

1 ***In vitro* testing of Hydroxychloroquine and Azithromycin on SARS-CoV-2 shows**
2 **synergistic effect**

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12

13 ***Abstract***

14 Human coronaviruses SARS-CoV-2 appeared at the end of 2019 and led to a pandemic with
15 high morbidity and mortality. As there are currently no effective drugs targeting this virus,
16 drug repurposing represents a short-term strategy to treat millions of infected patients at low
17 costs. Hydroxychloroquine showed an antiviral effect *in vitro*. *In vivo* it also showed efficacy,
18 especially when combined with azithromycin in a preliminary clinical trial. Here we
19 demonstrate that the combination of hydroxychloroquine and azithromycin has a synergistic
20 effect *in vitro* on SARS-CoV-2 at concentrations compatible with that obtained in human
21 lung.

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26 **Background**

27 Since the end of 2019, the world has encountered epidemic conditions attributable to a novel
28 Coronavirus SARS-CoV 2 (1-3). This is the 7th Coronavirus identified to infect Human
29 population (1;4;5) and the first one that had pandemic potential in non-immune populations in
30 the 21st century (6). Finding therapeutics is thus crucial, and it is proposed to do so by
31 repurposing existing drugs (7-9). This strategy presents the advantages that safety profiles of
32 such drugs are known and that they could be easily produced at relatively low cost, thus being
33 quicker to deploy than new drugs or a vaccine. Chloroquine, a decades-old antimalarial agent,
34 an analog of quinine, was known to inhibit the acidification of intracellular compartments
35 (10) and has shown *in vitro* and *in vivo* (mice models) activity against different subtypes of
36 Coronaviruses: SARS-CoV-1, MERS-CoV, HCoV-229E and HCoV-OC43 (11-16). In 2004 it
37 was tested *in vitro* against SARS-CoV 1 (17) and caused a 99% reduction of viral replication
38 after 3 days at 16 μ M. Moreover, tests *in vitro* have shown inhibition of viral replication on
39 SARS-CoV 2 detected by PCR and by CCK-8 assay (18). Hydroxychloroquine
40 (hydroxychloroquine sulfate; 7-Chloro-4-[4-(N-ethyl-N-b-hydroxyethylamino)-1-
41 methylbutylamino]quinoline sulfate) has shown activity against SARS-CoV2 *in vitro* and
42 exhibited a less toxic profile (19). This drug is well known and currently used mostly to treat
43 autoimmune diseases and also by our team to treat Q fever disease (20;21) and Whipple's
44 disease (22;23). In those clinical contexts, concentrations obtained in serum are close to 0.4-1
45 μ g/mL at the dose of 600 mg per day over several months (24). Clinical tests of chloroquine
46 and hydroxychloroquine to treat COVID-19 are underway in China (25), with such trials
47 using hydroxychloroquine in progress in the US (ClinicalTrials.gov Identifier:
48 NCT04307693) and in Europe with the Discovery Trial. In this drug repurposing effort,
49 antibacterial components have also been tested. Teicoplanin, a glycopeptide, was
50 demonstrated *in vitro* to inhibit cellular penetration of Ebola virus (26) and SARS-CoV 2

51 (27). Azithromycin (azithromycin dehydrate), a macrolide, N-Methyl-11-aza-10-deoxo-10-
52 dihydroerythromycin A, has shown antiviral activity against Zika (28-30) . Azithromycin is a
53 well-known and safe drug, widely prescribed in the US, for example, with 12 million
54 treatment courses in children under 19 years of age alone. (31). A recent study has identified
55 these two compounds (azithromycin and hydroxychloroquine) among 97 total potentially
56 active agents as possible treatments for this disease (32).

57 In a preliminary clinical study, hydroxychloroquine and, with even greater potency, the
58 combination of hydroxychloroquine and azithromycin were found effective in reducing the
59 SARS-CoV-2 viral load in COVID-19 patients (33). Since the beginning of the epidemic in
60 the Marseille region we isolated numerous strains and we tested one of them, the SARS-CoV-
61 2 IHUMI-3, using different concentrations of hydroxychloroquine and azithromycin, alone
62 and in combination, with Vero E6 cells.

63 **Materials and Methods**

64 **Viral isolation procedure and viral stock**

65 The procedure of viral isolation of our SARS-Cov 2 strain IHUMI-3 was detailed elsewhere
66 (33). The viral production was done in 75 cm² cell culture flask containing Vero E6 cells
67 (American type culture collection ATCC® CRL-1586™) in MEM with 4% of fetal bovine
68 serum and 1% glutamine. Cytopathic effect was monitored daily under an inverted
69 microscope (Figure 1). After nearly complete cell lysis (approximately 96 hours), viral
70 supernatant was used for inoculation on 96-wells plate.

71 **Testing procedure for drugs**

72 Briefly, we prepared 96-well plates with 5.10⁵ cells/mL of Vero E6 (200µL per well), using
73 Minimum Essential Media (Gibco, ThermoFischer) with 4% of fetal bovine serum and 1%
74 glutamine. Plates were incubated overnight at 37°C in a CO₂ atmosphere. Drug concentrations
75 tested were 1, 2 and 5 µM for hydroxychloroquine and 2, 5 and 10 µM for azithromycin. We

76 also tested combinations of these agents at these concentrations, each test done at least in
77 triplicate. Four hours before infection, cell culture supernatant was removed and replaced by
78 drugs diluted in the culture medium. At t=0, virus suspension in culture medium was added to
79 all wells except in negative controls where 50 μ L of the medium was added. We tested
80 different multiplicities of infection (MOI) at 2.5 and at 0.25. RT-PCR was done 30 minutes
81 post-infection in one plate and again at 60 hours post-infection on a second plate. For this,
82 100 μ L from each well was collected and added to 100 μ L of the ready-use VXL buffer from
83 QIAcube kit (Qiagen, Germany). The extraction was done using the manual High Pure RNA
84 Isolation Kit (Roche Life Science), following the recommended procedures. The RT-PCR was
85 done using the Roche RealTime PCR Ready RNA Virus Master Kit. The primers were
86 designed against the E gene using the protocol of Amrane et al. (34) in the Roche
87 LightCycler® 480 Instrument II.

88 **Results**

89 No cytotoxicity was associated with drugs alone or in combination in controls wells
90 (without viruses). We detected RNA viral production from 24 to 16 cycle-thresholds (Ct,
91 inversely correlated with RNA copy numbers) for the positive control that was associated
92 with cell lysis. In all cases, cell lysis at 60 hours was correlated with viral production as
93 compared to control (Figure 2). At low MOI, azithromycin or hydroxychloroquine alone had
94 no or low impact on the viral production compared to the positive control. We observed only
95 a moderate effect for hydroxychloroquine at 5 μ M in 2 of the 3 replicates (Figure 2a). For the
96 combination of azithromycin and hydroxychloroquine, we observed inhibition of viral
97 replication for wells containing hydroxychloroquine at 5 μ M in combination with
98 azithromycin at 10 and 5 μ M (Figure 2b). Moreover, no cytopathic effect was observed at 60
99 hours post infection in these wells (Figure 3). At high MOI, neither drug showed any effect.

100 The unique observed effect was with the combination of hydroxychloroquine at 2 μ M and
101 azithromycin at 10 μ M, leading to total inhibition of viral replication.

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103 **Discussion**

104 In this present work, we could confirm a moderate effect of hydroxychloroquine alone on
105 SARS-CoV2 at low MOI as previously observed with the lowest concentrations used in a
106 prior study (19). The most striking observation was the synergistic effect of the combination
107 of hydroxychloroquine and azithromycin. As compared to other studies testing
108 hydroxychloroquine for which viral growth was evaluated at 48h, our conditions with
109 prolonged incubation time of 60 hours showed that this effect remained observable. As for
110 MOI, even at the higher MOI of 2.5, as compared to the data of Liu et al. where the highest
111 MOI was of 0.8, the effect of the combination to inhibit viral growth was observable.
112 Hydroxychloroquine has been demonstrated in vitro to inhibit replication of SARS-CoVs 1
113 and 2 (17;19). Concentrations of drugs for our study were based on the known cytotoxicity
114 drugs (50% of cytotoxicity, EC 50) and their effect on microorganisms (50% inhibitory
115 concentration, IC50). With Zika virus, azithromycin showed activity with an IC 50 range
116 from 2.1 to 5.1 μ M depending MOI (28) without notable effect on EC 50 at high
117 concentration (29). On Vero E6 it was shown that for hydroxychloroquine, EC 50 is close to
118 250 μ M (249.50 μ M), which is significantly above the concentrations we tested herein (19).
119 Against SARS-CoV 2, the IC 50 of hydroxychloroquine was determined to be 4.51, 4.06,
120 17.31, and 12.96 μ M with various MOI of 0.01, 0.02, 0.2, and 0.8, respectively.

121 One of the main criticisms of previously published data was that drug concentrations for viral
122 inhibition used in vitro are difficult to translate clinically due to side effects that would occur
123 at those concentrations. The synergy between azithromycin and hydroxychloroquine that we
124 observed herein is at concentrations achieved in vivo and detected in pulmonary tissues (35-

125 37). Our data are thus in agreement with the clinical efficacy of the combination of
126 hydroxychloroquine and azithromycin demonstrated by Gautret et al. (33). They support the
127 clinical use of this drug combination, especially at the early stage of the COVID-19 infection
128 before the patients have respiratory distress syndrome with associated cytokine storm and
129 become less treatable by any antiviral treatment.

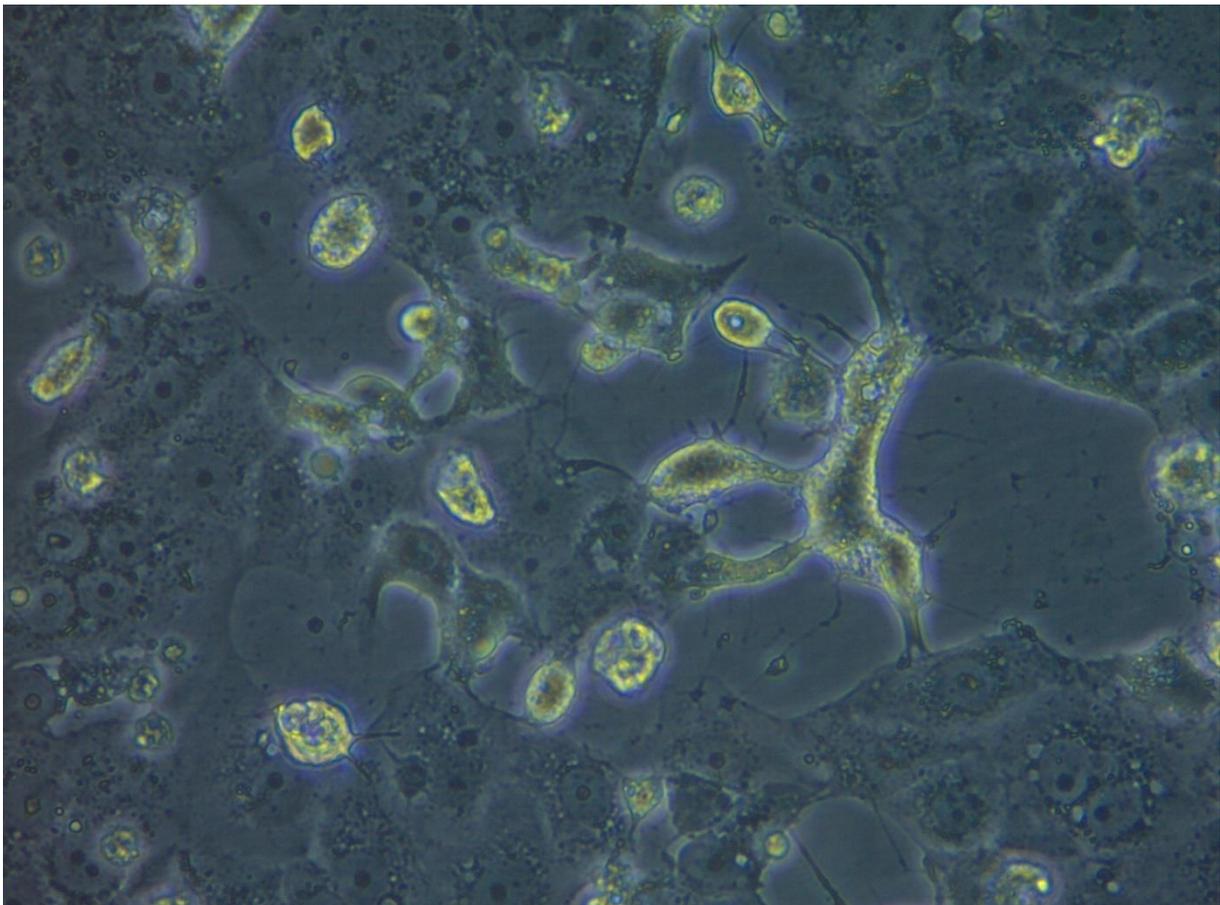
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132 **Figure 1: Observations of infected Vero E6 Monolayer.**

133 Observation was done 48 hours post infection by the SARS-CoV 2 strain IHUMI-3.

134 Magnitude X400. The picture was captured on ZEISS AxioCam ERC 5s.



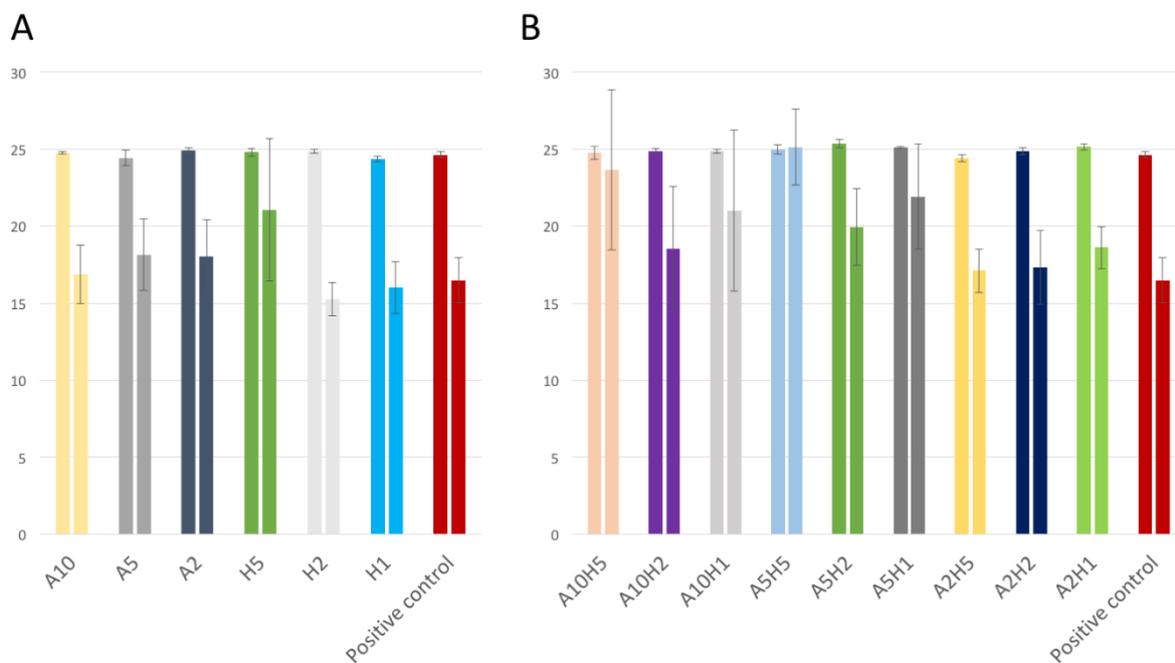
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137 **Figure 2: RNA viral quantification between 0 and 60 hours post infection.**

138 For each condition, the first histogram represents average RNA cycle-thresholds
 139 quantification at H0, and the second histogram represents average RNA viral quantification
 140 60 hours post-infection. Standard deviation scales are present for each condition (n=3 for all
 141 conditions and n=4 for the positive control).

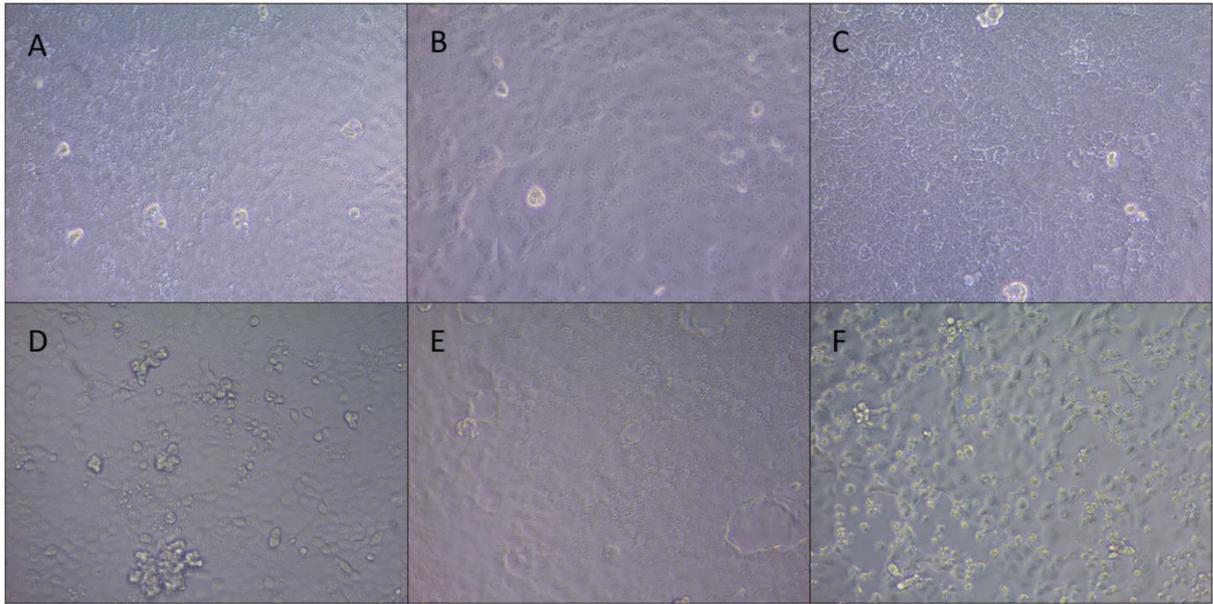
142 **2A.** represents molecules tested alone, A10 is for azithromycin at 10 μ M, A5 at 5 μ M, A2 is
 143 at 2 μ M, H5 is for hydroxychloroquine at 5 μ M, H2 for 2 μ M, H1 for 1 μ M. **2B.** represents
 144 the combination of molecules tested.



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146 **Figure 3: Observations of infected cells resistant or not to viral replication.**

147 Picture were captured on ZEISS AxioCam ERC 5s, 58 hours post infection by the SARS-CoV
 148 2 strain IHUMI-3. Magnitude X200. **3A-B-C.** overview of the monolayer in each well for the
 149 condition of Azithromycin 5 μ M associated with hydroxychloroquine at 5 μ M, **3D.** shows a
 150 cytopathic effect observed in one well in the condition Azithromycin 10 μ M combined with
 151 hydroxychloroquine at 2 μ M **3E.** negative control well and **3F.** positive control well.



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161 **Conflicts of Interest:**

162 The authors declare no conflict of interest.

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166 **References**

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