High-efficiency electroporation of chytrid fungi

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Two species of parasitic fungi from the phylum Chytridiomycota (chytrids) are annihilating global amphibian populations. These chytrid species—Batrachochytrium dendrobatidis and B. salamandrivorans—have high rates of mortality and transmission. Upon establishing infection in amphibians, chytrids rapidly multiply within the skin and disrupt their hosts' vital homeostasis mechanisms. Current disease models suggest that chytrid fungi locate and infect their hosts during a motile, unicellular ‘zoospore’ life stage. Moreover, other chytrid species parasitize organisms from across the tree of life, making future epidemics in new hosts a likely possibility. Efforts to mitigate the damage and spread of chytrid disease have been stymied by the lack of knowledge about basic chytrid biology and tools with which to test molecular hypotheses about disease mechanisms. To overcome this bottleneck, we have developed high-efficiency delivery of molecular payloads into chytrid zoospores using electroporation. Our electroporation protocols result in payload delivery to between 75 and 97% of living cells of three species: B. dendrobatidis, B. salamandrivorans, and a non-pathogenic relative, Spizellomyces punctatus. This method lays the foundation for molecular genetic tools needed to establish ecological mitigation strategies and answer broader questions in evolutionary and cell biology.

Derived from a node basal to the multicellular Dikarya that includes both Ascomycota (e.g. sac fungi) and Basidiomycota (e.g. mushrooms) (Fig. 1A)1–3, zoosporic fungi have retained many ancestral phenotypes including motile, unicellular, flagellated cell types called zoospores4,5. Crucial for dispersal, zoospores likely rely on complex sensory-motor coordination to locate and settle at nutrient-rich sites6–9. Settled zoospores quickly grow into sporangia, inside of which dozens to hundreds of new zoospores develop to begin the cycle anew (Fig. 1B).

Zoosporic fungi in the phylum Chytridiomycota (chytrids) include ecologically important pathogens. For example, two species in the genus Batrachochytrium have been recognized as virulent amphibian pathogens causing mass amphibian die-offs as far back as the 1970s10. While not all chytrids are pathogenic—Spizellomyces punctatus (Sp) is non-pathogenic—Batrachochytrium dendrobatidis (Bd) and Batrachochytrium salamandrivorans (Bsal) infections result in mortality rates that have driven multiple species extinct11–15, and are the first known example of a chytrid-driven epidemic in vertebrates10,16. Although exact numbers are currently under analysis, Bd alone has led to a catastrophic global decline of amphibians and Bsal is now considered an emerging global pathogen17,18. Exacerbated by climate change and anthropogenic distribution, it is unclear how much additional amphibian diversity will be lost to chytrid infection without intervention19,20.

A significant hurdle to effective management of chytrid infections is the lack of basic knowledge about their biology. With no way to study the molecular mechanisms underlying host identification, infection, and growth in parasitic chytrids, the field is struggling to develop mitigation strategies for even a single host species. While tools are just emerging for non-parasitic relatives21, developing molecular genetic tools for parasitic chytrids will remove key roadblocks hindering our efforts to curb the pathogens’ ecological impact. Additionally, these tools will leverage the unique diversity and evolutionary history of zoosporic fungi to better understand broader principles underlying the origins of parasitism, transitions to pathogenicity, and evolutionary cell biology.

Molecular genetic tools generally require driving molecular payloads across the cell membrane. To deliver molecules to the cytoplasm of chytrid cells, we have optimized one such method, electroporation, because of its widespread use in fungi and the flexibility of its downstream applications22–24. Electroporation exposes cells to tightly controlled electrical pulses that create holes in the cell membrane. The now-permeable membrane allows for co-incubated compounds to passively enter the cell before the membrane is sealed (Fig. 1C). Successful electroporation depends largely on physical parameters such as cell size, pulse profile, buffer conductance, and electrode contact area22,25, each of which must be optimized for a particular cell type. The comparatively
small role that biological parameters play in the success of electroporation make it applicable—given enough persistence—to cells from almost any taxa.

Although the initial process of developing electroporation for a new species is time-intensive, this investment yields a general template that can be quickly expanded to closely related species and optimized for delivery of a wide variety of payloads. Electroporation is widely used to introduce heterologous DNA into cells and perform CRISPR/Cas9-mediated gene editing, although these methods require significant development beyond electroporation. Conversely, other tools such as antisense oligonucleotides, siRNAs, and small molecule inhibitors can be used almost immediately after electroporation has been optimized.

To study the biology of infectious chytrids, we have developed a reliable, high-efficiency electroporation protocol for chytrid fungi. We optimized this protocol for two pathogenic chytrids (B. dendrobatidis, B. salamandrivorans), and a non-pathogenic chytrid (S. punctatus). This protocol opens the door for further studies using molecular genetics to address both ongoing and future chytrid epidemics while laying the foundation for comparative studies in these unique and diverse lineages.

Results
Voltage for optimal electroporation varies between chytrid species. To track the success of molecular payload delivery during electroporation, we used fluorescein-labelled dextrans (hereafter referred to simply as dextrans) and measured delivery using both microscopy and flow cytometry. To optimize electroporation, one must explore a large parameter space that includes the following variables: pulse shape, pulse time, number of pulses, contact distance, contact surface area, buffer conductivity, and pulse voltage. To quickly explore this parameter space, we conducted single-replicate trials looking for parameter combinations that resulted in at least 15% of single cells loaded with dextrans. This wide exploration led us to an unoptimized protocol using two three-millisecond square-wave pulses, five seconds apart yielding 10–20% efficiency (data not shown). To optimize this protocol for Bd, we adjusted pulse voltage alone testing 750 V, 1000 V, and 1250 V. Over this range, we found efficiency peaked at 1000 V, with ~95% of zoospores loaded with dextrans (Fig. 2B,C) and a 41–71% survival rate (Fig. 2D). We compared this efficiency to controls for autofluorescent shift due to pulse exposure (electroporation without dextran incubation) and dextran ingestion or non-specific binding to cell walls/coats (dextran incubation without electroporation). These comparisons revealed the fluorescence of dextran-incubated, electroporated cells is due to successful dextran delivery into the cytosol. When translating this optimized protocol to other species we found the optimal voltages were 1250 V for Bsal and 1000 V for Sp, with no other adjustments (Figs. 2, 3, 4).

Cell walls of encysted spores accumulate fluorescent molecules. When analyzing flow cytometry data, we used non-electroporated, no dextran controls to set a ‘positive’ gate which excluded 99.5% of all control
cells. To obtain the percent of positive cells under other treatments, we transposed this gate onto each experimental and control treatment within the experimental replicate. Unexpectedly, non-electroporated spores incubated with dextrans showed 3–10% of cells in the treatment were positive (Figs. 2, 3, 4, 5). To understand why

**Figure 2.** Electroporation efficiency and survival of *B. dendrobatidis* at 750 V (green), 1000v (blue), and 1250 V (purple). (A) Representative images of cells after electroporation at 1000v showing live cells adhered to the slide and attempting amoeboid movement. (B) Representative flow cytometry data from a single replicate showing the fluorescence intensity of single cell events for non-electroporated, no dextran controls (grey); non-electroporated, dextran incubated controls (red); and electroporated, dextran incubated treatments (colors vary by voltage). (C) Percent of single cell events with fluorescence intensities above the non-electroporated, no dextran control relative to all single cell events for three independent biological replicates. (D) Percent survival of cells in electroporated, dextran incubated controls in three independent biological replicates. Counts were normalized to the respective non-electroporated, no dextran controls.
incubation with dextrans leads to an increase of positive cells in the absence of electroporation, we inspected these samples using fluorescence microscopy. We observed cells with a bright ring of fluorescence on the periphery of a small percentage of cells in all samples incubated with dextrans regardless of their exposure to electric shock (Fig. 5), suggesting the small percent of positive cells in non-electroporated samples is due to non-specific binding of dextrans to the cell-coat or cell-wall.

Electroporation efficiency is strongly affected by dextran manufacturer. To explore if cheaper sources of 3000 MW dextrans could substitute our initial choice (Thermo-Fisher), we tested an alternative manufacturer of 3000 MW dextrans (Sigma-Aldrich). In the lots and concentrations we tested (stored under recommended storage conditions), we find a 30–70% difference in loading efficiency between two dextrans manufacturers (Fig.S2). In side-by-side comparisons, Thermo Fisher anionic 3000 MW dextrans (Thermo, D3305) yield 75–97% loading efficiency while the cheaper option (Fig. S1A), Sigma 3000 MW dextrans (Sigma, FD4) results in a lower and more variable efficiency (Fig. S1B). After electroporation, Sigma FD4 dextrans are confined to a small point in the majority of cells but do not stain the cell wall (Fig. S1B). In contrast, Thermo-Fisher dextrans appear evenly distributed throughout the cytoplasm in the majority of cells with occasional staining of the cell wall (Fig. S1A).

Figure 3. Electroporation efficiency of *B. salamandrivorans* for 750 V (green), 1000 V (blue), and 1250 V (purple). (A) Representative images (DIC, FITC, Merged) from a single replicate at 1000 V showing positive cells with dextran inside the cell and not only stuck to the cell wall/coat. (B) Representative flow cytometry data from a single replicate showing the fluorescence intensity of single cell events for non-electroporated, no dextran controls (grey); non-electroporated, dextran incubated controls (red); and electroporated, dextran incubated treatments (colors vary by voltage). (C) Percent of single cell events with fluorescence intensities above the non-electroporated, no dextran control relative to all single cell events for three independent biological replicates.
Synchronizing spores is not required for high efficiency. Without synchronizing cultures, chytrid zoospores are different ages and represent a very heterogeneous population of cells ranging from just released to about to encyst. Because zoospore age could impact on electrocompetency, for our initial experiments we synchronized cultures to limit the confounding variables. Once we achieved reproducible and effective electroporation with synchronized cultures, we then tested the effect of zoospore synchrony on electroporation efficacy. We find that although synchronization of zoospores may slightly increase efficiency, it is not required to reach >80%
loading of all cells (Fig. S2). Use of synchronized spores, however, may facilitate downstream applications of electroporated zoospores.

Methods

Culture and synchronization of zoospores. We cultured Batrachochytrium species in tissue culture treated flasks (Celltreat 229,330) and S. punctatus in petri dishes. Strains were thawed and then routinely passed according to the following species-specific protocols: We grew B. dendrobatidis (JEL423; from the highly virulent Bd-GPL global pandemic strain) in 1% Tryptone for 3 days at 24 °C, B. salamandrivorans (AMFP1) in ½ strength TGHL for 4 days at 15 °C, and S. punctatus (ATCC 48,900) on K1 plates at 30 °C for 1–2 days. After the fungi were mature, we collected zoospores and seeded the next passage.

For Batrachochytrium species, we synchronized zoospores by discarding growth media and gently washing older zoospores away from the sporangia adhered to the tissue culture flask three times with fresh growth media. After two hours of incubation with fresh growth media, we collected synchronized zoospores for use. For S. punctatus, we checked for mature sporangia after 19–32 h of growth while ensuring the surface of the agar had a thin film of liquid around each sporangium. As soon as sporangia began releasing zoospores, we washed plates three times with 2 mL of K1 media and let the sporangia sit under another 2 mL for 30 min before collecting synced zoospores.

Electroporation. We have made the detailed electroporation protocol available on protocols.io (dx.doi.org/10.17504/protocols.io.88xhzxn) and in a notebook-ready checklist format (File S1). In short, zoospores are brought to a specific concentration, co-incubated on ice with dextrans, electroporated using high-voltage square-wave pulses, then allowed to recover on ice. Unless specifically noted, all solutions were sterilized and we synchronized zoospores prior to collection. To achieve proper buffer conductance during electroporation, we transferred zoospores into SM buffer (5 mM KCl, 15 mM sodium phosphate buffer pH 7.2, 15 mM MgCl₂, 25 mM sodium succinate dibasic hexahydrate, 25 mM D-Mannitol) after collection by repeating the following wash steps three times: centrifuge zoospores for 5 min at 2,500 rcf, remove supernatant, then resuspend in 5 mL of SM buffer. We then brought zoospores to a final concentration of 1 × 10⁷ zoospores/mL. In preparation for electroporation, we loaded 100 uL of this 1 × 10⁷ zoospores/mL suspension into a 0.2 cm electroporation cuvette (Bulldog Bio 12,358–346) along with either: 100 uL of 2 mg/mL 3000MWCO fluorescein-labelled dextrans (Thermo-Fisher, Cat. no. 3305) in SM buffer for experimental treatments, or 100 uL SM buffer for control treatments. We then chilled all cuvettes on ice for 10 min. Immediately before electroporation, we gently mixed each cuvette by pipetting up and down to resuspend chilled zoospores. To electroporate cells, we used two, three millisecond, square-wave pulses five seconds apart (Biorad GenePulser Xcell + CE module) for all samples. We

Figure 5. Although dextran staining of cell wall/coat occurs without electroporation, stained and electroporated cells can be distinguished via microscopy. Representative images of each treatment were taken from a single replicate performed on subsets of spores from the same population. Fluorescence intensity was normalized across all images. Non-electroporated cells incubated with dextrans (non-electroporated, + dextran) show weak staining in a pattern that suggests binding to the cell wall. This same pattern can be observed in electroporated samples (electroporated, + dextran) where stained cells (right) co-occur with positive cells (left) in which dextran fluorescence is limited to the inside of the cell.
tested 750 V, 1000 V, and 1250 V for all species. We immediately placed each cuvette on ice for 10 min after electroporation then gently added 200 µL of 1% tryptone and allowed cells to rest for another 10 min on ice.

**Cytometry and microscopy measurement of electroporation efficiency & mortality.** We measured parameter efficiency using flow cytometry and used microscopy to verify that cells were loaded, rather than just externally stained. These steps are not necessary to grow electroporated cells for downstream applications and can be applied to a subset of cells, or even skipped after optimizing the protocol for new payloads. To prepare cells for flow and microscopy, we transferred all zoospores into 5 mL of chilled media. To remove extracellular dextrans and autofluorescent media, we performed a series of 6 wash steps. For each step, we centrifuged zoospores at 2,500 rpm for 5 min at then discarded the supernatant and gently resuspended the zoospore pellet in 5 mL of the indicated solution. For the first 3 wash steps, we resuspended zoospores in chilled growth media; for the next 2 wash steps we resuspended in room-temperature SM buffer. For the final wash step in SM buffer, we resuspended the zoospores using only 600 µL of SM buffer. We then transferred 100 µL of zoospores to a separate tube for imaging and fixed the remaining volume by adding an equal volume of paraformaldehyde (PFA) based fixation buffer, final concentrations: 4.8% PFA, 9 mM sucrose, 50 mM Sodium Phosphate Buffer pH 7.2. We incubated the zoospores and fixation buffer on ice for 20 min, after which we centrifuged them for 5 min at 2,200 rpm, discarded the supernatant, and resuspended in 400 µL of unchilled SM buffer.

We used flow cytometry of the fixed cells to measure the overall ratio of loaded to unloaded zoospores. Using a BD LSR Fortessa 3 Laser cytometer and BD FACSDiva software, we captured 10,000 events and analyzed the resulting data using FlowJo v10.1. Cells were gated for enhanced FITC in single-cell events relative to control treatments. To corroborate flow cytometry measurements, we imaged live cells on a Nikon Ti-E inverted microscope equipped with a 100 × PlanApo objective and sCMOS 4mp camera (PCO Panda) using NIS Elements software to capture images in FITC (LED illuminator at 488 nm) and DIC (white LED transmitted light) images. *Bd* zoospores were adhered to the slide by pre-treating it with concanavalin-A31, all other spores were imaged without treating the slide. To estimate the percent of cells that survive the protocol, we counted the number of live spores in electroporated, dextran- incubated treatments and normalized to the non-electroporated, no dextran control. We only collected this data for *B. dendrobatidis* because other species could not be adhered to the slide and we could not ensure zoospore behavior would remain identical across treatments, potentially skewing counts. Qualitative observations, however, suggest that survival rates of other species are similar to that of *Bd*.

**Discussion**

Developing efficient, reproducible molecular tools for emerging species remains a crucial step towards unraveling the general principles, origins, and evolution of basic cell biology. We present a well-optimized electroporation protocol for the zoospores of infectious chytrid fungi, with between 75 and 97% of cells demonstrably loaded with a fluorescent dextran payload. This range of efficiencies reported here matches and exceeds what is expected based on well-established electroporation protocols for mammalian cells (e.g. C3H/10 T 1/2 cells32). The efficiencies we report are likely conservative because electroporated cells are remarkably fragile, and the stringent wash steps we used to quantify the percent of cells loaded likely resulted in additional cell death, which could be avoided in typical applications.

Electroporation provides a fast, cost-effective method to deliver the material necessary to use genetic approaches to determine the molecular mechanisms that enable disease. The protocol we have developed allows delivery of antisense oligonucleotides, siRNAs, and small molecule inhibitors that allow direct testing of molecular hypotheses of chytrid development and pathogenesis. This approach can be extended to developing transformation protocols that allow transgenic lines and the expression of genetically encoded reporters. With the ability to examine the molecular mechanisms driving the chytrid infection process and life cycle, it is likely that researchers will uncover new avenues for disease control for current and future chytrid epidemics.

Chytrid fungi occupy an important phylogenetic position derived from one of the most basal nodes in the fungal kingdom, making comparative studies between chytrids useful for understanding early fungal evolution. We have, therefore, developed this protocol with comparative study in mind. Our successful electroporation of the non-pathogenic *S. punctatus* shows that this method is likely to be successful in many other chytrids with little modification. We anticipate that researchers will be able to quickly optimize electroporation protocols for a wide array of chytrids and zoosporic fungi by using this protocol as a guide. The development of electroporation and transformation in additional, diverse zoosporic fungi will be a boon to future comparative studies seeking to understand the molecular toolkit in ancestral opisthokonts and the processes that influenced its evolution in fungi.

**Potential pitfalls and difficulties.** Despite the general convenience of electroporation, we find that there are a number of qualitative variables which can have a large effect on the replicability of this protocol. For instance: using a cheaper source of what should be identical dextrans results in a 30–70% drop in efficacy but using asynchronized cells produces no major effect on efficacy. In an effort to help future researchers, we discuss a number of pitfalls and difficulties we believe will help make applications of this protocol successful:

*Zoospores are fragile.* We observed a large drop in efficacy and single cell events measured by flow cytometry if cells were pipetted or poured too roughly at any stage of the protocol. Thus, the use of any shaker/vortexer is not recommended at any point in this protocol before cell fixation. Researchers attempting this protocol should also exercise caution and be sure to pipette gently when resuspending spores. We also noted a qualitative increase in cell death at the end of this protocol if cells experienced large temperature fluctuations and advise that cuvettes...
are initially loaded with room-temperature reagents, then everything be placed and kept on ice after the cuvettes have been loaded to allow the cells to chill gradually.

**Cell walls bind dextrans.** Qualitative image analysis suggests that “stained” cells all lack a visible flagellum and therefore have likely begun the encystment process. This process is also marked by rapid synthesis of cell wall, endocytosis, and growth. We find it likely that dextrans in the external environment binds to the newly synthesized chitin cell wall similar to the binding of wheat germ agglutinin to saccharides in fungal cell walls. Without confirmation and the establishment of methods to quench externally bound FITC molecules, the use of “no electroporation” controls incubated with dextrans are necessary to estimate the percentage of cells which are ‘positive’ but do not have dextrans in the cytoplasm.

**Loss of fluorescence in fixed cells after 6 h.** After approximately 6 h, dextran loaded cells were not detectable by flow cytometry, even when re-sampling previously fluorescent samples stored in the dark at 4 °C overnight. This is likely a result of using non-fixable fluorescein. If users elect to use the same dextran as we did in these experiments (D3305) we urge them to avoid permeabilization steps and to proceed as quickly as possible from fixation to flow cytometry. Alternatively, users may wish to try Thermo Cat. No. D3306, a lysine-fixable 3000 MW fluorescein-labelled dextran if they would like to leave fixed samples overnight or include additional stains that require permeabilization.

**Viability after electroporation.** Most downstream applications of this protocol will require cells to remain alive after electroporation. We confirmed that spores loaded with dextrans were alive, active, and crawling at least 4 h after electroporation. However, the survival rate of electroporated spores is likely to be heavily dependent on the molecules introduced into the cell. While not necessary to assess the efficacy of electroporation itself, any future application of this protocol should assess the halflife of their introduced molecule through flow cytometry, interrupted phenotypes, or antibiotic resistance.

**Conclusions**

Developing molecular tools for chytrids allows researchers access to unique, diverse lineages for molecular studies into the origin, evolution, and fundamental processes underlying cell biology and parasitism. Our protocol is a first step towards building a suite of molecular tools allowing us to study chytrid fungi in depth. We anticipate this protocol will give rise to a multitude of studies using morpholinos, small molecule inhibitors, siRNAs, and eventually DNA editing to investigate the mechanisms and evolution behind chytrid biology. Furthermore, our results suggest that the protocol presented here is likely universal within Chytridiomycota and may be used as a starting template for other zoosporic species. Zoosporic fungi are now poised to quickly advance solutions to current ecological problems and provide context for broader questions in evolutionary and sensory cell biology.

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**Author contributions**

All authors contributed to experimental design and performed experiments. A.S. and L.F.-L. produced the manuscript.

**Competing interests**

The authors declare no competing interests.

**Additional information**

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