomes apparent in *P. californica* (Fig. 1; Gerisch 1959). This alga shows a partial germ–soma division of labor: all cells initially differentiate into biflagellate cells, but then only cells of the posterior hemisphere redifferentiate as gonidia, while the cells of the anterior hemisphere remain as sterile somatic cells and eventually die.

In *V. carteri*, cell differentiation and germ–soma dichotomy have been fully established, i.e., this alga shows a complete germ–soma division of labor between ~2,000 somatic cells and ~16 germline cells (Fig. 1). During embryogenesis of volvocine species with a complete germ–soma division of labor, some of the cleavage divisions are asymmetric, producing large–small sister-cell pairs (Fig. 1) (Kirk 1995, 1998). In *V. carteri*, asymmetric cell division is accompanied by a bifurcation of the cell division program (Fig. 1; Kirk 2005): *V. carteri* embryos first cleave symmetrically five times to form a 32-cell embryo with identical cells, and then 16 cells divide asymmetrically to produce one large gonidal cell initial and one small somatic cell initial each (Kirk and Kirk 2004). These gonidial initials divide asymmetrically two more times and produce additional somatic initials at each division. The gonidial initials then temporarily stop any cleavage activity, while the somatic initials divide symmetrically about three more times. At the end of embryogenesis in *V. carteri*, the volume of the gonidal initials is about 30-fold larger than that of the somatic initials (Figs. 1, 7); however, the cells are only different in size. Then, by an as-yet-unknown mechanism, the size of each sister cell leads to the activation of either a somatic or germline program (Fig. 7) (Kirk et al. 1993). Thus, small cells develop as biflagellate somatic cells for ECM biosynthesis, motility, and phototaxis, and large cells develop as nonmotile, germline gonidia specialized for growth and reproduction.

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**Fig. 6** The switch from asexual to sexual reproduction in *V. carteri*. Asexual development (see Fig. 5) and the phenotypes of males and females are indistinguishable. **a** Photomicrograph of an asexual male. **b** Photomicrograph of an asexual female. When the sex inducer is present, the gonidia of both sexes undergo a modified embryogenesis, which is also different in males and females. **c** In males, embryonic cleavage divisions (including asymmetric cell division) result in juveniles with somatic cells and sperm packets in a 1:1 ratio. **d** In females, juveniles with ~32 eggs arise. Mature sperm packets (e) are released and swim around until they contact a sexual female with mature eggs. The sperm packets then break up into individual sperm, and the sperm penetrate the ECM of the female and fertilize the eggs (f). **g** The resulting diploid zygotes develop an orange coloration and secrete a thick, crenellated cell wall. When the zygote germinates, it undergoes meiosis to produce either a single haploid asexual male or female.
(Figs. 1, 5, 6a, b, 7; Kirk et al. 1993). The fact that the gonidia develop directly without first developing functional flagella is the criterion for “complete” germ–soma division of labor.

Mutational analysis of the genetic program for germ–soma differentiation in V. carteri identified three types of key genes: gonidialess A (glsA), regenerator A (regA), and late gonidia (lag) genes. Based on the mutational and molecular genetic results, a minimal model for the genetic program of germ–soma differentiation was established by David Kirk (Fig. 7; Kirk 1998, 2001, 2005; Kirk and Kirk 2004). In this model, the glsA gene acts during cleavage to permit asymmetric division and production of the large–small sister-cell pairs (Miller and Kirk 1999). In the small (somatic) cells, the regA gene acts as a transcriptional repressor to prevent all aspects of reproductive development (e.g., by repressing chloroplast biogenesis), while somatic genes are transcribed (Meissner et al. 1999; Kirk et al. 1999). The lag genes act as transcriptional repressors in the large cells (gonidia) to prevent the formation of somatic characteristics such as flagella and eyespots, while gonidial genes are transcribed (Kirk 1998). It is likely that the difference in cell size at the end of cleavage determines whether regA or the lag genes will be activated. However, it remains unclear how cell size is signaled to the regulatory program, which other components are involved and how the identified key genes fit into a larger regulatory network.

Several paralogous genes that encode proteins resembling regA have been identified both in V. carteri [e.g., the regA-like sequence D (rlsD) gene] and C. reinhardtii [e.g., RLS1] (Kirk et al. 1999; Duncan et al. 2006, 2007). However, Chlamydomonas does not have a gene that is orthologous to regA (Duncan et al. 2007). The closest homolog of Volvox regA in C. reinhardtii is RLS1, but the orthologous gene of C. reinhardtii RLS1 in Volvox is rlsD (Nedelcu 2009). These findings might be an indication of gene duplications in the common unicellular ancestor of Volvox and Chlamydomonas and then divergence of one of the gene copies, regA. It was speculated that the ancestor of regA might have acted in a stress-activated pathway that led to the repression of growth and cell division in response to energy or nutrient deprivation (Nedelcu 2009; Miller 2010). The paralogous genes of regA both in Volvox and Chlamydomonas might still have this function, while the function of regA might have changed and then was used to co-opt the entire pathway to repress growth and division in a developmental context (Nedelcu 2009; Miller 2010). In contrast to the regA gene, the glsA gene of V. carteri has an orthologous gene in C. reinhardtii, GARI (Cheng et al. 2003). GARI was shown to function just like glsA: When transformed into Volvox glsA mutants, it rescued the wild-type phenotype (Cheng et al. 2003). In the evolution of the volvocine lineage, glsA obviously was adopted for its novel function in germ–soma differentiation with no significant changes.

**Key evolutionary innovations in the sexual life cycle**

In the case of C. reinhardtii, the usual trigger for the switch to the sexual mode of reproduction is a lack of nitrogen (Sager and Granick 1954), indicating nutrient deficiency; in the case of V. carteri, the transition can be triggered by heat shock (Kirk and Kirk 1986), indicating drying up of the pond. The sexual cycle produces heavily walled, dormant zygotes (zygospores) that can resist tough conditions like drought, heat, and cold for a long period of time.
Volvocine algae are haplonts, and the only diploid stage of development is the zygote.

In the unicellular alga *C. reinhardtii*, haploid vegetative cells of both sexes, the plus and minus mating types, differentiate into gametes in response to nitrogen starvation, and the isogamous gametes of opposite mating types can agglutinate and fuse to form zygotes (Harris et al. 2009). The zygote further matures into a dormant zygospore. At the time of spore germination, meiosis occurs to form four haploid progeny cells, each of which is able to enter the asexual life cycle (Fig. 4).

In the multicellular alga *V. carteri*, heat shock causes the production and release of a sex-inducer glycoprotein (Starr 1970; Starr and Jaenicke 1974; Kirk and Kirk 1986; Tschochner et al. 1987; Mages et al. 1988), which is one of the most potent biological effector molecules known because it can act at concentrations as low as 10⁻¹⁶ M (Gilles et al. 1984; Starr 1970; Sumper et al. 1993). Reproductive cells of asexually grown individuals (Fig. 6a, b) that have been exposed to the sex-inducer switch to the sexual mode (Hallmann et al. 1998). In sexually induced female embryos, the first asymmetrical cell division is postponed from the sixth to the seventh division cycle (Starr 1969, 1970; Hallmann et al. 1998). After the asymmetric cell division, the somatic cell initials undergo further cleavage (as in asexual embryos). The large reproductive cell initials develop into eggs, such that the sexual female ends up with ~32 eggs and ~2,000 somatic cells (Fig. 6d), compared to ~16 gonidia and ~2,000 somatic cells in asexual females (Fig. 6b). Sexually induced male embryos exhibit yet another pattern of asymmetric division (Starr 1969, 1970; Hallmann et al. 1998): the asymmetric cell division is postponed from the sixth to the eighth division cycle, and thereafter, the somatic cell initials no longer divide. Then about a day later, the large reproductive cell initials each undergo a new round of up to seven symmetric divisions, to form sperm packets containing up to 128 sperm. The sexual male thus ends up with 128 sperm packets and 128 somatic cells (Fig. 6c), compared with ~16 gonidia and ~2,000 somatic cells in asexual males (Fig. 6a) (Starr 1969, 1970; Hallmann et al. 1998). Eventually, the sperm packets, consisting of biflagellate sperm (Fig. 6e), are released from the spheroid. As soon as a swimming sperm packet makes contact with a sexual female, which seems to happen by chance rather than by chemotaxis (Kirk 1998), the packets break up into individual sperm, and the sperm penetrate the ECM of the female to reach the eggs inside the sphere (Fig. 6f) and fertilize them. Following fertilization, the resulting diploid zygotes mature and build tough cell walls (Fig. 6g). After a rest period, favorable conditions cause the zygotes to undergo meiosis and germination to form only a single viable germling and three nonviable small polar bodies (Starr 1975). The germling will produce a haploid female or male, depending on its genetic sex, which then reproduces asexually.

The sexual dimorphism, i.e., male–female dichotomy and anisogamy (oogamy), observed in *V. carteri* (Fig. 6) and many other volvocine species with an increased level of organismal complexity (Fig. 1) evolved from a *Chlamydomonas*-like, isogamous unicellular ancestor. Phylogenetic analyses indicate that either oogamy evolved twice or it evolved once and was lost once in the *Eudorina-Pleodorina-Volvox* lineage (Kirk 2006). Oogamy also evolved independently in almost all the lineages of eukaryotes that evolved multicellularity, e.g., in other Chlorophyta (green algae), Embryophyta (land plants), Rhodophyta (red algae), Phaeophyceae (brown algae), fungi, and animals (Kirk 2006). However, except for one lineage, there are no molecular genetic data relating the sex-determining loci of oogamous organisms to the mating types of their isogamous ancestors. The only lineage for which we have such data is the volvocine lineage.

Sexual development in *V. carteri* and *C. reinhardtii* is controlled by a large, multigenic, haploid sex-determining locus (mating-type locus) that is located on homologous chromosomes in the two species and that segregates as a single Mendelian trait in each. In *C. reinhardtii*, the mating-type locus is a 200- to 300-kb region within which the order of genes is rearranged between sexes (plus and minus mating type); thus, meiotic recombination is suppressed in this region (Ferris et al. 2002; Merchant et al. 2007). Relative to *C. reinhardtii*, the sex-determining locus of *V. carteri* has undergone a remarkable divergence in both sexes and has expanded fivefold (Kianianmomeni et al. 2008; Ferris et al. 2010; Prochnik et al. 2010).

Only the sex-specific genes *MID* (Ferris and Goodenough 1997) and *MTD1* (Lin and Goodenough 2007) from *C. reinhardtii* have clear homologs in *V. carteri*, namely in the male mating-type locus (Nozaki et al. 2006; Ferris et al. 2010). The *MID* gene of *C. reinhardtii* was shown to be both necessary and sufficient to cause cells to differentiate as gametes of the minus mating type (Ferris and Goodenough 1997). Thus, the MID homologs indicate that maleness in the oogamous *V. carteri* might have evolved from the minus mating type of its isogamous unicellular ancestor (Nozaki et al. 2006; Nozaki 2008).

Further analysis of the mating-type locus of *V. carteri* identified several male- and female-specific genes without detectable homologs in *C. reinhardtii* (Ferris et al. 2010), but their functions remain to be investigated.

A key protein in sexual development seems to be the gender-specific retinoblastoma-related protein *RBR1* (Kianianmomeni et al. 2008; Hallmann 2009a). The *RBR1* gene maps to the sex-determining locus and exhibits divergent evolution in females and males; in addition,
several splice variants of the RBR1 transcript exist (Kianianmomeni et al. 2008). An RBR1-related homolog exists not only in C. reinhardtii, where it is called MAT3 (Gillham et al. 1987; Armbrust et al. 1995; Umen and Goodenough 2001), but also in almost all eukaryotes (Hallmann 2009b), which highlights the importance of RBR1.

The information about the function of the genes described above is promising, but clearly more work is required to understand the evolution and functionality of the male–female dichotomy at a molecular level.

**Conclusions and prospects**

The gradual evolutionary transition in volvocine algae from unicellular species to multicellular species with a full division of labor between germline and somatic cells and with male–female dichotomy not only exemplifies the evolution of cellular cooperation from cellular autonomy but is also a prime example of Darwin’s notion that complex traits arise via evolution by small steps (Darwin 1872). However, the observed disappearance of innovations in the evolution of volvocine algae also indicates that evolution is not a one-way street from simple to complex organisms.

On the basis of the above-mentioned traits characteristic of volvocine algae, and taking into account previous considerations by Hoops (1997), Herron and Michod (2008) and Kirk (2005), the transition to differentiated multicellularity in asexual volvocine algae encompasses the following nine developments:

1. an increase in the number of cleavage divisions and an incomplete cytokinesis that generates a cytoplasmic bridge network between cells;
2. inversion of the (spheroidal) cell sheet to correct a maladaptive arrangement produced during cleavage; this inversion is the solution to a special problem found only in volvocine embryos;
3. development and expansion of a multifunctional ECM, which displays a dynamic and multifunctional interface; the ECM holds cells in the appropriate places on the surface of the spheroid, and developmentally controlled ECM-specific enzymes allow for ECM remodeling during ontogenesis;
4. modification of the breaststroke-type swimming motion of unicellular species to a beating mode in which the effective strokes of both flagella of each cell beat in the same direction, which is accomplished by rotation of the basal bodies;
5. modification of flagellar beating direction in such a way that it causes rotation of the spheroid, which is required for phototaxis, rather than rotation of the cell;
6. invention of organismal polarity with gradients in cell size, cell-to-cell distance, eyespot size, and light sensitivity along the anterior–posterior axis;
7. invention of partial germ–soma division of labor; all cells first differentiate into motile biflagellate (somatic-like) cells and then later at least some cells redifferentiate and turn into nonmotile gonidia, while all the other cells remain as motile biflagellate cells and eventually die;
8. evolution of full germ–soma division of labor between somatic and germline cells (germ–soma dichotomy); gonidia differentiate without first developing functional flagella and contributing to the motility of the organism; all the other cells develop as sterile somatic cells and eventually die;
9. evolution of asymmetric cell division producing large and small daughter cells and bifurcation of the cell division program.

The stepwise transition to multicellularity with differentiated cell types in volvocine algae likely has parallels in the transitions to multicellularity that occurred in other eukaryotic lineages (Fig. 2). The frequent and convergent evolution of differentiated multicellularity seems to indicate that it can be achieved relatively easily. This speculation is corroborated by a recent mathematical model, which shows that complete germ–soma differentiation can be achieved easily and quickly (within a million generations) via the evolution of developmental plasticity (Gavrilets 2010).

Volvocine algae that exhibit a partial or complete division of labor in the asexual cycle exhibit a similar division of labor in the sexual cycle. Thus, for example, in *V. carteri* asexual gonidia, eggs and sperm-forming cells are all set apart from somatic cells by asymmetric cleavage divisions, whereas *Volvox* species that form gonidia without benefit of asymmetric division form prospective eggs and sperm-forming cells by similar mechanisms. Sex seems to provide the following advantages for volvocine algae (Maynard Smith 1978; Dawes 1981; Coleman 1983; Bernstein et al. 1985; Kirk and Kirk 1986; Goodenough et al. 1995; Kirk 1998; Burt 2000; Cavalier-Smith 2002; Colegrave et al. 2002; Colegrave 2002; Kaltz and Bell 2002; Hallmann 2003b):

a. survival during life-threatening conditions by the production of stress-resistant zygotes (zygospores);
b. creation of genetic variation, which is required for evolution;
c. recombinational DNA repair during meiosis because homologous chromosomes pair at that time.

Like differentiated multicellularity in asexual reproduction, different cell types in sexual males and females...
with male–female dichotomy were probably also achieved relatively easily in evolution. In the first phase of the sexual life cycle of *V. carteri* females, only the first asymmetrical cell division had to be postponed from the sixth to the seventh division cycle. The large reproductive cell initials develop into haploid eggs, which resemble haploid gonidia, albeit somewhat smaller; in fact, after a waiting period, unfertilized eggs can develop into gonidia and enter the asexual cycle. Likewise, in the first phase of the sexual life cycle of *V. carteri* males, only the asymmetrical cell division had to be postponed from the sixth to the eighth division cycle. However, these large reproductive cell initials later undergo six additional symmetric division cycles to form small haploid biflagellate sperm cells. Such traits and cleavage programs related to fertilization and reproduction are known to evolve extremely rapidly (Rice 1998; Palumbi 1998; Howard et al. 1998; Gavrilets 2000). After fertilization, the formation of dormant, heavily walled, diploid zygotes/zygospores in multicellular species is quite similar to that observed in unicellular species.

Taken together, the evolution of multicellularity, development of sterile somatic cells, and generation of a male–female dichotomy are clearly among life’s greatest innovations. However, we learn from volvocine algae that the transition to such complex life is probably much easier to achieve than might be commonly believed. *Volvox* and its relatives seem to be ideal model organisms for addressing fundamental issues in the evolution of organismic complexity and for discovering universal rules that characterize this transition.

The post-genomic era is just beginning for *Chlamydomonas* and *Volvox*. Future comparative genomics studies of gene expression and gene regulation in volvocine algae of different organismic complexities promise further insights into how the ancestral genes of unicellular species have been changed and co-opted to play modified roles and allow for the development of colonial and multicellular descendants.

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