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JMX0207, a niclosamide derivative with improved pharmacokinetics, suppresses Zika virus infection both *in vitro* and *in vivo*

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Abstract

Flaviviruses causes significant human disease. Recent outbreaks of the Zika virus highlight the need to develop effective therapies for this class of viruses. Previously we identified niclosamide as a broad-spectrum inhibitor for flaviviruses by targeting the interface between viral protease NS3 and its cofactor NS2B. Here, we screened a small library of niclosamide derivatives and identified a new analogue with improved pharmacokinetic properties. Compound **JMX0207** showed improved efficacy in inhibition of the molecular interaction between NS3 and NS2B, better inhibition of viral protease function, and enhanced antiviral efficacy in cell-based antiviral assay. The derivative also significantly reduced Zika virus infection on 3D mini-brain organoids derived from pluripotent neural stem cells. Intriguingly, the compound significantly reduced viremia in a ZIKV animal model. In summary, a niclosamide derivative **JMX0207** was identified which shows improved pharmacokinetics and efficacy against Zika virus both *in vitro* and *in vivo*. **Keywords: Flavivirus, Zika virus, Dengue virus, antiviral, protease inhibitor**

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The genus *Flavivirus* is composed of more than 70 viruses. Many flaviviruses, including the four serotypes of the Dengue virus (DENV1-4), the West Nile virus, and the Zika virus (ZIKV or ZK) cause serious human diseases. Recently, significant ZIKV outbreaks have occurred worldwide.¹⁻⁵ ZIKV is transmitted to humans primarily through bites from infected *Aedes* species mosquitoes, but may also be transferred prenatally or through sexual activities or blood transfusions.⁶⁻⁹ ZIKV infection is associated with devastating diseases, including Guillain-Barré syndrome, a rare neurological syndrome, and congenital Zika syndrome, which is manifested by microcephaly, brain abnormalities, and other central nervous system malformations.¹⁻⁵ In addition, ZIKV persists in neuronal and/or reproductive tissues of animals for up to 60 days after viral infection, leading to severe damage.¹⁰⁻¹² Currently there is neither a safe and effective vaccine nor a specific therapy for ZIKV. Although a DENV vaccine is recently approved, it is effective in children between 9 and 16-year-old, and posts increased risk for naïve children due to antibody-dependent enhancement.¹³ Therefore, there is an unmet medical need to develop direct antivirals against flaviviruses.

The flavivirus genome encodes a single open reading frame, from which a polyprotein precursor is expressed.¹⁴⁻¹⁶ The polyprotein precursor is post-translationally processed into 10 mature proteins, including three structural and seven non-structural proteins, by viral and host proteases in a sequential manner.¹⁴⁻¹⁵ The viral protease is a heterocomplex composed of viral NS3 protein and its viral co-factor NS2B.¹⁴ The viral protease has been considered a highly promising drug target.^{14, 17-19} Various strategies have been used to develop protease inhibitors. The majority of efforts have been focused on the viral protease active site,^{13,16-18} but with limited success in finding inhibitors with *in vivo* efficacy.²⁰ Recently, alternative strategies to target the NS2B-NS3

interaction²¹⁻²² and allosteric sites²³⁻²⁵ have led to identification of potent protease inhibitors with *in vivo* efficacy against Zika virus.

Previously, we developed a split luciferase complementation (SLC)-based high throughput screening strategy to identify inhibitors targeting the NS2B-NS3 interface of DENV2 (DN2).²¹ Niclosamide was found to be a broad-spectrum inhibitor for flaviviruses by targeting the interface between viral protease NS3 and its co-factor NS2B, leading to accumulation of non-functional viral polyprotein precursor.²¹ In addition, niclosamide was found to interfere with virus entry²⁶ and endosomal acidification.²⁷ Moreover, niclosamide was found as a broad-spectrum antiviral against multiple viruses including SARS Cov-2, the causing agent of current COVID-19 pandemic.²⁸ Unfortunately, niclosamide did not show appreciable *in vivo* antiviral efficacy towards ZIKV (data not shown), possibly due to its poor pharmacokinetic properties.^{26, 29}

In this work, we screened a small library of niclosamide derivatives. We found that one derivative, **JMX0207**, showed improved efficacy in inhibition of the NS2B-NS3 interaction, better inhibition of viral protease function, and enhanced antiviral efficacy in cell-based antiviral assay, compared to niclosamide. **JMX0207** was also found to significantly reduce ZIKV infection on a 3D mini-brain organoid model derived from pluripotent neural stem cells. Intriguingly, **JMX0207** significantly improved the pharmacokinetic properties and markedly reduced viremia in a ZIKV animal model.

Results

Synthesis and screening of Niclosamide derivatives.

Niclosamide was originally used as an anthelmintic drug. It was also found to have anticancer and antibacterial activities.³⁰⁻³⁴ We previously developed a small library composed of 62 niclosamide derivatives for other purposes.³³⁻³⁸ These niclosamide analogs have been

resynthesized in-house following our published procedures for antiviral evaluation. To identify niclosamide derivatives that have equal or better antiviral efficacy and improved pharmacokinetic properties, we used our SLC-based NS2B-NS3 interaction assay to screen this library, with niclosamide as a control.

Compound ^b	Structure	IC _{50-SLC-} (DN2)	IC _{50-pro} (DN2)	<i>EC</i> _{50-DN2} (A549)	<i>EC</i> _{50-ZK} (А549)	CC ₅₀ (A549)	TI ^c (DN2)	TI (ZK)
Niclosamide		2.0	21.6	0.55	0.48	4.8	8.7	10.0
HJC0114		18.4	33.6	>30	>30	>200		
HJC0125	HN O C C HCI HCI	16.3	44.5	0.21	0.15	50.2	239	335
HJC0308	H_2N O O CI NO_2 CI	4.9	39.7	1.4	6.2	17.7	12.6	2.8
HJC0365		13.2	>60	6.4	6.0	177	27.6	29.5
HJC0381		10.8	25.9	1.2	3.9	91.1	76	23.3
НЈС0390	NH ₂ Cl	6.0	46.5	0.43	<0.1	30.1	70	>300
HJC0431	H ₂ N CI	3.1	34.0	2.2	0.36	28.4	13	79
НЈС0129	CI NH2	9.6	11.3	>30	>30	176		
HJC0140		9.4	25.5	>30	>30	>200		
HJC0149	OH OH NH OF OF	18.6	20.3	0.9	1.1	241	268	219
JMX0207	OH O NO2	1.3	8.2	0.31	0.30	31.9	103	106

Table 1. <i>IC</i> ₅₀ , ^a <i>EC</i> ₅₀ ,	, <i>CC</i> 5θ (all in μM), ε	ind therapeutic index	x (TI) of niclosamide	e derivatives.
---------------------------------------------------------------------------	-------------------------------	-----------------------	-----------------------	----------------

^a *IC*_{50-SLC}/*IC*_{50-pro}/*EC*₅₀/*CC*₅₀: Compound concentrations required to reduce 50% of the split luciferase complementation (SLC) signal, protease activity, virus production, and cell viability, respectively. ^bOriginal

compound names from literature and this work. According to their structures, these compounds can be classified into three groups, colored as red, magenta, and blue, respectively. ^c TI, therapeutic index (TI) defined as CC_{50}/EC_{50} . All values are in micromolar (μ M) except TI. DN2, Dengue virus serotype 2; ZK, Zika virus.



Figure 1. Inhibition of the NS2B-NS3 interactions and protease activity. (A) Dose-dependent inhibition of SLC upon binding of NLuc-NS2B₄₉₋₆₆ to GST-CLuc-NS3 by JMX0207. N=3. (B) Dose-response inhibitions of the DENV2 His-NS2B/His-MBP-NS3 protease activity by JMX0207. N=3. In both (A) and (B), DMSO control was set as 100%. The values represent means \pm standard deviation (S.D.) in all panels.

Eleven out of 62 niclosamide derivatives showed dose-responsive inhibition of the interaction between viral NS3 protease and its cofactor NS2B, with IC_{50-SLC} values lower than 30 μ M, the highest concentration tested ($IC_{50-SLC}/IC_{50-pro}/EC_{50}/CC_{50}$: compound concentrations required to reduce 50% of the split luciferase complementation signal, protease activity, virus production, and cell viability, respectively) (**Table 1, Figure 1**). The derivatives identified included seven compounds with *O*-alkylamino chains, one compound with 3-NO₂ instead of 5-Cl on the salicylic ring, and three compounds with modifications on the nitro group of the aniline moiety. Of these active derivatives, compounds **HJC0308** with 3-aminopropoxy moiety, **HJC0431** with 4-aminobutoxy moiety, and **JMX0207** with 3-NO₂ group showed comparable or

Inhibition of the NS2B-NS3 protease activity.

We next evaluated the inhibition efficacy of these positive derivatives against the viral NS2B-NS3 protease function as previously described.²¹ Previously, we generated two versions of the NS3 protease, the NS3 protease domain refolded from inclusion bodies and a soluble version of the NS3 protease domain fused to maltose-binding protein (MBP).²¹⁻²² Our data showed that the refolded and MBP-tagged NS3 proteins have indistinguishable protease activities when transactivated by co-factor NS2B. In this work, we used both versions to characterize the inhibitors and found that the IC_{50} values determined from both versions were similar. Therefore, only values for the refolded NS3 were shown. As shown in Table 1, most active compounds in SLC assay except **HJC0365** still showed inhibition against the viral NS2B-NS3 protease activity. However, we found that there seemed to be no strict correlation between IC_{50-SLC} and IC_{50-pro} values. This is not surprising because niclosamide is a known luciferase inhibitor.³⁹ The SLC inhibition is an indirect readout of the NS2B-NS3 interactions. Direct inhibition of luciferase activity and/or direct inhibition of luciferase complementation in addition to inhibition of NS2B-NS3 interaction may account for the poor correlation between IC_{50-SLC} and IC_{50-pro} values. While IC_{50-SLC} is a poor indicator of inhibition efficacy of the NS2B-NS3 interactions, it remains a valuable indicator to eliminate compounds if they failed to inhibit SLC.

Among these derivatives, compounds **HJC0140** with 4'-acetamido group ($IC_{50-pro} = 25.5$ μ M), **HJC0149** with benzo[*b*]thiophene 1,1-dioxide moiety ($IC_{50-pro} = 20.3 \mu$ M) and **HJC0381** with (1-methylpiperidin-4-yl)oxy moiety ($IC_{50-pro} = 25.9 \mu$ M) maintained the same level of potency, while compounds **HJC0129** with 4'-NH₂ group ($IC_{50-pro} = 11.3 \mu$ M) and **JMX0207** with

3-NO₂ group ($IC_{50-pro} = 8.2 \ \mu\text{M}$) showed nearly or more than 2-fold improvement in inhibition of the viral NS2B-NS3 protease activity, compared to niclosamide ($IC_{50-pro} = 21.6 \ \mu\text{M}$) (**Table 1**, **Figure 1**). Taken together, the three compounds **HJC0129** ($IC_{50-SLC} = 9.6 \ \mu\text{M}$ and $IC_{50-pro} = 11.3 \ \mu\text{M}$), **HJC0140** ($IC_{50-SLC} = 9.4 \ \mu\text{M}$ and $IC_{50-pro} = 25.5 \ \mu\text{M}$), and **JMX0207** ($IC_{50-SLC} = 1.3 \ \mu\text{M}$ and $IC_{50-pro} = 8.2 \ \mu\text{M}$) showed potent inhibitory activity against the NS2B-NS3 interactions in both NS2B-NS3 SLC and NS2B-NS3 protease assays.



Figure 2. JMX0207 inhibits DENV. (A) Dose-dependent inhibition of DENV2 infectivity by **JMX0207**. N=3. (**B**) Cell viability assay. A549 cells were incubated with various concentrations of **JMX0207** and then assayed for viability at 48 hrs post-incubation. N=3. Error bars in both panels represent the standard deviations at each concentration.

Inhibition of DENV2 viral replication.

A viral plaque reduction assay was performed next to evaluate the antiviral efficacy of these derivatives. Human lung carcinoma A549 cells were infected with DENV2 in the presence of a concentration series of derivatives or a dimethyl sulfoxide (DMSO) control, and viral titers were measured at 48 hrs post-infection, as described previously.^{21-22, 40} Our results indicated that eight out 11 active derivatives showed appreciable antiviral efficacy with EC_{50-DN2} less than 30 μ M (**Table 1**). The best derivative, **JMX0207**, showed slightly improved antiviral efficacy (EC_{50-DN2} of 0.31 μ M), compared to niclosamide (EC_{50-DN2} of 0.55 μ M) (**Figure 2A**).

The observed antiviral activity could result from compound's cytotoxicity. To address this concern, we next measured the compound cytotoxicity CC_{50} using a WST-8 cell proliferation assay, as we described previously.²¹ As shown in **Table 1** and **Fig. 2B**, the majority of these derivatives showed improved CC_{50} and therapeutic index (defined as CC_{50}/EC_{50}), compared to niclosamide.



Figure 3. JMX0207 inhibits ZIKV. (A) Dose-dependent inhibition of ZIKV infectivity. N=3. (**B**) qRT-PCR analysis of inhibition of viral RNA from ZIKV-infected A549 cells by **JMX0207**. N=3. Error bars in panels (**A**) and (**B**) represent the standard deviations at each concentration. (**C**) Immunofluorescence assay of inhibition of viral protein production by **JMX0207**, using pan-flavivirus anti-E 4G2 antibody (green) (ATCC).

Inhibition of ZIKV replication.

Our previous studies indicated that niclosamide is a broad-spectrum inhibitor against multiple flaviviruses²¹. To determine the antiviral spectrum of the derivatives, we evaluated the antiviral potency of the derivatives against ZIKV. Our results showed that the derivatives were also potent inhibitors against ZIKV, with EC_{50-ZK} values comparable to those for DENV2 (**Table 1, Figure 3A**). Similarly, **JMX0207** showed better inhibition efficacy towards ZIKV than niclosamide (**Figure 3A**).

JMX0207 treatment leads to reductions in viral RNA yield and protein production.

Among the derivatives, **JMX0207** showed improved efficacy in almost all categories (**Table 1**), compared to niclosamide. Therefore, we chose **JMX0207** to further characterize whether **JMX0207** treatment led to inhibition of viral RNA synthesis and viral protein production, using qRT-PCR and immunofluorescence assay (IFA), respectively. Our results indicated that **JMX0207** significantly reduced the ZIKV RNA copy number in a dose-dependent manner (**Figure 3B**). Using a pan flavivirus anti-E antibody 4G2, we conducted an IFA to demonstrate that **JMX0207** greatly reduced ZIKV antigen production in A549 cells in a dose-dependent manner (**Figure 3C**), presumably because of inhibition of viral replication.

Overall, these experimental results indicated that **JMX0207** showed improved antiprotease, antiviral, and cytotoxicity properties, compared to niclosamide. **JMX0207** treatment results in inhibition of viral infectivity, viral RNA replication, and viral protein production, and is a broad-spectrum antiviral for flaviviruses.

JMX0207 inhibits viral production in cells relevant to ZIKV pathogenesis.

We next used human induced pluripotent stem cell (iPSC)-derived neural progenitor cell (HNPC) ^{21, 41} to investigate whether **JMX0207** is an effective inhibitor in human primary cells demonstrated to be relevant to ZIKV pathogenesis.⁴²⁻⁴⁶ As shown (**Figure 4A & 4B**), **JMX0207** effectively inhibited ZIKV protein expression and RNA synthesis in HNPC in dose-dependent manner. Overall, these experiments demonstrate that **JMX0207** is an effective antiviral in neural progenitor cells relevant to ZIKV infection.



Figure 4. Inhibition of ZIKV in cells relevant to ZIKV. (A) Immunofluorescence assay (IFA) of inhibition of viral protein expression in human neural progenitor cells by **JMX0207**, using panflavivirus anti-E 4G2 antibody (green) (ATCC). Nuclei (blue) was stained in all IFA assays by the Hoechst stain solution. (**B**) qRT-PCR analyses of inhibition of viral RNA from ZIKV-infected human neural progenitor cell (HNPC) by **JMX0207**. N=3 (**C**) ZIKV organoid infected with ZIKV-Venus. The 3D organoids were infected with PBS (Mock), or ZIKV untreated (DMSO), or ZIKV treated with **JMX0207**, or Mock treated with **JMX0207**. Upper panel, bright field image of intact organoids. Lower panel, Venus fluorescence image (excitation 515 nm, emission 528 nm) of the intact 3D organoids. (**D**) Forebrain reginal specification of organoids. Organoids were stained

positive for forebrain identity markers PAX6 (green, upper panel), FOXG1 (green, lower panel) and SOX2 (Red (Magenta after merge with DAPI (Blue)), upper panel) at 20 days. The sections were stained positive for general neuronal marker TUJI (cyan, upper and lower panels) and were negative for SOX10 (Red, lower panel). DAPI-merged data were shown. Nuclei (DAPI, Blue); scale bar 200µm. (E) Slices of organoid infected with ZIKV PRVABC59. The 3D organoids were infected with PBS (Mock), or ZIKV untreated (DMSO), or ZIKV treated with JMX0207. Upper panel, IFA using anti-E 4G2 antibody (green); blue, DAPI. Lower panel, details of signature rosette region of the 3D organoids (Mock) or infected region (DMSO/ZIKV and JMX0207/ZIKV). Red: Pax6. (F) ZIKV production from the 3D organoids at 5 dpi. Culture supernatants were collected, and virus production was quantified by PFU assay. N=3. Error bars in panels (B) and (F) represent the standard deviations at each concentration.

JMX0207 protects 3D mini-brain organoid from ZIKV infection.

Newborns from mothers with ZIKV infection during pregnancy have significantly increased risk of developing microcephaly, a birth defect where a baby has smaller than normal head circumference. Babies born with microcephaly may develop many brain-related or other neurological problems. Recently, neural stem cell-derived 3D cerebral organoids were used to dissect ZIKV pathogenesis and anti-viral development.^{45, 47-49} Compared to the cultured 2D monolayer cells, the 3D brain organoids can better represent the composition, diversity and organization of cell types found in the developing human brain.

Therefore, we used the 3D mini-brain organoid model to further investigate whether **JMX0207** can protect against ZIKV-associated neurological damage. Induced pluripotent stem cells (iPSC) derived from a healthy control (Male, Caucasian) were differentiated using an

 established protocol to generate region-specific organoids patterned to resemble the dorsal forebrain (**Figure 4C**).⁵⁰ Organoids were stained positive for forebrain identity markers PAX6 (dorsal forebrain progenitors; upper panel), FOXG1 (lower panel) and SOX2 (neural ectoderm marker, upper panel) at 20 days. The sections were stained positive for general neuronal marker TUJI (neuron-specific class III β -tubulin; upper and lower panels) and were negative for SOX10 (Neural crest, Red, lower panel).

To evaluate if **JMX0207** protects organoids from ZIKV infections, we first generated a full-length infectious ZIKV clone expressing Venus fluorescent protein (ZIKV-Venus) (details to be published elsewhere). At day 20, the organoids displayed signature features of forebrain, including neural rosettes (**Figure 4C-E**), and were pre-treated with **JMX0207** (1.5 μ M) or DMSO control. At day 20, the organoids were infected with either Mock or ZIKV strain PRVABC59 or ZIKV-Venus, at an estimated multiplicity of infection (MOI) of 1 in the presence of 1.5 μ M **JMX0207** or DMSO control. ZIKV infection was quantified by a plaque forming unit (PFU) assay at 5 dpi and by fluorescence imaging at 7 dpi, respectively. Our results showed that **JMX0207** at 1.5 μ M concentration did not have any toxic effect on the 3D organoids, which remained intact and unchanged in morphology (**Figure 4D**). Compared to DMSO control, **JMX0207** treatment completely abolished ZIKV infection in the 3D organoid (**Figure 4D**).

The 3D mini-brain organoids were sliced for immunostaining to investigate the antiviral effects of **JMX0207** (**Figure 4E**). As shown, untreated organoids (DMSO) were infected with ZIKV throughout all layers (**Figure 4E**). In contrast, **JMX0207** treatment nearly completely protected the organoids from ZIKV infection (**Figure 4E**). Moreover, ZIKV production was also significantly inhibited by **JMX0207** treatment (**Figure 4F**). Collectively, the results indicated that



JMX0207 is an effective inhibitor to protect developing human cortical tissue from ZIKV



Figure 5. In vivo antiviral activity of JMX0207 against ZIKV. (A) Pharmacokinetic study of niclosamide and JMX0207. Adult female B6 mice were given a single oral dose of either niclosamide or **JMX0207** at 40 mg/kg. Plasma was obtained at various times after dosing. Both niclosamide and JMX0207 were extracted from the plasma and then analyzed by LC-MS/MS as described in the Methods. N=4. (B) Pharmacokinetic study of JMX0207 at different doses. N=4-5. The values in panels (A) and (B) represent means \pm S.D. (C) Viremia was detected by a plaque forming unit (PFU) assay on day 3 post-infection of ZIKV with a dose of 1.7×10^5 PFU/mouse in four-week-old A129 mice, which were treated with vehicle or JMX0207 though oral gavage. Difference between JMX0207 (N=10) or vehicle (N=10) treatment was analyzed by using the unpaired, two-tailed t-test. *, P=0.0081. Error bars represent data range of median with 95% confidence interval.

JMX0207 shows improved pharmacokinetic properties.

We next evaluated whether JMX0207 has favorable pharmacokinetic properties using a mouse model. As shown in Table 2 and Figure 5A, JMX0207 displayed much better pharmacokinetic properties than the lead compound niclosamide. In contrast to niclosamide, which had a short $T_{1/2}$ and low C_{max} (0.6 μ M), when given to B6 mice orally at 40 mg/kg JMX0207 had

an excellent pharmacokinetic profile with C_{max} of 145 μ M (**Table 2**), which was much higher than the *EC*₅₀ (0.3 μ M) required to inhibit ZIKV. The longer T_{1/2} (~11 h) was also more favorable to ensure less frequent drug administration to infected animals. Further pharmacokinetic study with different doses of JMX0207 indicated that the plasma drug concentration remained at the same level after the dose was reduced to 20 mg/kg (**Figure 5B**). More importantly, mice with repeated oral dosing at 40 mg/kg/day for 7 days did not display any sign of toxicity (N=6). These results indicate good bioavailability and low toxicity for the derivative **JMX0207**.

 Table 2. Pharmacokinetic properties of niclosamide and JMX0207

	$T_{max} \pm S.D.^{a}(h)$	$C_{max} \pm S.D. (\mu M)$	$T_{1/2} \pm S.D.$ (h)	$AUC_{0\to\infty} \pm S.D.$
Niclosamide	0.7±0.4	0.6±0.1	4.6±1.0	3.5±0.8
JMX0207	1.2±0.5	145±18	11±6.8	2719±1018

^a S.D., standard deviation

JMX0207 reduces viremia in a ZIKV mouse model.

Finally, we evaluated the *in vivo* antiviral efficacy using a viremia mouse model, as we described previously.⁵¹⁻⁵² Our data showed that **JMX0207** treatment at 20 mg/kg/day resulted in a significant reduction in ZIKV-induced viremia in the A129 mice inoculated with 1.7x10⁵ PFU PRVABC59 ZIKV/mouse compared to the vehicle control (**Figure 5C**). Overall, the results indicated that **JMX0207** not only inhibited viral replication *in vitro* but also significantly reduced viremia in an *in vivo* animal model.



Figure 6. Mechanism of action studies. (**A**) Time of addition. **JMX0207** (0.75 μ M) was added at indicated time points post-infection. Viral titers were quantified using plaque forming assay 48 hrs post-infection. DMSO was added to control. N=3. ***, p<0.001. (**B**) Dose-dependent inhibition of DENV2 replicon by niclosamide and **JMX0207**. N=3. (**C**) WB analysis of dose response inhibition of ZIKV NS3 production by **JMX0207** treatment (left). Right, Band intensities of PP and NS3 normalized to the GAPDH control (N=3). Pre-seeded A549 cells in 6-well plate were treated with **JMX0207** or DMSO control and infected with ZIKV with MOI of 0.1, as described previously.²¹ At 48 hrs post-infection, cells were washed, harvested, and protected by protease inhibitor cocktail prior to lysis by SDS-PAGE loading buffer. Upon incubation at 95 °C for 10 min, sample was subjected to Western blot analysis with anti-ZIKV NS3 (GTX133309, GeneTex, Inc.) and anti-GAPDH (CB1001, EMD Millipore) as primary antibodies. Error bars in all panels represent the standard deviations at each condition.

JMX0207 is effective post-infection.

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Antiviral inhibitors are generally divided into two categories, entry and post-entry (replication), respectively. To determine the mode of action, we first performed a time of addition experiment by adding JMX0207 at different time points post-infection (**Figure 6A**). Our results showed that **JMX0207** was equally effective in reducing ZIKV titer even at 24 hrs post-infection. The results indicated that **JMX0207** inhibited viral replication instead of viral entry, which agrees with our hypothesis that **JMX0207**, as a protease inhibitor, attenuates viral production in the post-entry replication stage.

JMX0207 inhibits viral replication using a DENV2 replicon cell line.

To further investigate the mechanism of action, we next used a DENV2 replicon cell line⁵³ to investigate if **JMX0207** inhibited flavivirus replication without an entry step. Our results indicated that both niclosamide and **JMX0207** showed dose-responsive inhibition of DENV2 replication using the DENV2 replicon cell line (**Figure 6B**). **JMX0207** has slightly improved antiviral efficacy (EC_{50-DN2} of 1.1 µM) compared to niclosamide (EC_{50-DN2} of 2.0 µM) (**Figure 6B**). Overall, the results indicate that **JMX0207** is an effective antiviral to inhibit viral replication instead of entry.

JMX0207 treatment inhibits viral polyprotein precursor (PP) processing

We carried out Western blot (WB) analysis using an antibody recognizing ZIKV NS3 protein (**Figure 6C**). Our results showed that **JMX0207** treatment led to dose-dependent reduction of viral NS3 protein production, presumably due to inhibition of viral production by **JMX0207**. In addition to reduced NS3 protein production, we observed dose-dependent increase of a protein with high molecular weight (MW) (**Figure 6C**). It was known from our previous studies that the high MW protein is the unprocessed viral polyprotein precursor (PP) which can be recognized by the anti-NS3 antibody^{21, 23}. The results are consistent with the mechanism by which **JMX0207**

inhibits viral protease function, leading to inhibition of viral PP processing, accumulation of nonfunctional viral PP, and finally virus reduction.

JMX0207 directly binds to the NS2B-NS3 interface.

As a niclosamide derivative, **JMX0207** is expected to directly bind to the viral NS3 protease, as did niclosamide.²¹ To demonstrate binding, we first carried out a protein thermal shift assay (PTSA). Our data indicated that the binding of **JMX0207** stabilized the viral NS3 protease, leading to a 0.75 °C increase in T_m of the viral protein. The data suggests that **JMX0207** directly binds the viral NS3 protease.



Figure 7: Docking and mutagenesis. (**A**) Predicted docking pose of **JMX0207** (magenta) docking into NS3pro of DENV2 (PDB Code: 2FOM). NS3pro is in blue ribbon representation and binding site key interaction residues are highlighted in stick presentation. π cation and π stacking are shown as cyan dotted lines, H-bond in purple and salt bridge in blue. (**B**) Predicted binding pose of Niclosamide (green) docked into NS3pro of DENV2. H-bond is shown as purple dotted lines. (**C**) **JMX0207** and Niclosamide superimposed at the predicted binding site, in surface representation.

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JMX0207 is shown as magenta sticks and Niclosamide in green. (D) Representative SPR analysis of **JMX0207** binding to the MBP-NS3 protease wild-type and mutant L58A.

To investigate the molecular interaction between JMX0207 and viral NS3 protease, we docked **JMX0207** to the NS3 protease structure of DENV2 (PDB ID: 2FOM) using Induced Fit Docking (IFD) protocol, as we describe previously²¹⁻²². The docked pose (**Figure 7A**) depicted **JMX0207** can be well docked into the same 2B53 pocket identified from previous study²¹. One NO₂ group of JMX0207 interacts with Y23 and F46 through π cations, while another NO₂ group forms H-bonds with Y33 and K61 and a salt bridge with K26. The phenyl ring on one side of JMX0207 interacts with L58 through hydrophobic interaction, and the other phenyl ring engages with Y33 with π - π interaction. The phenol forms an H-bond with H60. The O atom on carbonyl group of JMX02027 forms an additional H-bond with K26 to further stabilize the binding. We superimposed the IFD docking pose of JMX0207 and the IFD docking pose of niclosamide. The overlay analysis demonstrates that **JMX0207** binds at the 2B53 pockets on NS3 protease of ZIKV in a similar manner to that of niclosamide (**Figure 7B,C**), while an additional nitro group of **JMX0207** also forms critical binding interactions with the protease.

To further investigate JMX0207 binding to viral NS3 protein, we used surface plasmon resonance (SPR) to measure the binding affinity of **JMX0207** to the viral NS3 protease. Because we previously generated several MBP-NS3 mutants,²² we used MBP-NS3 fusion protein in the SPR analyses for better comparison. As shown in **Figure 7D**, **JMX0207** bound the viral NS3 protease with a binding affinity of 1.1 μ M. Collectively, our data demonstrated that **JMX0207** directly binds to the viral NS3 protease, with an affinity better than that for niclosamide (6.4 μ M).²¹

In a previous study, we generated five NS3 mutants, some of which had significant impact on binding of small molecule inhibitors to the NS2B-NS3 interface and on viral protease activity²². We measured the binding affinity of **JMX0207** to these mutants as well. Our data showed that L58A abolished the binding of **JMX0207** (**Table 3**, **Figure 7D**). Two other mutations, I25A and H60A reduced the binding affinity 2 to 3 folds, whereas mutations Y23A and F46A do not have any effect.

Table 3. Binding affinity of JMX0207 to MBP-NS3 wild-type (WT) or mutant

	WT	Y23A	I25A	F46A	L58A	H60A
$K_{D}(\mu M)$	1.1	1.2	2.3	1.2	NDa	2.9

ND^a, no detectable binding.

Discussion

Dengue and Zika viruses cause significant human disease, for which there are no specific vaccines or therapies. Therefore, development of effective antivirals is urgent. Previously we identified an existing FDA-approved drug, niclosamide, as a potent antiviral against DENV and ZIKV. However, the poor pharmacokinetic properties of niclosamide prevent its clinical use to treat DENV and ZIKV infections.

In this study, we screened a small in-house compound library of niclosamide derivatives and identified **JMX0207** as a candidate inhibitor with improved properties. **JMX0207** inhibited the viral NS2B-NS3 protease activity of DENV2 with better efficacy *in vitro* than niclosamide. In cell culture, **JMX0207** significantly inhibited the growth of representative flaviviruses, including DENV2 and ZIKV, with EC_{50} in nanomolar range. By using IFA, WB, and qRT-PCR analyses, we showed that **JMX0207** also significantly inhibited viral RNA synthesis and protein production. More importantly, compared to niclosamide, **JMX0207** has reduced cytotoxicity profile towards

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human cells, indicating a larger therapeutic window towards these flaviviruses. In addition, we showed that JMX0207 could rescue ZIKV-relevant neural progenitor cells from viral infection in a dose-dependent manner. Moreover, JMX0207 could almost completely protect 3D cortical organoids originated from neural stem cells from ZIKV infection. The findings establish the efficiency of **JMX0207** to eliminate infection of ZIKV from human neural progenitor cells, and provide a path forward to minimize the risk of fetal acquired microcephaly resulting from ZIKV infection of pregnant women.⁵⁴ Furthermore, we showed that **JMX0207** displayed significantly improved pharmacokinetic properties, compared to the parent compound niclosamide. The improved pharmacokinetics allowed us to carry out *in vivo* antiviral efficacy studies using an animal model for ZIKV infection. Our data showed that JMX0207 treatment markedly reduced viral viremia in the A129 ZIKV mouse model infected with clinical strain ZIKV PRVABC59. Mechanistically, we showed that **JMX0207** directly binds to the viral NS3 protease with improved binding affinity, compared to its lead drug niclosamide. We also showed that JMX0207 treatment led to dose-dependent inhibition of viral polyprotein precursor processing, a direct consequence of inhibition of viral protease activity.

Conclusion

We conclude that the niclosamide derivative **JMX0207** is a valuable candidate for further study, due to its demonstrated ability to inhibit several activities required for ZIKV propagation. When paired with its reduced cytotoxicity profile and superior pharmacokinetic properties, these positive attributes indicate that **JMX0207** might effectively be used to combat viral progression following initial infection with ZIKV. Successful development of this compound would be a valuable tool in the prevention of negative ZIKV-related outcomes, including Guillain-Barre

syndrome, neuropathy and myelitis, as well as pregnancy-related outcomes including miscarriage, preterm birth, and congenital Zika syndrome.

Methods

Synthesis of niclosamide derivatives.

The synthetic route of compound **JMX0207** was described as below (**Scheme 1**), and the synthesis of other niclosamide derivatives was reported in our previous publications.^{34-36, 38} The structures and purity of all synthesized compounds were confirmed by 1H and 13C NMR, HRMS and HPLC analysis, and all biologically evaluated compounds are >95% pure.

JMX0207 Synthesis.



Scheme 1. JMX0207 synthesis route.

Reagents and conditions: (a) CH₃I, K₂CO₃, DMF, 50 °C, 24 h, 95%. (b) NaOH, H₂O/MeOH, r.t., 1 h, 93%. (c) 2-chloro-4-nitroaniline, POCl₃, pyridine, DCM, 0 °C to r.t., 8 h, 67%. (d) BBr₃, DCM, 0 °C to r.t., 12 h, 96%.

Methyl 2-methoxy-3-nitrobenzoate (2).

CH₃I (3.9 g, 27.3 mmol) was added to a solution of 3-nitrosalicylic acid (1.0 g, 5.5 mmol) and K_2CO_3 (2.3 g, 16.4 mmol) in 20 mL of DMF. The mixture was stirred at 50 °C for 24 hrs and then diluted with 300 mL of AcOEt. The resulting mixture was washed with water (3 × 70 mL) and

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brine (50 mL), dried over Na₂SO₄, and concentrated *in vacuo*. The residue was purified by silica gel column (Hexane/EtOAc = 2/1) to afford compound **2** (1.1 g, 95%) as a yellow oil. ¹H NMR (300 MHz, CDCl₃) δ 8.01 – 7.89 (m, 1H), 7.85 – 7.78 (m, 1H), 7.23 – 7.09 (m, 1H), 3.91 (s, 3H), 3.87 (s, 3H). ¹³C NMR (75 MHz, CDCl₃) δ 164.6, 153.1, 145.4, 135.5, 128.3, 127.2, 123.7, 64.1, 52.6.

2-Methoxy-3-nitrobenzoic acid (3).

NaOH (1.1 g, 27.3 mmol, in 6 mL of H_2O) was added to a solution of compound 2 (1.1 g, 5.5 mmol) in 10 mL of MeOH. The mixture was stirred at r.t. for 1 h, and then the pH value was adjusted to 5~6 with 1 M HCl (aq.). The mixture was extracted with EtOAc (2 × 120 mL) and washed with water (80 mL) and brine (60 mL), dried over Na₂SO₄, and concentrated to give acid **3** (1.0 g, 93%) as a light yellow solid for direct use in the next step.

N-(2-Chloro-4-nitrophenyl)-2-methoxy-3-nitrobenzamide (4).

POCl₃ (3.9 g, 25.4 mmol) was added slowly at 0 °C to a solution of compound **3** (1.0 g, 5.1 mmol), 2-chloro-4-nitroaniline (876 mg, 5.1 mmol) and pyridine (8.0 g, 101.4 mmol) in 50 mL of DCM. After addition, the mixture was stirred at r.t. for 8 h and then poured into 200 mL of ice-water. Compound **4** (1.2 g, 67%) was isolated by filtration as a yellow solid. ¹H NMR (300 MHz, CDCl₃) δ 10.58 (s, 1H), 8.93 (d, *J* = 9.3 Hz, 1H), 8.46 (dd, *J* = 7.8, 1.8 Hz, 1H), 8.37 (d, *J* = 2.7 Hz, 1H), 8.24 (dd, *J* = 9.3, 2.4 Hz, 1H), 8.06 (dd, *J* = 8.1, 1.8 Hz, 1H), 7.46 (t, *J* = 8.1 Hz, 1H), 4.12 (s, 3H).¹³C NMR (75 MHz, CDCl₃) δ 161.9, 151.7, 144.4, 143.6, 140.7, 137.0, 129.9, 128.2, 125.3, 125.1, 123.8, 123.1, 121.1, 64.9.

N-(2-Chloro-4-nitrophenyl)-2-hydroxy-3-nitrobenzamide (JMX0207).

BBr₃ (7.4 ml, 7.40 mmol, 1M in DCM) was added dropwise at 0 °C to a solution of compound 4 (1.3 g, 3.70 mmol) in 250 mL of DCM. The mixture was stirred at r.t. for 2 h. Then the mixture

was poured into 200 mL of ice water. The yellow precipitate **JMX0207** was isolated by filtration. The organic layer was separated and concentrated. Then 50 mL of MeOH was added and the mixture was stirred at r.t for 20 min. The yellow solid was isolated by filtration. The two parts of yellow solids were combined to afford 1.2 g (96%) of compound **JMX0207** in total. HPLC purity 99.9% ($t_R = 18.75 \text{ min}$). ¹H NMR (300 MHz, DMSO- d_6) δ 12.10 (s, 1H), 8.65 (d, J = 9.3 Hz, 1H), 8.42 (d, J = 2.4 Hz, 1H), 8.34 – 8.24 (m, 2H), 8.13 (d, J = 8.1 Hz, 1H), 7.03 (t, J = 7.8 Hz, 1H). ¹³C NMR (75 MHz, DMSO- d_6) δ 164.2, 154.1, 143.1, 141.3, 139.1, 136.3, 129.8, 124.8, 124.1, 123.6, 122.5, 122.4, 117.3. HRMS (ESI) calcd for C₁₃H₉ClN₃O₆ 338.0180 (M + H)⁺, found 338.0172.

Cloning, Expression and Purification.

All clones and proteins were generated as previously described.²¹⁻²²

Split Luciferase Complementation (SLC) Assay.

The SLC assay was performed as previously described.^{21-22, 55}

Protease Inhibition Assay.

The protease inhibition assay was performed using the refolded DENV2 NS3 fusion protein (50 nM) and a fluorescence resonance energy transfer (FRET) peptide substrate (100 μ M) (Abz-RRRRSAG-nTyr (NeoBiolab)), as described previously.^{21-23, 55} Substrate cleavage was monitored at excitation/emission wavelengths of 360 nm/420 nm (Abz substrate) by a BioTek Flx800. The rate of increase in relative fluorescence unit (RFU) over time was calculated in the linear range and normalized as a percent of the DMSO control. The $IC_{50}/CC_{50}/EC_{50}$ was determined by fitting the dose-responsive curve with a non-linear regression function using the GraphPad Prism 8 (San Diego, CA). All experiments were performed in triplicates.

Cytotoxicity Assay.

Cytotoxicity was measured by a WST-8 cell proliferation assay kit (Dojindo Molecular Technologies, Inc.) as previously described.²¹ All experiments were performed in triplicates.

Viral Titer Reduction Assay.

A viral titer reduction assay was used to determine the compounds' effect on DENV2 and ZIKV strains, as described previously²¹⁻²². Human A549 lung carcinoma cells and human primary neural progenitor cells were used, as described previously.²¹⁻²² All experiments were performed in triplicates.

Immunofluorescence Assay.

The immunofluorescence assay was performed using ZIKV-infected cells treated with DMSO or **JMX0207**, as described previously.²¹⁻²² A mouse monoclonal pan anti-E antibody 4G2 (ATCC) and a DyLight® 488 goat anti-mouse IgG (ImmunoReagents, Inc.) were used to monitor viral E protein expression, as described previously.²¹⁻²²

Quantitative qRT-PCR.

Quantitative qRT-PCR was performed as described previously.²¹⁻²² ZIKV primers CCGCTGCCCAACAACAAG and CCACTAAYGTTCTTTTGCAGACAT with ZIKV probe Cy5-AGCCTACCT/TAO/TGACAAGCAGTCAGACACTCAA-IAbRQSp were used. The $2-\Delta\Delta$ CT ("delta-delta Ct") method was used to quantify samples. All experiments were performed in triplicates.

Western Blot.

Western blot was performed using anti-ZIKV NS3 (GTX133309, GeneTex, Inc.) and anti-GAPDH (CB1001, EMD Millipore) antibodies, as described previously.^{21,23} All experiments were performed in triplicates.

DENV2 Replicon Assay.

BHK-21 cells stably expressing DENV2 replicon with a *Renilla luciferase (Rluc)* reporter gene⁵³ were seeded into white 96-well plate at a density of 2 x 10⁵ cells per well. The cells were incubated at 37°C in a humidified incubator with 5% CO₂ for hrs in a 100 µl medium containing Minimal Essential Medium (MEM) supplemented with 10% fetal bovine serum (FBS), 100 I.U./ml penicillin, 100 µg/ml streptomycin, and 300 µg/ml G418. Upon 24 hrs incubation, culture medium was discarded. Fresh culture medium of 100 μ l with a concentration series of compounds or DMSO control was added to the cells in triplicates. The culture was further incubated for 48 hours. The cells were then washed twice with 100 µl phosphate-buffered saline (PBS), followed by addition of 25 µl of a lysis buffer containing 1X PBS and 1% Triton X-100. The mixture was incubated at room temperature with gentle shaking (50 rpm) for 30 minutes. Assay buffer (125 μ l) 1X PBS, 0.05% (3-((3-cholamidopropyl) containing dimethylammonio)-1propanesulfonate)(CHAPS), and 0.1% BSA was then added to each well. In the meantime, a fresh 4X working substrate coelenterazine was prepared by diluting 1 µl of stock coelenterazine (2.4 mM) dissolved in ethanol into 10 ml assay buffer (10,000-fold dilution). Finally, 50 µl 4X substrate coelenterazine was added to each well with a final substrate concentration of 0.06 µM, using a Veritas luminometer. The luminescence was recorded immediately using the Veritas luminometer. The luminescence data was normalized to DMSO control. The EC_{50} was determined by nonlinear regression fitting of normalized experimental data in GraphPad Prism 8.0 (San Diego, CA). All experiments were performed in triplicates.

Docking.

The crystal structure of NS3pro of DENV2 (PDB Code: 2FOM) was downloaded from RCSB PDB bank. After excluding the cofactor NS2B peptide, the structure was preprocessed and optimized with Schrödinger Protein Preparation Wizard using default settings. The 3D structure

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of JMX0207 and niclosamide were created using Schrödinger Maestro and prepared with LigPrep to generate a low energy conformation suitable for docking. The Induced Fit Docking (IFD) protocol of Schrödinger Small-Molecule Drug Discovery Suite was employed in this docking study. The grid box for docking was centered on the 2B53 binding site. The box size was set to 20 Å on each side. Selected side chains of R24, K26, and M59 were temporarily trimmed (the equivalent of being mutated to alanine) during the initial IFD process and were restored later. The receptor-ligand complexes structures generated from IFD docking were imported into Schrödinger Maestro for visualization and analysis of binding site interactions.

Surface Plasmon Resonance.

Surface plasmon resonance (SPR) was used to determine the affinity and kinetic analyses of the interactions between JMX0207 and the MBP-NS3 proteins at 25 °C using a ProteOn XPR36 SPR instrument (Bio-Rad). The MBP-NS3 wild-type or mutant proteins were immobilized onto a ProteOnTM GLH sensor chip (~9,000 RU) (Bio-Rad). A 2-fold dilution series of compounds were injected as the analytes. A blank surface blocked by ethanolamine was used as the control surface. The experiment was carried out at a flow rate of 100 µl/min using a PBSTD buffer containing 1x PBS, 0.005% surfactant P20, and 5% DMSO. Association (k_a) and dissociation (k_d) rates, as well as the dissociation constant (K_D), were obtained by global fitting of the SPR data from multiple concentrations to a simple 1:1 Langmuir binding model, using the ProteOn Manager software suite (Bio-Rad).

Culturing of human iPSCs.

Human iPSC F11350.1 culture was maintained in feed-free conditions using mTeSR1 medium (StemCell Technologies #85851) with 5X supplement (StemCell Technologies #85852).

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Cells were maintained on six-well plates coated with human embryonic stem cell-qualified Matrigel at a 1:60 dilution (Corning Catalog #354277) and passaged with ReLeSR (StemCell Technologies #05872). Cortical 3D organoids were generated based on the protocol described by Yoon et al.⁵⁰ Briefly, iPSCs were dissociated with accutase (ThermoFisher #A1110501) and then plated at a density of 3x10⁶ cells per well in AggreWell 800 24-well plate (StemCell Technologies #34811) in Essential 8^{TM} medium (ThermoFisher #A1517001) supplemented with 10µM rock inhibitor (Tocris #Y27632). The plate was centrifuged at 100 x g for 3 min and incubated at 37°C overnight. The next day (culture day 0) the organoids were transferred to a low attachment 10cm plate (Corning #3262) and maintained in 10mL of neural induction medium (Essential 6TM medium, Life Technologies, #A1516401) supplemented with two SMAD pathway inhibitors, 10 µM SB431541 (R&D #1614/50) and 2.5 µM dorsomorphin (Tocris #3093), and Wnt pathway inhibitor 2.5 µM XAV-939 (Tocris #XAV-939). Media were exchanged daily. On day 6, the medium was changed to neural expansion medium (Neuralbasal A medium, ThermoFisher #10888) supplemented with 2% B27 (without Vit A; ThermoFisher #12587001), 1% antibioticantimycotic (ThermoFisher #15240062), 1% glutaMAX (ThermoFisher # 35050061), 20 ng/mL FGF2 (R&D #233-FB-500) and 20 ng/mL EGF2 (PeproTech #AF-100-15). Media were exchanged daily.

Cryopreservation.

Organoids were fixed in 4% paraformaldehyde (Santa Cruz) overnight at 4°C. They were then washed three times in PBS and transferred to 30% sucrose solution for 72 h. Organoids were then transferred into plastic cryomold (10 X 10 X 5mm; Tissue Tek cat. no. 4565) with embedding medium (Tissue-Tek OCT compound, Sakura Finetek 4583) and stored at -80° C. For

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immunohistochemistry, 20 μ m thick sections were cut with a Leica cryostat (model CM3050S).

Immunohistochemistry.

Cryosections were blocked in 3% bovine serum albumin (BSA), 10% normal goat serum (NGS), and 0.3% Triton X–100 diluted in PBS for 1 hour at room temperature. The sections were incubated overnight at 4°C with primary antibodies diluted in blocking solution. Sections were washed three times with PBS and then incubated with appropriate secondary antibodies diluted in blocking solution for 1 hour. The following primary antibodies were used for immunohistochemistry: anti-Pax-6 (1:200; Biolegend #901301), anti-Sox2 (1:100; Santa Cruz #sc365823), anti-FoxG1 (1:500; Takara #M227), Sox10 (1: 100; Santa Cruz # sc369692), and anti-tubulin III (1:1000; Sigma #T8660).

Pharmacokinetics.

Two- to three-month-old female B6 mice were used for the study. Four to five mice per group were given niclosamide at 40 mg/kg (in 1% DMSO and 0.5% carboxymethylcellulose) or JMX0207 in the same vehicle at the indicated dose by oral gavage. Sample preparation and liquid chromatography with tandem mass spectrometry (LC-MS/MS) detection of niclosamide and JMX0207 were essentially the same as described in a previous study and were performed using a Sciex 4000 Q-Trap mass spectrometer (AB SCIEX, Framingham, MA) with the Agilent 1200 high-performance liquid chromatography system (Agilent Technologies, Santa Clara, CA).²⁹ JMX0207 was monitored at m/z 336/171. Declustering potential, entrance potential, collision energy, and collision cell exit potential were optimized for detection and quantification of JMX0207 at -45, -10, -35, and -14V, respectively. Data from 4 or 5 mice at each time point were averaged and used to calculate pharmacokinetic parameters, using a pharmacokinetic solver (Microsoft, Redmond, WA) by assuming a noncompartmental model. Statistical significance of

various data comparisons was determined with the use of GraphPad Prism (GraphPad Software,

La Jolla, CA). Student's t-test was used. P values < 0.05 were considered statistically significant.

In vivo protection efficacy.

All animal studies involving infectious ZIKV were conducted at an Animal Biosafety Level 2 (ABSL-2) facility at the Wadsworth Center with Institutional Biosafety and Animal Welfare Committee approval. The *in vivo* antiviral activity of **JMX0207** was evaluated in a viremia animal model.

A group of four-week-old A129 mice were infected by subcutaneous injection with 1.7×10^5 PFU of the PRVABC59 strain. Then, the infected mice were administered **JMX0207** at 20 mg/kg of body weight (n=10) or with vehicle control (n=10) every day for 3 consecutive days post-infection (dpi). Mice were observed daily for signs of illness and mortality. Viremia on day 3 post-infection (pi) was determined by plaque forming assay, and statistical analysis was performed using the unpaired, two-tailed t-test (Prism).

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Notes

The authors declare no competing financial interests.

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ABBREVIATIONS

ABSL-2, Animal Biosafety Level 2; BSA, bovine serum albumin; CHAPS ((3-((3-cholamidopropyl) dimethylammonio)-1-propanesulfonate)); DENV, dengue virus; DMSO, dimethyl sulfoxide; DN2, dengue virus serotype 2; dpi, days post-infection; FBS, fetal bovine serum; FRET, fluorescence resonance energy transfer; HNPC, human neural progenitor cells; IFA, immunofluorescence assay; iPCS, induced pluripotent stem cell; LC-MS/MS, liquid chromatography with tandem mass spectrometry; MEM, Minimum Essential Medium; MOI, multiplicity of infection; NGS, normal goat serum; PBS, phosphate-buffered saline; PFU, plaque forming unit; PTSA, protein thermal shift assay; qRT-PCR, quantitative reverse transcription polymerase chain reaction; RFU, relative fluorescence unit; RLU, relative luminescence unit; SLC, split luciferase complementation; SPR, surface plasmon resonance; WB, Western blot; ZIKV/ZK, Zika virus

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Figure 1. Inhibition of the NS2B-NS3 interactions and protease activity. (A) Dose-dependent inhibition of SLC upon binding of NLuc-NS2B49-66 to GST-CLuc-NS3 by JMX0207. N=3. (B) Dose-response inhibitions of the DENV2 His-NS2B/His-MBP-NS3 protease activity by JMX0207. N=3. In both (A) and (B), DMSO control was set as 100%. The values represent means ± standard deviation (S.D.) in all panels.

101x48mm (300 x 300 DPI)



Figure 2. JMX0207 inhibits DENV. (A) Dose-dependent inhibition of DENV2 infectivity by JMX0207. N=3. (B) Cell viability assay. A549 cells were incubated with various concentrations of JMX0207 and then assayed for viability at 48 hrs post-incubation. N=3. Error bars in both panels represent the standard deviations at each concentration.

118x41mm (300 x 300 DPI)





Figure 3. JMX0207 inhibits ZIKV. (A) Dose-dependent inhibition of ZIKV infectivity. N=3. (B) qRT-PCR analysis of inhibition of viral RNA from ZIKV-infected A549 cells by JMX0207. N=3. Error bars in panels (A) and (B) represent the standard deviations at each concentration. (C) Immunofluorescence assay of inhibition of viral protein production by JMX0207, using pan-flavivirus anti-E 4G2 antibody (green) (ATCC).

185x49mm (300 x 300 DPI)



Figure 4. Inhibition of ZIKV in cells relevant to ZIKV. (A) Immunofluorescence assay (IFA) of inhibition of viral protein expression in human neural progenitor cells by JMX0207, using pan-flavivirus anti-E 4G2 antibody (green) (ATCC). Nuclei (blue) was stained in all IFA assays by the Hoechst stain solution. (B) qRT-PCR analyses of inhibition of viral RNA from ZIKV-infected human neural progenitor cell (HNPC) by JMX0207. N=3 (C) ZIKV organoid infected with ZIKV-Venus. The 3D organoids were infected with PBS (Mock), or ZIKV untreated (DMSO), or ZIKV treated with JMX0207, or Mock treated with JMX0207. Upper panel, bright field image of intact organoids. Lower panel, Venus fluorescence image (excitation 515 nm, emission 528 nm) of the intact 3D organoids. (D) Forebrain reginal specification of organoids. Organoids were stained positive for forebrain identity markers PAX6 (green, upper panel), FOXG1 (green, lower panel) and SOX2 (Red (Magenta after merge with DAPI (Blue)), upper panel) at 20 days. The sections were stained positive for general neuronal marker TUJI (cyan, upper and lower panels) and were negative for SOX10 (Red, lower panel). DAPI-merged data were shown. Nuclei (DAPI, Blue); scale bar 200µm. (E) Slices of organoid infected with ZIKV PRVABC59. The 3D organoids were infected with PBS (Mock), or ZIKV untreated (DMSO), or ZIKV treated with JMX0207. Upper panel, IFA using anti-E 4G2 antibody (green); blue, DAPI. Lower panel, details of signature rosette region of the 3D organoids (Mock) or infected region (DMSO/ZIKV and JMX0207/ZIKV). Red: Pax6. (F) ZIKV production from the 3D organoids at 5 dpi. Culture supernatants were collected, and virus production was quantified by PFU assay. N=3. Error bars in panels (B) and (F) represent the standard deviations at each concentration.

171x137mm (300 x 300 DPI)



Figure 5. In vivo antiviral activity of JMX0207 against ZIKV. (A) Pharmacokinetic study of niclosamide and JMX0207. Adult female B6 mice were given a single oral dose of either niclosamide or JMX0207 at 40 mg/kg. Plasma was obtained at various times after dosing. Both niclosamide and JMX0207 were extracted from the plasma and then analyzed by LC-MS/MS as described in the Methods. N=4. (B) Pharmacokinetic study of JMX0207 at different doses. N=4-5. The values in panels (A) and (B) represent means ± S.D. (C) Viremia was detected by a plaque forming unit (PFU) assay on day 3 post-infection of ZIKV with a dose of 1.7 × 105 PFU/mouse in four-week-old A129 mice, which were treated with vehicle or JMX0207 though oral gavage. Difference between JMX0207 (N=10) or vehicle (N=10) treatment was analyzed by using the unpaired, two-tailed t-test. *, P=0.0081. Error bars represent data range of median with 95% confidence interval.

169x46mm (300 x 300 DPI)



Figure 6. Mechanism of action studies. (A) Time of addition. JMX0207 (0.75 μM) was added at indicated time points post-infection. Viral titers were quantified using plaque forming assay 48 hrs post-infection. DMSO was added to control. N=3. ***, p<0.001. (B) Dose-dependent inhibition of DENV2 replicon by niclosamide and JMX0207. N=3. (C) WB analysis of dose response inhibition of ZIKV NS3 production by JMX0207 treatment (left). Right, Band intensities of PP and NS3 normalized to the GAPDH control (N=3). Pre-seeded A549 cells in 6-well plate were treated with JMX0207 or DMSO control and infected with ZIKV with MOI of 0.1, as described previously.21 At 48 hrs post-infection, cells were washed, harvested, and protected by protease inhibitor cocktail prior to lysis by SDS-PAGE loading buffer. Upon incubation at 95 °C for 10 min, sample was subjected to Western blot analysis with anti-ZIKV NS3 (GTX133309, GeneTex, Inc.) and anti-GAPDH (CB1001, EMD Millipore) as primary antibodies. Error bars in all panels represent the standard deviations at each condition.

136x83mm (300 x 300 DPI)



60



Figure 7: Docking and mutagenesis. (A) Predicted docking pose of JMX0207 (magenta) docking into NS3pro of DENV2 (PDB Code: 2FOM). NS3pro is in blue ribbon representation and binding site key interaction residues are highlighted in stick presentation. π cation and n stacking are shown as cyan dotted lines, H-bond in purple and salt bridge in blue. (B) Predicted binding pose of Niclosamide (green) docked into NS3pro of DENV2. H-bond is shown as purple dotted lines. (C) JMX0207 and Niclosamide superimposed at the predicted binding site, in surface representation. JMX0207 is shown as magenta sticks and Niclosamide in green. (D) Representative SPR analysis of JMX0207 binding to the MBP-NS3 protease wild-type and mutant L58A.

172x93mm (300 x 300 DPI)