

1 **Isolation of nearly 4000 SARS-CoV-2 shows increase of**
2 **contagiousness associated with alpha then delta variant**

3 Celine Boschi^{1,2,3,4}, Sarah Aherfi^{1,2,3,4}, Philippe Colson^{1,2,3,4}, Didier Raoult^{1,2,3,4} and Bernard La
4 Scola^{1,2,3,4*}

5

6 **Affiliations:**

7 ¹ Microbes, Evolution, Phylogeny and Infection (MEPHI), UM63, Institut de Recherche pour
8 le Développement (IRD), Assistance Publique - Hôpitaux de Marseille (AP-HM), Aix-
9 Marseille University, Marseille, France

10 ² IHU Méditerranée Infection, Marseille, France

11

12 *** Corresponding author:**

13 Bernard La Scola: bernard.la-scola@univ-amu.fr

14 IHU - Méditerranée Infection, 19–21 boulevard Jean Moulin, 13005 Marseille, France. Tel.:

15 +33 413 732 401, Fax: +33 413 732 402.

16

17 **Abstract**

18 Culture inoculation of 6722 nasopharyngeal samples since February 2020 allowed
19 isolation of 3637 SARS-CoV-2 and confirmed that isolation rate is correlated to viral load,
20 regardless symptomatology or vaccination status. Moreover, the delta variant is associated
21 with higher viral loads and therefore higher rates of viral isolation, explaining its greater
22 contagiousness.

23

24 **Introduction**

25 Since February 2020 and diagnosis of first cases of Covid-19 in south of France, our
26 BSL3 laboratory was involved in massive SARS-CoV-2 isolation attempt with up to 3790
27 samples inoculated and 1941 strains isolated [1,2]. As a consequence, we early established a
28 clear correlation between viral load, assessed by cycle threshold values (Ct) obtained by real
29 time reverse transcriptase-PCR (qPCR) and isolation of the virus in cell culture, which is an
30 indirect marker of patient contagiousness. In relation with our studies, it was suggested that
31 asymptomatic SARS-CoV-2-infected patients would have a lower viral load and/or would be
32 less contagious [3]. Furthermore, since the introduction of Covid-19 vaccination, it has been
33 also suggested that patients infected post-vaccine immunization would be less contagious [4].
34 In order to evaluate these two aspects, we analyzed different groups of patients for whom
35 these data were available in our collection. The present study covers different periods of time
36 corresponding each to the predominance in our country of a SARS-CoV-2 genotype:
37 February-September 2020 during which close Wuhan-Hu-1 strain derivatives predominated;
38 October-December 2020 during which the Marseille-4 (a.k.a 20A.EU2 (Nextstrain clade
39 (<https://clades.nextstrain.org/>)) or B.1.160 (Pangolin classification ([https://cov-
41 lineages.org/resources/pangolin.html](https://cov-
40 lineages.org/resources/pangolin.html)))) variant predominated; January-June 2021 during
41 which the alpha variant (Pangolin lineage B.1.1.7) predominated; and most recently, since
42 July 2021, the delta variant (Pangolin lineage B.1.617) predominated.

43 **Materials and methods**

44 Between February 2020 and the 27th of July 2021, a total of 8930 samples have been
45 inoculated in our BSL3 laboratory, including 6731 nasopharyngeal samples. qPCR and
46 culture were performed as previously described [1,2]. In the present work, only
47 nasopharyngeal samples were analyzed, and samples with a Ct>35 at time of diagnosis were
48 excluded because they were associated with <3% of positive culture and were at greater risk

49 of false positive results [2,5]. The information on the symptomatic or asymptomatic status of
50 patients, when available, was collected from the anonymized database of patients tested in our
51 laboratory. For vaccinated people, the information was collected since January 2021, and as
52 part of the present study we defined vaccinated patients as those diagnosed ≥ 15 days after
53 their first injection of any of the four vaccines available in Europe
54 (<https://www.ecdc.europa.eu>).

55 Statistical analysis was performed on GraphPad prism 5.03 using One-way Anova or
56 Mann-Whitney tests. SARS-CoV-2 culture was done as previously described, except for
57 samples inoculated the last study's week during which cultures were only observed for 2
58 weeks (we considered the third week could be omitted as it accounted for only 1% of isolates)
59 [2].

60 **Results**

61 We could isolate 3637 (54%) SARS-CoV-2 from the 6722 patients' samples
62 inoculated. Culture positivity was inversely proportional to Ct at diagnosis as Ct was
63 significantly lower for patients with a positive than negative culture (mean \pm standard
64 deviation: 23.2 \pm 4.83 versus 28.3 \pm 4.9, respectively; $p < 0.0001$) (Figure 1a). Symptomatic or
65 asymptomatic status was known for 3761 and 543 patients respectively. Mean Ct was
66 significantly lower in asymptomatic than symptomatic patients (23.1 \pm 5.9 versus 26.1 \pm 5.4,
67 respectively; $p < 0.0001$) (Figure 1b). Regarding culture isolation, it was positive for 50%
68 (1882/3761) of symptomatic patients compared to 69% (372/543) of asymptomatic patients
69 ($p < 0.0001$) (Figure 1c). Since January 2021, we inoculated 309 samples from patients having
70 received ≥ 1 dose of Covid-19 vaccine. Vaccine administered was Pfizer-BioNTech,
71 AstraZeneca, Moderna, and Janssen in 162, 33, 8 and one patient, respectively; vaccine
72 administered was unknown for 105 patients. In the same period of time, we also inoculated
73 433 samples for unvaccinated patients. Ct at diagnosis were significantly lower for vaccinated

74 patients than unvaccinated patients (21.5 ± 4.5 versus 23.4 ± 5.4 , respectively) (Figure 1d)
75 ($p < 0.0001$). We isolated 80% (249/309) of SARS-CoV-2 among vaccinated patients versus
76 66% (287/433) among unvaccinated patients ($p < 0.0001$) (Figure 1e). For the 150 vaccinated
77 patients for whom this information was available, 134 (89%) were symptomatic and 16 (11%)
78 were asymptomatic (Figure 1f). For the 209 unvaccinated patients for whom this information
79 was available, 167 (80%) were symptomatic and 42 (20%) were asymptomatic. The
80 proportion of symptomatic patients was statistically different from that among vaccinated
81 patients ($p = 0.008$) (Figure 1f). Finally, we tested if the predominant SARS-CoV-2 variant at
82 time of diagnosis was associated with a different Ct and culture positivity rate. We studied
83 four periods: (i) February-September 2020 (Wuhan-Hu-1 close virus derivatives (French
84 original viruses)), (ii) October-December 2020 (predominance of 20A.EU2 variant, in
85 1684/2135 (79%) of the samples with viral genotype available); (iii) January-June 2020
86 (predominance of alpha variant; 9060/14495 (63%) samples with viral genotype available);
87 and (iv) July 2021 (delta variant, 1932/2031 (95%) samples with viral genotype available).
88 There was no significant difference between mean Ct of the periods with the French original
89 SARS-CoV-2 and during which the 20A.EU2 variant predominated (25.6 ± 5.5 versus
90 25.7 ± 5.5 respectively). In contrast, mean Ct was significantly lower for periods during which
91 the alpha (22.6 ± 5.2) then delta variants (19.7 ± 3.4) predominated than for the two former
92 periods, and between January-June than in July ($p < 0.0001$) (Figure 1g). In addition, the
93 culture isolation rate was inversely correlated with the Ct (Figure 1h) ($p < 0.0001$).

94 **Discussion**

95 This work allowed us to still confirm a relationship between qPCR Ct at diagnosis and
96 SARS-CoV-2 isolation [1,2]. The same correlation between culture positivity rate and Ct (as a
97 proxy of viral load) was observed by others authors despite culture viral isolation sensitivity
98 could vary dramatically according to the procedure [6,7]. In our work, such correlation was

99 independent of the period of study, of the presence or absence of clinical symptoms, and of
100 the vaccination status of the patients. Unexpectedly, we observed that patients symptomatic at
101 SARS-CoV-2 diagnosis had lower viral loads and culture isolation rate than asymptomatic
102 patients, regardless they were vaccinated or not (Figures 1b-c, 1f). We believe that this could
103 be due to the fact that asymptomatic patients may test earlier than symptomatic patients in the
104 course of infection, when viral loads are highest. Still more surprising are greater viral loads
105 and culture isolation rates in vaccinated patients compared to those unvaccinated (Figures 1d-
106 e). A bias in selection of patients coming to our institute to be tested for SARS-CoV-2
107 infection is possible, but such correlation has already been observed among healthcare
108 workers immediately after vaccination for whom the absolute risk of testing SARS-CoV-2-
109 positive was increased without obvious explanation [8]. Rate of culture isolation was
110 correlated with the Ct in nasopharyngeal samples and the predominantly circulating SARS-
111 CoV-2 genotype. Indeed, Ct during periods during which for the alpha variant largely
112 predominated was lower than those during which French original SARS-CoV-2 and 20A.EU2
113 variants predominated. This is in agreement with a study of 341 patients infected between
114 November and December 2020 in London that reported that those infected with the alpha
115 variant (n=198 (58%)) had a lower Ct than those infected with a non-alpha variant virus (29
116 versus 32; $p < 0.0011$) [9]. Despite a short period of time (only July 2021) and a small effective
117 of patients (n=94) were analyzed, the delta variant was notwithstanding associated with a
118 lower mean Ct than the alpha variant, an observation in agreement with other recent reports
119 [10]. Indeed, two recent US studies reported similar conclusion, as patients infected with the
120 delta variant had higher viral loads than those infected with former variants, regardless they
121 were vaccinated or not [11]. The lack of information about the proportion of fully-vaccinated
122 patients (2-dose course of vaccine ≥ 14 days before SARS-CoV-2 infection) could be
123 considered a weakness in our work. However, Ct were also similar among samples from

124 patients fully-vaccinated or not in the study by Brown et al. [12] who enrolled 74% of patients
125 fully-vaccinated and 90% infected with the delta variant. Overall, the present work confirms
126 that higher viral loads observed with the delta variant are correlated to higher positivity rate of
127 culture virus isolation, and therefore to higher contagiousness of patients, regardless of the
128 vaccination status of the patients and/or the presence or absence of clinical symptoms.

129

130 **Funding:** This research was funded by the French Government under the “Investissements
131 d’avenir” (Investments for the Future) program managed by the Agence Nationale de la
132 Recherche (ANR, French National Agency for Research), (reference: Méditerranée Infection
133 10-IAHU-03).

134

135 **Conflict of Interest:** The others authors declare no conflict of interest.

136

137 **Ethical approval :** The protocol was approved by the ethical committee of the University
138 Hospital Institute Méditerranée Infection (No.: 2020-029 and 2021-023).

139

140 **Informed consent:** All subjects provided informed consent in accordance with the
141 Declaration of Helsinki.

142

143 **References**

- 144 [1] La Scola B, Le Bideau M, Andreani J, Hoang VT, Grimaldier C, Colson P, et al. Viral
145 RNA load as determined by cell culture as a management tool for discharge of SARS-CoV-2
146 patients from infectious disease wards. *Eur J Clin Microbiol Infect Dis* 2020;39:1059–61.
147 <https://doi.org/10.1007/s10096-020-03913-9>.
- 148 [2] Jaafar R, Aherfi S, Wurtz N, Grimaldier C, Van Hoang T, Colson P, et al. Correlation
149 Between 3790 Quantitative Polymerase Chain Reaction-Positives Samples and Positive Cell
150 Cultures, Including 1941 Severe Acute Respiratory Syndrome Coronavirus 2 Isolates. *Clin*
151 *Infect Dis* 2021;72:e921. <https://doi.org/10.1093/cid/ciaa1491>.
- 152 [3] Rivett L, Sridhar S, Sparkes D, Routledge M, Jones NK, Forrest S, et al. Screening of
153 healthcare workers for SARS-CoV-2 highlights the role of asymptomatic carriage in COVID-
154 19 transmission. *ELife* 2020;9:e58728. <https://doi.org/10.7554/eLife.58728>.
- 155 [4] Levine-Tiefenbrun M, Yelin I, Katz R, Herzel E, Golan Z, Schreiber L, et al. Initial
156 report of decreased SARS-CoV-2 viral load after inoculation with the BNT162b2 vaccine.
157 *Nat Med* 2021;27:790–2. <https://doi.org/10.1038/s41591-021-01316-7>.
- 158 [5] Chang MC, Hur J, Park D. Interpreting the COVID-19 Test Results: A Guide for
159 Physiatrists. *American Journal of Physical Medicine & Rehabilitation* 2020;99:583–5.
160 <https://doi.org/10.1097/PHM.0000000000001471>.
- 161 [6] Huang C-G, Lee K-M, Hsiao M-J, Yang S-L, Huang P-N, Gong Y-N, et al. Culture-
162 Based Virus Isolation To Evaluate Potential Infectivity of Clinical Specimens Tested for
163 COVID-19. *J Clin Microbiol* 2020;58:e01068-20. <https://doi.org/10.1128/JCM.01068-20>.
- 164 [7] Wölfel R, Corman VM, Guggemos W, Seilmaier M, Zange S, Müller MA, et al.
165 Virological assessment of hospitalized patients with COVID-2019. *Nature* 2020;581:465–9.
166 <https://doi.org/10.1038/s41586-020-2196-x>.
- 167 [8] Keehner J, Horton LE, Pfeffer MA, Longhurst CA, Schooley RT, Currier JS, et al.

168 SARS-CoV-2 Infection after Vaccination in Health Care Workers in California. *N Engl J Med*
169 2021;384:1774–5. <https://doi.org/10.1056/NEJMc2101927>.

170 [9] Genomic characteristics and clinical effect of the emergent SARS-CoV-2 B.1.1.7
171 lineage in London, UK: a whole-genome sequencing and hospital-based cohort study - *The*
172 *Lancet Infectious Diseases* n.d. [https://www.thelancet.com/journals/laninf/article/PIIS1473-](https://www.thelancet.com/journals/laninf/article/PIIS1473-3099(21)00170-5/fulltext)
173 [3099\(21\)00170-5/fulltext](https://www.thelancet.com/journals/laninf/article/PIIS1473-3099(21)00170-5/fulltext) (accessed August 5, 2021).

174 [10] Li B, Deng A, Li K, Hu Y, Li Z, Xiong Q, et al. Viral infection and transmission in a
175 large, well-traced outbreak caused by the SARS-CoV-2 Delta variant. *MedRxiv*
176 2021:2021.07.07.21260122. <https://doi.org/10.1101/2021.07.07.21260122>.

177 [11] Riemersma KK, Grogan BE, Kita-Yarbro A, Jeppson GE, O'Connor DH, Friedrich
178 TC, et al. Vaccinated and unvaccinated individuals have similar viral loads in communities
179 with a high prevalence of the SARS-CoV-2 delta variant. *MedRxiv*
180 2021:2021.07.31.21261387. <https://doi.org/10.1101/2021.07.31.21261387>.

181 [12] Brown CM. Outbreak of SARS-CoV-2 Infections, Including COVID-19 Vaccine
182 Breakthrough Infections, Associated with Large Public Gatherings — Barnstable County,
183 Massachusetts, July 2021. *MMWR Morb Mortal Wkly Rep* 2021;70.
184 <https://doi.org/10.15585/mmwr.mm7031e2>.

185

186

187 **Figure 1.** Correlation between viral load evaluated by RT-PCR Ct (a, b, d, g), symptomatology (b, c, f), vaccine status (d, e, f), time of sampling
188 (g, h) and isolation yield (c, e, h).

