

1 ***Short-form paper***

2 **Niclosamide shows strong antiviral activity in a human airway model of SARS-CoV-2 infection and**
3 **a conserved potency against the UK B.1.1.7 and SA B.1.351 variant**

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21 *Running title: Niclosamide is active against SARS-CoV-2 variants*

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24

25 **Abstract**

26 SARS-CoV-2 variants are emerging with potential increased transmissibility highlighting the great unmet
27 medical need for new therapies. Niclosamide is a potent anti-SARS-CoV-2 agent that has advanced in
28 clinical development. We validate the potent antiviral efficacy of niclosamide in a SARS-CoV-2 human
29 airway model. Furthermore, niclosamide is effective against the D614G, B.1.1.7 and B.1.351 variants. Our
30 data further support the potent anti-SARS-CoV-2 properties of niclosamide and highlights its great potential
31 as a therapeutic agent for COVID-19.

32

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34 **Keywords:** COVID-19, small molecule, niclosamide, HAE model, variants of concern, SARS-CoV-2

35

36 **Main Body**

37 Since its emerge in 2019, coronavirus disease 2019 (COVID-19) caused by severe acute respiratory
38 syndrome coronavirus 2 (SARS-CoV-2) led to over 3.1 million deaths worldwide as of April 26, 2021 (1).
39 A tremendous joint research effort led to the approval of several vaccines at unprecedented speed yet anti-
40 viral treatment options remain limited. At the same time, several viral variants harboring mutations in the
41 N-terminal (NTD) and receptor-binding domain (RBD) of the spike protein gene, such as the B.1.1.7 (also
42 named 20I/501Y.V1), B.1.351 (also named 20H/501Y.V2) variants, are causing global concern as they have
43 been associated with enhanced transmissibility and possible resistance to vaccines and antibody
44 neutralization (2–6). The B.1.1.7 and B.1.351 lineages have been linked to a ~50% increased transmission
45 of SARS-CoV-2 infection and the vaccine efficacy of ChAdOx1 nCoV-19 has been reported to be reduced
46 to 10.4% against the B.1.351 variant (6–9). Thus, despite the recent vaccine roll-out, there remains a high
47 unmet need for novel therapeutics against SARS-CoV-2, which should be effective against circulating and
48 potentially emerging variants of concern of SARS-CoV-2.

49
50 Niclosamide has been identified as a potent inhibitor of SARS-CoV-2 *in vitro* and *in vivo* and its optimized
51 formulation for intranasal application and inhalation, was well-tolerated in healthy volunteers in a Phase 1
52 trial (10–13). Herein, we sought to further characterize the anti-viral properties of niclosamide by
53 determining its potency in a human epithelial airway model of SARS-CoV-2 infection and tested its efficacy
54 against several variants of concern of SARS-CoV-2.

55
56 To strengthen the existing data on the potent antiviral activity of niclosamide with a preclinical model
57 resembling the human respiratory tract, we employed a trans-well bronchial human airway epithelium
58 (HAE) model infected with SARS-CoV-2. HAE cultured at an airway-liquid interface has been extensively
59 used as an *in vitro* physiological model mimicking the human mucociliary airway epithelium to validate the
60 effectivity of antivirals on infections in conducting airways (14–16). The effect of niclosamide on the

61 replication of SARS-CoV-2 in the HAE bronchial model (Eptihelix) was determined as previously described
62 by Touret *et al.* (17) and Pizzorno *et al.* (14).

63 Briefly, human bronchial epithelial cells were apically infected with the European D614G strain of SARS-
64 CoV-2 (BavPat1/2020; obtained from EVA-GLOBAL) at a MOI of 0.1 and cultivated in basolateral media
65 that contained different concentrations of niclosamide (in duplicates) or no drug (virus control) for up to 4
66 days. Media was renewed daily containing fresh niclosamide. Remdesivir was used as experimental positive
67 control and non-treated samples as negative control. On day 4, samples were collected at the apical side and
68 the viral titer was estimated with a TCID₅₀ assay. Then, cells were lysed, and the intracellular viral RNA
69 was extracted and quantified by qRT-PCR. The viral inhibition was calculated with the infectious titers by
70 normalizing the response, having the bottom value as 100% and top value as 0%. The IC₅₀ was determined
71 using logarithmic interpolation ($Y=100/(1+10^{((LogEC50-X)*HillSlope)})$) in GraphPad Prims 7. Statistical
72 analysis was performed using the Ordinary One-way Anova with Dunnett's multiple comparisons test.

73
74 Niclosamide exhibited anti-SARS-CoV-2 activity by reducing the infectious titer and intracellular RNA
75 levels in the HAE model in a dose-responsive manner. Niclosamide treatment with concentrations $\geq 1 \mu\text{M}$
76 significantly reduced the infectious titer to below the level of detection at Day 4 post-infection, yielding an
77 IC₅₀ of 0.96 μM (Fig. 1A and 1C). Furthermore, treatment with concentrations $\geq 1 \mu\text{M}$ of niclosamide
78 significantly reduced the intracellular viral RNA level reaching a maximum effect of a 3-fold reduction on
79 Day 4 (Fig. 1B). These data validate the substantial anti-SARS-CoV-2 effect of niclosamide in a
80 reconstituted human airway model.

81
82 We then tested the activity of niclosamide against several variants of concern of SARS-CoV-2, including
83 the BavPat1 strain (D614G), SARS-CoV-2 201/501YV.1 (UVE/SARS-CoV-2/2021/FR/7b; lineage
84 B.1.1.7, ex UK), SARS-CoV-2 Wuhan D614, and SARS CoV-2 SA lineage B.1.351 (UVE/SARS-CoV-
85 2/2021/FR/1299-ex SA) in VeroE6 TMPRSS2 cells (ID 100978, CFAR). All viruses were obtained through
86 EVA GLOBAL. The IC₅₀ were determined by RT-qPCR as previously described by Touret *et al.* (18).

87 Briefly, eight 2-fold serial dilutions of niclosamide in triplicate were added to the cells 15 min prior to viral
88 infection and incubated for 2 days at 37°C. Remdesivir was used as experimental positive control and non-
89 treated samples as negative control. The viral genome was quantified by real-time RT-qPCR from the cell
90 supernatant (17). The IC₅₀ was calculated as described above. All data associated with this study are present
91 in the paper.

92
93 Niclosamide inhibited replication of the SARS-CoV-2 original strain (Wuhan D614) in VeroE6 TMPRSS2
94 cells with an IC₅₀ of 0.13µM and IC₉₀ of 0.16 µM which is in accordance with previous studies (10, 11).
95 Importantly, niclosamide also blocked the replication of the European BavPat D614G, UK B.1.1.7 and SA
96 B.1.351 variant with an IC₅₀ of 0.06 µM, 0.08 µM and 0.07 µM, respectively (Fig. 2). Thus, niclosamide is
97 effective against all tested variants of SARS-CoV-2 having a similar potency across the different strains
98 compared to the original Wuhan D614 strain.

99
100 These data are in line with the host-targeted mode of action of niclosamide, which has been described to
101 interfere with basic cellular mechanisms involved in SARS-CoV-2 replication, such as autophagy, the
102 endosomal pathway and the TMEM16A chloride channel (11, 19–21). Accordingly, niclosamide is a potent
103 antiviral therapeutic agent against SARS-CoV-2 and its variants. The molecule will also deserve further
104 investigations to assess its potential role in the chemotherapeutic armamentarium required for future
105 emerging infectious disease preparedness.

106
107 Taken together, our findings support niclosamide's therapeutic potential as a potent anti-viral agent against
108 SARS-CoV-2, including its variants of concern. Trials in patients with COVID-19 are needed to substantiate
109 future clinical use.

110

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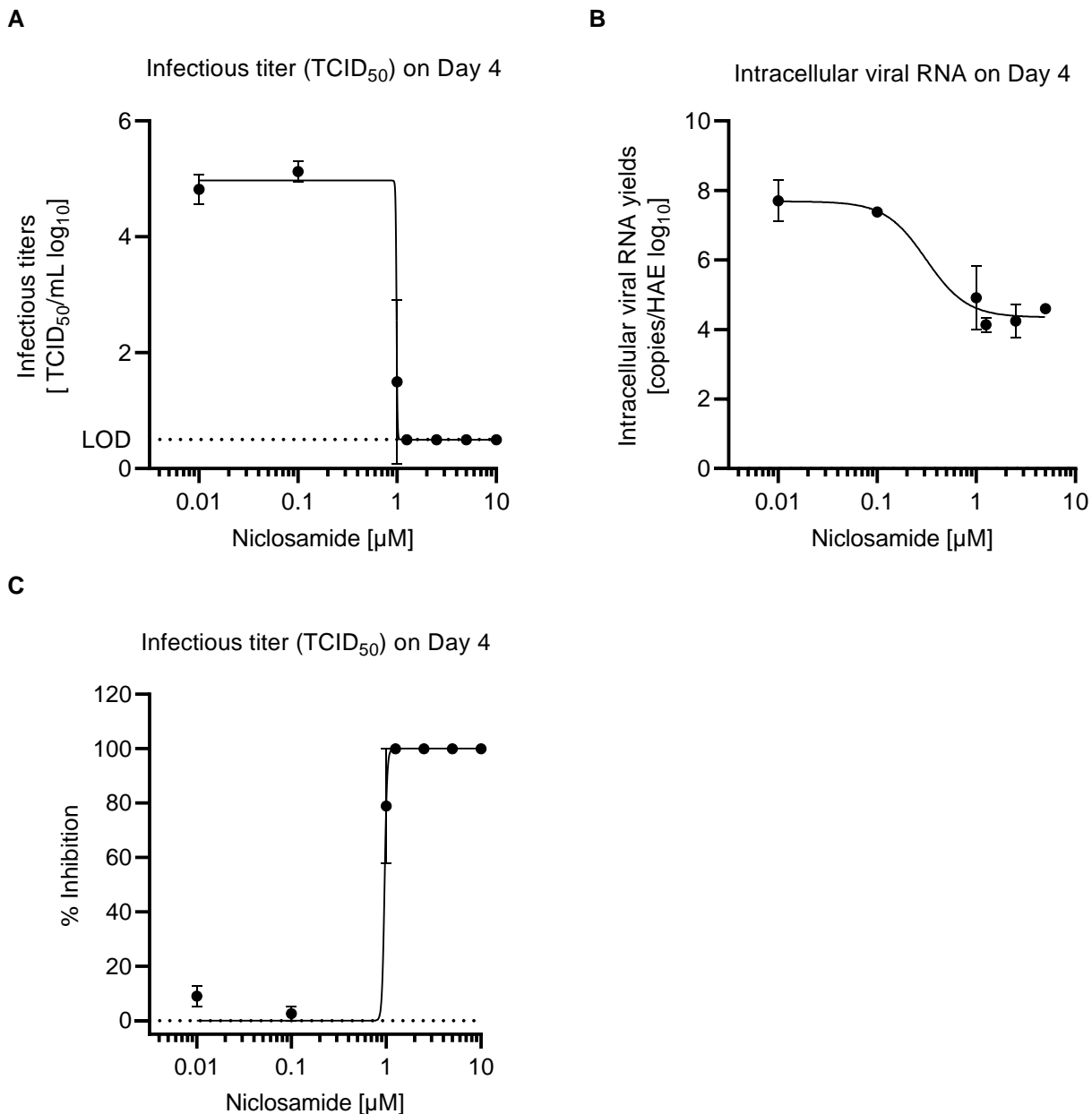
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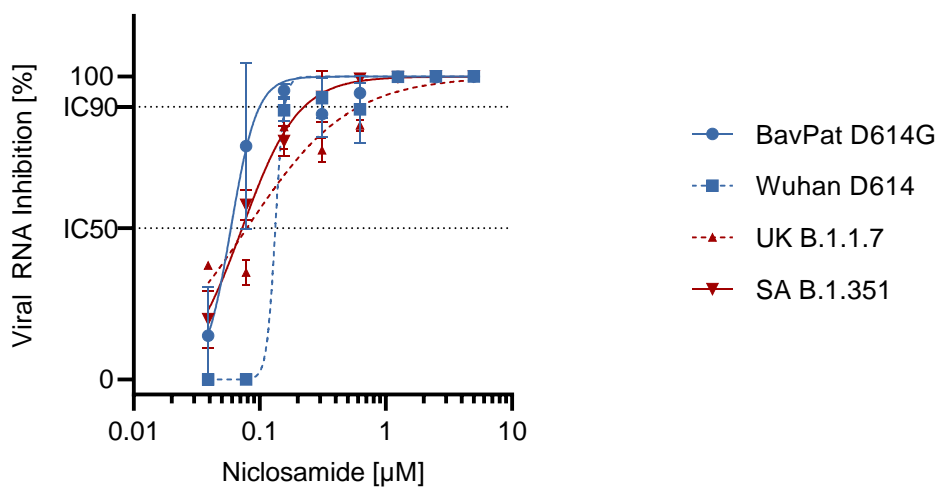
193 **Figures**

194



195
196 **Figure 1: Antiviral efficacy of niclosamide in a trans-well model of human bronchial epithelium**
197 **infected with SARS-CoV-2.** Dose-dependent effects of niclosamide on infectious titer of SARS-CoV-2
198 (A) and intracellular viral RNA levels (B) on Day 4 post-infection. The reduction of infectious titer and
199 intracellular RNA was significant for concentrations $\geq 1 \mu\text{M}$ niclosamide (infectious titer: $1 \mu\text{M} = p < 0.05$,
200 $1.25 - 10 \mu\text{M} = p < 0.0001$; intracellular viral RNA: $1, 2.5, 5 \mu\text{M} = p < 0.01$, $1.25 = p < 0.001$ compared to

201 non-treated control; Ordinary One way Anova with Dunnett's multiple comparisons test). The IC₅₀ based on
202 the infectious titer on Day 4 was 0.96 μM (C). N = 2



	BavPat D614G	Wuhan D614	UK B.1.1.7	SA B.1.351
IC50 [µM]	0.06	0.13	0.08	0.07

203
 204 **Figure 2: Effect of niclosamide on SARS-CoV-2 variants, including UK B.1.1.7 and SA B.1.351 in**
 205 **VeroE6 TMPRSS2 cells. IC = Inhibitory concentration. The origin of the tested variants is available at**
 206 **EVA-GLOBAL. N = 3**