

Synthesis and Evaluation of 9-Aminoacridines with SARS-CoV-2 Antiviral Activity

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acridine scaffold against SARS-CoV-2 using traditional medicinal chemistry to identify promising new analogs. Our studies identified several novel analogs possessing potent *in vitro* activity in U2-OS ACE2 GFP 1-10 and 1-11 (IC₅₀ < 1.0 μ M) as well as moderate cytotoxicity (CC₅₀ > 4.0 μ M). Compounds such as 7g, 9c, and 7e were more active, demonstrating selectivity indices SI > 10, and 9c displayed the strongest activity (IC₅₀ ≤ 0.42 μ M, CC₅₀ ≥ 4.41 μ M, SI > 10) among them, indicating that it has potential as a new lead molecule in this series against COVID-19.

INTRODUCTION

As of September 20, 2023, the COVID-19 pandemic has ended based on the WHO recommendation. However, there are very few small molecule treatments developed for this disease^{1,2} and preparation for future outbreaks of this virus or related viruses is still of great importance.³ COVID-19 is an infectious disease caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). The pandemic caused unprecedented economic and social hardships, affecting over 770 million people and causing over 6.9 million deaths worldwide.⁴ Infection with SARS-CoV-2 leads to a broad range of clinical symptoms including respiratory distress, cough, immune system disruption, and loss of smell and taste.^{5,6} While the development of various COVID-19 vaccines has been successful at preventing infection and decreasing symptom severity,⁷ there is a recognized need for complementary antiviral agents, which have direct activity against circulating SARS-CoV-2 strains. Remdesivir^{8,9} was the only FDA-approved antiviral drug for the treatment of COVID-19, while molnupiravir and paxlovid both obtained emergency use authorizations in 2021 for treating infections with this virus in the USA,¹⁰ and the latter obtained full approval from the FDA more recently.

Current treatment strategies for COVID-19 involve either prevention of viral attachment to the host cell or disruption of viral replication.¹ SARS-CoV-2 enters the host cell via a spike protein attachment to the host angiotensin converting enzyme 2 (ACE2) receptor.¹¹ After entry, viral RNA is translated by the host ribosome to generate two polyproteins, pp1a and pp1ab. The virus encodes two cysteine proteases, papain-like protease (PL^{pro}) and the coronavirus main protease (M^{pro}), which cleave pp1a and pp1ab, yielding nonstructural proteins that form complexes with the host membrane.¹² These proteases are essential for viral replication, making them attractive targets for antiviral therapeutics.¹³ There are currently a number of M^{pro} inhibitors that have advanced to clinical trials; however, none have yet reached the market.¹⁴

A hot topic in the identification for antiviral drug candidates is the repurposing of existing drugs for new targets.¹ The above-mentioned remdesivir was originally developed for the treatment of hepatitis C and was later repositioned for COVID-19.¹⁵ Several existing drugs with a variety of biological activities have been tested against SARS-CoV-2,¹⁶ and we and others have recently found that an antimalarial agent pyronaridine (Figure 1) displays activity against a number of viruses including SARS-CoV-2.^{17–19} In subsequent studies, we

Received: August 10, 2023 Accepted: October 4, 2023 Published: October 19, 2023





Figure 1. Structures of pyronaridine and quinacrine.

have shown that pyronaridine reduces viral load in the lungs of SARS-CoV-2-infected mice.²⁰ The antiviral effect of pyronaridine is most likely based on inhibition of replication because it was found to selectively inhibit SARS-CoV-2 PL^{pro} activity *in vitro*.²⁰ Another antimalarial drug popular outside of the U.S., quinacrine (Figure 1), which has similar structural motifs, also displays comparable *in vitro* activity against SARS-CoV-2.^{21,22} We have shown that quinacrine possesses activity against SARS-CoV-2 in A549 + ACE2 cells (pretreatment, IC₅₀ = 0.19 μ M, CC₅₀ = 9.24 μ M).¹⁹ These findings prompted us to study the structure–activity relationships of pyronaridine and quinacrine scaffolds against SARS-CoV-2 using a traditional medicinal chemistry approach.

RESULTS AND DISCUSSION

Compound Synthesis. Derivatives and analogs based on the pyronaridine core were synthesized according to Schemes 1 and 2. The aryl carboxylic acids 3a-3h were obtained under Ullmann reaction conditions from commercial aryl chlorides 1a-1e and arylamines 2a-2c (Scheme 1). Subsequent cyclization of 3a-3h with phosphorus oxychloride led us to afford the desired intermediate acridines and naphthyridines 4a-4h in high yields. Further, some compounds 4a-4h were reacted with the corresponding amines listed in Scheme 2 in DMF medium at reflux or 100 °C to give the final products 7a-7g and 8a-8g.

Structure–Activity Relationship Study of Pyronaridine and Quinacrine Analogs against SARS-CoV-2. The antiviral activity against SARS-CoV-2, cytotoxicity in U2-OS ACE2 GFP cells, and selectivity index (SI) for the synthesized compounds are summarized in Tables 1 and 2. The quinacrine derivatives in Table 1 were more active when bearing a 2chloro substituent compared to a 3-chloro substituent (for example, quinacrine is less active than 9a and 9c). We did not see a significant difference between the substituents at R₃ in this series. For the symmetric pyronaridine derivatives in Table 2 ($R_4 = Py$), compounds bearing trifluoromethyl substituents at the 2-position (R_1 as drawn) were more active than the 2chloro compounds and had comparable antiviral activity to that of the 3-chloro derivatives (R_2 as drawn). We did not observe a significant relationship regarding the 2- and 3-chloro derivatives with respect to activity. For the asymmetric pyronaridine derivatives (Table 2, $R_4 = H$), 2-chloro substituents were generally more active than the 3-chloro derivatives, and the more electronegative fluoro- and trifluoromethyl groups did not increase the activity to a significant degree. The presence of a chloro or methoxy substituent at R₃ did not have a significant effect on activity. Last, the introduction of an additional nitrogen atom into the ring system did not affect antiviral activity significantly but did result in lower cytotoxicity. In summary, there were few general trends for the SAR of the pyronaridine/quinacrine derivatives, as antiviral activity was dependent on both the identity of the side chain at the acridine 9-position and the halogenation pattern around the central ring system. The evaluation of additional derivatives in future studies may shed more light onto the specific substitution patterns, which affect potency against SARS-CoV-2.

Overall, compounds 7e, 7g, and 9c were the most active compounds, which displayed high selectivity indexes (SI > 10). Compounds 8a, 9a, and 7b were also comparable in antiviral activity but displayed higher cytotoxicity and lower selectivity indexes (SI < 10). Subsequent studies of these compounds in other common cells lines such as A549-ACE2, Huh-7, and Caco-2 will be performed in the future, as cell-type specific activity against SARS-CoV-2 has been demonstrated previously and should be addressed.^{19,23} However, the most active compounds are still less active than remdesivir (IC₅₀ = 0.02-0.05 μ M),^{8,9} suggesting the need to continue searching for compounds with increased antiviral activity. Nonetheless, these findings suggest that compounds 7e, 7g, and 9c may support further investigation of the repurposing of pyronaridine and quinacrine as antivirals against SARS-CoV-2 and illustrate how they can be used to generate new analogs with improved properties (compared to quinacrine).

While there has been considerable progress made on COVID-19 drug discovery in over 3 years since the pandemic began with preclinical and clinical molecules showing





Reagents and conditions: a) K₂CO₃, Cu, DMF, reflux then rt; b) POCl₃, (C₂H₅)₃N·HCl, reflux

Scheme 2. Synthesis of 9-Aminoacridines and Analogs 7a-7g and 8a-8g



Table 1. In Vitro Activity IC_{50} against SARS-CoV-2 Strain BetaCoV/France/IDF0372/2020, Cytotoxicity CC_{50} in U2-OS ACE2 GFP Cells, and Selectivity Index SI (Ratio of CC_{50} to IC_{50}) of Quinacrine Derivatives

	$ \begin{array}{c} $							
compound	Y	R ₁	R ₂	R ₃	R_4	IC ₅₀ (µM)	CC ₅₀ (µM)	SI
quinacrine	СН	Н	Cl	OMe	5-NEt ₂ pentyl	2.57	2.43	0.95
5	СН	Н	Cl	OMe	morpholine	>50.0	>50.0	1.0
9a	СН	Cl	Н	OMe	5-NEt ₂ pentyl	0.41	2.41	5.87
9b	СН	CF ₃	Н	OMe	5-NEt ₂ pentyl	1.81	7.26	4.02
9c	СН	Cl	Н	Cl	5-NEt ₂ pentyl	0.42	4.41	10.50
9d	Ν	Н	Cl	OMe	5-NEt ₂ pentyl	1.27	8.25	6.50

activity,^{1,2} there is still a need to identify promising lead compounds.³ This may be particularly important as they may be needed as the virus mutates and overcomes existing small-molecule antivirals. The 9-aminoacridines may provide a scaffold for further optimization as antivirals.^{17–19}

MATERIALS AND METHODS

Chemistry. All reagents and solvents were purchased from commercial suppliers (AlfaAesar, Acros, and Chimmed) and used without further purification. The ¹H and ¹³C spectra were recorded on Bruker AC-300 (200 MHz, ¹H) or a Bruker AC-200 (50 MHz, ¹³C) NMR spectrometers. Chemical shifts were

measured in DMSO- d_6 or CDCl₃ using tetramethylsilane as an internal standard and reported as ppm values. The following abbreviations are used to indicate multiplicity: s, singlet; d, doublet; t, triplet; m, multiplet; dd, doublet of doublets; brs, broad singlet. The mass spectra were recorded on a Finnigan MAT INCOS 50 quadrupole mass spectrometer (Thermo Finnigan, San Jose, CA USA) (EI, 70 eV) with direct injection. The purity of the final compounds was analyzed by analytical high-performance liquid chromatography (HPLC) on an Elute HPLC system (Bruker Daltonik, Heidelberg, Germany) equipped with an Azura UVD 2.1S UV detector (Knauer, Berlin, Germany) with a wavelength at 254 nm and acquisition

Table 2. In Vitro Activity IC ₅₀ a	gainst SARS-CoV-22 Strair	n BetaCoV/France/IDF0372/2020,	Cytotoxicity CC ₅₀ in U2-OS
ACE2 GFP Cells, and Selectivit	y Index SI of Pyronaridine	Derivatives	



compound	Х	Y	R ₁	R ₂	R ₃	R_4	IC_{50} (μM)	CC_{50} (μM)	SI
6 ^{<i>a</i>}	CH	CH	Н	Cl	OMe	Н	10.20	>50.0	>4.90
7a	СН	СН	Cl	Н	OMe	Ру	1.88	5.19	2.75
7b	СН	СН	CF ₃	Н	OMe	Ру	0.44	4.36	9.91
7c	СН	СН	F	Н	OMe	Ру	1.84	3.74	2.03
7 d	СН	СН	Cl	Н	Cl	Ру	1.76	5.46	3.09
7e	СН	СН	Н	Cl	Cl	Ру	0.57	8.86	15.50
7 f	Ν	СН	Н	Н	OMe	Ру	1.95	26.34	13.48
$7g^b$	СН	СН	Н	Cl	OMe	Ру	0.42	4.70	11.18
8a	CH	CH	Cl	Н	OMe	Н	0.48	3.20	6.67
8b	СН	СН	CF ₃	Н	OMe	Н	1.90	10.08	5.32
8c	СН	СН	F	Н	OMe	Н	1.91	4.03	2.11
8d	СН	СН	Cl	Н	Cl	Н	1.75	4.26	2.44
8e	СН	СН	Н	Cl	Cl	Н	1.37	19.30	14.41
8f	Ν	СН	Н	Н	OMe	Н	1.96	30.77	15.7
8g	СН	Ν	Н	Cl	OMe	Н	1.96	3.62	1.85
³ NMe ₂ instead of pyrrolidine rings, ^b Pyronaridine.									

rate at 1 Hz. Chromatographic separation was carried out on an Acquity HSS T3 column (2.1 \times 100 mm, 1.3 μ m, 100 Å) at 30 °C and a sample injection volume of 2.0 μ L. A mobile phase consisting of 0.1% formic acid in water (A) and 0.1% formic acid in acetonitrile (B) was programmed with gradient elution of 30–95% at a flow rate of 250 μ L/min. Mass spectrometric detection was operated in the positive ion mode. Data were processed using Compass DataAnalysis 5.1 software (Bruker Daltonik). All final compounds were $\geq 95\%$ pure. Elemental analysis (% C, H, and N) was performed on a EURO EA elemental analyzer (HEKAtech, Wegberg, Germany). Melting points were determined on an Electrothermal 9001 melting point apparatus (Electrothermal, UK) (10 °C per min) and were uncorrected. Merck KGaA silica gel 60 F₂₅₄ plates were used for analytical thin-layer chromatography. Spots were detected with a UV lamp. Column chromatography was performed using silica gel, Merck 60 (70-230 mesh). Yields refer to purified products, and they were not optimized.

4-Amino-2,6-bis((dimethylamino)methyl)phenol (b), 4amino-2,6-bis(pyrrolidin-1-ylmethyl)phenol (c), and 4amino-2-(pyrrolidin-1-ylmethyl)phenol (d) (see Scheme 2) were prepared according to the procedures as described.^{24,25} The synthetic procedures are described in the Supporting Information. Compounds 5, 6, 7g, and 8g were synthesized as previously described,²⁰ and their physicochemical properties are identical to those described in Puhl et al.²⁰

In Vitro **Testing.** We have previously reported our method of generating SARS-CoV-2-infected US-OS ACE2 GFP cells.²⁶ In short, U2-OS ACE2 GFP cells were split 1 day before infection. Compounds were resuspended in DMSO or H₂O at 20 or 5 mM. These concentrations were sufficient to solubilize the molecules. The experimental compounds were incubated for 2 h at 37 °C in 100 μ L of growth medium (DMEM, 10% FCS, and 1% PS). Ten microliters of SARS-CoV-2 virus (strain BetaCoV/France/IDF0372/2020) was then added to each

well (final MOI, 0.1). Cells were incubated at 37 °C for 20 h, fixed with 8% PFA for 30 min at RT, and washed with PBS. To stain the nuclei and measure the viability, 100 μ L of Hoechst solution was added. The plates were read with an automated confocal microscope (Opera Phoenix), which measures the number of infected cells (GFP signal) and viability (Hoechst signal). The compounds were tested from 50 μ M to 0.64 nM with serial 5-fold dilutions.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsomega.3c05900.

Synthetic procedures, ¹H and ¹³C NMR spectra for all new compounds, and representative HPLC traces (PDF)

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Author Contributions

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Author Contributions

T.J.: data curation, writing, and review and editing; N.M.: investigation, data curation, and review and editing; F.G.-B.: investigation; A.L.: investigation and data Curation; T.B.: investigation; O.S.: investigation, methodology, and resources; T.R.L.: editing; A.C.P.: editing; V.M.: methodology, resources, supervision, and review and editing; S.E.: writing, resources, and review and editing.

Funding

S.E. kindly acknowledges NIH funding R44GM122196-02A1 from NIH NIGMS 1R43AT010585-01 and from NIH/ NCCAM.

Notes

The authors declare the following competing financial interest(s): SE is CEO of Collaborations Pharmaceuticals, Inc. TJ, TRL and ACP are employees at Collaborations Pharmaceuticals, Inc. Collaborations Pharmaceuticals, Inc. has obtained FDA orphan drug designations for pyronaridine, tilorone and quinacrine for use against Ebola. CPI has also filed a provisional patent for use of these molecules against Marburg and other viruses. The other authors declare that they have no conflict of interest.

ABBREVIATIONS

ACE2: angiotensin converting enzyme 2; CC_{50} : concentration of cytotoxicity 50%; COVID-19: coronavirus disease of 2019; DMEM: Dulbecco's modified eagle medium; DMF: *N*,*N*dimethylformamide; DMSO: dimethylsulfoxide; Et: ethyl; FCS: fetal calf serum; FDA: Food and Drug Administration; GFP: green fluorescent protein; IC_{50} : inhibitory concentration 50%; Me: methyl; MOI: multiplicity of infection; M^{pro}: coronavirus main protease; PBS: phosphate buffered saline; PFA: paraformaldehyde; PL^{pro}: papain-like protease; PS: phosphatidylserine; rt: room temperature; SARS-CoV-2: severe acute respiratory syndrome coronavirus 2; SI: selectivity index

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