- 1 Brequinar and Dipyridamole in Combination Exhibits Synergistic Antiviral
- 2 Activity Against SARS-CoV-2 in vitro: Rationale for a host-acting antiviral
- 3 treatment strategy for COVID-19
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- 17 SARS-CoV-2

18 ABSTRACT

19 The continued evolution of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has 20 compromised the efficacy of currently available vaccines and monoclonal antibody (mAb)-based treatment options for COVID-19. The limited number of authorized small-molecule direct-acting 21 antivirals present challenges with pill burden, the necessity for intravenous administration or 22 23 potential drug interactions. There remains an unmet medical need for effective and convenient 24 treatment options for SARS-CoV-2 infection. SARS-CoV-2 is an RNA virus that depends on 25 host intracellular ribonucleotide pools for its replication. Dihydroorotate dehydrogenase 26 (DHODH) is a ubiquitous host enzyme that is required for de novo pyrimidine synthesis. The 27 inhibition of DHODH leads to a depletion of intracellular pyrimidines, thereby impacting viral replication in vitro. Brequinar (BRO) is an orally available, selective, and potent low nanomolar 28 29 inhibitor of human DHODH that has been shown to exhibit broad spectrum inhibition of RNA 30 virus replication. However, host cell nucleotide salvage pathways can maintain intracellular 31 pyrimidine levels and compensate for BRQ-mediated DHODH inhibition. In this report, we show that the combination of BRQ and the salvage pathway inhibitor dipyridamole (DPY) 32 exhibits strong synergistic antiviral activity in vitro against SARS-CoV-2 by enhanced depletion 33 34 of the cellular pyrimidine nucleotide pool. The combination of BRQ and DPY showed antiviral 35 activity against the prototype SARS-CoV-2 as well as the Beta (B.1.351) and Delta (B.1.617.2) 36 variants. These data support the continued evaluation of the combination of BRQ and DPY as a 37 broad-spectrum, host-acting antiviral strategy to treat SARS-CoV-2 and potentially other RNA 38 virus infections.

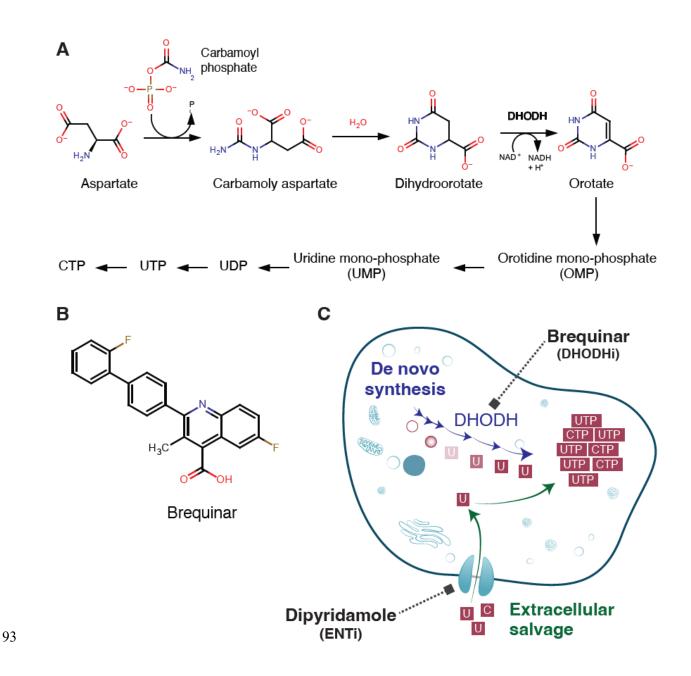
39 INTRODUCTION

40 As of March 2022, there have been more than 6-million reported deaths and greater than 400-41 million cases of coronavirus disease 2019 (COVID-19) worldwide as suggested by the World Health Organization (WHO; https://covid19.who.int/). Furthermore, the number of infections 42 with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), the causative agent of 43 44 COVID-19, likely exceeds the number of reported cases with an estimated excess mortality of 45 18.2 million [1]. Despite the availability of multiple prophylactic vaccines, SARS-CoV-2 46 continues to evolve, compromising the efficacy of vaccines and monoclonal antibody (mAb)-47 based treatment options. Furthermore, there are only a handful of small-molecule antivirals 48 including two RNA-dependent RNA polymerase (RdRp) inhibitors, remdesivir and molnupiravir, and one inhibitor of the SARS-CoV-2 main protease (Mpro), nirmatrelvir 49 50 (nirmatrelvir also requires the pharmacologic booster ritonavir in order to achieve sufficient 51 plasma drug levels). There remains therefore a high unmet medical need for safe, efficacious, 52 and patient-friendly treatments for SARS-CoV-2 infection. 53 While the therapeutic small molecules described above are direct-acting antivirals (DAAs) that 54 target virus-specific proteins, an alternative treatment strategy is to develop host-acting antivirals 55 (HAAs) that target host pathways essential for the viral lifecycle. A host-based mechanism of 56 action may provide unique advantages over DAAs including a greater likelihood of broad-57 spectrum antiviral activity against several families of RNA viruses. In addition, HAAs may 58 possess an inherently higher barrier to the development of resistance when compared to DAAs as 59 host-targets typically remain unchanged in contrast to the rapid emergence of viral variants 60 containing mutations that decrease the efficacy of DAAs.

61	Dihydroorotate dehydrogenase (DHODH) is a host enzyme that is essential for <i>de novo</i>
62	pyrimidine synthesis and has emerged as a candidate target of HAAs. Brequinar (BRQ) is an
63	orally available, selective, and potent low nanomolar human DHODH inhibitor (DHODHI)
64	shown to deplete intracellular uridine, cytidine, and thymidine levels in vitro and in vivo [2, 3, 4].
65	As an antiviral approach, DHODHIs such as BRQ block the host production of cellular
66	pyrimidine nucleotide triphosphates (NTPs) required by viruses for replication [5, 6, 7, 8]. To
67	date, BRQ has been studied in multiple human clinical studies that have included more than 1000
68	subjects with viral infection, hematologic malignancies, or autoimmune disorders [Clear Creek
69	Bio data on file; 2, 3, 9, 10].
70	DHODHIs exhibit potent in vitro activity against several RNA viruses such as SARS-CoV-2,
71	influenza, Zika (ZKV), Dengue (DENV), respiratory syncytial virus (RSV), and Ebola (EBOV)
72	[5, 6, 7, 8]. Despite this promising <i>in vitro</i> data, the successful translation of DHODHI
73	monotherapy showing clinical benefit is lacking. While the DHODHI NITD-982 exhibited in
74	vitro antiviral activity against DENV, the in vivo treatment of infected mice had no effect on
75	viremia [11]. This is likely due to other pathways that can compensate for the reduction of
76	pyrimidine NTPs by DHODH inhibition such as the host nucleoside transporters (equilibrative
77	nucleoside transporters, ENT) that facilitate salvage via extracellular uridine and cytidine [12,
78	13]. The concentration of extracellular uridine in mammalian plasma may range up to 10 μ M
79	[12] and may have contributed to limited clinical efficacy of DHODH inhibition in the oncology
80	setting [2].

A combination of DHODH and nucleotide salvage pathway inhibitors may therefore be required for optimal therapeutic efficacy (**Figure 1**). Consistent with this hypothesis, the antiviral activity

- 83 of a DHODHI against DENV was enhanced by the addition of cyclopentenyl uracil, an inhibitor
- of uridine salvage [14]. BRQ has also been shown to synergize with the ENT1/2/4 inhibitor
- 85 dipyridamole (DPY) which blocks the transport of extracellular pyrimidines needed for *in vitro*
- growth of tumor cell lines [13]. DPY, FDA approved in 1961
- 87 [https://www.accessdata.fda.gov/drugsatfda_docs/label/2019/012836s061lbl.pdf], also inhibits
- 88 platelet aggregation and has widespread clinical use in combination with aspirin for the
- 89 secondary prevention of stroke
- 90 [https://www.accessdata.fda.gov/drugsatfda_docs/label/2012/020884s030lbl.pdf]. We chose to
- 91 evaluate the *in vitro* pharmacologic and antiviral activity of the combination of BRQ and DPY
- 92 (BRQ+DPY) in uninfected and SARS-CoV-2-infected A549/ACE2 cells.



94 Figure 1: Rationale for combined DHODH and salvage pathway inhibition

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95 RESULTS

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96	DPY potentiates the suppression of pyrimidine NIPs by BRQ without apparent cytotoxicity
97	To study the effect of BRQ+DPY on the concentration of cellular pyrimidine nucleotides (CTP

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and UTP), uninfected A549/ACE2 cells were treated either with BRQ, DPY, or in combination,

99 over an 8-hour time course. The A549/ACE2 cell line was derived from the parental lung

100 carcinoma A549 and engineered to express human ACE2, the entry receptor for SARS-CoV-2,

101 for use in antiviral assays. Cells were harvested after treatment at each time point (0, 1, 2, 4, and

102 8 hours) and the NTP concentrations were compared to the control group (T=0, DMSO treated)

103 (**Figure 2**).

104 The concentrations of purine nucleotides (ATP and GTP) were not decreased by any of the

105 treatments. As a single agent, DPY (1 µM) had no impact on either UTP or CTP levels relative

106 to control at time 0 (baseline). Single agent BRQ ($1 \mu M$) exhibited a reduction in pyrimidine

107 NTPs (CTP and UTP) concentrations starting at 4 h (16% and 33% decrease for CTP and UTP,

respectively) and the effect was more pronounced at 8 h (77.5% and 83.32% decrease for CTP

and UTP, respectively) (Figure 2). The combination of BRQ $(1 \mu M) + DPY (1 \mu M)$

110 demonstrated a potentiated effect compared to either agent alone. At 2 h, BRQ+DPY reduced

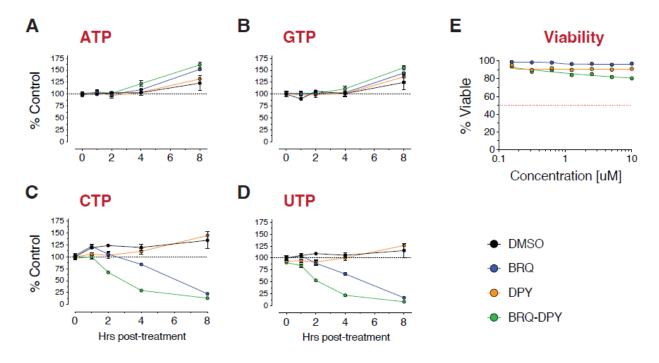
111 CTP and UTP from baseline by 33.6% and 47.3%, respectively. At 4 h, BRQ+DPY reduced CTP

and UTP concentrations from baseline by 71% and 79%, respectively; this was similar to what

113 was seen with BRQ alone at 8 h. At 8h, BRQ+DPY reduced CTP and UTP concentrations from

- baseline by 86.6% and 91.6%, respectively. The effect of BRQ+DPY on pyrimidine NTP levels
- 115 was significant relative to DMSO controls and BRQ as a single agent (Supplementary Tables).

We queried whether the effect of BRQ+DPY on pyrimidine NTP concentrations was driven by general cytotoxicity. The CC₅₀ values for BRQ, DPY, or BRQ+DPY (1:1 ratio) were >10 μ M even after three days of exposure. Notably, these concentrations of BRQ and DPY are 10-fold higher than those used to assay the inhibition of pyrimidine biosynthesis and nucleoside salvage, respectively.



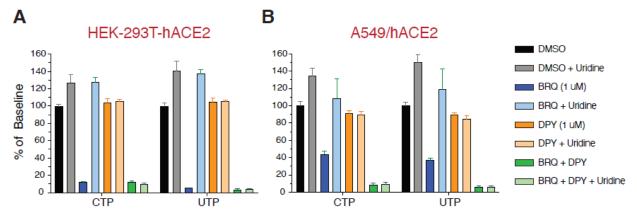
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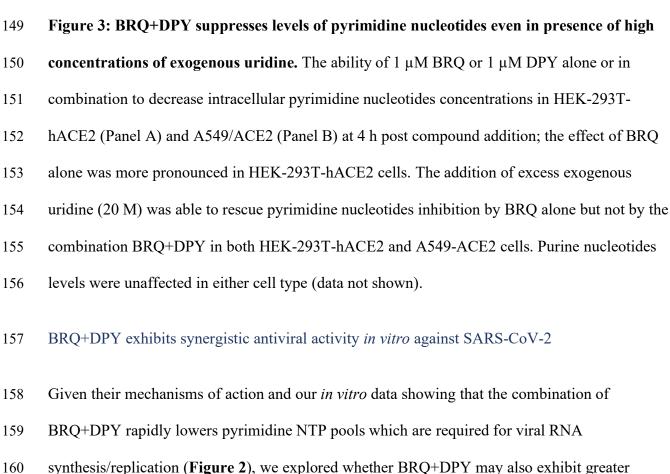
122 Figure 2. DPY potentiates the suppression of pyrimidine NTPs by BRQ without

123 **cytotoxicity. a-d)** Changes in cellular nucleotide pools over 8 hours in response to the treatment 124 of brequinar (BRQ; 1 μ M) and dipyridamole (DPY; 1 μ M) for single and combination treatment. 125 Free NTP concentrations were normalized to the vehicle control group at 0 hours post treatment 126 (n=3 per time point, per group). **e**) A549/ACE2 cell viability following treatment of BRQ and 127 DPY for three days. Cell viability was measured with CellTiter-Glo and compared to the vehicle 128 control group (DMSO treated).

129 BRQ+DPY suppresses pyrimidine nucleotides even with high exogenous uridine

130	Given the mechanism of action as a nucleoside transport inhibitor, we asked whether DPY could
131	potentiate the BRQ-mediated decrease in concentration of pyrimidine NTPs even in the presence
132	of higher concentrations of exogenous uridine. To address this, we repeated the experiment in the
133	presence or absence of uridine (20 μ M) at 4 h post compound addition in HEK-293T-hACE2
134	(Figure 3A) and A549/ACE2 cells (Figure 3B). Pyrimidine NTP levels were substantially
135	reduced with single agent BRQ (1uM) in both cell lines: HEK-293T-hACE2 (88% and 94%
136	decrease for CTP and UTP, respectively) and A549/ACE2 (56% and 63% decrease for CTP and
137	UTP, respectively). The combination of BRQ (1 μ M) and DPY (1 μ M) exhibited greater
138	reduction of pyrimidine NTP levels than BRQ alone in both HEK-293T-hACE2 (87% and 96%
139	decrease for CTP and UTP, respectively) and A549/ACE2 (91% and 94 % for CTP and UTP,
140	respectively) cells; the latter being consistent with our previous experiment (Figure 2).
141	The addition of excess uridine rescued cellular CTP and UTP concentrations in cells treated only
142	with BRQ. In contrast, and consistent with the pharmacological action of DPY, rescue of CTP
143	and UTP levels with excess uridine was not observed in cells treated with BRQ+DPY. Purine
144	NTP levels were not impacted by BRQ or DPY or the combination (data not shown). This result
145	confirms that blocking pyrimidine salvage potentiates the BRQ effect on intracellular pyrimidine
146	NTP concentrations in cells even in the presence of excess concentrations of extracellular
147	uridine.





- 161 antiviral activity than either agent alone.

148

162 BRQ+DPY demonstrates synergistic antiviral activity

163	The antiviral activity of BRQ+DPY inhibition was assessed in A549/ACE2 cells infected with
164	SARS-CoV-2 Beta variant (B.1.351) (Figure 4). DPY treatment alone had no impact on SARS-
165	CoV-2 infection (data not shown). The antiviral activity of BRQ was enhanced in a dose-
166	dependent manner by the addition of DPY (Figure 4A and Table 1). The decrease in EC_{50}
167	values was observed across pharmacologically relevant concentrations of DPY (0.78 μ M or
168	1.563 μ M; [15]); decreases from 2.67 μ M for BRQ alone to 0.80 μ M (3.3-fold lower) and to
169	0.59 μ M (4.5-fold lower), respectively. Similar enhancement of BRQ antiviral activity with the
170	addition of DPY was also observed in separate experiments using Vero cells infected with
171	geographically distinct Wuhan-related SARS-CoV-2 strains (data not shown). The antiviral
172	activity of BRQ+DPY was not attributable to cellular cytotoxicity as cytotoxicity of BRQ and
173	DPY alone or as a combination of BRQ (5.0 μM max) and either DPY at 0.78 μM or 12.5 μM
174	was <50% in the MTT assay (Supplemental Figure S1).
175	To determine whether BRQ+DPY had additive, synergistic, or antagonistic interactions, the
176	antiviral data were analyzed using a combination of Loewe and Bliss models [16]. The 2-drug

177 combination of BRQ+DPY exhibited a strong synergistic interaction (Figure 4B and Table 1).

178 The average synergy score from three replicates was 22.6, with the most synergistic area

179 covering 75.2 arbitrary square units. These data confirm synergistic antiviral activity of

180 BRQ+DPY in the treatment of SARS-CoV-2 at pharmacologically relevant drug concentrations.

181 Exogenous uridine supplementation partially abrogates the synergistic effect

182 The effect of excess exogenous uridine on the antiviral activity of BRQ+DPY was more

183 pronounced at DPY concentrations $<3 \mu$ M where substantial or complete reversal of antiviral

184 activity (e.g., EC_{50} > maximum concentration tested) was observed. DPY concentrations at or

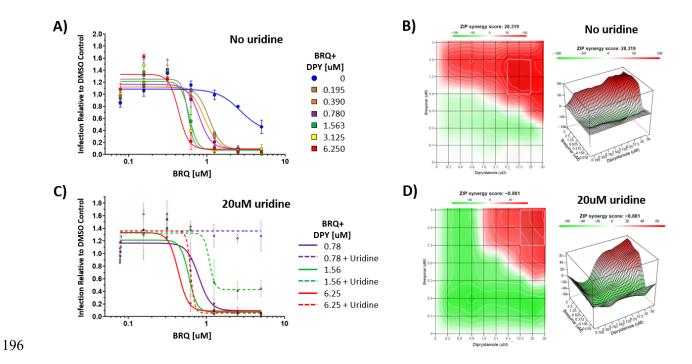
185	above 6.25 μ M decreased the impact of excess uridine (Table 1, examples in Figure 4C).
186	Consistent with the increase in EC $_{50}$ values for BRQ+DPY in the presence of 20 μM uridine,
187	synergy analysis of infection data revealed a shift in synergistic dose combinations to higher
188	concentrations of both drugs (Figure 4D). Furthermore, the synergy score in the presence of
189	excess exogenous uridine was -4.7, which falls in the range of additivity. The additive score is in
190	part confounded by the apparent antagonism of excess uridine at low concentrations of DPY and
191	by the lesser effect at higher DPY concentrations. The physiological relevance of 20 μ M uridine,
192	some 2- to 9-fold higher than physiologic levels [12], is not clear.

Treatment	EC ₅₀ [uM] No Added Uridine	Fold-Change vs BRQ Alone	EC ₅₀ [uM] With 20uM Uridine	Fold-Change vs BRQ Without Added Uridine
DPY Alone	>50	N/A	>50	N/A
BRQ Alone	2.67	1.00	>5	>1.87
BRQ + 0.195uM DPY	1.06	-2.51	>5	>4.70
BRQ + 0.390uM DPY	0.94	-2.85	>5	>5.34
BRQ + 0.789uM DPY	0.80	-3.35	>5	>6.27
BRQ + 1.563uM DPY	0.59	-4.50	1.09	1.84
BRQ + 3.125uM DPY	0.57	-4.66	0.90	1.57
BRQ + 6.250uM DPY	0.44	-6.12	0.63	1.44
BRQ + 12.5uM DPY	0.32	-8.24	0.40	1.23
BRQ + 25uM DPY	0.28	-9.48	0.28	1.00
BRQ + 50uM DPY	0.28	-9.53	0.27	1.00
Synergy				
Synergy Score (Ave)*	22.57	Synergy	-4.66	Additivity
Most Synergistic Area	75.15		58.36	

193 Table 1. BRQ+DPY Synergy vs SARS-CoV-2 Beta (B.1.351)

¹⁹⁴ *Determined with ZIP (Bliss+Loewe); Synergy scores: >10.0 indicates synergy; -10 to 10

195 indicates additivity; < -10 indicates antagonism; N/A: Not Applicable.

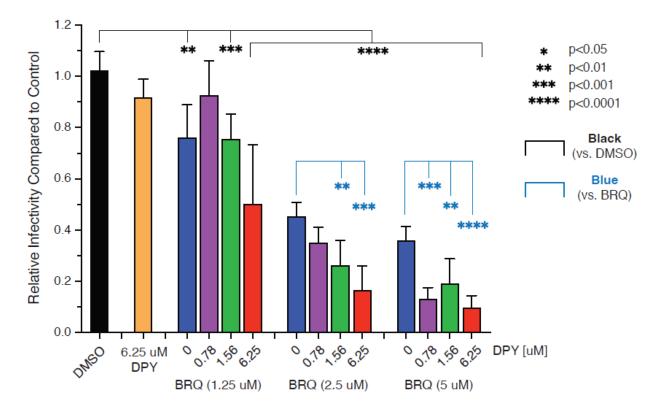


197 Figure 4: BRQ+DPY exhibits strong synergistic antiviral activity that is partially 198 abrogated by supplementation of extracellular uridine. A) The antiviral activity of BRQ 199 alone was enhanced in a dose-dependent manner by the addition of DPY. BRQ at concentrations 200 of $0.078 - 5.0 \mu$ M alone or in combination with DPY at concentrations of $0.195 - 50.0 \mu$ M were 201 evaluated. B) The 2-drug combination of BRQ+DPY exhibited strong synergy with highest 202 effect observed at BRO $0.6 - 2.5 \mu$ M and DPY $6.2 - 25 \mu$ M. C) The addition of excess 203 exogenous uridine reduces the dose-dependent effect of BRO+DPY. D) Addition of 20 µM uridine shifts the area of greatest synergy towards higher concentrations of both BRQ and DPY, 204 205 and synergy scores suggest that uridine addition abrogates synergy still observed at higher DPY 206 concentrations, rendering the effect of the drug combination overall as additive. A549-ACE2 207 cells were dosed with combinations of drugs and challenged with SARS-CoV-2 infection for 208 48 h. Infection was quantified from IFA images stained for SARS-CoV-2 N protein, and synergy 209 was calculated using SynergyFinder2.0 [16] under the LL4 curve and ZIP models. Dark red areas 210 and peaks in the contour plot indicate strong synergistic interaction, with the white box

- 211 representing the area of strongest synergy. Synergy scores greater than 10 indicate a synergistic
- interaction, -10 to 10 suggests additivity, and less than -10 supports antagonism.
- 213 BRQ antiviral activity against SARS-CoV-2 Delta Variant of Concern (B.1.617.2) is enhanced
- 214 *with low concentrations of DPY*

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- 215 Based on the findings with SARS-CoV-2 Beta (B.1.351), the antiviral activity of BRQ+DPY
- 216 was evaluated in the same assay system using the SARS-CoV-2 Delta variant of concern (VOC)
- 217 (B.1.617.2) (Figure 5). BRQ+DPY also exhibited enhanced antiviral activity against SARS-
- 218 CoV-2 Delta (B.1.617.2), relative to DMSO control as well as single agent BRQ.





with low concentrations of DPY. The antiviral activity of BRQ alone was enhanced in a dosedependent manner by the addition of low concentrations of DPY. BRQ at concentrations of 1.25,

223 2.5 and 5.0 μM alone or in combination with DPY at concentrations of 0.78, 1.56 and 6.25 μM
224 were evaluated. A549-ACE2 cells were dosed with combinations of drugs and challenged with
225 SARS-CoV-2 infection for 48 h. Infection was quantified from IFA images stained for SARS226 CoV-2 N protein.

227 DISCUSSION

There remains a significant unmet medical need for safe and convenient treatments for people 228 229 infected with SARS-CoV-2. The most recently authorized antivirals have challenges with pill 230 burden, inconvenient routes of administration, or potential drug interactions. The potential 231 efficacy of DHODHI monotherapy for treatment of viral infection in humans remains to be 232 proven. Clinical experience in the oncology space suggests that the impact of available 233 extracellular uridine and the activity of nucleoside transporters may limit the efficacy of 234 DHODHI monotherapy. The in vitro antiviral activity of BRQ alone against SARS-CoV-2 has 235 been reported by others [17, 18, 19]. Given the growing evidence suggesting that DHODH inhibition alone *in vitro* may not translate to *in vivo* activity, we explored a host-target antiviral 236 237 strategy with the combination BRQ and DPY.

The addition of DPY enhanced the inhibitory effect of BRQ on the concentration of intracellular pyrimidine NTPs *in vitro*. This effect was consistent in the HEK-293T-hACE2 and A549/ACE2 cell lines (**Figs 2-3**) as well as multiple other cell lines and primary cells (data not shown). The depletion of pyrimidine NTP pools by BRQ could be overcome with the addition of excess exogenous uridine (20 μ M) whereas the inhibitory activity of BRQ+DPY was preserved in the setting of high concentrations of extracellular uridine. Furthermore, the inhibitory activity associated with low concentrations of BRQ and DPY was not driven by apparent cytotoxicity.

Within a cell infected with SARS-CoV-2, more than 50% of RNA transcripts are viral RNA 245 transcripts, and all nucleotides are host-derived [20]. Thus, potent reduction of pyrimidine NTP 246 247 pools by the combination of BRQ+DPY should effectively limit viral replication. In A549/ACE2 248 cells (Figure 4), DPY alone had no evidence of antiviral effect against SARS-CoV-2 Beta 249 (B.1.351) and BRQ had marginal single agent activity consistent with published reports [17, 18, 250 19] and Clear Creek Bio data (not shown). Consistent with the enhanced in vitro reduction of 251 pyrimidine NTP levels in uninfected cells with 1 µM BRQ (Figure 2), the addition of DPY 252 increased the antiviral activity of BRQ alone in a dose-dependent manner (Figure 4A and Table 253 1). It is worth noting that as with the *in vitro* analyses of pyrimidine NTP levels, the antiviral activity of BRQ+DPY was evident at concentrations with potential pharmacologic relevance, 254 255 $<2 \mu$ M each [Clear Creek Bio data on file; 9, 15] and was not driven by apparent cytotoxicity 256 even at high concentrations. The analysis of 2-drug antagonism or additivity/synergy using a 257 combination of Loewe and Bliss models demonstrated strong synergy for the combination of 258 BRQ+DPY.

The addition of excess exogenous uridine abrogated the antiviral activity of BRQ in combination with lower concentrations of DPY in SARS-CoV-2 infected A549/ACE2 cells though this abrogation was less pronounced at the higher DPY concentrations (**Figure 4C and Table 1**). The synergy analysis in the presence of 20 μ M uridine shifted from synergy to additivity. These data highlight the importance of extracellular uridine in maintaining cellular pyrimidine NTP levels and supports the concept that inhibiting both *de novo* and salvage pyrimidine pathways with BRQ+DPY merits further exploration.

266	A major area of concern is the development of viral resistance, as seen with selective pressures
267	by DAAs, or viral escape from immune pressures that lead to the emergence of novel SARS-
268	CoV-2 VOCs, as has been observed after vaccination and the development of natural immunity.
269	As BRQ and DPY target host rather than viral proteins, the antiviral activity of BRQ+DPY
270	should have limited liability with respect to viral escape or development of resistance. Our data
271	demonstrated comparable antiviral activity of BRQ+DPY against SARS-CoV-2 strains including
272	those like the original Wuhan-1 as well as Beta and Delta variants of concern.
273	Given the <i>in vitro</i> synergy against multiple SARS-CoV-2 strains in different cell types, a similar
274	enhancement of BRQ antiviral activity may be observed in vivo with the host-based combination
275	treatment of BRQ+DPY. Furthermore, given this synergy and the inability of excess exogenous
276	uridine to reverse this, the HAA combination of BRQ + DPY may be expected to present a high
277	barrier to the development of clinically relevant resistance relative to DAAs. This needs to be
278	formally assessed in clinical trials.
279	In this report, we demonstrated that the combination of BRQ+DPY significantly reduces
280	pyrimidine NTP levels which translates to synergistic antiviral activity against SARS-CoV-2
281	variants in vitro. The antiviral activity observed with BRQ+DPY at pharmacologically relevant
282	concentrations supports continued investigation of this combination as an oral treatment
283	approach for COVID-19. A small, outpatient Phase 2 clinical trial is currently underway to
284	evaluate this concept. [https://clinicaltrials.gov/ct2/show/NCT05166876]. Finally, if this
285	approach is successful, investigation of the use of BRQ+DPY in other RNA viral infections may
286	be warranted.

287 MATERIALS AND METHODS

288 Test Articles

- 289 Brequinar (Selleck Chemicals, Cat# S6626) and dipyridamole (Sigma Aldrich Prod# D9766)
- 290 were provided by Clear Creek Bio. Uridine was obtained from Sigma (Cat#: U3003-5G) and

Alfa Aesar/Fisher (AAA1522706) for the NTP and antiviral assays, respectively.

292 Cell lines and virus

- A549/ACE2 and HEK-293T-hACE2 (BEI resources, NR-53821 and NR-52511) cells were
- cultured in DMEM media supplemented with 10% FBS and 2 mM L-glutamine. Cells were
- 295 passed twice in a week and maintained at 37° C with 5 % CO₂. The absence of mycoplasma
- 296 contamination was validated regularly with a PCR-based method (ATCC, Universal
- 297 Mycoplasma Detection Kit, 30-1012K).

298 In vitro determination of BRQ+DPY effect on pyrimidine NTPs

299 Cells were plated in 6-well plates (200,000 cells/well) one day prior to experiment. Test 300 compounds were dissolved in DMSO as a stock and then diluted in culture media before testing. 301 The final concentration of DMSO was kept at 0.25%. For drug treatment, cell supernatant was 302 removed and replaced with media containing test compounds (n=3 per group). At timepoints 303 denoted, cell lysates were prepared according to a protocol published [21] with minor 304 modifications. In brief, cells were washed with PBS and treated with 0.75 mL of with 0.4 N 305 perchloric acid (PCA) on ice and harvested. Extracts were centrifuged (1500 rpm) and 306 supernatants were combined with a second 0.25 ml extraction. Extracts were combined and

307	neutralized with 10 N and 1 N KOH. Neutralization was determined using pH paper. Samples
308	were stored at -20°C until. The NTP concentrations of samples were analyzed with HPLC.
309	PCA extracts were analyzed using either a Waters 2695e HPLC with a Waters 2489 UV/Visible
310	detector or a Waters 2695 HPLC with a Waters 2487 Dual λ Absorbance detector. A Partisil 10
311	SAX column separated nucleoside triphosphates at a flow rate of 1.5 ml/min with a 50-minute
312	concave gradient curve (curve 8) from $60\% 0.005$ M NH4H2PO4 (pH 2.8) and $40\% 0.75$ M
313	NH4H2PO4 (pH 3.8) to 100% 0.75 M NH4H2PO4 (pH 3.8). Standard ribonucleotides were used
314	to create a standard curve, which was used to quantitate nucleotide pools [21].
315	Analysis of Variance with Dunnett's Multiple Comparisons Test was used to test for significant
316	differences from the DMSO Control at Time 0 vs all other conditions or DMSO vs BRQ, DPY,
317	or BRQ+DPY treatment at each timepoint (Supplemental Tables S1 and S2).
318	For assessment of potential cytotoxicity, cells were seeded in 96-well white plates at a density of
319	12,000 cells per well and cultured overnight. The next day, test compound diluted in cell culture
320	media were added to the cells at the final concentration from 50 μ M to 0.4 μ M, 8 points by 2-
321	fold serial dilution. After three days of incubation, cell viability was evaluated with CellTiter-
322	Glo, measuring luminescence with Synergy 4 (Biotek). 1% of Triton-X100 and 0.25% DMSO
323	were used as the positive and negative control, 0 and 100% cell viability, respectively.
324	BRQ+DPY antiviral experiments
325	A549/ACE2 cells were plated in 96-well plates at a density of 10,000 cells per well in RPMI
326	supplemented with 10% FBS and allowed to adhere overnight. The following day cells were

327 treated with 2-fold dilution series in triplicate of brequinar (ranging 5 μ M – 0.0781 μ M) and

328	dipyridamole (ranging 50 μ M – 0.195 μ M) in a matrix format allowing each concentration pair
329	to be tested. Dilution series of each compound alone were included on each plate, as well as
330	10 μ M remdesivir positive control and DMSO negative control wells. Exogenous uridine
331	(20 μ M) was added to an additional three plates to test whether brequinar inhibition of DHODH-
332	catalyzed uridine synthesis can be overcome by addition of excess uridine. Following one hour
333	of incubation with compounds, cells were challenged with approximately 400 FFU of SARS-
334	CoV-2 Beta (B.1.351; hCoV-19/USA/MD-HP01542/2021) and incubated at 37 °C for 48 h. In a
335	separate experiment using A549/ACE2 cells, dilution series of brequinar and dipyridamole alone
336	or brequinar + 0.78 μ M or 12.5 μ M dipyridamole were tested with the SARS-CoV-2 Delta VOC
337	(B.1.617.2). Pre-treatment time, infection dose, and length of infection were the same as with
338	Beta variant. Analysis of Variance with Dunnett's Multiple Comparisons Test was used to test
339	for significant differences from the DMSO Control vs BRQ or BRQ+DPY treatment or between
340	BRQ alone vs BRQ+DPY.

341 Immunofluorescence detection of SARS-CoV-2 infection efficiency

Following 48 h of infection plates were fixed in 10% formalin for at least 6 h before removal 342 from the high containment laboratory. Plates were washed in PBS, permeabilized in 0.1% Triton 343 344 X-100 for 15 min at room temperature and blocked in 3.5% BSA for at least 1 h at room 345 temperature. To detect infection, plates were incubated overnight at 4 °C with a rabbit anti-N 346 protein monoclonal antibody (SioBiological 40143-R004) diluted 1:20,000. Plates were washed 347 and treated with an AF488-conjugated goat anti-rabbit secondary antibody for 2 h at room 348 temperature. Finally, Hoechst33342 was added to visualize cell nuclei. Plates were imaged on a 349 Cytation 1 Multimode Plate Reader (BioTek) using a 4X objective lens. Infection efficiency,

- defined as GFP-positive cells divided by total nuclei, was calculated for each image using a
- 351 CellProfiler pipeline.
- 352 Synergy calculations
- 353 Drug synergy was calculated with SynergyFinder2.0 [16] using inhibition readout, LL4 curve
- 354 fitting, and ZIP model parameters.

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