RESEARCH ARTICLE



Efficacy of APX2039 in a Rabbit Model of Cryptococcal Meningitis

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ABSTRACT Cryptococcal Meningitis (CM) is uniformly fatal if not treated, and treatment options are limited. We previously reported on the activity of APX2096, the prodrug of the novel Gwt1 inhibitor APX2039, in a mouse model of CM. Here, we investigated the efficacy of APX2039 in mouse and rabbit models of CM. In the mouse model, the controls had a mean lung fungal burden of 5.95 log₁₀ CFU/g, whereas those in the fluconazole-, amphotericin B-, and APX2039-treated mice were 3.56, 4.59, and 1.50 log₁₀ CFU/g, respectively. In the brain, the control mean fungal burden was 7.97 log₁₀ CFU/g, while the burdens were 4.64, 7.16, and 1.44 \log_{10} CFU/g for treatment with fluconazole, amphotericin B, and APX2039, respectively. In the rabbit model of CM, the oral administration of APX2039 at 50 mg/kg of body weight twice a day (BID) resulted in a rapid decrease in the cerebrospinal fluid (CSF) fungal burden, and the burden was below the limit of detection by day 10 postinfection. The effective fungicidal activity (EFA) was $-0.66 \log_{10}$ CFU/mL/day, decreasing from an average of 4.75 log₁₀ CFU/mL to 0 CFU/mL, over 8 days of therapy, comparing favorably with good clinical outcomes in humans associated with reductions of the CSF fungal burden of $-0.4 \log_{10}$ CFU/mL/day, and, remarkably, 2-fold the EFA of amphotericin B deoxycholate in this model ($-0.33 \log_{10}$ CFU/mL/day). A total drug exposure of the area under the concentration-time curve from 0 to 24 h (AUC $_{0-24}$) of 25 to 50 mg · h/L of APX2039 resulted in near-maximal antifungal activity. These data support the further preclinical and clinical evaluation of APX2039 as a new oral fungicidal monotherapy for the treatment of CM.

IMPORTANCE Cryptococcal meningitis (CM) is a fungal disease with significant global morbidity and mortality. The gepix Gwt1 inhibitors are a new class of antifungal drugs. Here, we demonstrated the efficacy of APX2039, the second member of the gepix class, in rabbit and mouse models of cryptococcal meningitis. We also analyzed the drug levels in the blood and cerebrospinal fluid in the highly predictive rabbit model and built a mathematical model to describe the behavior of the drug with respect to the elimination of the fungal pathogen. We demonstrated that the oral administration of APX2039 resulted in a rapid decrease in the CSF fungal burden, with an effective fungicidal activity of $-0.66 \log_{10}$ CFU/mL/day, comparing favorably with good clinical outcomes in humans associated with reductions of $-0.4 \log_{10}$ CFU/mL/day. The drug APX2039 had good penetration of the central nervous system and is an excellent candidate for future clinical testing in humans for the treatment of CM.

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ryptococcal Meningitis (CM) is a significant source of global morbidity and mortality. Current estimates are 220,000 cases and 181,000 deaths per year in individuals living with HIV/AIDS (1). The majority of infections are in Africa, but a significant number of cases continue to occur outside Africa due to progressively enlarging populations of immunocompromised hosts. Even in resource-rich health care systems, up to 15% of CM cases result in death (2), and this rate is much higher in resource-poor health care systems (1). The current treatment guidelines from the WHO include 7 days of intravenous amphotericin B deoxycholate (AMB) combined with oral flucytosine, followed by consolidation and then maintenance therapy with fluconazole (FLU) (3). Recently, with the report of the AMBITION study, the WHO also recommended induction therapy using a single dose of AmBisome (liposomal amphotericin B) at 10 mg/kg of body weight followed by 14 days of flucytosine at 100 mg/ kg/day and fluconazole at 1,200 mg/day (4). In contrast, in resource-rich countries, a combination of a polyene plus flucytosine for the treatment of CM for 2 weeks remains the standard of care (5). Both amphotericin B and fluconazole target portions of the ergosterol pathway of Cryptococcus, while flucytosine inhibits RNA/DNA synthesis. Although effective, this combination poses problems of accessibility, compliance, toxicity, and clinical failures. Thus, there is an urgent need for new antifungal therapies that can effectively treat CM and especially oral drugs that can be more readily administered for induction, consolidation, and maintenance therapy.

Manogepix (APX001A) is the first compound in the new gepix class that demonstrates broad-spectrum *in vitro* antifungal activity (6). The *N*-phosphonooxymethylene prodrug fosmanogepix (APX001) demonstrated significant activity in a wide array of fungal infection models and is currently in clinical development for the treatment of *Candida, Aspergillus,* and rare mold infections (ClinicalTrials.gov identifiers NCT04148287, NCT03604705, and NCT04240886) (6). The cellular target of manogepix is the Gwt1 protein, which catalyzes an early step in glycosylphosphatidylinositol (GPI) anchor biosynthesis (7), from which the new class name "gepix" is derived. This process is important for anchoring proteins to the cell walls of fungi. Manogepix, the active moiety of the prodrug fosmanogepix, does not inhibit PIGW, the closest human homolog, even at concentrations >100-fold higher than the 50% inhibitory concentration (IC₅₀) for inhibition of the *Candida albicans* and *Aspergillus fumigatus* Gwt1 proteins (7).

We previously reported on the efficacy of fosmanogepix and the analogous N-phosphonooxymethylene prodrugs of two other Gwt1 inhibitors, APX2020 and APX2039 (prodrugs APX2097 and APX2096, respectively), in murine models of cryptococcosis (8). In those studies, APX2096 was dosed at 60 mg/kg intraperitoneally (i.p.) 2 h after the administration of 100 mg/kg of 1-aminobenzotriazole (ABT), a pan-cytochrome P450 inhibitor that has been shown to increase systemic APX2039 exposure. In experiments where treatment with APX2096 was initiated 1 h after infection, the Cryptococcus neoformans H99 fungal burden was reduced by approximately 7 log₁₀ CFU/g and 5 log₁₀ CFU/g in the brain and lung tissue, respectively, after 7 days of therapy. Similarly, in a delayed (24 h postinfection) treatment model, the brain and lung fungal burdens were reduced by averages of 6.8 log₁₀ CFU/g and 4.6 log₁₀ CFU/g, respectively. The prodrug APX2097 (active moiety APX2020) was not effective in these models, which is consistent with the 8-fold-higher MIC of the active moiety APX2020 than APX2039 against C. neoformans var. grubii H99 (8). Although the mouse model is robust, the rabbit CM model offers an environment that more closely mimics that of the typical CM patient. The rabbits are immunosuppressed with hydrocortisone, which removes the anticryptococcal lymphocyte response, and the infection is established directly in the central nervous system (CNS) (9-11), whereas the mouse model utilizes immunocompetent mice and relies on systemic intravenous (tail vein) infection. Here, we present compelling evidence for the superior efficacy of orally administered APX2039, first in the mouse model and then in the rabbit model of CM, compared to FLU, AMB, and fosmanogepix.



FIG 1 Mouse delayed-treatment model (\log_{10} CFU per gram of mouse brain and lung tissue). Each point represents data from a single mouse. Data were analyzed with a Kruskal-Wallis test followed by a Wilcoxon rank sum test with Holm's correction. Dosing groups were as follows: control (vehicle [p.o.]) (n = 14), ABT (50 mg/kg ABT [p.o., QD]) (n = 11), FLU (150 mg/kg fluconazole [p.o., QD]) (n = 9), AMB (3 mg/kg amphotericin B deoxycholate [i.p., QD]) (n = 10), and 60 mg/kg APX2039 (p.o., QD) plus 50 mg/kg ABT administered 2 h prior to dosing with APX2039 (n = 7). (A) Data from brain tissue. (B) Data from lung tissue.

RESULTS

APX2039 reduces the fungal burden in mice. The efficacies of 60 mg/kg of APX2039 plus 50 mg/kg of ABT (orally [p.o.], once a day [QD]) administered 2 h prior to dosing with APX2039, 3 mg/kg of AMB (i.p., QD), and 150 mg/kg of fluconazole (p.o., QD) were evaluated in a 24-h delayed-treatment model of disseminated cryptococcal disease. Mice were infected via tail vein injection with approximately 5 \times 10⁴ cells of *C. neoformans* H99, and treatments were initiated at 24 h postinfection. Animals were euthanized on day 9 after 7 days of treatment. Vehicle controls with and without ABT were evaluated, and no significant differences were found between the two groups (Fig. 1). The fungal burden (log₁₀ CFU per gram of tissue) in the lung and brain was significantly lower in all treatment groups. The reductions in the fungal burdens (mean log₁₀ CFU/g control - mean log₁₀ CFU/g treated) in the lung versus the vehicle control were 2.38, 1.36, and 4.45 log₁₀ CFU/g for FLU, AMB, and APX2039, respectively. The reductions in the brain burden versus the vehicle control were 3.32, 0.82, and 6.53 log₁₀ CFU/g for FLU, AMB, and APX2039, respectively. Treatment with APX2039 was significantly better than that with FLU or AMB in terms of reductions in fungal burdens in both the brain (P < 0.001) and lung ($P \le 0.002$). Of note, a fungal burden below the limit of detection was observed only for APX2039. These data are consistent with a combination of a very potent direct antifungal effect of the drug as measured by a very low MIC and a favorable CNS pharmacokinetic/pharmacodynamic (PK/PD) profile (C. neoformans H99 MIC values previously observed for APX2039 [0.008 μ g/mL] versus manogepix [0.25 μ g/mL], AMB $[0.25 \,\mu g/mL]$, and FLU $[2 \,\mu g/mL]$ [8]).

APX2039 is effective in a rabbit model of cryptococcal meningitis. To evaluate efficacy in the rabbit CM model, male New Zealand White rabbits were inoculated with *C. neoformans* H99 (\sim 1.4 \times 10⁶ CFU) directly into the cisterna magna under sedation. Rabbits were immunosuppressed with daily hydrocortisone acetate. Treatments were initiated on day 2 postinfection and continued through day 14. Cerebrospinal fluid (CSF) was removed via a cisternal tap on days 2, 7, 10, and 14 postinfection to quantify the fungal burden. Animals were euthanized either on day 14 or when clinical symptoms required euthanasia, and the fungal burden in the brain tissue was assessed.

The primary endpoint was the fungal burden in the CSF. To assess this, we collected CSF 2 days after the initial inoculation prior to treatment to determine the baseline yeast burden and then repeated the CSF collections on days 7, 10, and 14 postinfection (Fig. 2A and B). Reductions in CFU per milliliter for all treatment groups were statistically different



FIG 2 Average \log_{10} CFU per milliliter of rabbit CSF over time. (A) CFU per milliliter of CSF data were averaged and then \log_{10} transformed. Error bars are SEM. Data are pooled from three independent experiments (control, n = 9 for day 2, n = 11 for day 7, n = 9 for day 10, and n = 5 for day 14; fluconazole, n = 5 for all time points; amphotericin B, n = 5 for all time points; forsmanogepix, n = 6 for all time points; APX2039 [25 mg/kg, QD], n = 5 for day 2, 7, and 10 and n = 4 for day 14; APX2039 [50 mg/kg, QD], n = 8 for day 2, day 7, and day 10 and n = 2 for day 14; APX2039 [50 mg/kg, BID], n = 8 for day 2, day 7, and day 10 and n = 7 for day 14; APX2039 [50 mg/kg, With each line corresponding to a single rabbit.

from those for the control group, with the exception of APX2039 at 25 mg/kg QD (see Table S1 in the supplemental material). Rabbits administered APX2039 at 50 mg/kg BID demonstrated a rapid decrease in the fungal burden by day 7, and the fungal burdens were undetectable on days 10 and 14.

Daily treatment with APX2039 at 50 mg/kg BID resulted in a mean total change of $-4.6 \pm 0.44 \log_{10}$ CFU/mL of CSF within 8 days of treatment. A dose response was observed for the APX2039 QD dosing groups, with a $-2.7 \pm 2.06 \log_{10}$ reduction by day 14 in rabbits treated with 75 mg/kg, whereas 50 mg/kg resulted in a $-0.72 \pm 1.37 \log_{10}$ CFU/mL reduction. Treatment with 25 mg/kg APX2039 was not effective and was equivalent to the vehicle control, as rabbits in this group had a small increase of $0.07 \pm 1.30 \log_{10}$ CFU/mL. Treatment with 100 mg/kg BID fosmanogepix was minimally effective and resulted in a $-0.34 \pm 0.86 \log_{10}$ CFU/mL reduction. Treatment with AMB and FLU resulted in -3.9 ± 0.72 and $-2.28 \pm 0.57 \log_{10}$ CFU/mL reductions, respectively. Rabbits that were either untreated or vehicle treated had an average increase of $1.17 \pm 0.46 \log_{10}$ CFU/mL versus day 2 values, starting at an average burden of 4.67 log₁₀ CFU/mL and reaching 5.65 log₁₀ CFU/mL by day 14.

Previous publications have reported the fungal clearance rate in \log_{10} CFU per milliliter per day using a variety of approaches, including manually calculated slopes, linear models, and linear mixed-effects models (12–25). To compare the rates of reduction that we observed in these experiments with those in the existing literature, we employed three approaches. For the manual linear calculation, we compared the log-subtracted differences of the baseline and the final collection across the groups (Table 1) [fungal clearance = (\log_{10} CFU/mL final – \log_{10} CFU/mL baseline)/interval in days].

Using this approach, APX2039 at 50 mg/kg BID demonstrated a remarkable reduction of -0.66 (standard deviation [SD], ± 0.15) log₁₀ CFU/mL/day, compared to -0.33 (SD, ± 0.06) log₁₀ CFU/mL/day for rabbits treated with AMB. We compared the means using a Kruskal-Wallis test followed by pairwise comparisons using a Wilcoxon rank sum test with Holm's adjustment, with a *P* value of 0.054 for comparisons of APX2039 at 50 mg/kg BID to both AMB and FLU (Fig. 3A).

Using a linear model, we examined the effect of treatment and day postinfection on the log₁₀ CFU per milliliter of CSF. We tested two models, one with an interaction of treatment group and day and one without the interaction, where treatment was a cofactor and day was a covariate. The model was created using all experiment data, and the coefficients within each model were then compared to the control values. We performed pairwise comparisons of the estimated marginal means for linear model 1 (log₁₀ CFU/mL \sim day + treatment) and linear model 2 (log₁₀ CFU/mL \sim day \cdot treatment [includes interaction]).

Treatment	Change in CFU/day				Total change in CFU during the study			
	Mean slope	No. of Rabbits	SD	SEM	Mean	SD	No. of Rabbits	SEM
Control	0.13	9	0.06	0.02	1.17	0.46	9	0.15
FLU (80 mg/kg, QD)	-0.19	5	0.05	0.02	-2.28	0.57	5	0.25
AMB (1 mg/kg, QD)	-0.33	5	0.06	0.03	-3.92	0.72	5	0.32
Fosmanogepix (100 mg/kg, BID)	-0.03	6	0.07	0.03	-0.34	0.86	6	0.35
APX2039 (25 mg/kg, QD)	0.02	5	0.13	0.06	0.07	1.30	5	0.58
APX2039 (50 mg/kg, QD)	-0.06	4	0.11	0.06	-0.72	1.37	4	0.69
APX2039 (75 mg/kg, QD)	-0.34	5	0.35	0.15	-2.70	2.06	5	0.92
APX2039 (50 mg/kg, BID)	-0.66	8	0.15	0.05	-4.60	0.44	8	0.16

TABLE 1 Change in CFU in log₁₀ CFU per milliliter of CSF

In the linear model without the interaction, the overall slope was -0.13 (95% confidence interval [CI], -0.17 to 0.09) log₁₀ CFU/mL/day. Treatment with AMB reduced the log₁₀ CFU per milliliter by -1.33 (95% CI, -2.88 to 1.66) units. Treatment with APX2039 at 50 mg/kg BID resulted in a far-larger effect, with a -3.47 (95% CI, -4.07 to -2.88)-log₁₀ CFU/mL reduction (FLU, $P \leq 0.0001$; AMB, P = 0.014 [by pairwise comparisons]). We then incorporated an interaction to more directly probe the slope of the groups and the effect of group for each unit change in days within the model (Fig. 3B). We calculated the estimated marginal means and then extracted the slopes. APX2039 at 50 mg/kg BID reduced the log₁₀ CFU/mL by -0.60 (95% CI, -0.69 to -0.50) units per day, compared to reductions of -0.19 (95% CI, -0.27 to -0.12) and -0.33 (95% CI, -0.40 to -0.26) log₁₀ CFU/mL/day for FLU and AMB, respectively (P < 0.0001 by pairwise comparisons of estimated marginal means for both FLU and AMB).

Next, to align our findings with the methodology employed previously by Molloy et al. (14), we used a linear mixed-effects model to model the fungal burden in CSF across treatment groups, with an interaction between day postinfection and treatment group as the fixed effects and individual variation in each rabbit as the random effect, along with random slopes for each rabbit [log₁₀ CFU/mL \sim day postinfection \cdot treatment + (0 + day|rabbit ID)]. Because we infected all rabbits at the same time with the same inoculum, we included a fixed intercept. Analyses of the models with a random intercept did not improve the model (Fig. 3C).

We calculated the estimated marginal means and then extracted the slopes. FLU and AMB accounted for daily decreases of -0.19 (95% Cl, -0.27 to -0.11) and -0.33(-0.41 to -0.25) \log_{10} CFU/mL for each day postinfection, respectively (Table 2). Oral APX2039 at 50 mg/kg BID showed a more rapid reduction than AMB or FLU, at -0.62(95% Cl, -0.71 to -0.54) \log_{10} CFU/mL/day (Table 2), which compares very favorably with a good clinical outcome in humans associated with a reduction of approximately $-0.4 \log_{10}$ CFU/mL/day in the CSF (14). Pairwise comparisons of the estimated marginal means between APX2039 at 50 mg/kg BID and FLU and AMB had *P* values of <0.0001 (Table 3).

One final statistical approach to examine the change in the CSF fungal burden used a linear mixed-effects model but did not include treatment as a fixed effect [log₁₀ CFU/mL \sim day postinfection + (0 + day postinfection|rabbit ID)] (Fig. 3D). This created a single slope for all of the data, and the only adjustments to the slope or intercept were from the individual rabbits.

The coefficients for each rabbit were then extracted and compared using one-way analysis of variance (ANOVA) (P < 0.001), and pairwise comparisons of means were then performed using Tukey's honestly significant difference test. Using this approach, APX2039 at 50 mg/kg BID had an average reduction of -0.53 (SEM [standard error of the mean] = 0.24) \log_{10} CFU/mL/day, compared to -0.15 (SEM = 0.1) \log_{10} CFU/mL/day for FLU and -0.27 (SEM = 0.03) \log_{10} CFU/mL/day for AMB. Regardless of the approach used to evaluate the rate of fungal clearance in the CSF, APX2039 at 50 mg/kg BID resulted in $-0.2 \log_{10}$ CFU/mL/day greater clearance than either AMB or FLU (Table 4).



FIG 3 Log₁₀ CFU per milliliter of rabbit CSF over time: plot of slopes and intercepts from a linear calculation, a linear model, and a linear mixed-effects model. Fungal clearance was calculated with four separate methods. Dots are log_{10} CFU per milliliter of CSF data collected during the study and are the same for each panel. Line labels are slopes of selected lines: control, FLU, AMB, and APX2039 (50 mg/kg, BID). (A) Slopes were manually calculated for each rabbit. The mean values of the slopes and intercepts were used to create the lines using the formula log_{10} CFU/mL = mean change in log_{10} CFU/mL/day · day postinfection + intercept mean. (B) Estimated marginal means from a linear model that includes an interaction between day postinfection and treatment (log_{10} CFU/mL ~ day postinfection and treatment). (C) Estimated marginal means from a linear mixed-effects model with day postinfection and treatment + (0 + day postinfection | rabbit ID)]. (D) Means of coefficients extracted from a linear mixed model with only day postinfection as the fixed effect and random slopes for each rabbit [log_{10} CFU/mL ~ day postinfection + (0 + day postinfection | rabbit ID)].

To examine the burden of yeasts in the rabbit brain, we collected brain tissue after euthanasia. Brains harvested on day 14 demonstrated mean reductions versus the control of -1.8 and $-3.4 \log_{10}$ CFU/g tissue for FLU and AMB, respectively, while a mean reduction of $-6.6 \log_{10}$ CFU/g was observed for APX2039 (Fig. 4). After the administration of APX2039 at 50 mg/kg BID, the burdens of yeasts in both the cerebellum/brainstem and cerebrum were below the limit of detection (data not shown). When comparing the total yeast burdens in the brain, treatment with APX2039 showed superiority over both FLU and AMB (adjusted P = 0.027 for both comparisons by a Wilcoxon rank sum test). Rabbits treated with APX2039 at 50 mg/kg BID had undetectable CFU in the vitreous humor, but the statistical comparisons to FLU and AMB did not have an adjusted P value of <0.05 (Fig. S1). Forsmanogepix was not effective in reducing the fungal burden in brain tissue at day 14 relative to the control (adjusted P = 0.06) and performed worse than either FLU or AMB (adjusted P = 0.082 for both by a Wilcoxon rank sum test). Forsmanogepix was able to reduce the fungal burden in the vitreous humor relative to the control but did not perform as well as FLU (Fig. S1).

PK analysis and drug concentrations. To investigate the PK of APX2039 in the rabbit model of CM, we performed serial blood collections after dosing on day 8 postinfection,

m	Bio

	Change in Log ₁₀ CFU/mL/day				
			Lower confidence	Upper confidence	
Treatment	Slope	SE	level	level	
Control	0.12	0.03	0.06	0.19	
Fluconazole	-0.19	0.04	-0.27	-0.11	
Amphotericin B	-0.33	0.04	-0.41	-0.25	
Fosmanogepix	-0.02	0.04	-0.09	0.06	
APX2039 (25 mg/kg, QD)	0.01	0.04	-0.07	0.09	
APX2039 (50 mg/kg, QD)	-0.04	0.05	-0.13	0.05	
APX2039 (75 mg/kg, QD)	-0.27	0.05	-0.36	-0.17	
APX2039 (50 mg/kg, BID)	-0.62	0.04	-0.71	-0.54	

TABLE 2 Slopes from a linear mixed-effects model for CSF in the rabbit model of cryptococcal meningitis

after 6 days of dosing. The PK parameters are shown in Table 5. The results across the groups can be seen in Fig. 5, with 50-mg/kg BID dosing having higher levels of APX2039 than with 25, 50, or 75 mg/kg QD. Although the PK index that best links drug exposure with antifungal efficacy has not been formally investigated for APX2039, the efficacy of the related Gwt1 inhibitor manogepix was previously shown to be the area under the concentration-time curve (AUC)/MIC for *C. albicans*, and this metric was used in the PK/PD analyses. The PK was linear. Levels of APX2039 in the CSF were well above the MIC for rabbits dosed at 50 mg/kg BID and 75 mg/kg QD and closer to the MIC for rabbits dosed with either 25 mg/kg QD or 50 mg/kg QD. This finding aligns with the lower fungal burdens in CSF for both the 50-mg/kg BID and 75-mg/kg QD dosing groups (Fig. 6).

PK/PD analysis. The PK-PD model fitted the data well. The coefficients of determination for the linear regression of observed-predicted values for the plasma and fungal density in the CSF were 0.94 and 0.85%, respectively, after the Bayesian step. The estimates of central tendency and dispersion for the parameters from the PK/PD model are summarized in Table 5.

The relationship between the average total drug AUC from 0 to 24 h (AUC₀₋₂₄) and the decline in \log_{10} CFU per milliliter as estimated by the area under the \log_{10} CFU per milliliter-versus-time plots is shown in Fig. 7. Near-maximal antifungal activity was achieved with a total drug AUC₀₋₂₄ of 25 to 50 mg \cdot h/L and a total drug AUC/MIC ratio of 3,125 to 6,250.

DISCUSSION

CM was uniformly fatal until treatment with amphotericin B deoxycholate was developed in the late 1950s (16). A second major improvement was identified with the successful use of amphotericin B deoxycholate and flucytosine in combination therapy, which reduced toxicity and shortened the treatment course (17). In the early 1990s, the approval of fluconazole allowed a safe, oral anticryptococcal treatment that was successful in managing some cases of CM, but it was clear that fluconazole monotherapy would not be consistently successful (18). Therefore, with three drugs but no uniformly successful monotherapy, a landmark article in 1997 delineated CM management into 3 stages: induction, consolidation, and suppression (19). Since that critical study, many resource-available health care systems have established specific guidelines using a combination of amphotericin B, or a lipid formulation of AMB, plus flucytosine followed by various doses of fluconazole. Over the last 3 decades, it has been clearly shown with the use of quantitative CSF yeast cultures that it is important to achieve rapid effective fungicidal activity (EFA) with treatment to efficiently reduce the fungal burden and, thus, improve patient outcomes (13, 14). Furthermore, it was shown that fluconazole is

TABLE 3 Selected pairwise contrasts of estimated marginal means

Contrast	P value
Control-APX2039 (50 mg/kg, BID)	< 0.000
Fluconazole-APX2039 (50 mg/kg, BID)	< 0.000
Amphotericin B-APX2039 (50 mg/kg, BID)	< 0.000
Control-APX2039 (75 mg/kg, QD)	< 0.000

TABLE 4 Selected pairwise comparisons of coefficients from a linear mixed-effects model

Comparison	P value
Control-APX2039 (50 mg/kg, BID)	<0.0001
Fluconazole-APX2039 (50 mg/kg, BID)	< 0.0001
Amphotericin B-APX2039 (50 mg/kg, BID)	0.0001
Control-APX2039 (75 mg/kg, QD)	<0.0001

most effective after the yeast burden is already substantially reduced. Mechanistically, this likely occurs because fluconazole acts as a fungistatic drug in the central nervous system, which may be due in part to the plasticity of the cryptococcal genome under stress and yeast drug selection. Fluconazole heteroresistance and tolerance frequently occur in monotherapy with high-burden yeast infections, and thus, it is not consistently effective as monotherapy (18).

There have been only small incremental improvements in the therapy of CM over the last 30 years, for instance, refinements in drug formulations such as lipid-based amphotericin B formulations, other combination therapies (amphotericin B and fluconazole, flucytosine and fluconazole, and triple therapy), and even a single dose of AmBisome (liposomal amphotericin B) on an azole and flucytosine base. Unfortunately, the search for an effective new anticryptococcal drug class has not been rewarding. For example, research on repurposed drugs like sertraline and tamoxifen failed to show a benefit in CM (20–22). Simply, this life-threatening common fungal CNS infection needs better therapies.

Our laboratory has studied many antifungal compounds in the immunosuppressed rabbit model, which uses the impact of drugs on fungicidal activity in the subarachnoid space as an endpoint, similar to human studies. By allowing the model to establish the infection over 2 days, at a yeast burden similar to that observed in heavily infected humans prior to treatment, the rabbit model presents a very high barrier to drug success. A similar endpoint of EFA in the subarachnoid space was adopted in human clinical trials, and the EFA measured early in the treatment course in rabbit and human CSF was correlated with the outcome (23, 24). The rabbit has thus been a reasonable surrogate for outcomes that might



FIG 4 Brain tissue fungal burden in the rabbit model of cryptococcal meningitis. Each point represents data from a single rabbit. Group effects were tested using a Kruskal-Wallis test. Pairwise comparisons were made using a Wilcoxon rank sum test with Holm's adjustment (control, n = 10; FLU, n = 5; AMB, n = 5; fosmanogepix, n = 6; APX2039 at 25 mg/kg QD, n = 5; APX2039 at 50 mg/kg QD, n = 5; APX2039 at 75 mg/kg QD, n = 5; APX2039 at 50 mg/kg BID, n = 8).

Table 5 Values for each parameter estimated from the rabbit population pharmacokinetic
pharmacodynamic-linked model

Parameter (Unit)	Mean	Median	SD
$\overline{K_a(h^{-1})}$	14.452	13.778	12.135
SCL (L/h)	5.869	4.644	4.654
V (L)	15.002	13.204	10.766
K_{cp} (h ⁻¹)	7.432	7.434	5.713
$K_{pc}(h^{-1})$	14.103	11.101	9.573
K _{gmax} (Log ₁₀ CFU/mL/h)	0.114	0.095	0.086
H _q	9.218	7.271	7.635
C _{50g} (mg/L)	5.841	5.748	3.387
Popmax (CFU/mL)	255386.168	202038.155	315859.857
K _{kmax} (Log ₁₀ CFU/mL/h)	0.321	0.235	0.244
H _k	16.733	16.081	11.843
C _{50k} (mg/L)	4.638	3.150	3.745
IC (CFU/mL)	540.496	510.244	329.735

occur in humans. For example, despite very low CSF levels of amphotericin B, monotherapy with this polyene can produce CSF fungicidal activity (11). In addition, a polyene can have a positive interaction with azole therapy in reducing CSF yeast counts (10). Furthermore, the rabbit model showed that amphotericin B lipid formulations could work in CM if there was careful attention paid to the dosing of the lipid formulation (25). With the ability to administer high-dose AMB lipid formulations, the hypothesis of whether liposomal amphotericin B could be effectively given in an intermittent dose schedule compared to daily dosing was tested, and it was found to possess similar efficacies with both schedules (26). These findings allowed investigators to test the efficacy of the intermittent dosing of liposomal amphotericin B in humans, and it culminated in the recent randomized controlled AMBITION trial with 844 patients that showed the noninferiority of one high dose of liposomal amphotericin B (10 mg/ kg) on a base of azole and flucytosine compared to daily polyene therapy (4, 27). The rabbit model predicted this outcome (26). Also, a series of azoles were tested in the rabbit model, including fluconazole (28), posaconazole (29), and isavuconazole (30). Similar to monotherapy in humans, these azoles all tend to be fungistatic or possess a slow fungicidal response in the rabbit. Thus, with serial CSF quantitative yeast measurements, a CNS site of infection, severe cell-mediated host immunosuppression, and a track record with drugs also used in human dis-



FIG 5 Plasma levels of APX2039. Each line shows data from a single rabbit. Plasma was collected from each rabbit for the following treatments: APX2039 at 25 mg/kg QD (n = 5) (top left), APX2039 at 50 mg/kg QD (n = 5) (top right), APX2039 at 75 mg/kg QD (n = 5) (bottom left), and APX2039 at 50 mg/kg BID (n = 8) (bottom right).



FIG 6 Levels of APX2039 in rabbit CSF. Concentrations of APX2039 in micrograms per milliliter of CSF were quantified by LC-MS. Each line shows data from a separate rabbit. All CSF samples were collected 1 to 2 h after dosing with APX2039. The black dotted line is the \log_{10} -transformed value of the MIC of APX2039 (0.008 μ g/mL).

ease that recapitulates or predicts outcomes in humans, this rabbit model can be a critical link between *in vitro* susceptibility testing and possible human disease outcomes for testing new potential anticryptococcal agents.

In the current rabbit study, we demonstrated that the oral administration of 50 mg/ kg APX2039 BID resulted in a rapid reduction of the fungal burden in the CSF, and we further showed the concomitant clearance of the yeast in brain tissue. These results provide strong evidence for oral APX2039, a member of the gepix class, as a potential new therapy for CM. We also examined the efficacy of fosmanogepix, which is presently in clinical trials for the treatment of *Candida, Aspergillus*, and rare mold infections. However, APX2039 was specifically chosen from a series of Gwt1 inhibitors because of its improved *in vitro* anticryptococcal activity compared to the active moiety, manogepix (0.008 μ g/mL versus 0.25 μ g/mL) (7). It was hypothesized that the higher potency of APX2039 would make it an ideal drug for the treatment of cryptococcal meningoencephalitis in the quest for rapid EFA in the host CNS site if



FIG 7 Average total drug AUC_{0-24} and decline in the log_{10} CFU per milliliter as estimated by the area under the fungal density (log_{10} CFU per milliliter)-versus-time curve for each rabbit.

the PK/PD were supportive, and this study confirmed this assumption.

APX2039 at 50 mg/kg BID produced a remarkable rate of CSF clearance of between -0.53 and $-0.66 \log_{10}$ CFU/mL/day using several calculation methods. In this model, it significantly outperformed fluconazole, amphotericin B, and fosmanogepix, for which dosing regimens were chosen to give rise to exposures in rabbits that are similar to those achieved clinically for each of the drugs. Furthermore, the rate of clearance was higher than the rate of successful yeast reduction of \log_{10} CFU per milliliter per day in humans, where the most potent regimen of amphotericin B and flucytosine was found to be approximately -0.4 to $-0.5 \log_{10}$ CFU/mL/day (13, 14). Impressively, by 8 days of treatment with oral APX2039 alone, both the CSF and brain tissue had undetectable levels of yeasts. This rate of fungal clearance has not been observed previously with azole or polyene treatments in our model, and even high-dose liposomal amphotericin B (5 mg/kg/day) for 3 days could not consistently sterilize the CSF in the model in a week (26).

Although the burden of yeasts was undetectable in the brain tissue at the time of necropsy after 50 mg/kg APX2039 BID, we did not determine how early this occurs or if the tissue was sterilized with no yeast survival. Future experiments may determine the kinetics of clearance of *Cryptococcus* from the brain and if the clearance effect persists after the cessation of the APX2039 therapy for several weeks without treatment. These findings might gauge how long induction therapy with APX2039 is needed for reliable tissue sterilization and help predict future studies in humans for the length of drug exposure prior to consolidation or suppression stages or even if these secondary stages are necessary if no viable yeasts survive after 10 to 14 days of treatment.

In summary, this is a new potential oral antifungal agent with extremely potent *in vitro* anticryptococccal activity against *Cryptococcus neoformans* that is clearly translated into rapid EFA in several mammalian models of cryptococcosis. Importantly, the impressive activity of APX2039, which inhibits the novel Gwt1 target, suggests that oral monotherapy may be a viable option, without the need for a second antifungal agent or prolonged therapies. There are now PK/PD relationships that can help predict initial and maintenance dosing goals in humans. These results have set the stage for studies of this drug in humans that could potentially revolutionize the treatment of CM.

MATERIALS AND METHODS

Animals. All animal experiments were performed in accordance with Duke University IACUC guidelines and approval.

Mice. Male CD-1 mice, weighing 22 to 24 g, were obtained from Charles River Laboratories International. They were infected with approximately 5×10^4 cells of *C. neoformans* (strain H99) via tail vein injection. Daily treatment was initiated 24 h after infection and continued for 7 days, with 50 mg/kg ABT (p.o.) being administered 2 h prior to each APX2039 (60 mg/kg, p.o.) dose, AMB (3 mg/kg, i.p.), fluconazole (150 mg/kg, p.o.), ABT alone (50 mg/kg, p.o.), or the vehicle (5% dextrose), as previously described (7). Mice were euthanized 24 h after the last dose, and the brain and left lung were used to quantify the fungal burdens.

Rabbits. Male New Zealand White rabbits, weighing between 2.3 and 3.1 kg, were purchased from Robinson Services Incorporated (Winston-Salem, NC). They were administered hydrocortisone acetate at 5 mg/kg intramuscularly (i.m.) daily, beginning 1 day prior to infection. If a rabbit showed disabling neurological signs, it was euthanized prior to the study endpoint.

The inoculum was prepared as previously described (31). The concentration was adjusted to approximately 3.3×10^6 cells/mL in phosphate-buffered saline (PBS), and the rabbits were inoculated with 0.3 mL of the suspension. Rabbits were sedated with ketamine (30 mg/kg) and xylazine (3 mg/kg) i.m. Next, 0.3 mL of the inoculum was injected into the cisterna magna. The rabbit was then administered yohimbine at 0.2 mg/kg intravenously (i.v.) (ZooPharm) to antagonize the anesthesia. Rabbits were recovered with fluids and supplemental heat.

APX2039, fosmanogepix, AMB, and FLU preparation and dosing. APX2039 was reconstituted to a 12.5-mg/mL solution suspended in 10% *N*,*N*-dimethylacetamide (DMA), 20% propylene glycol (PG), and 40% polyethylene glycol 400 (PEG 400) (pH 2 to 3). When precipitate was visible, aliquots were sonicated for 20 min in a 37°C water bath immediately prior to dosing. Fosmanogepix was supplied as a dry powder and stored at -20° C. The powder was reconstituted to a 25-mg/mL solution in 5% dextrose, with a pH of between 7.5 and 8.0, and aliquoted. Fluconazole (Rising) was purchased as a dry powder for oral administration at a concentration of 40 mg/mL. The powder was reconstituted according to the manufacturer's instructions. Fresh aliquots were prepared each day.

The fosmanogepix, APX2039, and fluconazole solutions were administered by oral gavage. Administration started on day 2 postinfection and continued through the day of euthanasia. For doses that occurred on CSF collection days, with the exception of day 2 postinfection, the drug was administered prior to CSF collection.

Amphotericin B deoxycholate (XGen Pharmaceuticals) was purchased as a lyophilized powder. The powder was reconstituted using sterile USP-grade water and adjusted to a final concentration of 2.5 mg/mL in 5% dextrose using a 50% dextrose stock. The rabbit's ear was numbed with lidocaine-prilocaine cream before dosing. The drug was administered through the marginal ear veins.

CSF collection and CFU quantification. CSF was collected on days 2, 7, 10, and 14 postinfection. If a rabbit needed to be euthanized before the study endpoint period of 14 days, then a final CSF sample was collected. For CSF collection, the sedation methods were identical to the methods used for the inoculations. The rabbits were given yohimbine and recovered according to the same protocol as that used for the inoculation.

Samples were plated onto yeast extract-peptone-dextrose (YPD) agar containing chloramphenicol (100 μ g/mL) in a petri dish. For dilutions, 0.1 mL of the CSF was serially diluted (1:10) in PBS. The petri dishes were incubated for 3 to 5 days at 30°C. The CFU were counted, and the CFU per milliliter were calculated by multiplying the colonies counted by the dilution factor/0.1 mL. The remaining CSF that was not used for CFU quantification was centrifuged at 2,000 relative centrifugal force (rcf). The supernatant was collected and stored at -80° C for CSF drug level measurements by liquid chromatography-mass spectrometry (LC-MS) analysis.

PK/PD blood collection and plasma processing. On day 8 postinfection, PK/PD blood collection was performed. The ears were numbed with lidocaine-prilocaine cream. The rabbits were lightly sedated with acepromazine at 1 to 2 mg/kg subcutaneously. A 1.5-in. 21- to 23-gauge catheter was then inserted into the ear artery. Blood samples were collected in K₂EDTA tubes (BD). The samples were centrifuged at 2,000 rcf for 15 min at 4°C and stored in 2-mL screw-cap tubes at -80° C for LC-MS analysis.

Euthanasia and tissue collection. Rabbits were sedated with ketamine at 40 mg/kg i.m. and xylazine at 5 mg/kg i.m. CSF was collected as described above. One milliliter of pentobarbital-phenytoin (Euthasol; Virbac, Westlake, TX, USA) was administered intravenously. A final blood sample was collected from the ear artery prior to the administration of pentobarbital-phenytoin or via cardiac puncture immediately after pentobarbital-phenytoin administration. The aqueous humor, vitreous humor, meninges, and brain were collected and either frozen on dry ice for later LC-MS analysis or homogenized in PBS for the quantification of the fungal burden.

The tissues from the right side were placed into 8-mL screw-cap tubes (Sarstedt) and homogenized using a Pro Scientific P200 homogenizer, or they were placed into tubes that were prefilled with homogenization beads (Precellys; Bertin Corp., France) and homogenized using a Precellys instrument at 6,500 rpm for 30 s.

LC-MS analysis. For all tissues, one of two internal standard solutions (ISSs) was used for homogenization or dilution: either 50% acetonitrile in water containing 10 nM warfarin and 1 μ M natamycin or 50% acetonitrile in water containing 10 nM warfarin, 1 μ M natamycin, and 20 μ M d₆-APX2039 internal standard.

Rabbit brain tissue samples were homogenized using Precellys lysing kits (Bertin, France) in the ISS at 10,000 rpm for 10 s for 3 cycles, with 60-s pauses between cycles, at 4°C. The homogenates were centrifuged at 13,000 rcf for 5 min at 4°C. The cleared lysates were stored at -80° C before analysis. The tissues were run in two batches, with curve ranges of 20 nM to 10 μ M and 2.2 nM to 2.22 μ M. Samples were diluted up to 1000-fold in 50% acetonitrile before the LC-tandem MS (MS/MS) analysis. The calibration curve was made using APX2039 dissolved in dimethyl sulfoxide (DMSO) and then diluted in either a naive rabbit brain homogenate or the ISS.

The CSF samples were mixed 10:1 with the ISS and then run with curve ranges of 1.06 nM to 20 μ M or 1 nM to 10 μ M. Samples were diluted up to 10-fold in the ISS. The plasma samples were diluted up to 400-fold in the ISS. The curves ranged from 20 nM to 10 μ M or 30 nM to 20 μ M. The calibration curves were made with the diluted ISS in PBS or rat plasma.

The analytical LC-MS/MS method, using selected reaction monitoring (SRM), was performed on an Acquity ultraperformance liquid chromatography (UPLC) system coupled to a Xevo TQ-S mass spectrometer. Mobile phase A consists of 5 mM ammonium acetate containing 0.2% formic acid (FA) and 0.025% heptafluorobutyric acid (HFBA) in water, while mobile phase B is acetonitrile with 0.2% FA. A 2.1-mm by 50-mm BEH Shield RP18 column (Waters) with a flow rate of 0.50 mL/min and a column temperature of 35°C was utilized. The electrospray source conditions were 2.5 kV, a desolvation temperature of 400°C, desolvation gas flow at 1,000 L/h, cone gas at 150 L/h, and 7.0 × 10⁵ Pa of nebulizer gas. MS/MS transitions for the most intense product ions were selected from an infusion experiment, and collision energy optimization was performed using LC-MS/MS and Skyline v20.0 (www.skyline.ms). Data analysis was performed with Skyline-daily 20.2 using the small-molecule mode, including peak extraction and quantification, linear regression, and concentration calculations. APX2039 was quantified versus a linear regression against a calibration curve with $1/x^2$ weighting, excluding calibration points that were outside 15% bias except at the lower limit of quantification (LLOQ) (20%).

Brain and eye tissue CFU quantification. The homogenates from the right eye and the right brain hemisphere were used for quantification. Samples were serially diluted in PBS, plated onto petri dishes containing YPD plus chloramphenicol (100 μ g/mL), and incubated for 3 to 5 days at 30°C. The CFU per milliliter were obtained by multiplying the CFU counted by the dilution factor and then dividing this value by 0.1 mL. The total volume was obtained by adding the net weight of the tissue (the density was assumed to be 1 g/mL) to the volume of PBS that was in the tube. These two values, CFU per milliliter and total milliliters, were multiplied to obtain the total CFU. This value was then divided by the net weight in grams to yield the CFU per gram.

Data analysis. Data were cleaned using tidyverse (version 1.3.1) (32) and stringr (version 1.4.0) (33). Statistical analyses were performed in R (34) version 4.1.0. For fungal clearance calculations, if a rabbit had CFU per milliliter equal to zero prior to day 14, and the counts remained at zero, then only the data through the first instance of the CFU per milliliter being equal to zero were used for calculations.

To calculate the linear slopes for the change in the CFU per milliliter of CSF, the numerator was calculated by subtracting the day 2 postinfection \log_{10} CFU per milliliter from the \log_{10} CFU per milliliter for the first day that the CFU counts equaled zero, or if the rabbit had detectable CFU counts at all time points, then the day of the last sample was used. If the day 2 postinfection CFU sample was not usable, the rabbit was excluded from the calculations in the table. The denominator was calculated by subtracting 2 from the day of the final CFU count.

Pairwise comparisons of the linear mixed-effects model with the interaction between day postinfection

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and treatment group, log₁₀ CFU/mL \sim day \cdot treatment + (0 + day|rabbit ID), were performed using the estimated marginal means across all groups in R with the emmeans package, and pairwise comparisons were then performed for all pairs using the *t* test with Holm's adjustment. To compare the slopes from the linear mixed-effects model [log₁₀ CFU/mL \sim day + (0 + day|rabbit ID)], coefficients for each rabbit were extracted and analyzed by ANOVA and then Tukey's honestly significant difference test.

The data for fungal burdens in the mouse lung and brain tissues and the rabbit brain tissue were analyzed using a Kruskal-Wallis test followed by pairwise comparisons using a Wilcoxon rank sum test with Holm's adjustment (R package rstatix version 0.7.0) (35). The linear and linear mixed-effect models and calculations of the estimated marginal means were constructed in R using the ImerTest (36) and emmeans (version 1.6.3) (37) packages. Figures were made using ggplot2 version 2.3.3.5 (38) and sjPlot (version 2.8.10) (39). The source code will be provided upon request.

The PK/PD of APX2039 were described using a population methodology. A rabbit was considered an "individual" contributing both plasma concentrations and serial estimates of fungal density from the CSF. The PK and PD data were described by the following four inhomogeneous differential equations:

$$XP(1) = -K_a \cdot X(1) \tag{1}$$

$$XP(2) = K_a \cdot X(1) - (K_{cp} + (SCL/V)) \cdot X(2) + K_{pc} \cdot X(3)$$
(2)

$$XP(3) = K_{cp} \cdot X(2) - K_{pc} \cdot X(3)$$
 (3)

$$XP(4) = K_{gmax} \cdot \left(1 - \left((X(2)/V) \cdot H_g/(C_{50g} \cdot H_g + (X(2)/V) \cdot H_g)\right)\right)$$

$$\left(1.D0 - (X(4)/popmax)\right) \cdot X(4) - kk_{max} \cdot (X(2)/V) \cdot H_k/(C_{50k} \cdot H_k + (X(2)/V) \cdot H_k) \cdot X(4)$$

(4)

where XP is the rate of change with parentheses showing the relevant compartment. For example, XP(1) is the rate of change of drug amount in compartment 1. K_a is the first-order transfer rate constant connecting the gut with the plasma; SCL is the first-order clearance of APX2039 from plasma; V is the volume of the central compartment; K_{cp} and K_{pc} are the first-order intercompartmental rate constants; K_{gmax} and K_{kmax} are the rates of fungal growth and drug-induced killing, respectively, in CSF; H_g and H_k are the slope functions for the effects of the drug on growth and killing, respectively; C_{s0g} and C_{s0k} are the plasma concentrations of APX2039 that result in half-maximal effects on growth and killing, respectively; and popmax is the maximum achievable fungal density in the CSF. An initial condition (i.e., the fungal density in the CSF immediately after inoculation) was estimated as a parameter (not shown in the differential equations).

Equation 1 describes the movement of APX2039 from the gut into the plasma, equation 2 describes the rate of change of the mass of APX2039 in the central compartment (i.e., plasma), equation 3 describes the rate of change of the mass of APX2039 in the peripheral compartment, and equation 4 describes the pharmacodynamics of APX2039 and contains terms that describe the capacity-limited fungal growth in CSF, the drug-induced suppression of growth, and drug-induced fungal killing. Two output equations were used, one to describe the plasma concentrations and the other to describe the time course of the fungal density in the CSF. The structural model was fitted to the combined PK and PD data set using the nonparametric population PK program Pmetrics (40). The adequacy of the fitting was assessed by visual inspection of the observed-predicted results and linear regression of the observed-predicted values for plasma concentrations and pharmacodynamics.

The PK/PD population model was used to link drug exposure with the antifungal effect. A number of pharmacodynamic measures were used, and these included (i) the area under the \log_{10} CFU per milliliter-time curve for each rabbit and (ii) the rate of decline of \log_{10} CFU per milliliter in CSF. The total drug average AUC₀₋₂₄ in the first 168 h was used as an integrative measure of drug exposure.

SUPPLEMENTAL MATERIAL

Supplemental material is available online only. FIG S1, TIF file, 1.4 MB. TABLE S1, DOCX file, 0.01 MB.

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