Evolution of Individuality: A Case Study in the Volvocine Green Algae

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While numerous criteria have been proposed in definitions of biological individuality, it is not clear whether these criteria reflect the evolutionary processes that underlie transitions in individuality. We consider the evolution of individuality during the transition from unicellular to multicellular life. We assume that "individuality" (however it is defined) has changed in the volvocine green algae lineage during the transition from single cells, to simple multicellular colonies with four to one hundred cells, to more complex multicellular organisms with thousands of differentiated cells. We map traits associated with the various proposed individuality criteria onto volvocine algae species thought to be similar to ancestral forms arising during this transition in individuality. We find that the fulfillment of some criteria, such as genetic homogeneity and genetic uniqueness, do not change across species, while traits underpinning other aspects of individuality, including degrees of integration, group-level fitness and adaptation, and group indivisibility, change dramatically. We observe that different kinds of individuals likely exist at different levels of organization (cell and group) in the same species of algae. Future research should focus on the causes and consequences of variation in individuality.

Keywords

volvocine algae • multicellularity • individuality • evolution of individuality • multilevel selection • division of labor

1 Introduction

All disciplines must define their basic units and core processes. In evolutionary biology, the core process is natural selection and the basic unit of selection and adaptation is the individual. To operationalize the theory of natural selection we must count individuals, as they are the bearers of fitness. While canonical individuals have often been taken to be multicellular organisms

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(Jeuken 1952; Hull 1976, 1980; Tauber 1991; Pepper and Herron 2008), the hierarchy of life (e.g., simple prokaryotic cells, eukaryotic cells, multicellular organisms, and eusocial groups) shows that new kinds of individuals have evolved (Buss 1987; Maynard Smith 1988; Maynard Smith and Szathmáry 1995; Michod and Roze 1997). A variety of criteria have been used to define biological individuality. Some criteria rely on the presence/absence of a particular property while others advocate an approach that reflects the process of natural selection. The plethora of approaches to classifying individuality has resulted in confusion regarding how to study individuality in a given taxon (Pepper and Herron 2008).

We investigate how the various criteria proposed for delineating individuality map onto a well-studied case of the evolution of multicellular individuality in the volvocine green algae. We assume that "individuality" (however it is defined) has changed in the volvocine algae lineage from being a property of the cell to being a property of the multicellular group. We begin by briefly presenting the various criteria for individuality and some of the issues that can arise in applying them. We then use the volvocine green algae to examine which traits have changed during the evolution of multicellularity and how these traits relate to and satisfy the various criteria used to define individuality (Table 1). We find that the traits associated with some criteria, such as genetic homogeneity and genetic uniqueness, probably do not change much in this lineage, while others change dramatically, including degrees of integration, group-level fitness and adaptation, and group indivisibility. We discuss how differences in the fulfillment of these criteria between genera inform our understanding of individuality and the inherently multi-level nature of the process of natural selection.

2 Individuality Criteria

2.1 Genetic homogeneity

Genetic homogeneity requires that parts within the individual have the same genotype. For a multicellular organism, this can be achieved via a unicellular reproductive bottleneck (Weismann 1904; Simpson et al. 1957; Dawkins 1982; Crow 1988; Maynard Smith 1988; Grosberg and Strathmann 1998). Genetic homogeneity inhibits the potential for lower-level competition via the reduction of genetic variation amongst cells (Buss 1985, 1987). Issues with this definition arise for example, when mutations occur during development in a multicellular organism. In many plants, somatic mutations accumulate between cell lineages (Whitham and Slobodchikoff 1981). Somatic mutations have also been reported in fungi (Peabody et al. 2000; Stenlid 2000), red algae (Santelices and Varela 1993; Meneses et al. 1999; Santelices et al. 1999), invertebrates (Rinkevich and Weissman 1987; Barki et al. 2002; Sommerfeldt et al. 2003), and vertebrates (De 2011).

2.2 Genetic uniqueness

Genetic uniqueness is the possession of a genotype different from that of other individuals and is primarily conferred by sexual reproduction. The amount of genetic uniqueness created by sex will depend on the mating system and degree of inbreeding. This definition clearly distinguishes an individual from other individuals of the same species (Weismann 1904; Huxley 1912; Santelices 1999) and facilitates selection amongst colonies (Buss 1985). In this view, organisms or ramets produced asexually from the same clone are not individuals, but instead represent the growth or expansion of a single biological individual, the genet. This definition is difficult to relate to facultatively sexual species (where two ramets may or may not be different individuals

Criteria of	Chlamydomonas	Tetrahaana	Conium	Dandorina	Fudorina	Plandoring	Volvov
Genetic	Facultatively sexual	Facultatively sexual	Facultatively sexual	Facultatively sexual	Facultatively sexual	Facultatively sexual	Facultatively sexual
Genetic	Dependent on details of	Unicellular hottleneck 2–4	Unicellular hottleneck: 8–16 cells	Unicellular bottleneck, 16–32 cells	Unicellular bottleneck, 32–48 cells	Unicellular bottleneck, 32–128 cells	Unicellular bottleneck, thousands of cells
Division of	formation Undifferentiated	cells Undifferentiated	Undifferentiated	Undifferentiated	Undifferentiated	Somatic differentiation	Somatic and germ
Indivisibility	Cells regularly leave the group	Cells leave the group in stressful	Cells leave the group in stressful	Cells do not leave the group	Cells do not leave the group	Cells do not leave group and some cells specialize	Cells do not leave group and all cells specialize in
		body rotation may reduce cell fitness outside the group context	rotation may reduce cell fitness outside the group context			the group	group
Physiological unity and integration	No clear examples, possibly secretion of ECM	Cytoplasmic bridges, basal body rotation, cell	Cytoplasmic bridges, basal body rotation, cell number control,	Cytoplasmic bridges, basal body rotation, cell number control,	Cytoplasmic bridges, basal body rotation, cell number control,	Cytoplasmic bridges, basal body rotation, cell number control,	Cytoplasmic bridges, basal body rotation, cell number control, organismal
C		number control	organismal polarity	organismal polarity	organismal polarity, increased ECM	organismal polarity, increased ECM, somatic	polarity, increased ECM, somatic cells, germ cells
Spatial/ temporal boundaries	Spatial boundaries do not persist long	ECM, single layer colony wall	ECM, single layer colony wall	ECM, multilayer colony wall	ECM, multilayer colony wall	ECM, multilayer colony wall	ECM, multilayer colony wall
Group-level adaptations	No clear examples	Cytoplasmic bridges, basal body rotation, cell number control	Cytoplasmic bridges, basal body rotation, cell number control, organismal polarity, inversion	Cytoplasmic bridges, basal body rotation, cell number control, organismal polarity, inversion	Cytoplasmic bridges, basal body rotation, cell number control, organismal polarity, inversion, increased ECM	Cytoplasmic bridges, basal body rotation, cell number control, organismal polarity, inversion, increased ECM, somatic cells	Cytoplasmic bridges, basal body rotation, cell number control, organismal polarity, inversion, increased ECM, somatic cells, germ cells
Multilevel Selection 2	Cell lineages exist	Cell lineages possible	Cell lineages possible	Group lineages only	Group lineages only	Group lineages only	Group lineages only
(MLS2)	Group fecundity additive, group survival likely emergent; depends on few colony traits	Group fecundity additive, group survival likely emergent; depends on few colony traits	Group fecundity additive, group survival likely emergent; depends on more colony traits	Group fecundity additive, group survival likely emergent; depends on more colony traits	Group fecundity additive, group survival likely emergent; depends on more	Both components are emergent, fitness depends on many colony traits	Both components are emergent, fitness depends on many colony traits

depending on which genet they belong to) (Huxley 1912; Janzen 1977). The example of genetic heterogeneity accumulating through somatic mutations raises issues with this definition as well (Stoner et al. 1999).

2.3 Reproductive division of labor

Another criterion is reproductive division of labor (Weismann 1885; Buss 1987), in which lowerlevel units specialize on the basic fitness components of the group (Maynard Smith and Szathmáry 1995; Michod 2005, 2006, 2007). Reproductive division of labor promotes cooperation and inhibits conflict amongst lower-level entities. By specializing on survival or reproduction of the group, lower-level units, such as cells, give up their own fitness were they to live outside of the context of the group (Michod 2007). At the same time, fitness emerges at the group level and a change in the Darwinian unit of fitness and adaptation from one level to another has occurred. In the evolution of multicellularity, division of labor often takes the form of germ-soma differentiation, in which cells specialize in either group survival or group reproduction. The sequestration of a germ line also reduces the per-generation mutation rate (Simpson et al. 1957; Buss 1987; Michod 1996), thereby reducing the potential for cell-level conflict. Due to the ubiquity and importance of germ-soma division of labor (Simpson 2012), some multicellular groups without this trait have been considered to lack individuality (Godfrey-Smith 2009). Reproductive division of labor simultaneously increases group-level fitness and decouples the group-level fitness from the average of lower-level fitnesses (Michod 2006; Michod et al. 2006; Hanschen et al. 2015). While the evolution of division of labor is an especially clear example of how fitness decoupling may occur, it is not the only means. Any trait that is costly at the lower level but beneficial at the group level enhances the fitness of the group at the expense of lower-level fitness and may therefore contribute to fitness decoupling and the emergence of indivisibility of the group. In addition, traits that evolve in the context of the group may no longer be optimal were cells to leave the group (Shelton and Michod 2014). This context-dependent evolution increases the fitness of the group and decreases the fitness of cells were they to leave the group.

2.4 Indivisibility

The Latin root of the word "individual" means indivisible (Table 1). Under this criterion, an individual cannot be divided into smaller units that still maintain the properties of the whole (Michod 1999, 2007). Indivisibility facilitates selection at the group level and inhibits lower-level selection. Groups that are indivisible can be subjected to novel and different selection pressures. If a group is divisible, lower-level units could leave the group and form lineages, allowing for heritable variation in fitness and selection at the cell level. Fitness decoupling, discussed above in regards to division of labor, also leads to indivisibility as cells cannot well survive and reproduce outside of the group (Michod 2007). A group is indivisibility may also arise when cells possess properties in the context of the group that would make them less fit were they to leave the group (Shelton and Michod 2014).

2.5 Autonomy and physiological unity

Huxley (1932) considered individuality to be characterized by autonomy and physiological unity (Table 1), which imply that the individual interacts as a whole with the external environment. Integration of lower-level component parts is closely related to physiological unity (Sober and Wilson 1994; Michod 2007). The definition of an individual as possessing physiological unity

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and integration is meant to reflect the interaction between parts of an individual, increasing the likelihood that heritable variation in fitness will exist amongst groups and not the parts (Clarke 2013). This interaction among parts can be thought of as affecting the fitness of the whole. Identifying specific traits and characters which constitute physiological unity and integration can be challenging, especially as parts of an individual may demonstrate substantial internal unity and integration themselves (Clarke 2010). For example, a cell of a multicellular organism displays unity and integration between its sub-cellular component parts (Clarke 2010). Similarly, the rapid and high level of coordination amongst European starlings doesn't necessarily make a murmuration an individual (Sterelny and Griffiths 1999).

2.6 Spatial and temporal boundaries

Other individuality criteria focus on the discrete and integrated nature of individuals (Table 1). Individuals have clear spatial and temporal boundaries and are localized in both time and space (Hull 1976, 1980; Buss 1987; Mishler and Brandon 1987). The criterion of being spatiotemporally localized emphasizes that selection acts upon discrete units (Hull 1976) and the spatial/temporal organization of integrated parts which comprise that individual (Hull 1980). Spatial/temporal boundaries increase the likelihood that heritable variation in fitness exists between groups, with selection acting on these autonomous groups (Clarke 2013).

2.7 Group-level adaptations

The presence of group-level adaptations is another criterion for individuality (Vrba 1984; Folse and Roughgarden 2010; Shelton and Michod 2014). This criterion captures the expected outcome of Darwinian evolution at the group level (Lewontin 1970). The criterion of grouplevel adaptations specifically excludes cross-level byproducts, which occur when lower-level and group-level traits covary resulting in apparent selection at the group level (Okasha 2006). Nevertheless, it remains a challenge to identify what is an adaptation at the group level (Clarke 2010). Determining whether group properties are aggregative or emergent properties of cells and understanding the relationship between group-level traits and fitness at the cell and group levels may help identify group-level adaptations (Damuth and Heisler 1988; Lloyd and Gould 1993; Wimsatt 1997). Shelton and Michod (2014) have proposed testable predictions, based on an explicit mathematical model, which help to determine whether traits are group-level adaptations including strong group-level fitness effects.

2.8 Multilevel selection

Finally, the criteria used to understand different paradigm cases of multilevel selection (so-called MLS1 and MLS2 [Heisler and Damuth 1987; Damuth and Heisler 1988]) have been used to understand whether individuality exists at the group or the lower level (Okasha 2006; Shelton and Michod 2009). Do the groups themselves form lineages or are lineages just properties of the units that make up the group? How is group fitness composed out of cell fitness? Is the fitness of the group an average or aggregate property of the fitnesses of the cells comprising the group or is group fitness a non-linear or even discontinuous function of cell properties? Okasha (2006) argues the transition from MLS1 to MLS2 may be thought of as a defining property of individuality at the group level. Initially group-level fitness is defined as the average of lower-level fitness (MLS1) as in the trait-group model of Wilson (1975). As the groups become more integrated, group lineages arise and selection transitions from MLS1 to MLS2 (Okasha 2006),

and individuality emerges at the group level. In this view, once MLS2 is relevant, groups form lineages and group fitness is a non-aggregative function of cell properties.

3 The Volvocine Green Algae

Multicellularity has evolved at least 25 times across the tree of life including in bacteria, archaea, and eukaryotes (Bonner 1998; Grosberg and Strathmann 2007). In the Chlorophyte green algae, one finds the volvocine algae, a particularly tractable model system for studying the evolution of multicellularity (Figure 1). First described by Antony van Leeuwenhoek (1700), the volvocine green algae are an informal clade of approximately 90 species that includes unicellular, undifferentiated multicellular, and differentiated multicellular species. The volvocine algae include numerous phenotypes along this gradient of complexity, making them an ideal model system for studying the evolution of multicellular individuality.

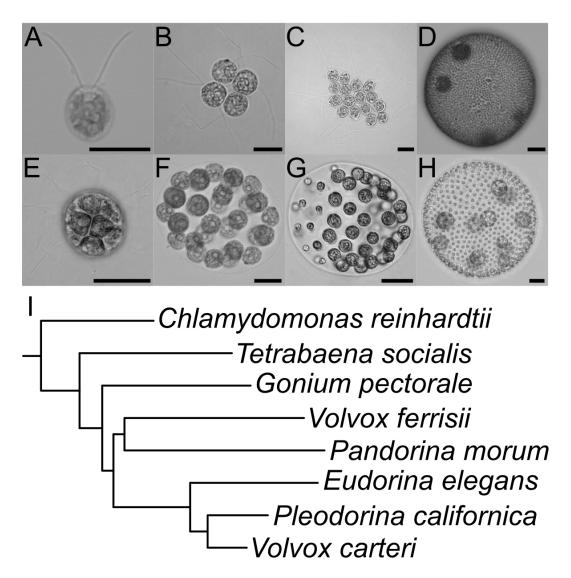


Figure 1: Micrographs of select volvocine genera (same order as in I.); A. *Chlamydomonas* (scale bar, 10 µm), B. *Tetrabaena* (10 µm), C. *Gonium* (10 µm), D. *Volvox ferrisii* (50 µm), E. *Pandorina* (25 µm), F. *Eudorina* (25 µm), G. *Pleodorina* (50 µm), H. *Volvox carteri* (50 µm). I. Phylogenetic tree of select volvocine species.

3.1 Overview of the volvocine green algae

The volvocine green algae (Figure 1) live in still bodies of freshwater such as puddles and ponds, germinating from desiccation-resistant diploid spores as conditions allow (Pocock 1933a,b). In permissive ecological conditions, volvocine algae germinate to form haploid colonies which reproduce asexually. When ecological conditions deteriorate, including due to nitrogen deprivation (Sager and Granick 1954; Kates and Jones 1964; Harris 2009) and heat shock (Kirk and Kirk 1986; Kirk 1998), colonies sexually differentiate and produce desiccation-resistant diploid spores (Janet 1912). They are photosynthetic but negatively buoyant, and thus need to maintain their position in the water column through flagellated motility. Motility plays a critical role in their ecology; some species can perform daily migrations through the water column, photosynthesizing at the water's surface during the day and migrating to the bottom of the pond, presumably in search of phosphorus and other nutrients, at night (Sommer and Maciej Gliwicz 1986). Flagellar action also serves to mix the surrounding water to aid in the uptake of nutrients and removal of waste products (Short et al. 2006; Solari et al. 2006a).

Species are globally distributed (Smith 1944; Coleman 2012) and range in size and complexity from the unicellular *Chlamydomonas reinhardtii* to undifferentiated species of four (e.g., *Tetrabaena socialis*), eight (e.g., *Gonium pectorale*), sixteen (e.g., *Pandorina morum*), or thirty-two cells (e.g., *Eudorina elegans*), to soma differentiated species (e.g., *Pleodorina californica*), to the germ-soma differentiated *Volvox carteri* with thousands of cells (Figure 1). The cells of undifferentiated species resemble *Chlamydomonas*, somatic cells in *Pleodorina* and *Volvox* resemble small *Chlamydomonas* cells, and germ cells in *Volvox* resemble *Chlamydomonas* but are much larger and not flagellated (Figure 1). These extant species, with diverse levels of complexity are hypothesized to be similar to ancestors during the evolutionary transition from unicellular to multicellular individuals. The volvocine algae do not have a multicellular ancestor and have evolved undifferentiated multicellularity once (Herron and Michod 2008; Leliaert et al. 2012). Their relatively recent radiation, approximately 200 million years ago, facilitates the identification of genetic and phenotypic changes associated with the evolution of multicellularity (Kirk 2005; Herron et al. 2009; Prochnik et al. 2010; Hanschen et al. 2016).

3.2 Individuality criteria applied to the volvocine algae

We now discuss how the previously introduced individuality criteria are satisfied throughout the volvocine algae. Regarding genetic homogeneity, mutations may arise in cell lineages during development, but given the relatively small number of rounds of cell division (approximately twelve in the case of Volvox carteri) compared to other multicellular taxa such as plants or animals, colony-level genetic homogeneity is probably high throughout this lineage. Still the opportunity for within colony variation and genetic conflict should correlate with the number of rounds of cell division and colony size (Table 1). All things being equal, genetic homogeneity (and hence individuality) should decrease with colony size. Regarding genetic uniqueness, all volvocine algae are facultatively sexual, so barring mutation, asexual clones arising in a lineage from a single germinated spore are not genetically unique. The degree of genetic uniqueness for colonies derived from different germinated spores will depend on the mating system, effective population size, and degree of inbreeding. Unfortunately, little is known about these factors in the volvocine algae. Regarding spatial/temporal boundaries, except for differences in how the colony wall is formed in these species (Kirk 2005), the volvocine algae have similar spatial/temporal boundaries, suggesting this criterion remains relatively constant in this lineage (Table 1). However, as discussed below, the elaboration of the colony wall may have implications for the indivisibility criterion and whether cells can leave the group.

We turn to the remaining criteria (physiological unity and integration, indivisibility, grouplevel adaptations, division of labor, and MLS2) and will discuss them one genus at a time. We have selected genera to represent observed variation in traits that appear most relevant to the evolution of individuality. Other genera exist but, with the exception of *Astrephomene* (which has cellular differentiation but not inversion [Iyengar and Desikachary 1981; Nozaki 1983]), they do not display different or unique combinations of the traits we discuss here. We present these genera in order of increasing body size, though some of the traits we discuss here have evolved multiple times in the volvocine green algae (Herron and Michod 2008). Based on the reconstructed evolutionary history in Herron and Michod (2008), it is hypothesized that many of these genera are similar to ancestral populations, as least with respect to the major traits related to the evolution of multicellularity as identified by Kirk (2005).

3.2.1 Chlamydomonas

Unicellular Chlamydomonas reinhardtii (Figure 1) forms multicellular clusters under certain conditions. One form of multicellular clusters, non-motile palmelloid clusters, are thought to form through the adhesion of daughter cells via an extracellular matrix or by failure of daughter cells to hatch out of the mother cell wall (Harris 2009; Khona et al. 2016). These clusters typically include 2, 4, 8, or 16 cells. Palmelloid clusters can facultatively form as a result of treatment inhibiting daughter cell hatching (e.g., calcium deprivation, chelating agents, high salt concentrations) and treatment resulting in cell wall aberrations (Iwasa and Murakami 1968; Nakamura et al. 1975, 1978; Harris 2009; Khona et al. 2016). More recently, multicellular clusters of Chlamydomonas have been used to investigate ecological and evolutionary questions. Of course, caution must be taken when using laboratory experiments to inform how selection operates on Chlamydomonas in nature. Lurling and Beekman (2006) demonstrated how palmelloid clusters facultatively form as a result of rotifer predator presence. These palmelloid clusters varied from 2 to 10+ cells and appeared to form as a result of adhesion of daughter cells via an extracellular matrix. Becks et al. (2010) used a Brachionus rotifer predator, palmelloid Chlamydomonas (evolved through six months of exposure to Brachionus), and unicellular Chlamydomonas to investigate community dynamics. Palmelloid clusters varied in size from 2 to 140+ cells held together by a thin extracellular matrix. Cells in these palmelloid clusters had a significant reduction in growth rate and a significant increase in predator survivorship, suggesting a trade-off between survival and reproduction when cells are in a palmelloid cluster. Ratcliff et al. (2013) experimentally evolved obligate multicellular Chlamydomonas clusters using centrifugation to select on cluster size. Evolved multicellular Chlamydomonas clusters contained over 100 cells descended from a single cell. Clusters were held together through secretion of an extracellular matrix (Ratcliff et al. 2013). Sathe and Durand (2015) used a Euglenoid predator (*Peranema trichophorum*) to demonstrate Chlamydomonas can aggregate (in contrast to forming clusters through daughter cells failing to leave the mother wall) in clusters of 10-100,000 cells. Using Chlamydomonas reinhardtii, C. moewusii, and C. debaryana, the formation of multi-species aggregations of approximately 100 cells was observed.

Multicellular *Chlamydomonas* clusters have clear spatial boundaries, but as cells are able to leave palmelloid clusters (Khona et al. 2016) we do not consider them indivisible. Cell clusters may form via aggregation between separate genetic strains, suggesting *Chlamydomonas* clusters are not always genetically homogenous (Sathe and Durand 2015); however, in the other studies discussed above, the clusters are considered to be clonally formed and genetically homogenous. Germ-soma division of labor is not present. The case of palmelloid clusters illustrates the challenges involved in distinguishing the level of selection leading to increased physiological unity

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and the presence of true group adaptations (see Clarke 2010, 2013). In our opinion, there are no clear examples of physiological unity or group-level adaptations in palmelloid clusters, although secretion of extracellular matrix upon which cluster formation is based is a potential candidate. Alternatively, the secretion of extracellular matrix may be a cell-level adaptation which increases the inclusive fitness of secreting cells. Shelton and Michod (2014) analyzed a model of simple clusters of genetically identical cells as a way to understand when group-level adaptations emerge. The conditions under which group adaptations emerge depend on how strongly group membership affects life history variables such as mortality and growth. While the beneficial effect of group membership on lowering predation rates is consistent with group adaptations being present in *Chlamydomonas* palmelloid clusters (Shelton and Michod 2014), it is not sufficient to clearly demonstrate their presence. The example of palmelloid *Chlamydomonas* serves as a null comparison for the undifferentiated colonial species in which group-level adaptation and integration are clearly present.

3.2.2 Tetrabaena

One of the simplest and smallest colonial volvocine algae, Tetrabaena socialis, consists of four undifferentiated Chlamydomonas-like cells (Figure 1, [Janet 1912]). During reproduction, each cell in a *Tetrabaena* colony undergoes two rounds of cell division, producing a four-celled colony. These cells are attached through cytoplasmic bridges which result from incomplete cytokinesis (Arakaki et al. 2013); these bridges likely serve to keep the cells in the group. Integration is enhanced by the existence of rotational asymmetry in the arrangement and rotation of basal bodies (Table 1). These basal bodies connect flagella to the cell wall, allowing for flagellar beating. Rotational asymmetry is unique and universal (in all species where it has been studied) to colonial/multicellular species (Kirk 2005; Arakaki et al. 2013), suggesting that this rotation plays an important role in colonial motility and thus may be considered a group-level adaptation. Finally, the ability to reproduce a four-celled colony (in contrast to a single Chlamydomonas-like cell that can grow and divide many more times) indicates that the number of cells in a colony is genetically modulated in Tetrabaena (Kirk 2005; Arakaki et al. 2013), again, demonstrating the presence of group-level adaptations (Shelton and Michod 2014). However, Tetrabaena does not fulfill other criteria of individuality such as germ-soma division of labor (Table 1) and indivisibility. Tetrabaena colonies break apart during conditions of stress (Harper 1912; Starr 1955) and so the group is divisible in the sense that cells are not completely committed to group living. However, basal body rotation may negatively affect swimming ability outside the context of the group; thus, *Tetrabaena* has a degree of indivisibility in the sense that it has group level adaptations that may reduce cell fitness outside of the context of a group.

Turning to the MLS criteria, we find that lineages exist at the group level, but may also exist at the cell level when cells leave the group. To understand how group fitness relates to cell fitness and cell properties, we consider each component of fitness (reproduction and survival). In *Tetrabaena*, as all cells produce a single daughter colony, the fecundity of the group is the sum of cell fecundity. Survival of the group is more complicated and difficult to relate to cell properties. Components of survivorship, such as predation resistance (Boraas et al. 1998), swimming speed (Solari et al. 2006b), and flagellar mixing of the environment (Solari et al. 2006a), may be aggregative properties of cells and may depend on group properties such as cell number. For example, flagellar force of the group (important for both swimming and mixing of the surrounding microenvironment) could be construed to be an aggregative property generated by cells. Likewise, group size (important for predation resistance) is likely an aggregative function of cell size but requires reproducing the number of cells in the group, which is a group trait. This

group trait likely depends on cell traits such as cell growth rate and cell size at which mitotic divisions commence. Ecologically, the predation resistance of the group could be a step function of cell number (Boraas et al. 1998; Kirk 1998), if the size of the group is above a certain threshold, groups could have high survival, while if the size of the group is below the threshold, the group may have little survivorship. Solitary cells, likely below any size threshold (Kirk 1998), might have little or no survivorship. Thus, the survival of the group (and consequently fitness, the product of survival and reproduction) may not be the average of cell-level fitnesses (Okasha 2006; Shelton and Michod 2009). Increased swimming speed and flagellar mixing of the environment (Solari et al. 2006a,b) also have complex relationships to group survivorship, and may not affect group survivorship until larger colony sizes evolve. Since group-level fitness is sensitive to changes in cell number and organization, group-level fitness is likely emergent (Wimsatt 1997).

In Table 1, we assume that survivorship starts out aggregative and depends on few colony traits. As colonies get larger and more morphologically complex, survivorship is less likely to be an aggregative property of cell properties; simultaneously, the number of group traits affecting survivorship increases. When group traits are no longer aggregative functions of cell traits we refer to them as emergent in Table 1.

3.2.3 Gonium

Gonium pectorale is a small colonial species in the Goniaceae family consisting of a plate of 4-32 Chlamydomonas-like cells (Figure 1). Like Tetrabaena, Gonium develops from a unicellular bottleneck without division of labor; all cells reproduce with no differential investment at the cell level in group survival or reproduction (Table 1). However, Gonium has several traits which satisfy individuality criteria not satisfied in *Tetrabaena*. First, development in *Gonium* includes inversion. A Gonium embryo is initially shaped like a shallow bowl, with the flagella pointed in towards the center of the bowl. During development, the colony inverts, reversing the curvature of the colony, so that flagella point outward (Stein 1965). Second, Gonium has a centerperipheral axis in which cells differ in their flagellar orientation (Gerisch 1959). Third, Gonium shares many genomic modifications with germ-soma differentiated Volvox including expanded cyclin D genes (genes which regulate the cell cycle) and structural modifications to retinoblastoma (RB), a cell cycle regulation gene (Hanschen et al. 2016). Given that the modifications to *RB* cause cell colonies to form and that similar expansions of cyclin D genes exist in other multicellular taxa, these changes are likely group-level adaptations (Hanschen et al. 2016). As with Tetrabaena, Gonium colonies break apart into smaller units composed of one or more cells in times of stress (Harper 1912). These smaller units can survive and reproduce (personal observation), raising the question as to whether Gonium colonies lack indivisibility. As with Tetrabaena, the fecundity of the group is the sum of the fecundity of the cells, while the survivorship of the group may or may not be an additive property of cell survivorship traits. Lineages typically exist at the group-level except when cells leave the group (Table 1).

3.2.4 Pandorina

Pandorina morum (Figure 1) consists of 8–16 cells and repeats many of the points already made, except *Pandorina* is spherical and has a novel trait potentially relevant to individuality not found in *Tetrabaena* or *Gonium*: a multilayer colony boundary. As far as we are aware (and we have tried growing them under stress), there are no reports of *Pandorina* (or any larger volvocine species) dissociating into single cells under stress. *Pandorina* appears to be committed to group living as cells do not leave the group (Table 1), which is likely the consequence of a multilayer

colony wall (Fulton 1978; Kirk 2005). This more complex colony wall is shared with larger species, which are also committed to group living.

3.2.5 Eudorina

Species of the genus *Eudorina* form spherical colonies of 32 to 64 cells (Figure 1) that are similar to *Pandorina* and repeat many of the points made so far with the exception that *Eudorina* has an expanded extracellular matrix (ECM) (Conrad 1913; Goldstein 1964; Kirk 2005) made of glycoproteins secreted by the cells (Hallmann 2006). The ECM is composed largely of secreted glycoproteins whose production likely imposes a metabolic cost on the cells, which secrete the ECM (Sumper and Hallmann 1998). This cost is likely overcome by key group-level benefits which may include protection against the external environment (Gerhart and Kirschner 1997) and nutrient storage (Bell 1985; Koufopanou and Bell 1993). The ECM is a shared resource that each cell contributes to, providing an environment in which cheaters, which benefit from the ECM but do not directly contribute to it, could flourish (Herron and Michod 2008). The ECM is a group-level adaptation that likely imposes costs at the level of the individual cell.

3.2.6 Pleodorina

The genus *Pleodorina* forms spherical colonies with 32–128 cells (Figure 1), and meets many of the same criteria as *Eudorina*, with the important novelty that 20–50% of these cells are terminally differentiated, non-reproductive, somatic cells located in the anterior hemisphere (Shaw 1894; Nozaki et al. 2006). These somatic cells indicate division of labor involving reproductive altruism is present in *Pleodorina* (Table 1). The rest of the cells in the colony are *Chlamydomonas*-like, general-purpose, flagellated cells contributing to both survival (flagellar action) and reproduction. Only the somatic cells are specialized in these species, which are specialized in survival and give up reproduction. The fitness costs of reproductive altruism incurred by somatic cells are compensated for by greater group-level benefits that are brought about by specialized somatic cells such as increased colony motility (Solari et al. 2006b). Cellular differentiation results in differences in cell-level fecundity, some cells do not reproduce at all and have zero fecundity. This means that group-level fitness is clearly not the average of cell-level fitnesses; thus group-level fitness is strongly decoupled from cell-level fitness (Table 1). The colony is indivisible in the sense that, if a colony were broken up, the somatic cells could not produce offspring.

3.2.7 Volvox

Volvox is a polyphyletic genus and member species fulfill nearly all the individuality criteria discussed (Table 1). *Volvox* is indivisible and has complete germ-soma division of labor. For example, *Volvox carteri* contains approximately 12 reproductive germ cells, which do not develop flagella, and approximately 2,000 somatic cells, which cannot reproduce but instead specialize in motility via flagellar action. Therefore both somatic and germ cells are specialized in either group-level survival or group-level reproduction. *Volvox* species have clear group-level adaptations such as greatly expanded ECM (Powers 1908; Starr 1969, 1970); greater than 99% of the volume of a *Volvox* colony is cell-secreted ECM (Hallmann 2003). Some species of *Volvox* have 50,000 cells and a corresponding increase in number of rounds of cell division, and hence, more opportunities for mutations to arise. *Volvox* species are therefore predicted to be marginally less genetically homogeneous than smaller species, all other factors being equal.

Volvox species display similar phenotypic characteristics with one key difference: the timing of reproductive cell flagellar loss. In the group of Volvox which includes Volvox ferrisii (represent-

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ing an independent evolution of the *Volvox* body plan from *Volvox carteri*; (Figure 1, [Herron et al. 2010]), reproductive cells are not specialized as they retain their flagella early in development (Pocock 1933b; Isaka et al. 2012). Could these *Volvox* species be considered to be less integrated than *Volvox carteri* because they do not have specialized germ cells? On the other hand, species such as *V. ferrisii* maintain cytoplasmic bridges between reproductive and somatic cells throughout their life cycle (Smith 1944; Isaka et al. 2012). Conversely, cytoplasmic bridges in *V. carteri* are broken down at the end of embryogenesis (Kirk 1998). While the function of these cytoplasmic bridges is unknown, they may serve to transport nutrients and waste or facilitate intercellular communication. If so, it could be argued that *V. ferrisii* is more integrated than *V. carteri* (Godfrey-Smith 2011). If traits such as the maintenance of cytoplasmic bridges or flagella in reproductive cells affect integration, or could be considered group-level adaptations, they affect the level of individuality in *Volvox* species. However, the differences of flagellar loss in reproductive cells and cytoplasmic bridges may be negligible compared to the morphological similarities they share.

4 Discussion

4.1 Summary of individuality in the volvocine algae

Our review of how various criteria for individuality relate to the volvocine algae shows that criteria differ in how they are fulfilled across genera that are thought to span the divide between unicellular and multicellular individuals. Some criteria, such as genetic uniqueness and genetic homogeneity, do not appear to change substantially in the extent to which they are satisfied from Chlamydomonas palmelloid clusters to Volvox. For other criteria, such as physiological unity and the presence of group-level adaptations, there is a gradient across genera in the extent to which the criteria are satisfied. Traits fulfilling group-level adaptations and physiological unity and integration criteria appear to change concurrently across the volvocine algae, as most of the traits that are possible group-level adaptations also promote physiological unity and integration. In contrast, the traits fulfilling other criteria do not change concurrently; for example, the pattern of gradual change in the traits underlying physiological unity and integration is very different from the step-like evolution in division of labor. Criteria such as indivisibility, MLS2, and division of labor are satisfied at different times and stages during the evolution of multicellularity in the volvocine green algae. These three criteria critically inform how selection likely acts at different levels in the different genera and so it may be expected that their underlying traits would change as the unit of selection changes levels. Indivisibility and division of labor reduce the potential for cell-level conflict and selection while MLS2 reflects how selection operates at both levels in the process of reducing the capacity for selection at the lower level, and enhancing selection at the higher level.

Based on the differences in how selection likely acts in different volvocine genera, we identify three types of volvocine multicellular individuals: *Tetrabaena* and *Gonium*, where selection can act on both cells and colonies; *Pandorina* and *Eudorina*, where selection acts on colonies that lack the mechanisms seen in other taxa that promote cell-level cooperation and inhibit the potential for cell-level conflict; and *Pleodorina* and *Volvox*, where selection clearly operates at the colony level and mechanisms which reduce the potential for cell-level conflict exist. The notion of multiple kinds of multicellular individuals is not novel (Santelices 1999) but our approach to delineating different types of individuality is.

Many definitions of individuality focus upon traits which reduce the possibility of withingroup conflict and selection at the lower level (Maynard Smith and Szathmáry 1995; Michod 1999; Gardner and Grafen 2009). Our analysis acknowledges the context-dependency and plasticity of individuality. For example, under certain conditions, *Gonium* multicellular groups reproduce as Darwinian individuals. Yet, when environmental conditions change, the group-level individual dies and selection operates on cell-level organisms. This represents a temporary shift from MLS2 (group lineages) to MLS1 (cell lineages). When conditions change again, these cells form multicellular organisms and selection again operates at the level of the group (MLS2). As reflected by the changes in Darwinian properties at different levels in different conditions, the levels at which selection acts can change temporally. Moreover, fitness interests between levels may not be aligned. If an organism lacks mechanisms to sufficiently suppress lower-level conflict, selection acting at one level may interfere with selection acting at another level. Alternatively, fitness interests across levels could be aligned. For example, imagine a mutation which increases the strength of flagella, thus increasing swimming speed and both cell-level and group-level fitness.

4.2 A multidimensional perspective on individuality

The level of attention in the literature dedicated to the concept of individuality is born out of necessity—in order to measure fitness and changing frequencies, researchers need to count individuals. To this end, many researchers have proposed a single criteria whose fulfillment is purportedly both necessary and sufficient for individuality (but see Santelices 1999; Pepper and Herron 2008; Clarke 2011, 2013), implying that individuality is a dichotomy and individuals are considered equivalent so long as the required criterion is present. Consistent with this interpretation, individuality is often thought to be present at a single hierarchical level at a time (Dawkins 1982; Maynard Smith 1988; Michod 1999; Gardner and Grafen 2009). Other researchers argue that individuality is not a dichotomy and that there are multiple different kinds of non-equivalent individuals based on variation in genetic uniqueness, genetic homogeneity, and autonomy/physiological unity (Santelices 1999). Queller and Strassmann (2009) compared species based on qualitative differences in the extent of cooperation and conflict, and suggested that continuous variation in individuality exists. Clarke (2013) proposed that definitions of individuality should not be mechanism-specific but should instead reflect both the suppression of selection acting at the lower level and the enhancement of selection acting at the group level, also suggesting group-level individuality may exist in degrees, the extent to which selection can act on group-level units.

A dichotomous formulation of individuality, in which individuality is only present at a single level and all multicellular individuals are conceptually equivalent, is not supported by our analysis and ignores the possibility of different kinds of individuals co-occurring in the same species. The presence of genetic homogeneity in *Chlamydomonas* clusters depends on the details of cluster formation. Similarly, the temporal variability of cell lineage formation in species such as *Tetrabaena* and *Gonium* and the environmentally determined, intermittent occurrence of sexual reproduction in all volvocine species demonstrates that the level and kind of individuality can change across time and space and can be dependent on the environmental context.

The context-dependency of individuality is not restricted to the volvocine algae. For instance, black spruce (*Picea mariana*) reproduce both sexually from seeds and asexually via clonal propagation (Legere and Payette 1981). On flat terrain at low elevations, sexual reproduction prevails and every ramet represents a unique genet. However, on subalpine slopes asexual reproduction is more common, and ramets do not represent a unique genet (Viktora et al. 2011). In this case, selection is likely operating both between genets and between ramets. Both ramet and genet are individuals, the genets composed of ramets, and individuality depends on environmental context. Similarly, when cancer develops within a canonical multicellular organism, selection is occurring between multicellular organisms but also within a multicellular organism, between cell lineages (Aktipis et al. 2015). Individuality resides at both the level of cell lineages and the multicellular organism, though only when mutation causes cancerous cell lineages to arise. Lastly, in the example of somatic mutation accumulation along a tree branch, genetic variation between branches could result in increased fruit production (increased reproduction) on one branch, beneficial at both the lower (branch) and higher (tree) levels (Whitham and Slobodchikoff 1981; Pineda-Krch and Lehtilä 2004). In this example, both the tree and tree branches are considered to be simultaneous Darwinian individuals.

While individuality is often assumed to be constant across members of a given species, the context-dependency of individuality should be considered and the evolutionary consequences investigated. We suggest researchers embrace a multidimensional approach based on multiple individuality criteria to determine the kinds of individuality present in their system and how selection may be affected. This approach may provide insight into individuality in taxa of interest, as well as how evolution is subsequently impacted. Our approach has utilized a multiplicity of individuality criteria. Mechanistic, trait-based definitions of individuality were critical to inform phenomenological, selection-based definitions of individuality, providing insight to how selection might be acting in the volvocine algae. This multidimensional approach to individuality, composed of trait-based definitions, selection-based definitions, and ecological context, may be informative in future research, particularly in the context of comparative studies.

This multidimensional approach may be fruitful when applied to the evolution of individuality during the evolution of multicellularity in other lineages or other evolutionary transitions such as the evolution of eukaryotes or eusociality. This would entail determining which individuality criteria participating taxa fulfill, as was carried out here with the volvocine algae. As appropriate, the ecological context affecting individuality needs to be taken into account. This mapping can then allow researchers to ask questions about individuality, macroevolutionary trends and patterns, and the capacity to respond to selection. This approach may be fruitful for understanding how different kinds of individuals across the tree of life differ in their evolutionary dynamics, how independent evolutions of one kind of individuality are similar, and how transitions between different levels of individuality are similar.

5 Conclusions

Our approach in this paper is to understand individuality by examining how traits that have changed during the evolution of multicellularity in the volvocine green algae relate to the various individuality criteria proposed in the literature. We find that the evolution of multicellular individuality from unicellular ancestors in the volvocine green algae likely involves minor changes in genetic homogeneity, genetic uniqueness, and spatial/temporal boundaries. While necessary for the initiation of group-level selection and evolution, these criteria do not appear to change substantially in the extent they are satisfied during the evolution of individuality in this lineage. Other individuality criteria, including division of labor, indivisibility, and the presence of multilevel selection, vary dramatically in how they are fulfilled. Examining these criteria in the context of ecology suggests three kinds of multicellular individuals (*Pandorina* and *Eudorina*), and committed, differentiated multicellular individuals (*Pleodorina* and *Volvox*). We suggest that a multidimensional approach to individuality in which researchers consider multiple criteria simultaneously may provide greater evolutionary insight via a more nuanced view that may better reflect the process of natural selection.

Although the evolution of individuality is often thought of as a singular and dichotomous event, the volvocine algae show that multicellular organisms can differ in which aspects of individuality they exhibit and multiple criteria may be fulfilled at the same time. The evolutionary implications of interspecific variation in how criteria are satisfied for individuality are unclear; fulfilling some but not all criteria may have consequences yet to be appreciated. Does a certain kind of individuality inhibit or enhance subsequent evolutionary change or speciation rates, allow for greater reversibility, lower degrees of adaptation, or lower complexity of the organisms involved? What selection pressures lead to the various kinds of individuals in this lineage? These questions remain unanswered and future research should focus on the causes and consequences of variation in individuality.

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