Toward establishing model organisms for marine protists: successful transfection protocols for *Parabodo caudatus* (Kinetoplastida: Excavata)

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Running title: Protist transfection

Originality-Significance Statement: This study is the first experimental evidence comparing three different electroporation methods for transfection of microbial eukaryotes. Our recently developed microfluidic transfection technology and square wave protocols resulted in higher efficiency and reproducibility compared to the traditional exponential decay electroporation. *Parabodo caudatus* has the potential to be a genetically tractable model protist organism because it grows quickly and relatively easily in the laboratory, and its transparent cell membrane facilitates observations of measurable phenotypic traits in genetically transformed cells. *P. caudatus* is abundant in marine and freshwater environments, and is of evolutionary interest for being one of the free-living close-relatives of parasitic trypanosomes. The development of transfection protocols for *P. caudatus* will help to further understand evolution, physiology, and ecological roles of *P. caudatus*.

Key words: Transfection; electroporation; microfluidic; micro-eukaryotes; reporter genes

Summary:

We developed protocols for, and demonstrated successful transfection of, the free-living kinetoplastid flagellate *Parabodo caudatus* with three plasmids carrying a fluorescence reporter gene (pEF-GFP with the EF1 alpha promoter, pUB-GFP with Ubiquitin C promoter, and pEYFP-Mitotrap with CMV promoter). We evaluated three electroporation approaches: 1) a square-wave electroporator designed for eukaryotes, 2) a novel microfluidic transfection system employing hydrodynamically-controlled electric field waveforms, and 3) a traditional exponential decay electroporator. We found the microfluidic device provides a simple and efficient platform to quickly test a wide range of electric field parameters to find the optimal set of conditions for

electroporation of target species. It also allows for processing large sample volumes (> 10 ml) within minutes, increasing throughput 100 times over cuvettes. Fluorescence signal from the reporter gene was detected a few hours after transfection and persisted for 3 days in cells transformed by pEF-GFP and pUB-GFP plasmids and for at least 5 days post-transfection for cells transformed with pEYFP-Mitotrap. Expression of the reporter genes (GFP and YFP) was also confirmed using reverse transcription-PCR (RT-PCR). This work opens the door for further efforts with this taxon and close relatives toward establishing model systems for genome editing.

Introduction

Protists are unicellular eukaryotes that are ubiquitous in the marine realm, their molecular signatures have been described from all marine habitats investigated, and they are recognized as pivotal members of aquatic microbial communities in models of carbon cycling (Worden et al., 2015). Phototrophic protists contribute to primary production, and heterotrophic protists shape organic matter pools and populations of their prokaryotic and eukaryotic prey, thus indirectly influencing activities at the foundation of microbially-driven major nutrient cycles (Azam and Malfatti, 2007). Free-living protists exhibit complex interactions with other protists, Metazoa, Bacteria, Archaea, and viruses. Our understanding of the extent of marine protist diversity has expanded tremendously over recent decades, due in significant part to information from high throughput sequencing approaches based on bulk extracted DNA/RNA. Protists also exhibit symbioses with prokaryotes and with other protists and Metazoa. Marine protist taxa with sequenced genomes reveal that microeukaryotes can have complex and large genomes that can be many times greater in size than even the human genome. The function of a significant portion of those genes remains unknown, and is referred to as "genetic dark matter" (Clark et al., 2013).

To uncover the scientific principles that govern the interactions of protists with other microbes and that mediate nutrient flow in the sea, an understanding of the function of this genetic "dark matter" facilitated by genetically tractable model representatives is required. Their development will allow us to systematically decipher the gene–gene and gene–environment interactions, and to understand processes underlying the roles of certain protists in biogeochemical cycling and the evolution and ecology of the microbial Eukarya. Low efficiency transfection protocols exist for a few marine protists; *Ostreococcus tauri*, *Phaeodactylum tricornutum*, *Amphidinium* sp. and *Symbiodinium microadriaticum* (Te et al., 1998; De Riso et al., 2009; van

73 Ooijen et al., 2012). Transfection of two dinoflagellates has been successful using silicon carbide
74 whiskers to deliver plasmid constructs (Te et al., 1998), and transfection of *Perkinsus marinus*, a
75 marine protist parasite, has been achieved using electroporation (Fernandez-Robledo et al., 2008;
76 Cold et al., 2016). Recent gene editing protocols have been established for few diatoms (Hopes et
77 al., 2016; Liu et al., 2016), and progress toward useful forward genetics approaches for
78 choanoflagellates have been established (Hoffmeyer and Burkhardt, 2016). Additional models are
79 needed for more widely distributed, ecologically important free-living lineages. Methods of gene
80 tagging and gene silencing using CRISPR/Cas9 have been developed for a few, mainly parasitic
81 protists (*Trypanosoma cruzi*, *Leishmania* spp., *Plasmodium* spp., *Cryptosporidium parvum*,
82 *Chlamydomonas reinhardtii*) (Lander et al., 2015; Peng et al., 2015; Lander et al., 2016a; Lander
83 et al., 2016b), but again, there is a lack of methods for most protist taxa. Genetic manipulation of
84 marine protists will make it possible to link genes of unknown function to cell behavior (e.g.,
85 colony formation, morphogenesis, cell-cell interactions), physiology (e.g., life cycle and
86 reproduction type), particular biogeochemical cycles, and processes of interest, such as, nitrogen
87 and carbon cycling, and production of climate active trace gases or initiation of harmful algal
88 blooms (Gong et al., 2017).

89 The first step in developing transgenic marine protists is the establishment of reliable and
90 reproducible transfection protocols. Transfection can be accomplished using chemical-based
91 methods (which include lipofection, calcium phosphate, etc.), electroporation, microinjection,
92 biolistic, laserfection/optoinjection, and virus-based methods (Kim and Eberwine, 2010).
93 Electroporation using pulsed electric fields is a technology that has become a powerful and mature
94 tool used for genetic engineering of prokaryotes and eukaryotes. Via electroporation, electrical
95 pulses of a specified voltage and duration transiently disrupt the membrane of cells and deliver

foreign DNA (i.e., plasmids) into the cells within milliseconds. With the advent of electroporation bacterial strains could be transformed with plasmids carrying marker genes (Teissie and Tsong, 1981; Neumann et al., 1982; Josenhans et al., 1998) and methods have evolved since then to include a wide range of tools including most recently, RNAi and CRISPR/Cas9 (Peng et al., 2015; Liang et al., 2017). Such approaches can allow researchers to gain an understanding of the role of a gene of interest in the physiology or behavior of an organism, and in turn, the role the gene plays in the organism's ecology.

While electroporation methods for some bacteria, such as *E. coli*, are technically simple and thus commonplace, the efficacy of electroporation, broadly speaking, depends on many variables. The critical parameters in electroporation are the electric field magnitude and pulse characteristics such as the shape, duration, and number of pulses (Weaver et al., 2012). Traditionally, electroporation is performed in plastic cuvettes outfitted with parallel plate metal electrodes. The separation distance between the plate electrodes is fixed at 1 mm, 2 mm, or 4 mm and the DNA-cell suspension is placed between the electrodes. Upon application of a voltage, an electric field is generated between the plates at a magnitude equal to the voltage-to-distance ratio (Corovic et al., 2007). Other experimental parameters affecting the outcome include the DNA concentration and the electrical conductivity (salt concentration) of the medium surrounding the cells (i.e., electroporation buffer). Buffers with lower ionic concentrations reduce the arcing potential (electrical charges that occur due to high or excess salt concentration), reduce deleterious heating, and generally increase the transfection efficiency (Kotnik et al., 2015). Notably, eukaryotic cells (including many protists) are highly sensitive to the experimental conditions that are optimal for electroporation, such as lower ionic concentrations. The low efficiency of

electroporation resulting from the myriad of technical challenges has hindered progress in developing electroporation-based methods appropriate for aquatic protists.

Accordingly, we aimed to develop an efficient method for electroporation of marine protists. For this effort, we selected *Parabodo caudatus*, a free-living biflagellate kinetoplastid (Parabodonida, Kinetoplastea, Euglenozoa), and free-living close relative of parasitic trypanosomatid flagellates. *P. caudatus* feeds on bacteria (e.g., *Klebsiella pneumoniae*, *Escherichia coli*, *Enterobacter aerogenes*, and others). *P. caudatus* cells divide by longitudinal binary fission. Members of this family have fast growth rates and are easy to culture and maintain in laboratory. They prey on bacterial cells and are ubiquitous in many environments including fresh and marine water columns and sediments, seawater from deep-sea hydrothermal vents, and as contaminants in food (von der Heyden et al., 2004; Tikhonenkov et al., 2016). The genome of a congener of *P. caudatus*, *Bodo saltans,* revealed that 60% of all coding genes had homologs in trypanosomatids, evolutionary close relatives to kinetoplastids (Jackson et al., 2008; Jackson et al., 2016; Opperdoes et al., 2016). The remainder of genes was found to be homologs of genes in other eukaryotes (i.e., fungi, animals, and plants) but not trypanosomatids, or *Bodo*-specific genes with no matches to sequences in public databases. These *Bodo*-specific genes were predicted as hypothetical proteins expressed on the cell surface (Jackson et al., 2008; Jackson et al., 2016; Opperdoes et al., 2016). These findings suggest that *Parabodo* and other free-living bodonids represent appealing model organisms for exploring potential roles of those unidentified genes.

To determine the efficacy of electroporation as a means of transforming marine protists, we tested three different transfection technologies. First, we evaluated a commercially available square-wave technology that has been successfully used to increase transfection rates by 2-3 times in living cells ranging from very fragile mammalian stem cells (Kaneko et al., 2014) to intact

141 vertebrate embryos (when compared to traditional exponential decay electroporation; (Sanders et
142 al., 2013)). Very few studies to date, however, have examined electroporation responses of free-
143 living microeukaryotes, which are markedly different than mammalian cell lines in their robustness
144 and their transfection amenability (Miyahara et al., 2013). While the exponential decay
145 electroporation methods enable voltage, resistance, and capacitance to be independently selected,
146 the square-wave generator enables additional parameters such as pulse duration, pulse number,
147 and polarity. The square-wave system may be used to apply two types of multi-pulse
148 electroporation protocols (Fig. 1b). The first sequence uses poring pulses, which are multiple high-
149 voltage, short duration (5 ms) pulses responsible for formation of the initial pores (membrane
150 defects) in the cell membrane. The second sequence uses transfer pulses, which are multiple low-
151 voltage, long duration (50 ms) pulses that deliver the target molecules into cells with minimal
152 damage. The low-voltage pulses are similar to those used in electrophoresis, which facilitates the
153 movement of charged molecules such as DNA into the cells.

154 The second method that we evaluated utilizes a novel microfluidic platform for identifying
155 critical electroporation conditions for successful transformation, recently developed by a subset of
156 the authors (Garcia et al., 2016). We further developed our microfluidic platform to conduct
157 continuous flow transformation of microorganisms (Garcia et al., 2017). This technology uses
158 microfluidic channels with geometric constrictions (see Experimental Procedures section for
159 physical dimensions) in order to amplify the electric field to achieve electroporation (Fig. 1a). In
160 our bilaterally converging microfluidic system, a single applied voltage results in a linear electric
161 field gradient along the length of the microchannel constriction and results in a maximum ~6x
162 amplification of the applied voltage (1 V = 6 V/cm). Thus, depending on the applied voltage and
163 the location within the constriction, the strength of the electric field will be able to induce cell

electroporation. Square wave pulses are delivered from electrodes with alternating polarity between the pulses to reduce electrolytic effects at the electrode-buffer interface (Fig. S1). For example, square wave pulses with 5 ms ON and 5 ms OFF cycles are applied to the microchannel through the dispensing needle, which results in 50 % of the cells experiencing the electric field during their transit through the channel. Increasing the duty cycle would result in a larger fraction of the cells being exposed to the electric field, but also more sample heating. In order to mitigate potential deleterious heating, the flow rate must be selected carefully in order to remove the heated sample from the constriction without compromising cell viability. The microfluidic device conveniently provides a simple platform to efficiently test transfection conditions and optimize parameters for genetic manipulation of recalcitrant organisms such as many protists.

The third system we investigated was a MicroPulser Bio-Rad (CN 165-2100) exponential decay system (Fig. 1). Among electroporation technologies, exponential decay technologies have been routinely used in laboratories the longest. Here we report successful transient transfection of the free-living kinetoplastid flagellate *Parabodo caudatus* with three plasmids carrying fluorescent protein (FP) reporter genes, using three electroporation approaches: 1) a new microfluidic transfection system using hydrodynamically-controlled waveforms, 2) a square-wave transfection system, and 3) traditional exponential decay electroporation. This study is the first experimental comparison of successful transient transfection of marine microeukaryotes employing three different electroporation methods. It lays the groundwork for future efforts aimed at stable transfection with a variety of gene targets, and genetic manipulation of this taxon and its close relatives.

Results

Cell Viability After Cytomix Buffer Incubation

Tolerance of *P. caudatus* to different electroporation buffers was tested. *P. caudatus* cells were viable after incubation in 50 % and 10 % cytomix (Knight and Scrutton, 1986) buffer concentrations for at least 15 minutes. However, *P. caudatus* incubated in 100 % cytomix were viable for about 10 minutes, after which increasing cell mortality was observed. Therefore, all our subsequent experiments were performed in 50 % cytomix buffer, 10% or 1% seawater, or MilliQ water in order to maintain sufficient cell viability.

Electroporation Buffer Conductivity

The exponential decay (MicroPulser Bio-Rad CN 165-2100) system resulted in arcing at the highest voltage tested (1000 V), because all cytomix buffer concentrations tested had relatively high electrical conductivity. However, when the voltage was reduced (800 V, 500 V, and 300 V) arcing was eliminated and the resulting pulses lasted between 0.7 ms and 1.2 ms. Buffers composed of 10 % seawater and 1 % seawater both resulted in pulse durations of 0.7 ms and 3.5 ms at all applied voltages between 1000 V and 300 V. These pulse durations are all shorter than the typical 5.0 ms that results when using low conductivity buffer, such as MilliQ water at all voltages tested.

For the square wave electroporation system (NEPA21 transfection system, Bulldog Bio), cytomix buffers at high or low concentrations and the 10 % seawater were too conductive, resulting in arcing at 150 V and 300 V during a continuous 5.0 ms square pulse (Miyahara et al., 2013). In contrast, when voltage strength of both poring and transfer pulses was reduced to 99 V, with multiple 5.0 ms square pulses, the treatment was successful in the 100 % cytomix, 50 % cytomix, 10 % cytomix, and 10 % seawater buffers (Table 2). Buffers with low conductivity, such

as, 1 % seawater and MilliQ water were also able to complete the entire treatment without arcing at any of the tested voltages (99 V, 150 V, and 300 V).

Post-Electroporation Cell Viability Quantification

Using the exponential decay system, *P. caudatus* cells did not survive exposure to 1000 V in any of the above mentioned electroporation buffers (Table S1). In contrast, 40-50 % of cells were viable post-electroporation when a single exponentially decaying pulse was applied at 800 V (E = 4,000 V/cm) in all tested electroporation buffers. When the maximum voltage was limited to 500 V (E = 2,500 V/cm) cell viability increased to between 60-70 %. Applied voltages of 300 V (E = 1,500 V/cm) resulted in the highest cell viability of about 80-90 %.

Using the square-wave system, we initially tested the same parameters that were successfully applied for transformation of diatoms (Miyahara et al., 2013). Electroporated cells exposed to 150 V (E = 750 V/cm) or 300 V (E = 1,500 V/cm) in MilliQ water or 1% seawater survived. However, these electric fields failed to successfully transform *P. caudatus* with plasmid DNA, potentially because the transfer pulses used very low voltage (8 V). We therefore tested several other poring and transfer pulse voltage combinations, pulse numbers, and durations. Cells of *P. caudatus* electroporated with a maximum applied voltage of 99 V for poring and transfer pulses (E = 500 V/cm) in any of the investigated electroporation buffers (cytomix, seawater, and MilliQ water) were viable with no observed cell damage or loss. These parameters were also successful for establishing plasmid DNA transfection for *P. caudatus*.

We did not specifically assess cell viability after applying the microfluidic platform since cells are exposed to different electric fields with a single applied voltage (Garcia et al., 2016). Additionally, depending on the duty cycle selected, some of the cells flowed through the device

without being exposed to any electric field. Therefore, we tested electric field parameters that resulted in high cell viability using the exponential decay platform in subsequent experiments in the microfluidic device.

Real-Time Permeabilization Confirmation with SYTOX® Post-Electroporation

We aimed to establish the first transfection protocols for marine protists using the microfluidic system, and to identify the critical electric field that is required for the onset of electroporation. Initially, the ability to permeabilize *P. caudatus* cells was tested using the intercalating dye SYTOX® blue nucleic acid stain, which fluoresces upon binding to intracellular DNA. We delivered a single pulse with applied voltages of 500 V (E_{max} = 3,000 V/cm) and 1,000 V (E_{max} = 6,000 V/cm) in the absence of flow to expose cells to a narrow range of electric fields. The fluorescence images depicted in Fig. 2e and 2f confirm the ability to electroporate the cells at electric fields ranging from 1,000 V/cm – 6,000 V/cm with a pulse duration of about 5 ms. The extremely high electroporation efficiency at the ideal electric field can be seen in Fig. 2f, in which the majority of cells in the microfluidic channel have been successfully permeabilized (Supplementary Video 1). However, because the dead cells are also labelled with SYTOX®, this electroporation assay does not inform on the upper limit of the electric field within the range where cells are labelled and still viable. Transfection with plasmid DNA using the microfluidic technology provided more conclusive evidence of transfection success.

Transformation of *P. caudatus* with Plasmids

Circular DNA plasmids pEF-GFP, pUB-GFP, and pEYFP-Mitotrap were introduced separately into *P. caudatus* using the three different electroporation systems with parameters presented in Tables 1 and 2. All of our plasmids were expressed in the cytoplasm of *P. caudatus*

252 cells, after they were transcribed in the host's nucleus. In all cases of successful transfection,
253 transformants were viable and their growth rate was similar to that of wild-type cells. No
254 morphological differences in the cell shape were detected between the transformed and wild type
255 cells (Figs. 3 and 4). Expression of the GFP gene, driven by either the EF1 alpha promoter or the
256 ubiquitin C promoter, and the YFP gene, driven by the CMV promoter, was documented using a
257 fluorescence microscope 12 hours post-electroporation (Figs. 3 and 4). Expressed GFP signal
258 levels decreased gradually over the 48 h post-electroporation, but YFP expression was maintained
259 for 5 days (the longest time that expression was monitored). Microscopy revealed that GFP
260 expression driven by the ubiquitin C promoter was stronger than the GFP expression pattern driven
261 by the EF1 alpha promoter. Reverse transcription-PCR performed using RNA isolated from *P.*
262 *caudatus* cells transfected using the microfluidic system and the pUB-GFP plasmid revealed the
263 presence of GFP transcripts 3 days post-transfection (Fig. 5). These results clearly indicate that the
264 pUB-GFP plasmid was delivered into *P. caudatus* nucleus by electroporation and was transcribed
265 to GFP mRNA *in vivo*.

266 The fluorescence signal resulting from transfection of *P. caudatus* with the pEYFP-
267 Mitotrap plasmid was stable for 5 days post-transfection. Transcription of the YFP gene was
268 confirmed by RT-PCR using RNA isolated 5 days post-transfection (Fig. 5). Stability was not
269 monitored past 5 days in this study. Given optimization of antibiotic selection markers was outside
270 the scope of this short-term project, post-transfection cultures were maintained in the absence of a
271 selection marker, and hence were not suitable for long-term observation or experiments to confirm
272 stable transfection.

273 The microfluidic technique was implemented in the transformation of *P. caudatus,* but
274 unlike the square-wave and exponential decay systems, this system does not employ a uniform

electric field. As cells flow through the microfluidic device, they are exposed to multiple electric fields, making it challenging to assess viability as a function of a specific, uniform electric field. Based on our electroporation assays, initial unsuccessful attempts were made using maximum electric fields of 6,000 V/cm or 9,000 V/cm at the constriction with a 20 % duty cycle (Table S2). In order to improve the probability of transfection, the duty cycle was increased to 50 % with maximum electric fields of 750 V/cm, 1,500 V/cm, or 2,250 V/cm. These experiments conducted with a 50 % duty cycle resulted in transfection efficiencies ranging between 20-30 %. Finally, we increased the duty cycle to 95 % to increase the fraction of treated cells with maximum electric fields ranging between 500 V/cm and 3,000 V/cm. Transfection efficiencies ranging between 30-50 % were also achieved with maximum electric fields of 1,500 V/cm and 2,250 V/cm using 5 ms pulses in MilliQ water. Additionally, we achieved transfection efficiencies ranging between 20-30 % using a maximum electric field of 1,000 V/cm with 20 ms pulses in 50 % cytomix buffer in a straight channel.

Transformation efficiencies (percentage of successfully transformed cells) were comparable for the microfluidic platform and the commercially available square-wave technology. The microfluidic platform was the most efficient method with 30-50 % of the cells successfully transformed (Table 1). The square-wave platform resulted in transformation efficiency of ≥ 40 % (Table 2). Finally, the exponential decay electroporation resulted in ≤ 5% transformation efficiency making it the least optimal transfection platform evaluated (Table 1).

Discussion

In this study we achieved successful transfection of *P. caudatus* using three electroporation systems; our microfluidic platform, a square-wave system, and traditional exponential decay

methods. The process of developing transient transfection protocols in *P. caudatus* involved initial testing and determination of proper electroporation buffers and parameters (voltage strength, pulse duration and number). Our results suggest that the type of electroporation buffer is critical for maintaining cell viability throughout the experiment and is essential for determining the optimum electric field range. We have demonstrated that successful electroporation conditions were different for the three electroporation systems utilized, and are largely dependent on the electric field strength, as well as the number and duration of pulses. Transient transfection was carried out using three plasmids, pUB-GFP, pEF-GFP, and pEYFP-Mitotrap, which utilize promoters that are recognizable to most eukaryotes, to determine and optimize electroporation parameters. We now know that all three promoters, CMV, ubiquitin C, and EF1 alpha, work successfully with *P. caudatus*. Our transient transfection experiments demonstrated the feasibility of introduction and expression of foreign DNA into *P. caudatus* using each of the three systems (microfluidic, square wave, and the exponential decay), and the optimal electroporation parameters to apply for future stable transfection of *P. caudatus*.

Although the pEYFP-Mitotrap plasmid includes neomycin resistance as a selection marker gene and Tom70p as a target gene for the mitochondrial outer membrane (Robinson et al., 2010), the long-term stability of transfection and the efficacy of transformant selection based on antibiotic resistance were not examined in our study. Assessing longer-term stability of transfection using selective marker genes requires initial screening with various antibiotics to determine the most effective antibiotic and concentration, and this was outside the scope of this project. Further, in order to confirm stable transfection, one should investigate integration of the plasmid genes into the host genome by Southern blot or PCR and sequencing methods. Given that expression of the YFP gene was detected 5 days post-transfection via RT-PCR (Fig. 5), which represents stability

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over at least 5-6 generations, it is possible that this plasmid integrated into the nuclear genome, but this would need to be confirmed with Southern blotting. Similarly, we established successful transfection of the choanoflagellate *Monosiga brevicolis* using the microfluidic system with the same plasmid, pEYFP-Mitotrap, which was expressed for at least 4 days post-transfection (data not shown). This was supported by RT-PCR but not microscopically, due to overlap between the strong cell autofluorescence signal and the reporter gene signal.

The square-wave system. The square wave system with the specific electric field conditions given in Table 2 successfully delivered pUB-GFP plasmid DNA into the *P. caudatus* cytoplasm and achieved transient GFP expression with a transfection efficiency of about 40 % (Fig. 4c and 4d). In Under these conditions, both poring and transfer pulses had an equal electric voltage strength of 99 V (E = 500 V/cm). In contrast, attempts to establish transgenic *P. caudatus* using the previously applied electroporation parameters for diatom transformation (Miyahara et al., 2013) with high poring pulses voltage (150 V or 300 V) and low transfer pulses voltage (8 V) were unsuccessful. These results suggested that even though *P. caudatus* and the diatom *Phaeodactylum tricornutum* are single-celled marine eukaryotes, they possess different cell characteristics and therefore a different electric field strength and pulse number are required for successful intracellular delivery of exogenous DNA.

The exponential decay system. The exponential decay electroporation system was also used successfully to establish transgenic *P. caudatus* using two plasmids: the pEF-GFP and the pEYFP-Mitotrap (Fig. 3a and 3b). Comparison of results for the square wave and the exponential decay systems shows the exponential decay system results in a lower transformation efficiency of 5 %, and that the survival rate of the electroporated cells after exponential decay pulses at 800 V (E = 4,000 V/cm) is less than 50 % (based on light microscopy observation of swimming cells).

Relative to the square wave system, these results clearly indicate the increased effectiveness of the lower voltages and multiple pulses produced by the square wave electroporation system for delivering the extracellular DNA to larger numbers of cells with minimum cell damage.

The microfluidic electroporation system. The microfluidic electroporation system resulted in the highest transfection efficiencies ranging from 20 % to 50 %. The applied electric fields were much smaller than the ones employed during the SYTOX® assay in order to increase cell viability. We demonstrated successful *P. caudatus* transfection employing electric fields of 1,500 V/cm, resulting in transformation efficiencies of 30-40 %, and 2,250 V/cm, resulting in transformation efficiencies of 40-50 % efficiency using 5 ms pulse durations in MilliQ water and the bilaterally constricting channel geometry. Additionally, by decreasing the electric field to 1,000 V/cm and by employing longer 20 ms pulses, we achieved 20-30 % transfection efficiencies in 50 % cytomix buffer using the straight channel constriction. These results demonstrate that different geometric constrictions can be used successfully to modulate the electric field that the cell is exposed to for successful transfection.

The major advantage of the microfluidic platform is that it allows continuous flow-through transfection in comparison to traditional, commercially available cuvette-based technologies, while achieving comparable or better transfection efficiencies. Additionally, since the flow-through transfection process is continuous in nature, there is flexibility in the sample volume. This has exciting implications for processing large sample volumes (> 10 ml) within minutes, increasing throughput by 100 times in comparison to cuvettes (Garcia et al., 2017). This has advantages for future genome editing applications including library generation, and the ability to transfect cells directly from aqueous environments.

Conclusions

Development of successful transfection protocols for marine protists will enable advances in our understanding of their ecology. Here we successfully transfected *P. caudatus* for the first time using three different electroporation-based transfection methods and three different DNA plasmids pEF-GFP, pUB-GFP, and pEYFP-Mitotrap. Between the two traditional cuvette-based technologies (exponential decay and square wave), multi-pulse square wave electroporation resulted in higher transformation efficiency and cell viability. The microfluidic electroporation system produced the highest transfection efficiency (20-50%) when the optimal combination of buffer, electric field, and flow rate (among those tested in this study) was employed. This implies that microfluidic transfection holds great promise for efficiently optimizing and conducting electroporation of a potentially wide range of microbial eukaryotes. The microfluidic system is economical and can be installed and easily used by researchers and academics. The device features hydrodynamically-controlled electric fields that allow cells to experience a time-dependent pulse waveform that is otherwise difficult to achieve using standard electronics. The ability to efficiently test a wide range of electroporation parameters, or to quickly transfect a target (or a collection of targets) with a range of genetic elements has significant advantages over cuvette-based methods for the field of genome editing. High-throughput transfection technologies such as our microfluidics system offer the possibility of parallel processing of multiple samples (cultures or environmental samples), making possible effective investigations into the ecological roles of protists.

Experimental Procedures

P. caudatus Strain and Growth Media

389 *Parabodo caudatus* culture (ATCC 50361) was used in this study. This ATCC strain was
390 isolated from a freshwater sediment location, but *Parabodo* is described from marine habitats
391 (Kopylov et al., 1980). Initially, *Parabodo caudatus* was grown in 50 % ATCC seawater 802
392 media. Subsequently, seawater was replaced with distilled water in order to reduce the high
393 electrical conductivity during the electroporation. Briefly, this is a cerophyl-based media enriched
394 with 3.5 mM sodium phosphate dibasic (Na_2HPO_4) and with *Klebsiella pneumoniae* (ATCC-BAA
395 1706) added as a food source. *K. pneumoniae* is a gram negative, rod-shaped facultative anaerobe
396 bacterium commonly found in animals and the environment, and routinely used as bacterial prey.
397 Cultures were incubated at 22 °C and sub-cultured weekly in fresh T-25 vented tissue culture flasks
398 (Falcon brand, Fisher Scientific) containing 30 ml of fresh media.

399 **Cell Viability Assay After Cytomix Buffer Incubation**

400 Cell viability of *P. caudatus* in cytomix buffers needed to be tested since they can be
401 cultured in either MilliQ water or ≤ 50 % seawater. Three replicates of *P. caudatus* cultures each
402 twenty-five ml (i.e., biological replicates of *P. caudatus*, defined as different starting culture
403 bottles, although it is noted that all originated from the same starting strain) in logarithmic growth
404 phase ($1x10^7$ cells to $1.3x10^7$) were harvested by centrifugation at 5000 x g for 30 seconds and re-
405 suspended in 200 µl of 100 % cytomix, 50 % cytomix, or 10 % cytomix. To evaluate survival in
406 these buffers, aliquots of 20 to 30 µl of the cell-buffer mixture were placed on a haemocytometer
407 every 5 minutes for 15 minutes and were imaged under bright field microscopy (Nikon) using a
408 20x objective. Survival was determined by counting the total number of swimming cells in the
409 haemocytometer and determining the fraction of live cells.

410 **Electroporation Buffer Conductivity**

411 A buffer with low electrical conductivity is recommended to minimize Joule heating during
412 electroporation. We evaluated the electrical conductivity for the following buffers: 100 % cytomix
413 (120 mM KCl; 0.15 mM $CaCl_2$; 10 mM KH_2PO_4; 25 mM HEPES; 2 mM EGTA; 5 mM $MgCl_2$;
414 pH adjusted to 7.6 with KOH), 50 % cytomix (in MilliQ water), 10 % cytomix (in MilliQ water),
415 10 % seawater, 1 % seawater, or 100% MilliQ water, at four different voltages (300 V, 500 V,
416 800 V, and 1000 V). Since the exponential decay system uses 2-mm gap cuvettes, the electric
417 fields result in 1,500 V/cm, 2,500 V/cm, 4,000 V/cm, and 5,000 V/cm, respectively, after
418 computing the voltage-to-distance ratio. Electric field amplitude and pulse duration were measured
419 for each electroporation event with parameters given in Table S1. For square wave we followed a
420 published protocol for diatom transformation by Miyahara *et al.* (Miyahara et al., 2013), which
421 uses 2-mm cuvettes with applied voltages of 150 V and 300 V. In the microfluidic device we only
422 tested combinations of *P. caudatus* cells in the presence of SYTOX® or DNA as outlined below.

423 **Electroporation Parameters Tested and Post-Electroporation Cell Viability Quantification**

424 Prior to electroporation of *P. caudatus* cells in the MicroPulser Bio-Rad (CN 165-2100)
425 exponential decay system (Fig. 1), cell pellets from 25 mL of replicate cultures were re-suspended
426 in 200 µl MilliQ water, 1 % seawater, 10 % seawater¸ 10 % cytomix, or 50 % cytomix and
427 transferred to 2-mm gap cuvettes. The cells were electroporated with applied voltages of 300 V
428 (E = 1,500 V/cm), 500 V (E = 2,500 V/cm), and 800 V (E = 4,000 V/cm). The pulse duration in
429 milliseconds (ms) after each electroporation was recorded (Table S1). The cells were immediately
430 transferred to a 1.5 ml Eppendorf tube containing 1 ml of fresh growth media (ATCC 802 medium
431 prepared with distilled water) for recovery. To determine cell viability, aliquots (20-30 µl) of
432 electroporated cells were quantified using microscopy for each electric field applied.

433 The NEPA21 transfection system (Bulldog Bio), which utilizes square wave pulses, was
434 used for electroporation of *P. caudatus* in 2-mm gap cuvettes with identical buffers as used for the
435 exponential decay experiments. We initially used the same electroporation parameters that were
436 successfully applied previously for transformation of diatoms (Miyahara et al., 2013). However,
437 these high applied voltages of 300 V or 150 V were found to compromise *P. caudatus* cell viability
438 so modifications were necessary with a lower applied voltage (Table 2). It is important to note that
439 transformation in *P. caudatus* was most successful when we employed 'poring' (t = 5 ms) and
440 'transfer' (t = 50 ms) pulses of the same amplitude (99 V) but with different pulse durations.

441 We recently developed a continuous flow system to transform microorganisms in high
442 throughput in a microfluidic device (Garcia et al., 2017). This system employs microfluidic
443 channels that contain a bilateral constriction between the inlet and outlet electrode connections
444 ($length$ = 3.0 mm, $width_{min}$ = 50 μm, $width_{max}$ = 2.0 mm, and $height$ = 100 μm). The constriction
445 amplifies the electric field under an applied voltage between the inlet and outlet electrodes to levels
446 sufficiently high to induce electroporation. As opposed to the previous two systems that deliver
447 uniform electric fields in static cuvettes, this system drives cells through the constriction, which is
448 the region of highest electric field. During *P. caudatus* transfection, the cells were driven through
449 the microfluidic device at flow rates of 50 μL/min and 500 μL/min, which correspond to residence
450 times (i.e., pulse durations) of 20 ms and 2 ms, respectively. Square wave pulses with, for example,
451 5 ms ON and 5 ms OFF cycles (50 % duty cycle) are applied to the microchannel through the
452 dispensing needle. Therefore, the cell viability cannot be accurately evaluated since only 50 % of
453 the cells experience the electric field. The pulses are delivered from electrodes with alternating
454 polarity between the pulses to reduce electrolytic effects at the electrode-buffer interface (Fig. 1
455 and Fig. S1). After flowing through the microchannel (See Supplementary Video 2), each 200 μL

cell sample is added to a 1.5 ml Eppendorf tube containing 1 ml of fresh growth media for cell recovery. The applied voltages we evaluated had amplitudes of 250 V (E_{max} = 1,500 V/cm), 375 V (E_{max} = 2,250 V/cm), and 500 V (E_{max} = 3,000 V/cm) for each polarity. The non-uniform constriction in the microfluidic devices generates a variable electric field that is capable of transfecting cells while minimizing exposure to the highest electric field.

Electroporation Protocol Optimization with SYTOX® Blue

We used the SYTOX® Blue dead cell stain (Thermo Fisher Scientific) to initially determine pulse parameters that induce electroporation for *P. caudatus*. The SYTOX® Blue dye cannot penetrate the plasma membrane of living cells, but easily penetrates compromised plasma membranes, such as those induced by electroporation. Thus, the only cells that fluoresce are those that are exposed to an electric field strength and duration within and above the cell-specific critical electroporation threshold. *P. caudatus* cultures at logarithmic growth phase ($1x10^7$ cells to $1.3x10^7$) were harvested by centrifugation at 5000 g for 30 s. Cells were re-suspended in 200 µL of MilliQ water, 1 % seawater, 10 % seawater¸ 10 % cytomix, or 50 % cytomix and mixed with SYTOX® Blue dead cell stain to a final concentration of 5 µM. Cells were incubated for 2 min, then electroporated with exponential decay or microfluidic systems using different electroporation parameters. Two to three biological replicates (i.e., cells mixed with one of the tested buffer) were used for each of the tested applied voltages (technical replicates). In the exponential decay system we applied voltages of 300 V (E_{max} = 1,500 V/cm), 500 V (E_{max} = 2,500 V/cm), and 800 V (E_{max} = 4,000 V/cm). For the microfluidic device we applied voltages of 500 V (E_{max} = 3,000 V/cm) and 1,000 V (E_{max} = 6,000 V/cm) (Fig. 2). The applied voltage and pulse duration were measured for each electroporated sample and are shown in Table 1. For the exponential decay, cell integrity was confirmed using a bright field microscope (Nikon) and 20x

objective (Fig. 2b-2d). The bright blue signal was detected using a fluorescence microscope equipped with DAPI filter set. For the microfluidic system we were able to confirm the conditions that lead to successful entry of the SYTOX® Blue dye in real-time (Fig. 2e and 2f).

Plasmid Selection and Preparation

Three plasmids were obtained from Addgene (www.addgene.org/). pEYFP-Mitotrap (CMV mammalian and yeast promoter, the Tom70p gene targeting the outer membrane of the mitochondria in yeast and mammalian cells, and the YFP reporter) was a gift from Margaret Robinson (Addgene plasmid # 46942; (Robinson et al., 2010)); pEF-GFP (EF1 alpha promoter from mammalian cells for expression of GFP) and pUB-GFP (mammalian Ubiquitin C promoter for expression of GFP) were gifts from Connie Cepko (Addgene plasmid # 11154 and # 11155, respectively; (Matsuda and Cepko, 2004)). These plasmids were used to assess the transcriptional activity of those promoters and pEYFP-Mitotrap was used to assess whether Tom70p would only be expressed within the kinetoplast (a dense DNA-containing granule within the cell's single mitochondrion). Plasmids were purified from 100 mL cultures grown overnight in standard Luria Bertani liquid medium (Cold Spring Harbor Protocols 2006) with the appropriate selection marker. Purification was done according to the manufacturer's protocol for the Plasmid Midi Kit (Qiagen, Germantown, MD), with the following modifications: 1) Each 100 mL culture was split into two 50 mL volumes and centrifuged at 4,500 rpm for 20 min at 4°C to pellet bacterial cells; 2) Each half went through the lysis steps separately, and the lysate was pooled after neutralization; 3) Pelleting of precipitated DNA was done by centrifugation at 4,600 rpm for 60 min at 4°C; 4) Each 2 mL volume of pellet (in 70 % ethanol wash) was split into two 1 mL volumes, centrifuged at 15,000 X g for 10 min at 4°C, and the supernatant decanted; and 5) Dried DNA pellets were re-suspended in 50 µL of nuclease-free water, and the two 50 µL volumes were combined for each

sample. Purified plasmid DNA was quantified using the Qubit fluorometer (Thermo Fisher Scientific, Waltham, MA) and stored at -20°C until use. The success of our plasmid preparations was confirmed by PCR prior to use in transfection experiments.

Transfection of *P. caudatus*.:

P. caudatus cells were grown to logarithmic phase ($1x10^7$ cells to $1.3x10^7$) and harvested by centrifugation at 5000 X g for 30 s, re-suspended in 200 µl cytomix (50 % in distilled water), mixed with 20 to 40 µg of plasmid, and then transferred into an electroporation cuvette (2.0-mm gap) for electroporation with the exponential decay system and the square wave electroporation system. For the microfluidic system, cells in cytomix buffer were aspirated into 1/16 inch tygon tubing (McMaster-Carr) prior to being delivered into the microchannel. We carried out a minimum of ten trials of each combination of electroporation conditions tested using the three platforms; however, only the successful transformation parameters are summarized in Tables 1 and 2. Electroporation parameters that were not successful are included in Table S1 for the exponential decay system and Table S2 for the microfluidic system.

RT-PCR Confirmation for Expression of Plasmids in *P. caudatus*

Total RNA was isolated from transformed *P. caudatus* cells using the RNEasy Mini Kit (Qiagen, Hilden, Germany). Cells were filtered onto a Durapore® PVDF 0.45 µm-pore size filter (EMD Millipore, Billerica, MA). The filter was placed in 500 µL RLT lysis buffer (RNA Isolation Kit, Qiagen, Hilden, Germany) with 143 mM β-mercaptoethanol and vortexed. Following 10 min incubation at room temperature, 350 µL of 100 % ethanol were added and the lysate was purified using RNEasy Mini Kit according to the manufacturer's instructions.

Purified RNA then underwent two rounds of DNAse treatment (Jones et al., 2007). First,

the Turbo DNA-*free*™ Kit (Ambion®, Thermo Scientific, Waltham, MA) was used with the following modifications: 1) A total of 2 µL DNAse was added, 1 µL each time, with each addition followed by a 30 min incubation at 37 °C; and 2) 0.2 µL volumes of DNAse inactivation reagent were used. Next, the RNAse-Free DNase Set was used in combination with the RNEasy Mini Kit (Qiagen, Hilden, Germany) to perform an on-column DNAse digestion, followed by column-based purification, according to the manufacturer's instructions.

First-strand cDNA synthesis and PCR amplification were performed using the One*Taq*® RT-PCR Kit (New England Biolabs, Ipswich, MA). The appropriate reverse primer (0.5 µM final conc.; See Table S3) and 5 µL RNA were used for reverse transcription. Control reactions were performed with water in place of the reverse transcriptase enzyme mix. cDNA was amplified in a 25 µL PCR reaction, with final primer concentrations of 0.2 µM. Thermocycling conditions were as follows: 30 sec at 95 °C; 30 cycles of 30 s at 94 °C, 30 s at 58 °C, and 30 s at 68 °C; and a final extension for 5 min at 68 °C. PCR primers targeting expression of the GFP or YFP reporter gene (Table S3) were used. PCR products were visualized by gel electrophoresis, with purified plasmid as a positive control. Amplified PCR products at the expected size of 367-bp for YFP and GFP genes were documented (Fig. 5).

Supplementary Methods:

Soft Lithography Protocol for Microfluidic Device Fabrication

Soft lithography is employed in order to fabricate devices with microscale features. This process creates a master stamp from photomasks that can be used to create devices repeatedly. The photomasks are designed in AutoCAD 2014 (Autodesk, San Rafael, CA) with bilaterally converging or straight geometries, and are printed by Fine-Line Imaging, Inc. (Colorado Springs,

546 CO). The microchannels are microfabricated using soft lithography techniques described by
547 Garcia *et. al* (Whitesides et al., 2001; Garcia et al., 2016). Briefly, SU-8 (SU-8 2050, Micro-Chem,
548 Westborough, MA) molds are patterned on silicon wafers with standard photolithography.
549 Afterwards, the surfaces of the SU-8 master mold are treated for 2 hours with tridecafluoro-1,1,2,2-
550 tetrahydrooctyl-1-trichlorosilane (Sigma Aldrich, St. Louis, MO) under vacuum before being used
551 for molding. Next, the SU-8 master mold polydimethylsiloxane (PDMS, Sylgard 184, Dow
552 Corning, Midland, MI) was used at a 10:1 ratio after 2-hour vacuum for removal of air bubbles in
553 the polymer. The PDMS devices are bonded to a glass substrate after a 45 second plasma treatment
554 and placed overnight in an oven at 75 °C prior to subsequent experiments.

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Figure 1: Electric field waveforms employed for transient and stable transfection of *Parabodo caudatus*. a) Three independent electroporation systems were used for reproducible transfection, including our microfluidic electroporation platform (Garcia et al., 2017), the NEPA21 square-wave transfection system (Bulldog Bio), and the MicroPulser™ exponential decay electroporator (Bio-Rad). b) The signature waveforms for the NEPA21 square wave transfection system include both 'poring' and 'transfer' pulses for electroporation. Note: The time scale in Fig. 1a is a zoomed-in version of the red-dashed box from Fig. 1b.

Figure 2: Permeabilization confirmation of *P. caudatus* cells with SYTOX® Blue. a) Phase contrast and b-d) fluorescence images of *P. caudatus* electroporated with 5 µM SYTOX® Blue nucleic acid stain at 300 V (E_{max} = 1,500 V/cm), 500 V (E_{max} = 2,500 V/cm), and 800 V (E_{max} = 4,000 V/cm), respectively, using the exponential decay electroporation system in 2-mm cuvettes. Fluorescence images e) before and f) after electroporation in the microfluidic system using a single 5 ms exponential decay pulse at 500 V (E_{max} = 3,000 V/cm).

Figure 3: Microfluidic transfection of *Parabodo caudatus*. a) *P. caudatus* (brightfield), b) transient pEYFP-Mitotrap transfection at 250 V (E_{max} = 1,500 V/cm), c) transient pUB-GFP transfection using 375 V (E_{max} = 2,250 V/cm), d) autofluorescence control for *P. caudatus*, e) transient transfection using pEF-GFP and 313 V (E_{max} = 1,000 V/cm) in the straight channel, and f) merged image of brightfield and fluorescence image from e) for visualizing cell morphology.

Figure 4: Exponential decay and square wave transfection of *Parabodo caudatus*. Fluorescence imaging confirmation of *P. caudatus* after a) transient transfection with pEF-GFP using the MicroPulser^{TM} exponential decay electroporator, after b) transient transfection with pEYFP-Mitotrap using 800 V (E_{max} = 4,000 V/cm) in the exponential decay electroporator, and after c) transient transfection with pUB-GFP using the NEPA21 square-wave transfection system at 99 V (E_{max} = 500 V/cm). Panel d) shows merged fluorescence image from c) with the brightfield image.

Figure 5: Gel electrophoresis image showing the RT-PCR results detecting reporter genes expression in *P. caudatus* transformants. (L) 1kb ladder (Invitrogen, cat. #10787018); (1) GFP expression profile in *P. caudatus* cells transformed with pUB-GFP plasmid at 375 V using the microfluidic electroporation system; (2) Control reaction was performed without addition of RT to verify the absence of DNA in the RNA preparations in *P. caudatus* cells transformed with pUB-GFP transient plasmid using 375 V with the microfluidic electroporation system; (3) pEYFP expression profile in *P. caudatus* cells transformed with pEYFP-Mitotrap plasmid at 250 V with the microfluidic electroporation system; (4) Control reaction was performed without addition of RT to verify the absence of DNA in the RNA preparations in *P. caudatus* cells transformed with withpEYFP-Mitotrap plasmid at 250 V with the microfluidic electroporation system; (5) PCR negative control; (6) PCR positive control using the GFP plasmid DNA. The PCR products were separated on a 1% agarose gel, visualized under UV light, and DNA fragments of both reporter genes were at the expected size of 367-bp.

Figure S1: Representative 5-ms square waveform delivered with alternating polarity in the microfluidic device at a 95 % duty cycle. This device geometry results in a ~6x amplification of the applied voltage in the narrowest portion of the constriction. Therefore, the applied voltage of 250 V presented here results in a maximum electric field (E_{max}) of 1,500 V/cm in the microfluidic device.

Figure S2: Graphical representation of the microfluidic device used in this study for continuous flow-through transfection of *P. caudatus*. The device exhibits inlet (green) and outlet (red) fluidic ports that also serve as electrodes to generate the electric field within the bilateral constriction.

Supplementary Video 1: Microfluidic electroporation of *Parabodo caudatus* in the presence of 5 μM SYTOX® Blue nucleic acid stain using an applied voltage of 1,000 V (20X magnification) demonstrates successful intracellular delivery of dye due to real-time fluorescence detection.

Supplementary Video 2: Microfluidic electroporation of *Parabodo caudatus* with pUB-GFP driven at 500 μL/min with an applied voltage of 375 V (E_{max} = 2,250 V/cm) and a 50 % duty cycle resulted in 20-30 % transfection efficiency.