

# **Hydroxychloroquine and Azithromycin as a Treatment of COVID-19: Results of an Open-Label Non-Randomized Clinical Trial: Response to criticisms**

## **SUPPLEMENTARY DATA**

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### **Supplementary Methods**

We conducted a meta-analysis of studies evaluating the effects of chloroquine derivatives against SARS-CoV-2 in groups of COVID-19 patients as compared to control groups of patients who did not receive chloroquine derivatives. In these studies, groups were expected to be similar with respect to demographics, chronic conditions, clinical presentation at enrolment and use of other antiviral drugs during the course of the disease. The keywords “hydroxychloroquine”, “chloroquine”, “coronavirus”, “COVID-19” and “SARS-Cov-2” were used in the PubMed, Google Scholar and Google search engines without any restrictions as to date (research updated on September, 9, 2020) or language. Preprints were also included. Open reviews and reviewer’s recommendations regarding preprints are available in the supplementary data. Articles published in peer-reviewed journals, pre-prints and articles available on the internet, even when not published on official websites, were included. An overview of most of the screened studies can be accessed at <https://c19study.com/>. The following outcomes were considered: death and persistent viral shedding as assessed by PCR.

Only studies comparing a group of COVID19 patients, mandatorily confirmed by PCR, treated with a chloroquine derivative to a control group without chloroquine derivatives were included. Studies must provide the number of treated and untreated individuals. Non-comparative (single arm) studies and studies comparing two groups treated with chloroquine derivatives at different dosages or with different delay of treatment were excluded. Studies

analyzing safety, efficacy as a prevention, data provided as a webpage without an article format (such as a tweet), were also excluded. Studies without confirmation of the diagnosis by RT-PCR were excluded. For the “mortality” outcome, studies without any death were excluded. For the “viral shedding” outcome, only studies reporting at least the proportion of positive PCR were included. Studies assessing only viral load without data on the proportion of positive samples were excluded.

Studies were classified as “big data” studies when conducted on electronic medical records extracted by public health specialists and epidemiologists who did not care COVID-19 patients themselves. Conversely, studies were classified as “clinical studies” when mentioning details of treatments (dosages, duration, contraindications, monitoring...) and conducted by authors physicians (infectious diseases and internal medicine specialists, and pulmonologists) who cared COVID-19 patients themselves.

The meta-analysis was performed with a randomized model using Comprehensive Meta-Analysis v3 (Biostat, Englewood, NJ, USA) as recommended by Borenstein *et al.* (1). This software made it possible to include dichotomous outcomes (number of events out of the total) and quantitative outcomes (mean in each group, sample size, p-value). The most adjusted effect size reflecting the greatest control for potential confounding factors was extracted. Heterogeneity was considered substantial when  $I^2 > 50\%$ . A p-value  $< 0.05$  was considered significant.

**Supplementary Table 1. Studies assessing the death outcome (at least one death) but excluded and reason for exclusion**

Study	Reason
<p>Ahmad, MedRxiv, 2020 (2)</p> <p><a href="https://www.medrxiv.org/content/10.1101/2020.05.18.20066902v1">https://www.medrxiv.org/content/10.1101/2020.05.18.20066902v1</a></p>	<p>Number of treated and untreated patients not provided</p>
<p>Ayerbe, J Thromb Thrombolysis, 2020 (3)</p> <p><a href="https://link.springer.com/article/10.1007%2Fs11239-020-02162-z">https://link.springer.com/article/10.1007%2Fs11239-020-02162-z</a></p>	<p>Possible duplicate with Mateos Gonzales, MedRxiv, 2020</p>
<p>Calik Basaran, Turk J Med Sci, 2020 (4)</p> <p><a href="https://pubmed.ncbi.nlm.nih.gov/32718127/">https://pubmed.ncbi.nlm.nih.gov/32718127/</a></p>	<p>Diagnosis not confirmed by PCR</p>
<p>Chowdhury, Researchsquare, 2020 (5)</p> <p><a href="https://www.researchsquare.com/article/rs-38896/v1">https://www.researchsquare.com/article/rs-38896/v1</a></p>	<p>Control group treated by doxycycline and ivermectin</p>
<p>Fried, Clin Infect Dis, 2020 (6)</p> <p><a href="https://academic.oup.com/cid/advance-article/doi/10.1093/cid/ciaa1268/5898276">https://academic.oup.com/cid/advance-article/doi/10.1093/cid/ciaa1268/5898276</a></p>	<p>Confounding by indication</p> <p>“Patients treated with hydroxychloroquine were more likely to be on mechanical ventilation compared to those who did not receive hydroxychloroquine (24.9% vs 12.2%).” (1054/4232 vs 913/7489, bilateral khi square test, <math>p &lt; 0.0001</math>)</p>
<p>Horby et al., MedRxiv, 2020 (7)</p> <p><a href="https://www.medrxiv.org/content/10.1101/2020.07.15.20151852v1">https://www.medrxiv.org/content/10.1101/2020.07.15.20151852v1</a></p>	<p>Toxic doses (2400 mg fir the first 24 hours), PCR confirmation was not mandatory</p>

<p>Kelly, Br Pharmacol Soc, 2020 (8)</p> <p><a href="https://bpspubs.onlinelibrary.wiley.com/doi/full/10.1111/bcp.14482">https://bpspubs.onlinelibrary.wiley.com/doi/full/10.1111/bcp.14482</a></p>	<p>Confounding by indication :</p> <p>No approach to control for confounding and treated group with higher CRP (81.5 vs 28, <math>p &lt; .0001</math>), higher FiO<sub>2</sub> requirement median day 0 (24% vs 21%, <math>p &lt; .0001</math>).</p>
<p>Magagnoli, Med, 2020 (9,10)</p> <p><a href="https://www.cell.com/med/pdf/S2666-6340(20)30006-4.pdf">https://www.cell.com/med/pdf/S2666-6340(20)30006-4.pdf</a></p>	<p>Lymphopenia more frequent in the treated group / HCQ started after intubation / Azithromycin given to 30% of control group</p>
<p>McGrail, MedRxiv, 2020 (11)</p> <p><a href="https://www.medrxiv.org/content/10.1101/2020.07.17.20156521v1">https://www.medrxiv.org/content/10.1101/2020.07.17.20156521v1</a></p>	<p>Confounding by indication</p> <p>“The latter two groups were significantly more ill than the untreated group”</p>
<p>Peters, MedRxiv, 2020 (12)</p> <p><a href="https://www.medrxiv.org/content/10.1101/2020.08.14.20173369v1">https://www.medrxiv.org/content/10.1101/2020.08.14.20173369v1</a></p>	<p>HCQ initiation when patients deteriorated</p>
<p>Rivera, Cancer Discovery, 2020 (13)</p> <p><a href="https://cancerdiscovery.aacrjournals.org/content/early/2020/07/21/2159-8290.CD-20-0941">https://cancerdiscovery.aacrjournals.org/content/early/2020/07/21/2159-8290.CD-20-0941</a></p>	<p>Confounding by indication</p>
<p>Sanchez Alvarez, Nefrologia, 2020 (14)</p> <p><a href="https://www.sciencedirect.com/science/article/pii/S201325142030050X">https://www.sciencedirect.com/science/article/pii/S201325142030050X</a></p>	<p>Number of treated and untreated patients not provided</p>

<p>Skipper, Annals of Internal Medicine, 2020 (15)</p> <p><a href="https://www.acpjournals.org/doi/10.7326/M20-4207">https://www.acpjournals.org/doi/10.7326/M20-4207</a></p>	<p>Only 58% of participants received SARS-CoV-2 testing because of severe U.S. testing shortages.</p>
<p>Synolaki, MedRxiv, 2020 (16)</p> <p><a href="https://www.medrxiv.org/content/10.1101/2020.09.05.20184655v1">https://www.medrxiv.org/content/10.1101/2020.09.05.20184655v1</a></p>	<p>Number of treated and untreated patients not provided for the different groups of severity</p>

**Supplementary Table 2. Chloroquine derivatives and COVID19 mortality – Data extracted (as of September 2020, 21)**

	<b>Country</b>	<b>N treated</b>	<b>N untreated</b>	<b>Data in the manuscript</b>	<b>Data entered in the software</b>
<b>CLINICAL STUDIES (POSSIBLE CONFLICT OF INTEREST) (Reference)</b>					
Abd-Elsalam, Am J Trop Med Hyg, 2020 (No) (17)	<b>Egypt</b>	<b>97</b>	<b>97</b>	Table 4. Univariate regression Hydroxychloroquine treatment OR 0.824 (0.243 - 2.797) P = 0.757	<b>Positive direction P = 0.757</b>
Alamdari, Tohoku J Exp Med, 2020 (No) (18)	<b>Iran</b>	<b>427</b>	<b>32</b>	Table 4. Therapies and outcomes. P = 0.028	<b>Negative direction P = 0.028</b>
Alberici, Kidney International, 2020 (No) (19)	<b>Italy</b>	<b>72</b>	<b>22</b>	Table 3   Univariate analyses of the association between clinical characteristics and the risk of ARDS or death in hemodialysis patients with SARS-CoV-2 infection. Hydroxychloroquine: outcome death OR 0.44 (0.16–1.24) p = 0.12	<b>Negative direction P = 0.12</b>
Arshad, Int J Infect Dis, 2020 (I.B. received speakers' bureau honoraria from Gilead) (20)	<b>USA</b>	<b>190 (propensity score matched patients)</b>	<b>190 (propensity score matched patients)</b>	Table 4. Propensity Matched Cox Regression Result for Mortality Prediction	<b>Negative direction P = 0.009</b>

				Given HCQ p-value = 0.009 **, Hazard Ratio 0.487 – (0.285 0.832)	
Cavalcanti, N Eng J Med, 2020 – HCQ alone (No) (21)	<b>Brazil</b>	<b>159</b>	<b>173</b>	Table 2. Primary and Secondary Outcomes (Modified Intention-to-Treat Population).* Death 5/159 vs 5/173	<b>5/159 vs 5/173</b>
Cavalcanti, N Eng J Med, 2020 – HCQ+AZ (No) (21)	<b>Brazil</b>	<b>172</b>	<b>173</b>	Table 2. Primary and Secondary Outcomes (Modified Intention-to-Treat Population).* Death 3/172 vs 5/173	<b>3/172 vs 5/173</b>
D’arminio Monforte, IJID, 2020 – HCQ alone (No) (22)	<b>Italy</b>	<b>197</b>	<b>92</b>	Table 1 Unadjusted and adjusted marginal relative hazards of in-hospital mortality Adjusted HR 0.66 (0.39, 1.11), p = 0.118	<b>Negative direction p = 0.118</b>
D’arminio Monforte, IJID, 2020 – HCQ+AZ (No) (22)	<b>Italy</b>	<b>94</b>	<b>92</b>	Table 1 Unadjusted and adjusted marginal relative hazards of in-hospital mortality Adjusted HR 0.44 (0.24, 0.82), p = 0.009	<b>Negative direction P = 0.009</b>
Goldman, N Eng J Med, 2020 (Funded by Gilead Sciences) (23)	<b>Multinational</b>	<b>109</b>	<b>288</b>	Table S3. Baseline Predictors of Time to Clinical Improvement (with p-values <0.2) / Patients who Died Before Achieving Clinical Improvement (Competing Risks)	<b>10/109 vs 34 / 288</b>

				N (%) / Received hydroxychloroquine yes 10 / 109 vs no 34 / 288	
Guerin, Asian J Med Health, 2020 (No) (24)	<b>France</b>	<b>20</b>	<b>34</b>	“One patient, a man of 82-year-old without comorbidities in the NST group died suddenly;”	<b>0/20 vs 1/34</b>
Heras, Researchsquare, 2020 – HCQ+AZ (No) (25)	<b>Andorra</b>	<b>70</b>	<b>21</b>	Table 3 Risk factors associated with COVID-19 mortality on multivariate analysis Treatment H+A OR 0.044 p = 0.004	<b>Negative direction P = 0.004</b>
Heras, Researchsquare, 2020 – HCQ alone (No) (25)	<b>Andorra</b>	<b>9</b>	<b>21</b>	Table 3 Risk factors associated with COVID-19 mortality on multivariate analysis Treatment H OR 0.32 p = 0.369	<b>Negative direction P = 0.369</b>
Lagier, Trav Med Infect Dis, 2020 (No) (26)	<b>France</b>	<b>503</b>	<b>199</b>	Table 5 Age stratified multivariable analyses adjusted on comorbidities and severity of the disease addressing associations between treatment (HCQ-AZ $\geq$ 3 days) and clinical outcomes/viral shedding clearance (n = 3,737). Weighted Cox regression on	<b>Negative direction P = 0.003</b>



				Unmatched sample (n = 702) Hazard ratio 0.49 (0.31–0.79), p = 0.0030	
Lauriola, Clinical Transl Sci, 2020 – HCQ alone (No) (27)	<b>Italy</b>	<b>17</b>	<b>63</b>	Table 2. Multivariable Cox proportional hazard regression analysis of factors associated with in-hospital death. HCQ (vs. no treatment) 1.108 (0.536-2.293) p = 0.782	<b>Positive direction</b> <b>P = 0.782</b>
Lauriola, Clinical Transl Sci, 2020 – HCQ+AZ (No) (27)	<b>Italy</b>	<b>297</b>	<b>63</b>	Table 2. Multivariable Cox proportional hazard regression analysis of factors associated with in-hospital death. HCQ + azithromycin (vs. no treatment) HR 0.265, 95%CI 0.171-0.412, p<0.001	Exact p-value calculated* : <b>p = 6.67924E-09</b>
Lecronier, Critical care, 2020 (No) (28)	<b>France</b>	<b>38</b>	<b>22</b>	Table 2 Primary and secondary outcomes - 28-day mortality, n (%) standard of care 9/22 vs Lopinavir/ritonavir 7/20 vs hydroxychloroquine 9/38, p = 0.35	<b>9/38 vs 9/22</b>
Ly, IHU preprints, 2020 (No) (29)	<b>France</b>	<b>116</b>	<b>110</b>	Table 3. Associations between multiple factors and SARS-CoV-2 death among 226 infected elderly	<b>Negative direction</b> <b>P = 0.026</b>

				residents (univariate and multivariate analysis) / HCQ/AZ treatment for at least 3 days (226) / Multivariate 0.39 [0.17-0.89] 0.026	
Mahevas, MedRxiv, 2020 (in the final corrected version of the MS published in BMJ : SG reports personal fees and non-financial support from Gilead Sciences / FXL has received personal fees from Gilead / RL reports non-financial support from Eumedica SA, non-financial support from Gilead Sciences / CO reports non-financial support from MSD, non-financial support from Janssen, non-financial support from CSL Behring, non-financial support from Gilead / JMP reports personal fees from Abbvie, personal fees from Gilead / FS reports personal fees from Gilead Sciences /) (30)	<b>France</b>	<b>92</b>	<b>89</b>	Supplementary data 4: Sensitivity analyses* Trimmed sample that was truncated at 10% of the extreme weights.	Events were recalculated and this is explained in : <a href="https://www.mediterranee-infection.com/correction-scientifique/">https://www.mediterranee-infection.com/correction-scientifique/</a> <b>3/92 vs 4/89</b>
Membrillo de Novales, Preprints, 2020 (No) (31)	<b>Spain</b>	<b>123</b>	<b>43</b>	Table 4. Significant outcomes of the multivariate analysis of survival - HCQ treatment P = 0,003 - Exp(B) 0,070 (0,012-0,402)	<b>Negative direction</b> <b>P = 0.003</b>

<p>Paccoud, Clin Infect Dis, 2020 (eurosfordocs reported several authors with conflict of interests particularly Vincent Calvez, Marc Antoine Valantin, Romain Palich – each of them received more than 10,000 euros from Gilead) (32)</p>	<p><b>France</b></p>	<p><b>43</b></p>	<p><b>46</b></p>	<p>Supplementary Data table 2: Results of sensivity analyses - Other sensivity analyses: results on the Secondary population - Time-to-event outcomes evaluated from admission – Death - IPTW-weighted analysis HR 0.52 [0.12; 2.29], p = 0.38</p>	<p><b>Negative direction</b> <b>P = 0.38</b></p>
<p>Pinato, Cancer Research, 2020 (MP has declared consulting/advisory role for Gilead and Bayer /) (33)</p>	<p><b>Multinational</b></p>	<p><b>182</b></p>	<p><b>446</b></p>	<p>Table 3. Model-adjusted risk of mortality complemented by restricted mean survival time analysis according to type of anti-Covid-19 therapy in patients with cancer and SARS-Cov-2 infection – Therapy Antimalarials only (n=182) vs no drug (n=446) / Restricted mean survival time (RMST) analyses: Cox proportional model : HR 0.41 (0.26-0.66) p&lt;0.0001</p>	<p><b>Negative direction</b> <b>P = 0.0001</b></p>
<p>Scholz, Preprints, 2020 (No) (34)</p>	<p><b>USA</b></p>	<p><b>141</b></p>	<p><b>377</b></p>	<p>Table 7. Clinical Outcome in the Treated Patient Group versus the Untreated Patient Group / All-cause death 1/141 vs 13/377</p>	<p><b>1/141 vs 13/377</b></p>

Yu, Sci China Life Sci, 2020 (No) (35)	<b>China</b>	<b>48</b>	<b>502</b>	Table 3 Univariable and multivariable cox proportional hazards model for 60-day fatality after HCQ treatment Adjusted HR (95% CI), 0.36 (0.18–0.75), p = 0.006	<b>Negative direction</b> <b>P = 0.006</b>
<b>BIG DATA STUDIES</b>					
Bernaola, MedRxiv, 2020 (No) (36)	<b>Spain</b>	<b>1498</b>	<b>147</b>	Table 2: Hazard ratio with 95% confidence intervals and Cohen's d for various treatments before and after propensity-score matching, for their effects on mortality rate. Propensity score matching Hazard ratios HCQ 0.84 ± 0.08	<b>Negative direction</b> <b>P = 0.00037 (*calculated from the ratio 0.84 and confidence interval 0.76-0.92)</b>
Catteau, Int J Antimicrob Agents, 2020 (No) (37)	<b>Belgium</b>	<b>4542</b>	<b>3533</b>	« Treatment with HCQ alone was in contrast independently associated with decreased risk of in-hospital mortality (Adjusted hazard ratio [HR] 0.684, 95% confidence interval [CI] 0.617–0.758) compared to the no-HCQ group »	<b>Negative direction</b> <b>P* = 1.96xE-12</b>
Di Castelnuovo, Eur J Intern Med, 2020 (No) (38)	<b>Italy</b>	<b>2634</b>	<b>817</b>	Table 2 Incidence rates and hazard ratios for death in COVID-19 patients, according to hydroxychloroquine use	<b>Negative direction</b> <b>P* = 8.66xE-05</b>

				Propensity score analysis, inverse probability weighting** (primary analysis) HR 0.70 (0.59 to 0.84)	
Gonzalez, MedRxiv, 2020 (No) (39)	<b>Spain</b>	<b>8448</b>	<b>1169</b>	Table 4. Multivariate analysis of mortality. The effect of each factor is expressed as an Adjusted Odds Ratios (CI 95%). Hydroxychloroquine Adjusted OR 0.662 (0.432 to 1.013) p = 0.057	<b>Negative direction</b> <b>P = 0.057</b>
Ip, MedRxiv, 2020 – Inpatients (No) (40)	<b>USA</b>	<b>1914</b>	<b>598</b>	“This retrospective observational cohort study of 2512 hospitalized COVID-19 patients within a 13- hospital network did not find the empirical use of hydroxychloroquine with or without co-treatment with azithromycin to be associated with a reduction in mortality (adjusted HR, 0.99 for any hydroxychloroquine during hospitalization [95% CI, 0.80-1.22]).”	<b>Negative direction</b> <b>P* = 0.93</b>
Ip, MedRxiv, 2020 – Outpatients (AHG reports being a study investigator for Genentech-Hoffman La Roche, during	<b>USA</b>	<b>97 (propensity score)</b>	<b>970 (propensity score)</b>	Table 1 Baseline characteristics and outcomes / Propensity-score-Matched patients (N=1077) / Death	<b>Negative direction</b> <b>p-value = 0.427</b>

the conduct of the study; research funding as study investigator from Acerta, AstraZeneca, Celgene, Kite Pharma, Elsevier's PracticeUpdate Oncology, Gilead) (41)		<b>matched patients)</b>	<b>matched patients)</b>	p-value = 0.427	
Mikami, J Gen Intern Med, 2020 (No) (42)	<b>USA</b>	<b>2077</b>	<b>743</b>	Table 3 Risk Factors Associated with In-Hospital Death Hydroxychloroquine use HR 0.53 (0.41–0.67), p < 0.001	<b>Negative direction</b> <b>P* = 6.6xE-07</b>
Roomi, J Med Internet Res, 2020 (No) (43)	<b>USA</b>	<b>144</b>	<b>32</b>	Table 3: HCQ regression analysis with the outcome Adjusted OR (95%CI) 1.6 (0.33-7.9) p = 0.54	<b>Positive direction</b> <b>P = 0.54</b>
Rosenberg, JAMA, 2020 – HCQ alone (Dr Dufort reported that her spouse has a Gilead Foundation-Focus HIV/HCV testing research grant.) (44)	<b>USA</b>	<b>271</b>	<b>221</b>	Table 3. Model-Adjusted Risk of In-Hospital Death, Cardiac Arrest, Arrhythmia / In-hospital death (hazard ratio) / Hydroxychloroquine alone vs neither drug HR 1.08 (0.63-1.85)	<b>Positive direction</b> <b>P* = 0.79</b>
Rosenberg, JAMA, 2020 – HCQ+AZ (Dr Dufort reported that her spouse has a Gilead Foundation-Focus HIV/HCV testing research grant.) (44)	<b>USA</b>	<b>735</b>	<b>221</b>	Table 3. Model-Adjusted Risk of In-Hospital Death, Cardiac Arrest, Arrhythmia / In-hospital death (hazard ratio) / Hydroxychloroquine + azithromycin vs neither drug HR 1.35 (0.76-2.40)	<b>Positive direction</b> <b>P* = 0.31</b>

Sbidian, MedRxiv, 2020 – HCQ alone (No) (45)	<b>France</b>	<b>623</b>	<b>3792</b>	Table 3. Primary and secondary outcomes according to study population and treatment group / HCQ alone vs. neither drug / AIPTW Estimate* (95%CI) / Whole population / Ratio in average treatment effect / 1.05 (0.77 to 1.33)	<b>Positive direction</b> <b>P* = 0.73</b>
Sbidian, MedRxiv, 2020 – HCQ+AZ (No) (45)	<b>France</b>	<b>227</b>	<b>3792</b>	Table 3. Primary and secondary outcomes according to study population and treatment group / HCQ plus AZI vs. neither drug / AIPTW Estimate* (95%CI) / Whole population / Ratio in average treatment effect / 1.40 (0.98 to 1.81)	<b>Positive direction</b> <b>P* = 0.031</b>
Singh, MedRxiv, 2020 (No) (46)	<b>USA</b>	<b>910 (propensity score matched patients)</b>	<b>910 (propensity score matched patients)</b>	Table 1: Comparison of patient characteristics and outcomes among hospitalized COVID-19 Hydroxychloroquine Treatment group and Control Group / Treatment Hydroxychloroquine vs Control (Matched Cohorts) / Mortality 30-Day / Relative risk (95%CI)	<b>Negative direction</b> <b>P = 0.72</b>

				0.95 (0.74-1.23) p = 0.72	
Sulaiman, MedRxiv, 2020 (No) (47)	<b>Saudi Arabia</b>	<b>1817</b>	<b>3724</b>	Adjusted OR “0.36 (0.16 - 0.8) 0.012”	<b>P = 0.012</b>

CQ: Chloroquine, HCQ: hydroxychloroquine, (H)CQ: chloroquine derivative (HCQ or CQ), OR: Odds ratio, HR: Hazard ratio, Positive direction : Ratio > 1 ((H)CQ associated with higher mortality, Negative direction : ratio < 1 : (H)CQ associated with lower mortality. In the software, the data entered were the number of patients with treatment, without treatment and the effect size data. \*Altman DG, Bland JM. How to obtain the P value from a confidence interval. BMJ. 2011;343:d2304. doi:10.1136/bmj.d2304. Bold: data entered in the CMA software



**Supplementary Table 3. Studies assessing the viral shedding outcome but excluded and reason for exclusion**

Study	Reason
Gautret, Int J Antimicrob Agents, 2020 (48)	Included in Lagier, 2020
Mitja, Clin Infect Dis, 2020 (49)	<p>“The viral load was provided in logarithmic scale; specimens with undetectable viral load at a given follow-up assessment were assigned a value of 3 log<sub>10</sub> copies per mL (i.e., lower limit of detection) for the purpose of statistical analysis.” As mentioned in our methods, we excluded studies that did not mention the proportion of positive. To our opinion, a negative PCR cannot be confused with a positive PCR with 3 log<sub>10</sub> copies DNA/mL.</p>

**Supplementary Table 4. Chloroquine derivatives and COVID19 Viral shedding – Data extracted (as of September 2020, 21)**

<b>Study (conflict of interest)</b>	<b>Country</b>	<b>N treated</b>	<b>N untreated</b>	<b>Data in the manuscript</b>	<b>Data entered in the software</b>
<b>BIG DATA STUDIES</b>					
An, MedRxiv, 2020 – HCQ (No) (50)	South Korea	<b>20 (matched patients)</b>	<b>20 (matched patients)</b>	Table 3. Associations between hydroxychloroquine use and time to viral clearance and symptom duration in crude analysis, multivariable analysis, and propensity-score matching compare to standard supportive therapy. (Conservative therapy is the reference) / Time to viral clearance / Cox regression with matched population (n=20) ** HR 1.53 (0.83-2.94) p = 0.184	<b>Positive direction P = 0.184</b>
<b>CLINICAL STUDIES</b>					
Chen CP, MedRxiv, 2020 – HCQ – RCT (No) (51)	Taiwan	<b>21</b>	<b>12</b>	Table 2. Proportions of negative rRT-PCR assessments on day 14 and median times to negative rRT-PCR results after randomization in the multicenter, open-label, randomized controlled trial / Median time to negative# (Days, 95% CI) P-value*2	<b>Negative direction P = 0.40</b>

				#Time to negative = Event date or censored date – start day / *2 Log-rank test stratified by clinical syndromes 5 (1,9) vs 10 (2,12), p = 0.40	
Chen CP, MedRxiv, 2020 – HCQ – Retrospective study (No) (51)	Taiwan	<b>16</b>	<b>28</b>	“The median times (ranges) to undetected virus were 15 (6–31) days for the HCQ group and 14 (7–22) days for the control group (p = 0.37)”	<b>Positive direction</b> <b>P = 0.37</b>
Chen L, MedRxiv, 2020 – CQ (No) (52)	<b>China</b>	<b>18</b>	<b>12</b>	“Compared with the control group [median day: 7.0 (IQR: 3.0-10.0) days], the chloroquine group [median day: 2.5 (IQR: 2.0-3.8) days] (...) had significant decreases in the number of days required to reach RT-PCR negativity (P=0.006 (...) by Logrank (Mantel-Cox) test, respectively) (Figure 2b).”	<b>Negative direction</b> <b>P = 0.006</b>
Chen L, MedRxiv, 2020 – HCQ (No) (52)	<b>China</b>	<b>18</b>	<b>12</b>	“Compared with the control group [median day: 7.0 (IQR: 3.0-10.0) days], (...) the hydroxychloroquine group [median day: 2.0 (IQR: 2.0-3.5) days] had significant decreases in the number of days required to reach	<b>Negative direction</b> <b>P = 0.010</b>

				RT-PCR negativity ((...) P=0.010 by Logrank (Mantel-Cox) test, respectively) (Figure 2b).”	
Chen J, J Zhejiang U, 2020 – HCQ – RCT (No) (53)	<b>China</b>	<b>15</b>	<b>15</b>	“On day 7, nucleic acid of throat swab was negative in 13 (86.7%) cases in the HCQ group and 14 (93.3%) cases in the control group (p > 0.05).”	<b>2/15 vs 1/15</b>
Huang, J Mol Cell Biol, 2020 – HCQ – RCT (No) (54)	<b>China</b>	<b>10</b>	<b>12</b>	“There were then steady increases in the number of patients turning negative, cumulating at Day 13 when all of the Chloroquine-treated patients became negative (Figure 1B, left panel; Supplementary Table S2). In comparison, patients in the Lopinavir/Ritonavir group only became SARS-CoV-2 negative after 3 days of dosing, and 11 out of 12 turned negative at Day 14.”	<b>0/10 vs 1/12</b>
Huang, MedRxiv, 2020 – CQ – Prospective observational study (No) (55)	<b>China</b>	<b>197</b>	<b>176</b>	Table 2. Outcomes in the overall population with confirmed SARS-CoV-2 infection§. Patients with undetectable viral RNA by Day 10, N (%) 180/197 vs 101/176	<b>Proportion of positive (17/197 vs 75/176)</b>

Kamran, MedRxiv, 2020 – HCQ – RCT (No) (56)	<b>Pakistan</b>	<b>151</b>	<b>349</b>	Table-2. Assessment of Effect of HCQ on RT-PCR status of study population RT-PCR at day 7 / TREATMENT / 167/349 vs 97/151, p = 0.001 (NB: difference in PCR is most important around day 7, see Fig. 3 Lagier, TMAID, 2020)	<b>(proportion of positive PCR at day 7) 167/349 vs 97/151</b>
Kim, MedRxiv, 2020 – HCQ+AZ+Cefixime (No) (57)	<b>South Korea</b>	<b>22</b>	<b>40</b>	“The length of time to viral clearance, which was indicated by negative conversion on PCR after initiation of treatment, was significantly shorter with HQ plus antibiotics than with (...) conservative treatments (HR, 0.44; 95% CI, 0.25 to 0.78).”	<b>Negative direction P* = 0.0047</b>
Lagier, Travel Med Infect Dis, 2020 – HCQ+AZ (No) (26)	<b>France</b>	<b>3119</b>	<b>618</b>	Table 5 Age stratified multivariable analyses adjusted on comorbidities and severity of the disease addressing associations between treatment (HCQ-AZ ≥ 3 days) and clinical outcomes/viral shedding clearance (n = 3,737). Viral shedding persistence ≥ 10 daysf / All patients (n =	<b>Negative direction P* = 3.9E-07</b>

				3,737) / 10.6% vs 20.6%, HR 1.29 (1.17–1.42) p <0.0001	
Lecronier, Crit Care, 2020 – HCQ (No) (28)	<b>France</b>	<b>38</b>	<b>22</b>	Table 4 Virological findings on admission and on day 7 / Respiratory RT-PCR at day 7 / Positive RT-PCR, n (%) 19/26 vs 12/14 (positive / samples analyzed)	<b>19/26 vs 12/14</b>
Shabrawishi, MedRxiv, 2020 – HCQ/CQ (No) (58)	<b>Saudi Arabia</b>	<b>45</b>	<b>48</b>	“The primary endpoint of the study is achieving negative SARS-CoV-2 nasopharyngeal PCR within five days or less from the start of the intervention. Secondary endpoint was achieving negative sample within 12 days or less from the first positive PCR result.” “In group A 73.3% (n= 33) achieved the primary endpoint and 84.4% (n= 38) achieved the secondary endpoint. Smaller percentage of patients 68.8 (n= 33) and 79.2% (n= 38) achieved the primary and secondary endpoints in group B.”	<b>HCQ 33/45 vs 33/48</b>
Tang, MedRxiv, 2020 – HCQ – RCT (No) (59)	<b>China</b>	<b>75</b>	<b>75</b>	“The median time to negative	<b>Positive direction P = 0.34</b>

				conversion was also similar in the SOC plus HCQ group (8 days, 95%CI 5 to 10 days) with that in the SOC group (7 days, 95%CI 5 to 8 days) (Hazard ratio, 0.846; 95%CI, 0.58 to 1.23; p=0.34 by log-rank test) (Figure 2)	
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CQ: Chloroquine, HCQ: hydroxychloroquine, (H)CQ: chloroquine derivative (HCQ or CQ), OR: Odds ratio, HR: Hazard ratio, Positive direction : Ratio > 1 ((H)CQ associated with higher mortality, Negative direction : ratio < 1 : (H)CQ associated with lower mortality. In the software, the data entered were the number of patients with treatment, without treatment and the effect size data. \*Altman DG, Bland JM. How to obtain the P value from a confidence interval. BMJ. 2011;343:d2304. doi:10.1136/bmj.d2304. Bold: data entered in the CMA software.

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