

## RESEARCH ARTICLE

# Identification of antibiotics for use in selection of the chytrid fungi *Batrachochytrium dendrobatidis* and *Batrachochytrium salamandrivorans*

Kristyn A. Robinson, Mallory Dunn, Shane P. Hussey, Lillian K. Fritz-Laylin \*

Department of Biology, The University of Massachusetts Amherst, Amherst, MA, United States of America

\* [lfritzlaylin@umass.edu](mailto:lfritzlaylin@umass.edu)



## OPEN ACCESS

**Citation:** Robinson KA, Dunn M, Hussey SP, Fritz-Laylin LK (2020) Identification of antibiotics for use in selection of the chytrid fungi *Batrachochytrium dendrobatidis* and *Batrachochytrium salamandrivorans*. PLoS ONE 15(10): e0240480. <https://doi.org/10.1371/journal.pone.0240480>

**Editor:** Louise A. Rollins-Smith, Vanderbilt University School of Medicine, UNITED STATES

**Received:** July 16, 2020

**Accepted:** September 25, 2020

**Published:** October 20, 2020

**Peer Review History:** PLOS recognizes the benefits of transparency in the peer review process; therefore, we enable the publication of all of the content of peer review and author responses alongside final, published articles. The editorial history of this article is available here: <https://doi.org/10.1371/journal.pone.0240480>

**Copyright:** © 2020 Robinson et al. This is an open access article distributed under the terms of the [Creative Commons Attribution License](https://creativecommons.org/licenses/by/4.0/), which permits unrestricted use, distribution, and reproduction in any medium, provided the original author and source are credited.

**Data Availability Statement:** All relevant data are within the manuscript and its Supporting Information files.

## Abstract

Global amphibian populations are being decimated by chytridiomycosis, a deadly skin infection caused by the fungal pathogens *Batrachochytrium dendrobatidis* (*Bd*) and *B. salamandrivorans* (*Bsal*). Although ongoing efforts are attempting to limit the spread of these infections, targeted treatments are necessary to manage the disease. Currently, no tools for genetic manipulation are available to identify and test specific drug targets in these fungi. To facilitate the development of genetic tools in *Bd* and *Bsal*, we have tested five commonly used antibiotics with available resistance genes: Hygromycin, Blasticidin, Puromycin, Zeocin, and Neomycin. We have identified effective concentrations of each for selection in both liquid culture and on solid media. These concentrations are within the range of concentrations used for selecting genetically modified cells from a variety of other eukaryotic species.

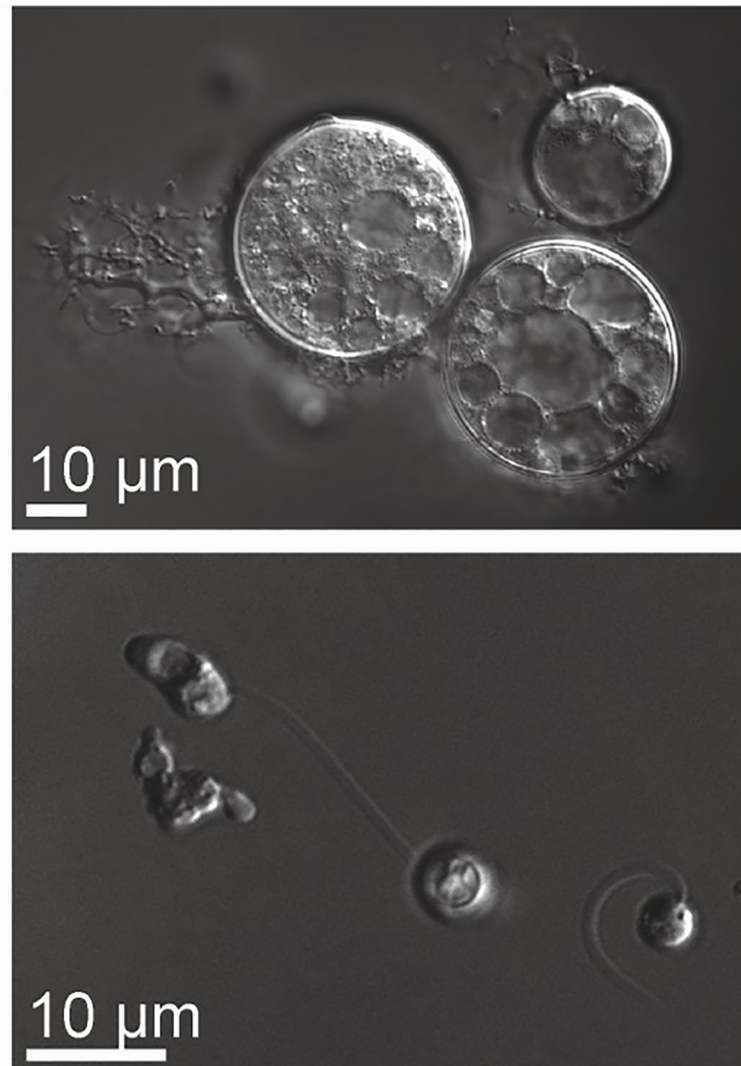
## Introduction

Chytrids are early diverging fungi that are commonly found in aquatic and moist environments [1]. They play key ecological roles, particularly by cycling carbon between trophic levels [2, 3]. Chytrids have a biphasic life cycle characterized by motile and sessile stages (Fig 1) [4–6]. They begin their life as motile “zoospores,” which use a flagellum to swim through water and, for some species, actin-based motility to crawl along surfaces [7, 8]. Zoospores then transition to a sessile growth stage by retracting their flagellum and building a cell wall in a process referred to as encystation. Encysted spores of many species develop into sporangia and develop hyphal-like structures called rhizoids and grow rapidly. Each sporangium produces many zoospores that exit via discharge papillae to begin the life cycle anew.

Many chytrids are pathogens that infect protists, plants, algae, fungi, and vertebrates [2]. The most infamous chytrids are the vertebrate pathogens *Batrachochytrium dendrobatidis* (*Bd*) and *B. salamandrivorans* (*Bsal*). Both pathogens cause chytridiomycosis, a skin disease plaguing amphibians worldwide [4, 6]. Recent estimates indicate that *Bd* has affected several hundred amphibian species and has been recorded on every continent except for Antarctica [9–11]. *Bsal* was more recently discovered in 2013 after a steep decline in fire salamander populations in Belgium [6].

**Funding:** This work was supported by the National Science Foundation (IOS 1827257), awarded to Lillian K Fritz-Laylin (LFL). [https://www.nsf.gov/funding/pgm\\_summ.jsp?pims\\_id=505480](https://www.nsf.gov/funding/pgm_summ.jsp?pims_id=505480) The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

**Competing interests:** The authors have declared that no competing interests exist.



**Fig 1. Life cycle of chytrid fungi.** As illustrated here with images of *Bsal*, chytrid fungi have a biphasic life cycle characterized by a stationary growth phase called a sporangium (top) and a motile dispersal phase called a zoospore (bottom). Images taken at 100X using differential interference contrast (DIC) microscopy.

<https://doi.org/10.1371/journal.pone.0240480.g001>

Management strategies for these pathogens have been developed and implemented in limited contexts, but implementation in real world settings remains a challenge. To develop better treatments, we need to understand the biology of chytrids in order to identify targets for drug development. However, studying the molecular mechanisms driving pathogenesis remains challenging due to the lack of genetic tools available for chytrid fungi. Electroporation protocols have been developed for *Bd* and *Bsal*, which could be used to deliver molecular payloads for genetics manipulation such as plasmids and/or CRISPR-Cas9 complexes [12]. The recent success in genetic manipulation of a related chytrid species, *Spizellomyces punctatus* (*Sp*), is a major breakthrough for our ability to study chytrid biology [8]. We and others are now striving to adapt this technology to *Bd* and *Bsal* to further our understanding of chytridiomycosis.

A key step to genetic tool development is the identification of methods for selection of successful transformants. The most commonly used selection method is antibiotic resistance:

incorporating a gene that provides specific drug resistance allows transformed cells to survive exposure to the antibiotic while all of the other cells are killed [13]. Distinct classes of antibiotics are commonly used for selection, each with their own molecular targets and corresponding organismal specificity. In addition to testing whether a given antibiotic kills cells of interest, it is important to pay attention to the effective concentration of each antibiotic. This is because a low concentration will not apply sufficient selective pressure and a high concentration could produce off-target effects and kill cells indiscriminately [14].

In this paper, we examine five antibiotics used in fungal and animal systems and identify the effective inhibitory concentration(s) necessary to prevent cell growth in liquid and solid media. Hygromycin, Blastidicin, and Puromycin inhibit protein translation in both bacterial and eukaryotic cells. Hygromycin inhibits protein synthesis by binding to the small ribosomal subunit and stabilizing the tRNA in the A site, preventing the progression of translation [15]. Blastidicin inhibits the terminating step of translation while Puromycin causes the ribosome to prematurely detach from mRNA [16, 17]. Although neomycin targets the prokaryotic 30S ribosomal subunit and causes codon misreading and mistranslation, it has been used in eukaryotes because of the similarity between mitochondrial and chloroplast ribosomes and bacterial ribosomes [18]. Zeocin intercalates in the DNA of both bacteria and eukaryotes and introduces double-stranded breaks, ultimately causing cell death [19].

## Results

To establish appropriate selection compounds for use with *Bd* and *Bsal*, we first identified antibiotics commonly used for selection with both mammalian and fungal systems. We chose five compounds (Hygromycin, Blastidicin, Puromycin, Zeocin, and Neomycin) to test based on the mechanism of action of each compound, their proven efficacy for use with both animal and fungal cells, and the availability of resistance genes (Table 1). We next tested the ability of these five compounds to inhibit the growth of *Bd* and *Bsal* cells in liquid culture. Although solid agar media is typically used for colony selection in chytrid and other fungi [8, 20, 21], we chose to use liquid culture to identify initial working concentrations because measuring zoospore release in liquid media is rapid and easily quantified.

To measure the effect of each antibiotic on *Bd* and *Bsal* growth, we added a wide range of antibiotic concentrations to cultures of age matched zoospores and allowed them to grow for one full life cycle: three (*Bd*) or four (*Bsal*) days. We then measured the concentration of released zoospores in each culture. Initial concentrations were selected based on known inhibitory concentrations for other organisms (Table 1) and spanned many orders of magnitude. Based on these preliminary experiments (not shown), we then identified possible working concentration ranges for each antibiotic in both species and tested intermediate concentrations using three biological replicates separated in time (Figs 2 and 3). To enable comparison of zoospore release from replicate experiments conducted on different days, we normalized counts for each replicate to its antibiotic-free control.

We identified antibiotic concentrations that consistently prevented growth in all three biological replicates—the successful concentrations are highlighted in orange in each figure. We found Hygromycin, Zeocin, Blastidicin and Neomycin could inhibit *Bd* growth in liquid culture (Fig 2), while all of the tested antibiotics inhibited *Bsal* growth (Fig 3). In *Bd*, Hygromycin has the lowest minimum inhibitory concentration (0.1 µg/ml), followed by Blastidicin (1 µg/ml), Zeocin (5 µg/ml), and Neomycin (600 µg/ml). Puromycin did not inhibit growth in *Bd* with the concentrations tested. In *Bsal*, Zeocin prevented growth at 1 µg/ml, followed by Blastidicin (2 µg/ml), Hygromycin (10 µg/ml), Puromycin (50 µg/ml), and Neomycin (250 µg/ml).

**Table 1. Antibiotic concentrations used to select for gene expression in select eukaryotes.** This table lists the key features of the antibiotics used in this study: the drug class, the target, known resistance genes, the current listed price per gram from Millipore Sigma, and the concentrations used in select eukaryotes. Species include representatives from plants (*Arabidopsis thaliana* and *Chlamydomonas reinhardtii*), protozoa (*Trypanosoma brucei*), amoebae (*Dictyostelium discoideum*), fungi (*Aspergillus spp.*, *Schizosaccharomyces pombe*, *Saccharomyces cerevisiae*), and animals (human) in addition to the two species tested in this study. The lowest concentrations of each antibiotic which inhibited growth in liquid and solid media for *Bd* and *Bsal* are listed from our findings in this study. These concentrations were used to calculate the cost per liter of growth media for both *Bd* and *Bsal*.

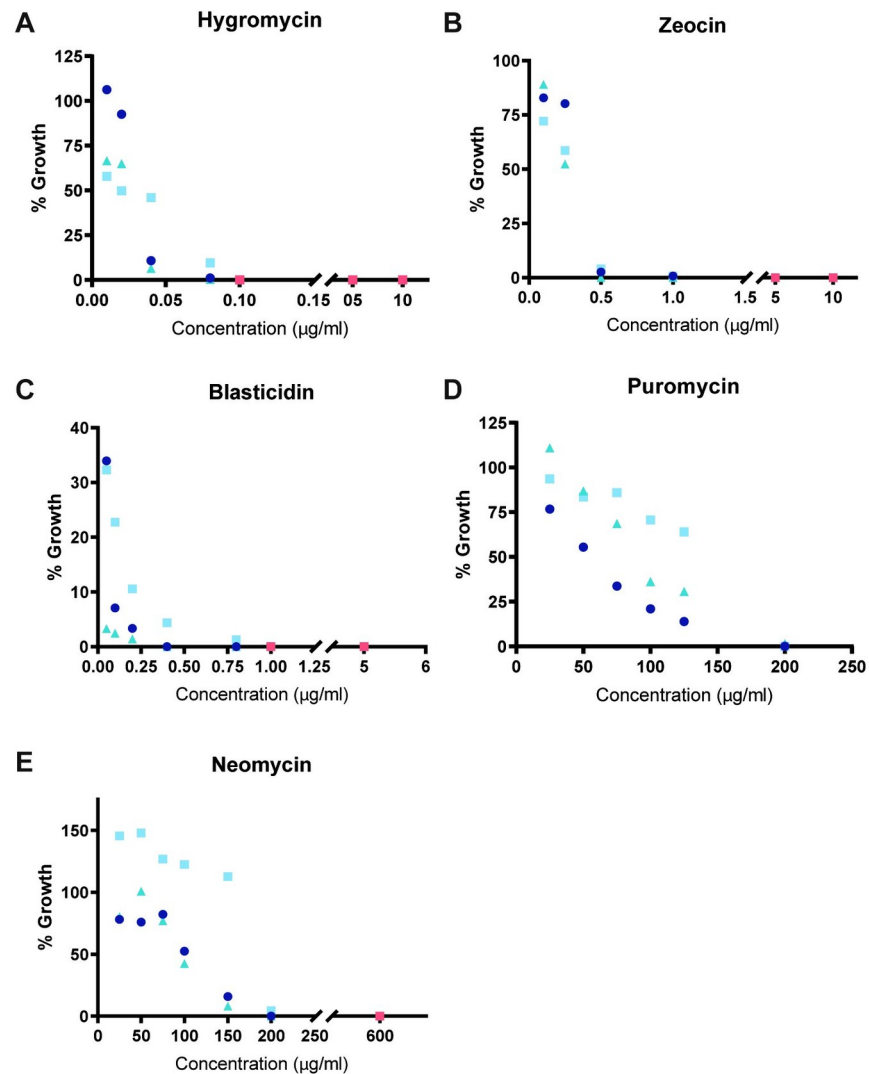
Drug Class	Target	Known Resistance Genes	List price per gram (MilliporeSigma)	Lowest drug conc. for growth inhibition for <i>Bd</i>	Cost per Liter for <i>Bd</i>	Lowest drug conc. for growth inhibition for <i>Bsal</i>	Cost per Liter for <i>Bsal</i>	Conc. for HeLa cells	Conc. for HESC	Conc. for Fibroblasts	Conc. for <i>Arabidopsis thaliana</i>	Conc. for <i>Dictyostelium discoideum</i>	Conc. for <i>Trypanosoma brucei</i>	Conc. for <i>Chlamydomonas reinhardtii</i>	Conc. for <i>Aspergillus spp</i>	Conc. for <i>S. pombe</i>	Conc. for <i>S. cerevisiae</i>
Neomycin	Ribosome [18]	<i>neo1</i>	\$1.93/g	Liquid: 600 µg/ml Solid: 1 mg/ml	Liquid: \$1.16/L Solid: \$1.93/L	Liquid: 250 µg/ml Solid: N/A	Liquid: \$0.48/L Solid: >\$193/L	-	-	-	-	-	-	300 µg/ml [22]	200–400 µg/ml [23]	0.375 g/L [24]	6.25 mM [25]
Hygromycin	Ribosome [15]	<i>hvg-1/hpt</i>	\$998/g	Liquid: 1 µg/ml Solid: 0.1 µg/ml	Liquid: \$1.00/L Solid: \$0.10/L	Liquid: 10 µg/ml Solid: 10 µg/ml	Liquid: \$9.98/L Solid: \$9.98/L	100–200 µg/ml [26,27]	40 µg/ml [28]	40 µg/ml [29]	15–50 µg/ml [30,31]	25–40 µg/ml [32]	5–50 µg/ml [33,34]	1–20 µg/ml [35]	100 µg/ml [36]	400 mg/L [37]	300 µg/ml [38]
Blasticidin	Ribosome [16]	<i>bsr, bbs, bsd</i>	\$6280/g	Liquid: 5 µg/ml Solid: 10 µg/ml	Liquid: \$31.25/L Solid: \$62.80	Liquid: 2 µg/ml Solid: 10 µg/ml	Liquid: \$12.56/L Solid: \$62.80/L	10–20 µg/ml [39,40]	2.0 µg/ml [41]	8 µg/ml [29]	10 µg/ml [42]	10 µg/ml [43,44]	2–10 µg/ml [45–47]	-	-	30 µg/ml [48,49]	10 mg/ml [50]
Puromycin	Ribosome [17]	<i>pac</i>	\$5340/g	Liquid: N/A Solid: 100 µg/ml	Liquid: >\$1068/L Solid: \$534/L	Liquid: 50 µg/ml Solid: N/A	Liquid: \$267/L Solid: >\$2670/L	1–2 µg/ml [51–53]	0.5–5 µg/ml [41,54,55]	2 µg/ml [29]	-	-	0.1 µg/ml [56]	-	-	-	*200 uM [57]
Zincin	DNA [19]	<i>ble</i>	\$177/g (Invitrogen)	Liquid: 10 µg/ml Solid: 10 µg/ml	Liquid: \$1.77/L Solid: \$1.77/L	Liquid: 1 µg/ml Solid: 10 µg/ml	Liquid: \$0.18/L Solid: \$1.77/L	50 µg/ml [58]	300 µg/ml [59]	800 µg/ml [29]	100 µg/ml [60]	100 mg/L [61]	-	5–15 µg/ml [62,63]	100–125 µg/ml [64]	150 mg/ml [65]	-

- no references were found.

\* the organism had to be made susceptible for the antibiotic to work.

‡ the *neo* resistance gene is also used for resistance to the drug G418 which was not tested in this study.

<https://doi.org/10.1371/journal.pone.0240480.t001>

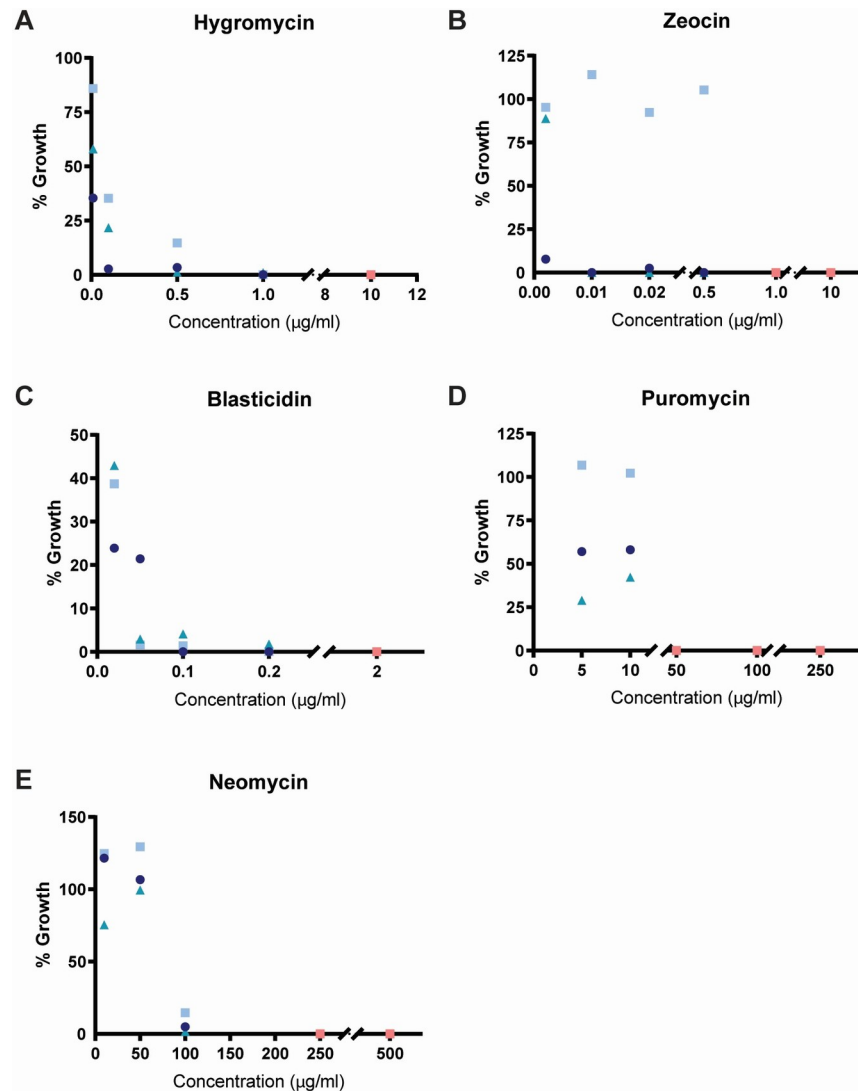


**Fig 2. Inhibition of *Bd* growth in liquid media.** Percent of *Bd* growth in liquid media supplemented with (A) Hygromycin, (B) Zeocin, (C) Blasticidin, (D) Puromycin, and (E) Neomycin as compared to an antibiotic free control for three temporally isolated replicates (circle, square, and triangle, shades of blue). Orange symbols indicate concentrations at which no growth occurred after three days in all three replicates.

<https://doi.org/10.1371/journal.pone.0240480.g002>

Having identified working concentrations of these compounds for use with liquid media, we next tested their efficacy on solid media. Growing cells on solid media allows for colony formation, which is useful for isolating successful and independent genetic transformants by “picking” colonies that grow under selection. To identify useful concentrations for selection on solid media, we inoculated zoospores on nutrient agar plates containing varying antibiotic concentrations. After a full growth cycle on selective media (three days for *Bd*, four days for *Bsal*), we compared zoospore release to antibiotic-free control cultures by flooding plates with water and looking for motile zoospores (S1 and S2 Videos). We defined successful concentrations as those which yielded no zoospore release in either replicate. We found at least one concentration for each antibiotic that prevented zoospore release in the timeframe of a typical growth cycle (Figs 4 and 5).

Because detection of colony formation often requires multiple growth cycles, we evaluated the efficiency of growth inhibition by growing plates with no zoospore release for 14 days. We



**Fig 3. Inhibition of *Bsal* growth in liquid media.** Percent of *Bsal* growth in liquid media supplemented with (A) Hygromycin, (B) Zeocin, (C) Blasticidin, (D) Puromycin, and (E) Neomycin as compared to an antibiotic free control for three temporally isolated replicates (circle, square, and triangle, shades of blue). Orange symbols indicate concentrations at which no growth occurred after four days in all three replicates.

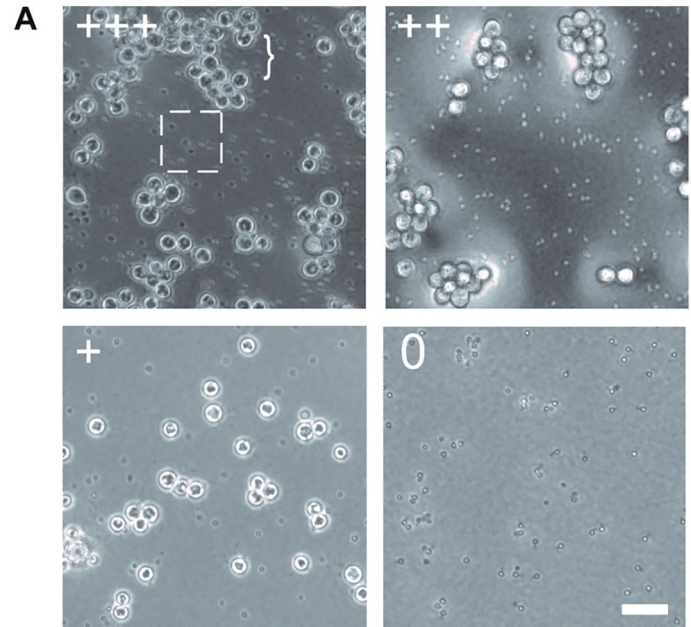
<https://doi.org/10.1371/journal.pone.0240480.g003>

found that all the tested antibiotics inhibited *Bd* growth on solid media, but only Hygromycin, Blasticidin and Zeocin inhibited growth in *Bsal*. For *Bd*, Hygromycin has the lowest minimum concentration at 0.1 µg/ml, with Blasticidin and Zeocin both following at 10 µg/ml, Puromycin at 100 µg/ml, and Neomycin at 1 mg/ml (Fig 4). In *Bsal*, Hygromycin, Blasticidin, and Zeocin all prevented growth for at least 14 days at a concentration of 10 µg/ml, while Puromycin and Neomycin did not prevent growth on solid media (Fig 5). The recommended concentrations for selection are highlighted in orange on the tables in both figures (Figs 4B and 5B).

## Discussion

This study identified drug concentrations that reproducibly inhibited *Bd* and *Bsal* growth in either liquid culture or on solid media. When a drug worked in both liquid culture and solid





**B**

Antibiotic	Concentration (µg/ml)	Rep 1	Rep 2
Hygromycin	0.01	++	++
	0.1	0	0
	1	0	0
	10	0	0
	100	0	0
	1000	0	0
Zeocin	0.1	+	++
	1	0	+
	10	0	0
	100	0	0
	1000	0	0
Blasticidin	0.01	++	++
	0.1	+	+
	1	0	0
	10	0	0
Puromycin	0.1	+++	+++
	1	+++	+++
	10	++	+++
	100	0	0
	1000	0	0
Neomycin	0.1	+++	+++
	1	+++	++
	10	+	+++
	100	+	+
	1000	0	0

**Fig 4. Inhibition of Bd growth on solid media.** (A) Examples of *Bd* growth after three days on antibiotic selection plates. The '+' demonstrates the relative zoospore activity of each plate compared to an antibiotic-free control plate. The box highlights zoospores, which appear as small dots while the bracket highlights sporangia. The zoospores in the '0' image are immotile (see [S1 Video](#)). Scale bar 50  $\mu\text{m}$ . (B) *Bd* growth on antibiotic selection plates. Concentrations highlighted in bold and orange are the lowest concentrations that prevent growth for at least 14 days post zoospore plating.

<https://doi.org/10.1371/journal.pone.0240480.g004>

media, the solid media typically required a higher concentration of antibiotic. This may be because of the additional minerals found in the agar not present in the liquid media [66]. Hygromycin, Zeocin, and Blastidicin worked well for both species and at concentrations within the typical range used for genetic selection in other species (Table 1). Puromycin and Neomycin were both able to inhibit growth of *Bd* and *Bsal*, but required higher concentrations than are used for animal cell lines. Although Hygromycin, Zeocin, and Blastidicin are all effective for preventing growth of *Bd* and *Bsal*, we recommend first using Hygromycin for genetic selection because it has been successfully used for selection of transformants in the nonpathogenic chytrid *Spizellomyces punctatus*, and is widely used for other fungal species [8, 36–38].

The ability to select for genetically transformed cells will allow for tractable genetic models to facilitate hypothesis testing in *Bd* and *Bsal*. The identification of useful selection agents and appropriate working concentrations is an important first step in developing genetic tools for use with *Bd* and *Bsal*. The natural step forward will be the design of selection cassettes, most commonly in the form of transformation plasmids. We look forward to the development of these and related molecular tools that will help us answer questions about the basic cell biology of chytrids, fungal evolution, and amphibian pathology.

## Methods

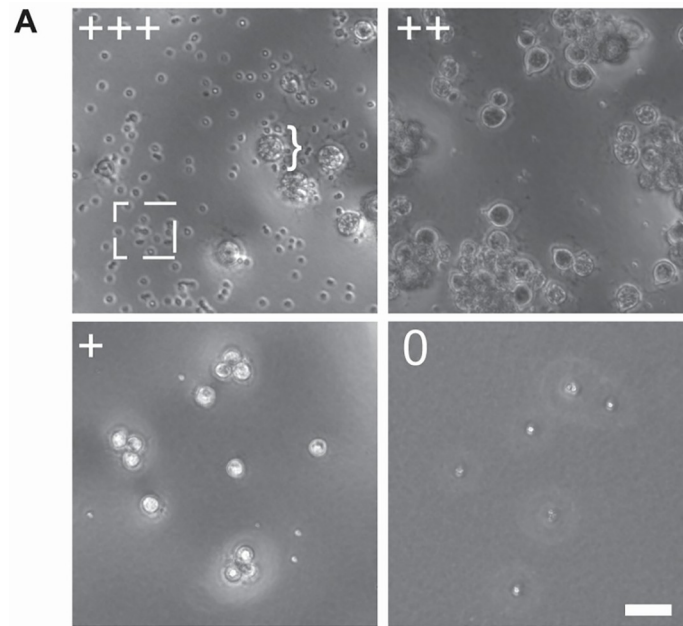
### Cell growth and synchronization

*Batrachochytrium dendrobatidis* (*Bd*) isolate JEL 423 was grown in 1% (w/v) tryptone (Apex Cat. 20–251) in tissue culture treated flasks (Cell Treat 229340) at 24°C for three days. *B. salamandrivorans* (*Bsal*) isolate AMFP 1 was grown in half-strength TGH liquid media (0.8% Tryptone, 0.2% gelatin hydrolysate, 0.1% lactose (w/v) in tissue culture treated flasks at 15°C for four days [67]. For both species, we synchronized the release of motile zoospores by gently washing the flask three times with fresh growth media and then incubating with 10 mL of media for 2 hours. Age matched zoospores were then collected by centrifugation at 2000 rcf for 5 mins, resuspended in media, counted, and used for experiments as outlined below.

### Drug treatments and quantitation for cells grown in liquid media

Neomycin (Fisher Cat. AAJ67011AE), Hygromycin B (Fisher Cat. AAJ60681MC), Blastidicin (Fisher Cat. BP2647100), Puromycin (Fisher Cat. BP2956100), and Zeocin (Fisher Cat. AAJ671408EQ), were screened for growth inhibition of *Bd* and *Bsal*. Cells were diluted to a starting concentration of  $5 \times 10^5$  cells/mL and 250  $\mu\text{L}$  of cells were added to each well of a sterile tissue culture treated 24-well plate (Cell Treat 229123). 250  $\mu\text{L}$  of appropriately diluted antibiotics and matched carrier controls were added to each well and mixed thoroughly. Plates were sealed with parafilm and grown at either 24°C for three days (*Bd*), or 15°C for four days (*Bsal*). For each of three biological replicates spaced in time, the concentration of released zoospores was estimated using the average of two independent hemocytometer counts. Zoospore concentrations were normalized to the no drug control and data plotted using Prism (GraphPad v8).





**B**

Antibiotic	Concentration µg/ml	Rep 1	Rep 2
Hygromycin	0.001	+++	+++
	0.01	+++	++
	0.1	++	+
	1	0	0
	10	0	0
Zeocin	0.01	++	++
	0.1	0	0
	1	0	0
	10	0	0
	100	0	0
Blasticidin	0.01	+	++
	0.1	+	0
	1	0	0
	10	0	0
	100	0	0
Puromycin	0.1	+++	+++
	1	++	+++
	10	++	+
	100	+	0
	500	0	0
Neomycin	0.1	++	++
	1	++	++
	10	++	+++
	100	+	+
	1000	0	0

**Fig 5. Inhibition of *Bsal* growth on solid media.** (A) Examples of *Bsal* growth after four days on antibiotic selection plates. The '+' demonstrates the relative zoospore activity of each plate compared to a no antibiotic control plate. The box highlights zoospores, which appear as small dots while the bracket highlights sporangia. The zoospores in the '0' image are immotile (see [S1 Video](#)). Scale bar 50  $\mu\text{m}$ . (B) *Bsal* growth on antibiotic selection plates. Concentrations highlighted in bold and orange are the lowest concentrations that prevent growth for at least 14 days post zoospore plating.

<https://doi.org/10.1371/journal.pone.0240480.g005>

## Drug treatments and quantitation for cells grown on solid media

We added 1% agar to 50 mL batches of 1% tryptone (w/v) and half-strength TGHl then autoclaved. Each antibiotic was added to a separate, pre-cooled, 50 mL batch of media, and 10 mL of the solution added to one of five 15 mm<sup>2</sup> plates (VWR 25384–090) and allowed to solidify. Equal volume of appropriate carrier liquid was added to the pre-cooled 50 mL batch of agar-media to create control plates. Plates were wrapped in parafilm and aluminum foil, and stored at 4°C. Plates were inoculated by evenly spreading  $5.0 \times 10^6$  zoospores across the agar and incubated at 24°C for three days (*Bd*) or 15°C for four days (*Bsal*). Three control plates were used per replicate to ensure a point of comparison if one were to be contaminated. Zoospore release was evaluated by imaging each plate for 20 seconds at one second intervals using a Nikon Ti2-E inverted microscope equipped with 10x PlanApo objective and sCMOS 4mp camera (PCO Panda) using white LED transmitted light. Approximate zoospore activity was assessed as: 0 (no visible zoospores), + (< 25% zoospore activity of control plates lacking antibiotic), ++ (~50% zoospore activity of control plates), or +++ (equivalent zoospore activity to control plates). To determine the lowest antibiotic concentration that could completely inhibit growth, plates that yielded "0" growth were allowed to grow for 14 days at the appropriate incubation temperature and reassessed as above.

## Supporting information

**S1 Video. *Bsal* zoospores with zero growth.** Zoospores grown on antibiotic selection plates are labeled "0" if no zoospores are released or zoospores showed no growth and are immotile. (MP4)

**S2 Video. *Bsal* zoospores with "+++" growth.** Zoospores grown on antibiotic selection plates are labeled "+++" if the zoospore release is comparable to the no antibiotic control. (MP4)

## Author Contributions

**Conceptualization:** Lillian K. Fritz-Laylin.

**Data curation:** Kristyn A. Robinson.

**Formal analysis:** Kristyn A. Robinson, Mallory Dunn.

**Funding acquisition:** Lillian K. Fritz-Laylin.

**Investigation:** Kristyn A. Robinson, Mallory Dunn, Shane P. Hussey.

**Methodology:** Mallory Dunn, Shane P. Hussey.

**Project administration:** Lillian K. Fritz-Laylin.

**Supervision:** Lillian K. Fritz-Laylin.

**Validation:** Kristyn A. Robinson.

**Visualization:** Kristyn A. Robinson.

**Writing – original draft:** Kristyn A. Robinson, Lillian K. Fritz-Laylin.

**Writing – review & editing:** Kristyn A. Robinson, Mallory Dunn, Shane P. Hussey, Lillian K. Fritz-Laylin.

## References

1. Grossart H-P, Wurzbacher C, James TY, Kagami M. Discovery of dark matter fungi in aquatic ecosystems demands a reappraisal of the phylogeny and ecology of zoosporic fungi. *Fungal Ecol.* 2016 Feb; 19:28–38.
2. Kagami M, Miki T, Takimoto G. Mycoloop: chytrids in aquatic food webs. *Front Microbiol* [Internet]. 2014 Apr 22 [cited 2020 Apr 24]; 5. Available from: <http://journal.frontiersin.org/article/10.3389/fmicb.2014.00166/abstract>
3. Gleason FH, Küpper FC, Amon JP, Picard K, Gachon CMM, Marano AV, et al. Zoosporic true fungi in marine ecosystems: a review. *Mar Freshw Res.* 2011 May 19; 62(4):383–93.
4. Longcore JE, Pessier AP, Nichols DK. *Batrachochytrium dendrobatidis* gen. et sp. nov., a chytrid pathogenic to amphibians. *Mycologia.* 1999 Mar; 91(2):219–27.
5. Berger L, Hyatt A, Speare R, Longcore J. Life cycle stages of the amphibian chytrid *Batrachochytrium dendrobatidis*. *Dis Aquat Organ.* 2005; 68:51–63. <https://doi.org/10.3354/dao068051> PMID: 16465834
6. Martel A, Spitzen-van der Sluijs A, Blooi M, Bert W, Ducatelle R, Fisher MC, et al. *Batrachochytrium salamandrivorans* sp. nov. causes lethal chytridiomycosis in amphibians. *Proc Natl Acad Sci.* 2013 Sep 17; 110(38):15325–9. <https://doi.org/10.1073/pnas.1307356110> PMID: 24003137
7. Fritz-Laylin LK, Lord SJ, Mullins RD. WASP and SCAR are evolutionarily conserved in actin-filled pseudopod-based motility. *J Cell Biol.* 2017 Jun 5; 216(6):1673–88. <https://doi.org/10.1083/jcb.201701074> PMID: 28473602
8. Medina EM, Robinson KA, Bellingham-Johnston K, Ianiri G, Laplante C, Fritz-Laylin LK, et al. Genetic transformation of *Spizellomyces punctatus*, a resource for studying chytrid biology and evolutionary cell biology. Rokas A, Baldwin IT, Stearns T, editors. *eLife.* 2020 May 11; 9:e52741. <https://doi.org/10.7554/eLife.52741> PMID: 32392127
9. Bellard C, Genovesi P, Jeschke JM. Global patterns in threats to vertebrates by biological invasions. *Proc R Soc B Biol Sci.* 2016 Jan 27; 283(1823):20152454.
10. Olson DH, Aanensen DM, Ronnenberg KL, Powell CI, Walker SF, Bielby J, et al. Mapping the Global Emergence of *Batrachochytrium dendrobatidis*, the Amphibian Chytrid Fungus. Stajich JE, editor. *PLoS ONE.* 2013 Feb 27; 8(2):e56802. <https://doi.org/10.1371/journal.pone.0056802> PMID: 23463502
11. Scheele BC, Pasmans F, Skerratt LF, Berger L, Martel A, Beukema W, et al. Amphibian fungal panzootic causes catastrophic and ongoing loss of biodiversity. *Science.* 2019 Mar 29; 363(6434):1459–63. <https://doi.org/10.1126/science.aav0379> PMID: 30923224
12. Swafford AJM, Hussey SP, Fritz-Laylin LK. High-efficiency electroporation of chytrid fungi. *Sci Rep.* 2020 Dec; 10(1):15145. <https://doi.org/10.1038/s41598-020-71618-2> PMID: 32934254
13. Smith HO, Danner DB, Deich RA. Genetic Transformation. *Annu Rev Biochem.* 1981; 50(1):41–68.
14. Stepanenko AA, Heng HH. Transient and stable vector transfection: Pitfalls, off-target effects, artifacts. *Mutat Res Mutat Res.* 2017 Jul 1; 773:91–103. <https://doi.org/10.1016/j.mrev.2017.05.002> PMID: 28927539
15. Borovinskaya MA, Shoji S, Fredrick K, Cate JHD. Structural basis for hygromycin B inhibition of protein biosynthesis. *RNA.* 2008 Jun 20; 14(8):1590–9. <https://doi.org/10.1261/rna.1076908> PMID: 18567815
16. Svidritskiy E, Ling C, Ermolenko DN, Korostelev AA. Blastocidin S inhibits translation by trapping deformed tRNA on the ribosome. *Proc Natl Acad Sci.* 2013 Jul 23; 110(30):12283–8. <https://doi.org/10.1073/pnas.1304922110> PMID: 23824292
17. Pestka S. Inhibitors of Ribosome Functions. 1971. 1971; 25:487–562.
18. Mehta R, Champney WS. Neomycin and Paromomycin Inhibit 30S Ribosomal Subunit Assembly in *Staphylococcus aureus*. *Curr Microbiol.* 2003 Sep 1; 47(3):237–43. <https://doi.org/10.1007/s00284-002-3945-9> PMID: 14570276
19. Chankova SG, Dimova E, Dimitrova M, Bryant PE. Induction of DNA double-strand breaks by zeocin in *Chlamydomonas reinhardtii* and the role of increased DNA double-strand breaks rejoining in the formation of an adaptive response. *Radiat Environ Biophys.* 2007 Oct 10; 46(4):409–16. <https://doi.org/10.1007/s00411-007-0123-2> PMID: 17639449
20. de Groot MJA, Bundock P, Hooykaas PJJ, Beijersbergen AGM. *Agrobacterium tumefaciens*-mediated transformation of filamentous fungi. 1998; 16:4.

21. Bundock P, den Dulk-Ras A, Beijersbergen A, Hooykaas PJ. Trans-kingdom T-DNA transfer from *Agrobacterium tumefaciens* to *Saccharomyces cerevisiae*. *EMBO J*. 1995 Jul 3; 14(13):3206–14. PMID: [7621833](https://pubmed.ncbi.nlm.nih.gov/7621833/)
22. Hasnain SE, Manavathu EK, Leung WC. DNA-mediated transformation of *Chlamydomonas reinhardtii* cells: use of aminoglycoside 3'-phosphotransferase as a selectable marker. *Mol Cell Biol*. 1985 Dec; 5(12):3647–50. <https://doi.org/10.1128/mcb.5.12.3647> PMID: [3018525](https://pubmed.ncbi.nlm.nih.gov/3018525/)
23. Dong Y, Cui C-B, Li C-W, Hua W, Wu C-J, Zhu T-J, et al. Activation of Dormant Secondary Metabolite Production by Introducing Neomycin Resistance into the Deep-Sea Fungus, *Aspergillus versicolor* ZBY-3. *Mar Drugs*. 2014 Jul 29; 12(8):4326–52.
24. Bureik M, Bruck N, Hubel K, Bernhardt R. The human mineralocorticoid receptor only partially differentiates between different ligands after expression in fission yeast. *FEMS Yeast Res*. 2005 Apr; 5(6–7):627–33. <https://doi.org/10.1016/j.femsyr.2004.12.007> PMID: [15780662](https://pubmed.ncbi.nlm.nih.gov/15780662/)
25. Shimma Y-I, Uno I. Isolation and characterization of neomycin-sensitive mutants in *Saccharomyces cerevisiae*. *J Gen Microbiol*. 1990; 136:1753–61.
26. Jiang H, Su ZZ, Lin JJ, Goldstein NI, Young CS, Fisher PB. The melanoma differentiation associated gene *mda-7* suppresses cancer cell growth. *Proc Natl Acad Sci*. 1996 Aug 20; 93(17):9160–5. <https://doi.org/10.1073/pnas.93.17.9160> PMID: [8799171](https://pubmed.ncbi.nlm.nih.gov/8799171/)
27. Buchschacher GL, Panganiban AT. Human Immunodeficiency Virus Vectors for Inducible Expression of Foreign Genes. *J VIROL*. 1992; 66:9.
28. Sakurai K, Shimoji M, Tahimic CGT, Aiba K, Kawase E, Hasegawa K, et al. Efficient integration of transgenes into a defined locus in human embryonic stem cells. *Nucleic Acids Res*. 2010 Apr; 38(7):e96. <https://doi.org/10.1093/nar/gkp1234> PMID: [20071742](https://pubmed.ncbi.nlm.nih.gov/20071742/)
29. Sato M, Ohtsuka M, Miura H, Miyoshi K, Watanabe S. Determination of the Optimal Concentration of Several Selective Drugs Useful for Generating Multi-Transgenic Porcine Embryonic Fibroblasts. *Reprod Domest Anim*. 2012; 47(5):759–65. <https://doi.org/10.1111/j.1439-0531.2011.01964.x> PMID: [22136322](https://pubmed.ncbi.nlm.nih.gov/22136322/)
30. Harrison SJ, Mott EK, Parsley K, Aspinall S, Gray JC, Cottage A. A rapid and robust method of identifying transformed *Arabidopsis thaliana* seedlings following floral dip transformation. *Plant Methods*. 2006; 2(1):19.
31. Rashid A. Comparison of a kanamycin versus hygromycin resistance gene in transgenic plant selection of *Arabidopsis thaliana* L. *J Cell Sci Mutat* [Internet]. 2017 [cited 2020 Apr 24]; 01(01). Available from: <http://www.alliedacademies.org/articles/comparison-of-a-kanamycin-versus-hygromycin-resistance-gene-in-transgenic-plant-selection-of-arabidopsis-thaliana-l.html>
32. Egelhoff TT, Brown SS, Manstein DJ, Spudich JA. Hygromycin resistance as a selectable marker in *Dicotylelium discoideum*. *Mol Cell Biol*. 1989 May; 9(5):1965–8. <https://doi.org/10.1128/mcb.9.5.1965> PMID: [2546056](https://pubmed.ncbi.nlm.nih.gov/2546056/)
33. Biebinger S, Elizabeth Wirtz L, Lorenz P, Christine Clayton. Vectors for inducible expression of toxic gene products in bloodstream and procyclic *Trypanosoma brucei*. *Mol Biochem Parasitol*. 1997 Mar; 85(1):99–112. [https://doi.org/10.1016/s0166-6851\(96\)02815-0](https://doi.org/10.1016/s0166-6851(96)02815-0) PMID: [9108552](https://pubmed.ncbi.nlm.nih.gov/9108552/)
34. Wirtz E, Leal S, Ochatt C, Cross GeorgeAM. A tightly regulated inducible expression system for conditional gene knock-outs and dominant-negative genetics in *Trypanosoma brucei*. *Mol Biochem Parasitol*. 1999 Mar; 99(1):89–101. [https://doi.org/10.1016/s0166-6851\(99\)00002-x](https://doi.org/10.1016/s0166-6851(99)00002-x) PMID: [10215027](https://pubmed.ncbi.nlm.nih.gov/10215027/)
35. Berthold P, Schmitt R, Mages W. An Engineered *Streptomyces hygroscopicus* aph 7" Gene Mediates Dominant Resistance against Hygromycin B in *Chlamydomonas reinhardtii*. *Protist*. 2002 Dec; 153(4):401–12. <https://doi.org/10.1078/14344610260450136> PMID: [12627869](https://pubmed.ncbi.nlm.nih.gov/12627869/)
36. Cullen D, Leong SA, Wilson LJ, Henner DJ. Transformation of *Aspergillus nidulans* with the hygromycin-resistance gene, *hph*. *Gene*. 1987 Jan; 57(1):21–6. [https://doi.org/10.1016/0378-1119\(87\)90172-7](https://doi.org/10.1016/0378-1119(87)90172-7) PMID: [3322945](https://pubmed.ncbi.nlm.nih.gov/3322945/)
37. Brown S, Lorenz A. Single-step Marker Switching in *Schizosaccharomyces pombe* Using a Lithium Acetate Transformation Protocol. *BIO-Protoc* [Internet]. 2016 [cited 2020 Jan 23]; 6(24). Available from: <https://bio-protocol.org/e2075>
38. Goldstein AL, McCusker JH. Three new dominant drug resistance cassettes for gene disruption in *Saccharomyces cerevisiae*. *Yeast*. 1999; 15(14):1541–53. [https://doi.org/10.1002/\(SICI\)1097-0061\(199910\)15:14<1541::AID-YEA476>3.0.CO;2-K](https://doi.org/10.1002/(SICI)1097-0061(199910)15:14<1541::AID-YEA476>3.0.CO;2-K) PMID: [10514571](https://pubmed.ncbi.nlm.nih.gov/10514571/)
39. Jin Q, Marsh J, Cornetta K, Alkhatib G. Resistance to human immunodeficiency virus type 1 (HIV-1) generated by lentivirus vector-mediated delivery of the CCR5Δ32 gene despite detectable expression of the HIV-1 co-receptors. *J Gen Virol*. 2008 Oct; 89(Pt 10):2611–21. <https://doi.org/10.1099/vir.0.2008/003624-0> PMID: [18796731](https://pubmed.ncbi.nlm.nih.gov/18796731/)

40. Cheng N, He R, Tian J, Ye PP, Ye RD. Cutting Edge: TLR2 Is a Functional Receptor for Acute-Phase Serum Amyloid A. *J Immunol*. 2008 Jul 1; 181(1):22–6. <https://doi.org/10.4049/jimmunol.181.1.22> PMID: 18566366
41. Moore JC, Atze K, Yeung PL, Toro-Ramos AJ, Camarillo C, Thompson K, et al. Efficient, high-throughput transfection of human embryonic stem cells. *Stem Cell Res Ther*. 2010 Jul 26; 1(3):23. <https://doi.org/10.1186/scri23> PMID: 20659329
42. Tamura K, Kimura M, Yamaguchi I. Blastidicin S Deaminase Gene (BSD): a new selection marker gene for transformation of *Arabidopsis thaliana* and *Nicotiana tabacum*. *Biosci Biotech Biochem*. 1995; 59(12):2336–8.
43. Thompson CRL, Kay RR. The Role of DIF-1 Signaling in Dictyostelium Development. *Mol Cell*. 2000 Dec; 6(6):1509–14. [https://doi.org/10.1016/s1097-2765\(00\)00147-7](https://doi.org/10.1016/s1097-2765(00)00147-7) PMID: 11163223
44. Li G, Alexander H, Schneider N, Alexander S. Molecular basis for resistance to the anticancer drug cisplatin in Dictyostelium. 2019; 9.
45. Brooks DR, McCulloch R, Coombs GH, Mottram JC. Stable transformation of trypanosomatids through targeted chromosomal integration of the selectable marker gene encoding blastidicin S deaminase. *FEMS Microbiol Lett*. 2000 May; 186(2):287–91. <https://doi.org/10.1111/j.1574-6968.2000.tb09119.x> PMID: 10802186
46. Roper JR, Güther MLS, MacRae JI, Prescott AR, Hallyburton I, Acosta-Serrano A, et al. The Suppression of Galactose Metabolism in Procyclic Form *Trypanosoma brucei* Causes Cessation of Cell Growth and Alters Procyclin Glycoprotein Structure and Copy Number. *J Biol Chem*. 2005 May 20; 280(20):19728–36. <https://doi.org/10.1074/jbc.M502370200> PMID: 15767252
47. Erben ED, Fadda A, Lueong S, Hoheisel JD, Clayton C. A Genome-Wide Tethering Screen Reveals Novel Potential Post-Transcriptional Regulators in *Trypanosoma brucei*. Tschudi C, editor. *Pathog PLoS*. 2014 Jun 12; 10(6):e1004178.
48. Zhang X-R, He J-B, Wang Y-Z, Du L-L. A Cloning-Free Method for CRISPR/Cas9-Mediated Genome Editing in Fission Yeast. *G3amp58 GenesGenomesGenetics*. 2018 Jun; 8(6):2067–77.
49. Kimura M, Kamakura T, Zhou Tao Q, Kaneko I, Yamaguchi I. Cloning of the blastidicin S deaminase gene (BSD) from *Aspergillus terreus* and its use as a selectable marker for *Schizosaccharomyces pombe* and *Pyricularia oryzae*. *Mol Gen Genet MGG*. 1994 Jan 1; 242(2):121–9. <https://doi.org/10.1007/BF00391004> PMID: 8159161
50. Fukuda H, Kizaki Y. A new transformation system of *Saccharomyces cerevisiae* with blastidicin S deaminase gene. 1999; 3.
51. Ukekawa R, Miki K, Fujii M, Hirano H, Ayusawa D. Accumulation of multiple forms of lamin A with down-regulation of FACE-1 suppresses growth in senescent human cells. *Genes Cells*. 2007 Mar; 12(3):397–406. <https://doi.org/10.1111/j.1365-2443.2007.01057.x> PMID: 17352743
52. Huang J, Dibble CC, Matsuzaki M, Manning BD. The TSC1-TSC2 Complex Is Required for Proper Activation of mTOR Complex 2. *Mol Cell Biol*. 2008 Jun 15; 28(12):4104–15. <https://doi.org/10.1128/MCB.00289-08> PMID: 18411301
53. Grueneberg DA, Degot S, Pearlberg J, Li W, Davies JE, Baldwin A, et al. Kinase requirements in human cells: I. Comparing kinase requirements across various cell types. *Proc Natl Acad Sci*. 2008 Oct 28; 105(43):16472–7. <https://doi.org/10.1073/pnas.0808019105> PMID: 18948591
54. Yang C, Przyborski S, Cooke MJ, Zhang X, Stewart R, Anyfantis G, et al. A Key Role for Telomerase Reverse Transcriptase Unit in Modulating Human Embryonic Stem Cell Proliferation, Cell Cycle Dynamics, and In Vitro Differentiation. *STEM CELLS*. 2008; 26(4):850–63. <https://doi.org/10.1634/stemcells.2007-0677> PMID: 18203676
55. Drobinskaya I, Linn T, Šarić T, Bretzel RG, Bohlen H, Hescheler J, et al. Scalable Selection of Hepatocyte- and Hepatocyte Precursor-Like Cells from Culture of Differentiating Transgenically Modified Murine Embryonic Stem Cells. *STEM CELLS*. 2008; 26(9):2245–56. <https://doi.org/10.1634/stemcells.2008-0387> PMID: 18556507
56. Niemirowicz GT, Cazzulo JJ, Álvarez VE, Bouvier LA. Simplified inducible system for *Trypanosoma brucei*. *PLoS ONE* [Internet]. 2018 Oct 11 [cited 2020 Apr 24]; 13(10). Available from: <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC6181392/>
57. Cary GA, Yoon SH, Torres CG, Wang K, Hays M, Ludlow C, et al. Identification and characterization of a drug-sensitive strain enables puromycin-based translational assays in *Saccharomyces cerevisiae*. *Yeast*. 2014; 31(5):167–78. <https://doi.org/10.1002/yea.3007> PMID: 24610064
58. Mesnil M, Piccoli C, Tiraby G, Willecke K, Yamasaki H. Bystander killing of cancer cells by herpes simplex virus thymidine kinase gene is mediated by connexins. *Proc Natl Acad Sci*. 1996 Mar 5; 93(5):1831–5. <https://doi.org/10.1073/pnas.93.5.1831> PMID: 8700844

59. Chow MZY, Geng L, Kong C-W, Keung W, Fung JC-K, Boheler KR, et al. Epigenetic Regulation of the Electrophysiological Phenotype of Human Embryonic Stem Cell-Derived Ventricular Cardiomyocytes: Insights for Driven Maturation and Hypertrophic Growth. *Stem Cells Dev.* 2013 Oct; 22(19):2678–90. <https://doi.org/10.1089/scd.2013.0125> PMID: 23656529
60. Van Hove J, Fouquaert E, Smith DF, Proost P, Van Damme EJM. Lectin activity of the nucleocytoplasmic EUL protein from *Arabidopsis thaliana*. *Biochem Biophys Res Commun.* 2011 Oct; 414(1):101–5. <https://doi.org/10.1016/j.bbrc.2011.09.031> PMID: 21945438
61. Nicolussi A, Dunn JD, Mlynek G, Bellei M, Zamocky M, Battistuzzi G, et al. Secreted heme peroxidase from *Dictyostelium discoideum*: Insights into catalysis, structure, and biological role. *J Biol Chem.* 2018 Jan 26; 293(4):1330–45. <https://doi.org/10.1074/jbc.RA117.000463> PMID: 29242189
62. Liu P, Lechtreck KF. The Bardet–Biedl syndrome protein complex is an adapter expanding the cargo range of intraflagellar transport trains for ciliary export. *Proc Natl Acad Sci.* 2018 Jan 30; 115(5):E934–43. <https://doi.org/10.1073/pnas.1713226115> PMID: 29339469
63. López-Paz C, Liu D, Geng S, Umen JG. Identification of *Chlamydomonas reinhardtii* endogenous genic flanking sequences for improved transgene expression. *Plant J.* 2017 Dec; 92(6):1232–44. <https://doi.org/10.1111/tbj.13731> PMID: 28980350
64. Han G, Shao Q, Li C, Zhao K, Jiang L, Fan J, et al. An efficient *Agrobacterium*-mediated transformation method for aflatoxin generation fungus *Aspergillus flavus*. *J Microbiol.* 2018 May; 56(5):356–64. <https://doi.org/10.1007/s12275-018-7349-3> PMID: 29721833
65. Benko Z, Zhao RY. Zeocin for selection of bleMX6 resistance in fission yeast. *BioTechniques.* 2011 Jul 1; 51(1):57–60. <https://doi.org/10.2144/000113706> PMID: 21781055
66. Tam JP, Lu Y-A, Yang J-L. Correlations of Cationic Charges with Salt Sensitivity and Microbial Specificity of Cystine-stabilized  $\beta$ -Strand Antimicrobial Peptides. *J Biol Chem.* 2002 Dec 27; 277(52):50450–6. <https://doi.org/10.1074/jbc.M208429200> PMID: 12399464
67. Robinson KA, Pereira KE, Bletz MC, Carter ED, Gray MJ, Piovia-Scott J, et al. Isolation and maintenance of *Batrachochytrium salamandrivorans* cultures. *Dis Aquat Organ.* 2020 Jun 18; 140:1–11. <https://doi.org/10.3354/dao03488> PMID: 32618283