



Since January 2020 Elsevier has created a COVID-19 resource centre with free information in English and Mandarin on the novel coronavirus COVID-19. The COVID-19 resource centre is hosted on Elsevier Connect, the company's public news and information website.

Elsevier hereby grants permission to make all its COVID-19-related research that is available on the COVID-19 resource centre - including this research content - immediately available in PubMed Central and other publicly funded repositories, such as the WHO COVID database with rights for unrestricted research re-use and analyses in any form or by any means with acknowledgement of the original source. These permissions are granted for free by Elsevier for as long as the COVID-19 resource centre remains active.



Real-life lack of evidence of viable SARS-CoV-2 transmission via inanimate surfaces: The SURFACE study



José Camilla Sammartino ^{a,1}, Marta Colaneri ^{b,1}, Cecilia Bassoli ^b, Mariaelena Ceresini ^b, Antonio Piralla ^a, Alessandro Ferrari ^a, Elena Percivalle ^a, Fausto Baldanti ^{a,c}, Raffaele Bruno ^{b,c}, Mario U. Mondelli ^{d,e,*}

^a Division of Microbiology and Virology, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy

^b Division of Infectious Diseases I, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy

^c Department of Clinical Surgical Diagnostic & Pediatric Sciences, University of Pavia, Italy

^d Department of Internal Medicine and Therapeutics, University of Pavia, Pavia, Italy

^e Division of Clinical Immunology and Infectious Diseases, Fondazione IRCCS Policlinico San Matteo, Pavia, Italy

ARTICLE INFO

Article history:

Received 6 February 2023

Received in revised form 8 March 2023

Accepted 16 March 2023

Keywords:

Inanimate surfaces

SARS-CoV-2

Contamination

Viable virus

ABSTRACT

Introduction: Although the potential role of inanimate surfaces in SARS-CoV-2 transmission has yet to be adequately assessed, it is still routine practice to apply deep and expensive environmental disinfection protocols. The aim of this study was to verify the presence of viable virus on different surfaces exposed to droplets released by coughing in SARS-CoV-2 RNA positive patients.

Methods: Patients admitted to hospital with a positive SARS-CoV-2 real-time (RT)-PCR swab were asked to cough on steel, cardboard, plastic and their hands. Surfaces were tested at baseline (T_0) and at different timepoints thereafter using swabs dipped in medium, and quickly seeded on VERO E6 cells that were checked every other day for cytopathic effect (CPE). Laboratory-propagated SARS-CoV-2 strains were examined at the same time points and on identical materials.

Results: Ten RNA-positive patients were enrolled into the study. The median cycle threshold value was 20.7 (range 13–28.3). Nasopharyngeal swabs from 3 of the patients yielded viable virus 2–10 days post-inoculation. However, in none of the patients was it possible to isolate viable SARS-CoV-2 from sputum under identical experimental conditions. A CPE was instead already visible using laboratory-propagated SARS-CoV-2 strains at 20', 60', 180' while an effect at 24 h required a 6-day incubation.

Conclusion: The evidence emerging from this real-life study suggests that droplets delivered by SARS-CoV-2 infected patients on common inanimate surfaces did not contain viable virus. In contrast, and in line with several laboratory-based experiments, *in vitro* adapted viruses could survive and grow on the same fomites.

© 2023 The Author(s). Published by Elsevier Ltd on behalf of King Saud Bin Abdulaziz University for Health Sciences. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Introduction

Severe acute respiratory Coronavirus 2 (SARS-CoV-2) is a human coronavirus first isolated in China in December 2019 that rapidly spread worldwide to become a pandemic in March 2020 as declared by the World Health Organization [1]. Since then, several public health interventions were implemented to contrast the pandemic

without tangible success [2]. The development and distribution of vaccines, and the implementation of barrier measures were instrumental in reducing SARS-CoV-2 spreading [3,4]; however, the virus keeps circulating in the community and in hospital settings [5]. A better understanding of the modes of transmission is of paramount importance to help reducing the spread of the infection [6]. One hypothetical transmission mode is *via* fomites since SARS-CoV-2 has been shown to survive for several hours on different surfaces under strict laboratory conditions. Indeed, most studies used aerosols of SARS-CoV-2 strains that were propagated and stocked in the laboratory and used at high concentrations [8–10]. Such potentially contaminated inanimate surfaces may be able to transmit the infection upon touching [7]. Sterilising with UV irradiation or high

* Correspondence to: SC Immunologia Clinica - Malattie Infettive, Fondazione IRCCS Policlinico San Matteo, Italy.

E-mail address: mario.mondelli@unipv.it (M.U. Mondelli).

¹ These authors have equally contributed.

temperatures is generally not possible in a communal setting, thus making these countermeasures unfeasible to contrast the spread of SARS-CoV-2. Last but not least, as the development of new variants hinders the efficacy of the vaccines changing the virus behaviour [11], it is even more important to assess the ability of SARS-CoV-2 to survive in the environment.

Materials and methods

Virus handling and SARS-CoV-2 isolation

All procedures of SARS-CoV-2 isolation, propagation and titration were performed by experienced personnel in a Biosafety Level 3 (BSL3) laboratory. The SARS-CoV-2 European Wild Type strain (D614G, B.1) and omicron variant (BA.1) used in this study for the *in vitro* evaluation of virus survival on fomites, were isolated from nasopharyngeal swabs of symptomatic patients, while, for the *ex vivo* evaluation of SARS-CoV-2 survival, COVID-19 patients' sputum was used. The patients were asked to cough on the designated surfaces, including their own hands, to produce enough sputum to cover about one cm², which was then sampled at each defined time point. Swabs were preserved in transport medium for a maximum of 24 h at 4 °C, then 200 µL from each sample were decontaminated with 50 µL of antibiotics and antifungals (penicillin [50 µg/mL], streptomycin [50 µg/mL], gentamicin [50 µg/mL], neomycin [50 µg/mL], amphotericin B [2.5 µg/mL], glutamine 0.1 %, foetal calf serum 0.05 %, NaHCO₃ 0.4 %) for 30 min at room temperature to remove contaminants. The decontaminated samples were then seeded on confluent VERO E6 cells (VERO C1008 (Vero 76, clone E6, Vero E6); ATCC® CRL-1586™) in a 24-well flat-bottomed tissue-culture microtitre plate (COSTAR, Corning Incorporated, Corning, NY, USA) and incubated at 33 °C in an atmosphere of 5 % CO₂ in air for 1 h. After incubation and inoculum removal, fresh Eagle's MEM (EMEM, Lonza Group Ltd, Basel, Switzerland) supplemented with 1 % v/v penicillin, streptomycin and glutamine (Euroclone SpA) and 0.1 % v/v trypsin, was added before incubation, under the same conditions. The plates were then incubated at 33 °C in 5 % CO₂ and checked every other day for the appearance of a cytopathic effect (CPE).

Whole genome sequencing

All variants for *in vitro* study were identified by complete genome sequencing [12] in order to confirm the presence of variant-defining mutations. Sequences were submitted to GISAID.

Virus titration for *in vitro* evaluation of SARS-CoV-2 survival

After propagation in a 25 cm² cell flask (Corning, NY14831, USA), the titre of each variant was defined as the 50 % tissue culture infectious dose (TCID₅₀) in six replicate wells of a 96-well flat-bottomed tissue-culture microtitre plate. Logarithmic dilutions of stock virus were incubated for 72 h in the presence of 3 × 10⁴ VERO E6 at 33 °C in 5 % CO₂. Cells were observed by light microscopy for CPE development and stained with Gram's crystal violet solution (Merck KGaA, Darmstadt, Germany) plus 5 % v/v formaldehyde 40 % m/v (Carlo Erba SpA, Arese, Italy). The value of TCID₅₀ mL⁻¹ was calculated with the Reed–Muench method [13].

Quantitative real-time PCR (RT-PCR)

To identify SARS-CoV-2 positive patients a RT-PCR was performed using in house developed primer sets as previously reported [14]. Briefly, total RNA was extracted using the QIAamp Viral RNA Mini Kit according to the manufacturer's instructions. The extracted RNA was subjected to a one-step RT-PCR using the SuperScript IV One-Step RT-PCR System (Thermo Fisher Scientific, Waltham, MA,

USA). Cycle threshold (Ct) values were used to screen eligible patients. All patients with a Ct below 30 were included in the study after providing their written informed consent.

Sampling

Patients were asked to provide a sputum sample, immediately after diagnosis of SARS-CoV-2 positivity. The sputum was subsequently distributed on different surfaces, including the palm of the patient's hand, a metal hospital food trail, a piece of cardboard and a piece of plastic. The contaminated surfaces, including the patient's hand, were then secured in plastic bags to preserve the sputum. At each defined time point (20', 60', 180' and 24 h) the residual sputum was sampled using a swab, dipped in 3 mL of medium and immediately transported to the BSL3 laboratory for further processing.

Viral isolation from sputum and fomites

At each time point, 200 µL from each sampled swab were used to verify SARS-CoV-2 viability as reported in the section 'SARS-CoV-2 isolation'. Every day, the plates were observed under an inverted microscope, 10 × magnification, to assess the appearance of CPE. Plates were kept for a maximum of 10 days, and fresh medium was added every 3–4 days to the wells in which no CPE was visible. For the *in vitro* analysis using in-lab propagated SARS-CoV-2 variants, at the same time points, 40 µL of 100, 1000 and 10,000 TCID₅₀ were placed on 2 × 4 cm² stainless steel, plastic and cardboard surfaces. These were kept inside sterile sealable containers and secured in the Glow-Box hood for the duration of the experiment without ventilation. At each time point the inoculum was resuspended in 400 µL Eagle's MEM, supplemented with 1 % v/v penicillin, streptomycin and glutamine (Euroclone SpA) and 0.1 % v/v trypsin, optimal for respiratory virus growth, of which 200 µL were used for SARS-CoV-2 isolation.

Results

A total of 10 patients (6 males and 4 females, median age 61.5 years, range 23.2–79.4 years) were enrolled in the study. Ct value range was 13–28.3, median 20.7. Viable SARS-CoV-2 was isolated from the nasopharyngeal swabs of three patients at T₀, but not from the remainder. The three nasopharyngeal swabs had Ct values equal to 13, 28.3 and 18.7 and were positive at 10-, 2- and 6-days post-inoculation, respectively. A SARS-CoV-2-like CPE was observed in only one patient at T₀ on cardboard, but molecular analysis on the sample failed to amplify viral RNA. In none of the patients was it possible to isolate viable SAR-CoV-2 from sputum at any of the predefined time points (Table 1). In contrast, laboratory-adapted SAR-CoV-2 strains, the European Wild Type strain D614G, B.1

Table 1
Overall results of SARS-CoV-2 Isolation from sputum on fomites.

| Ex vivo Isolation | | | | | |
|-------------------|--------|--------|---------|------|--------------------|
| Time | 20 min | 60 min | 180 min | 24 h | Days post-inoculum |
| Plastic | 0/10 | 0/10 | 0/10 | 0/10 | 1–5 |
| | 0/10 | 0/10 | 0/10 | 0/10 | 6–10 |
| | 0/10 | 0/10 | 0/10 | 0/10 | > 10 |
| Paper | 0/10 | 0/10 | 0/10 | 0/10 | 1–5 |
| | 0/10 | 0/10 | 0/10 | 0/10 | 6–10 |
| | 0/10 | 0/10 | 0/10 | 0/10 | > 10 |
| Metal | 0/10 | 0/10 | 0/10 | 0/10 | 1–5 |
| | 0/10 | 0/10 | 0/10 | 0/10 | 6–10 |
| | 0/10 | 0/10 | 0/10 | 0/10 | > 10 |
| Skin | 0/10 | 0/10 | 0/10 | 0/10 | 1–5 |
| | 0/10 | 0/10 | 0/10 | 0/10 | 6–10 |
| | 0/10 | 0/10 | 0/10 | 0/10 | > 10 |

Table 2
In vitro isolation of lab-adapted SARS-CoV-2 Wild type B.1 strain from fomites.

| Time | 20 min | | | 60 min | | | 180 min | | | 24 h | | | Days Post inoculum | | |
|-----------|--------|-----|------|--------|-----|------|---------|-----|------|--------|-----|------|--------------------|--------|---|
| | TCID50 | 100 | 1000 | 10,000 | 100 | 1000 | 10,000 | 100 | 1000 | 10,000 | 100 | 1000 | | 10,000 | |
| Plastic | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 | |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | 3/3 | 3/3 | - | - | - | 3 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | 3/3 | 4 | |
| | - | - | - | - | - | - | - | 3/3 | - | - | - | 3/3 | 3/3 | - | 5 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 6 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | - | 7 |
| Cardboard | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 | |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | - | - | 3 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 4 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | 3-3 | 5 | |
| | - | - | - | - | - | - | - | - | - | - | 2/3 | 3/3 | - | 6 | |
| | - | - | - | - | - | - | - | - | - | - | 1/3 | - | - | 7 | |
| Metal | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 | |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | - | - | 3 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 4 | |
| | - | - | - | - | - | - | - | - | - | - | 3/3 | 1/3 | 3/3 | 5 | |
| | - | - | - | - | - | - | - | - | - | - | - | 2/3 | - | 6 | |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 7 | |

Table 3
In vitro isolation of lab-adapted SARS-CoV-2 BA.1 strain from fomites.

| Time | 20 min | | | 60 min | | | 180 min | | | 24 h | | | Days Post inoculum | |
|-----------|--------|-----|------|--------|-----|------|---------|-----|------|--------|-----|------|--------------------|--------|
| | TCID50 | 100 | 1000 | 10,000 | 100 | 1000 | 10,000 | 100 | 1000 | 10,000 | 100 | 1000 | | 10,000 |
| Plastic | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | 3/3 | 3/3 | - | - | - | 3 |
| | - | - | - | - | - | - | - | 3/3 | - | - | - | - | 3/3 | 4 |
| | - | - | - | - | - | - | - | - | - | - | 3/3 | 3/3 | - | 5 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 6 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 7 |
| Cardboard | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | - | - | 3 |
| | - | - | - | - | - | - | - | - | - | - | - | - | 3/3 | 4 |
| | - | - | - | - | - | - | - | - | - | - | 3/3 | 3/3 | - | 5 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 6 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 7 |
| Metal | - | - | - | - | - | - | - | - | - | - | - | - | - | 1 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 2 |
| | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | 3/3 | - | - | - | 3 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 4 |
| | - | - | - | - | - | - | - | - | - | - | - | - | 2/3 | 5 |
| | - | - | - | - | - | - | - | - | - | - | 3/3 | 3/3 | 1/3 | 6 |
| | - | - | - | - | - | - | - | - | - | - | - | - | - | 7 |

(Table 2) and omicron subvariant BA.1 (Table 3) remained viable and produced CPE in each of the in vitro experimental conditions at different time points after inoculation.

Discussion

In this study, we provide definitive real-life evidence that contamination of inanimate surfaces with viable SARS-CoV-2 should be regarded as exceedingly rare and very short lasting. This is in contrast with data using laboratory-generated viruses, which showed robust and long-lasting virus viability.

Patient enrolment began in March 2022 when the prevalence of the omicron variant reached 98.3 %. The omicron variant (subvariants BA.1, BA.2, BA.2.12.1, BA. 4 and BA.5) involved new mutations in the spike protein, most of which are in the receptor binding domain (RBD), increasing its transmissibility which resulted in a decreased monoclonal antibody and vaccine response due to reduced fusogenesis [15]. Moreover, omicron shows a wider diffusion potential than previously isolated variants of concern (VOCs) and

does not preferentially affect the lower respiratory tract, being characterised by a non-TMPRSS-dependent cellular entry mechanism [15]. Furthermore, it is usually associated with milder symptoms than other VOCs in immunocompetent individuals [16–18].

Previous studies found that the recovery of infectious virus from cell culture inversely correlated with the presence of SARS-CoV-2 specific IgG in the respiratory tract, specific for both prior VOCs or monovalent vaccines [19]. Moreover, recovery of infectious virus was more frequent from samples with lower mean Ct values, regardless of the subvariant whereas, when Ct values < 20 were considered, BA.5 showed a statistically higher recovery rate of infectious virus compared with other subvariants [20]. In our study, we did not find viable virus on different surfaces, even if a low Ct was obtained from nasal swabs of the enrolled individuals. We can speculate that vaccine-induced or hybrid immunity may have played a role, even though previous findings failed to show viable virus on inanimate surfaces in hospital settings, before the advent of mRNA vaccines

[21]. Indeed, our patients were all vaccinated, enjoying robust protection by hybrid immunity.

Another interesting hypothesis regarding reduced virus viability in real-life conditions is provided by a recent study showing that mucin inhibits the infection of cells by human betacoronavirus OC43 in a concentration- and glycan-dependent manner [21]. This observation is consistent with low rates of fomite transmission of SARS-CoV-2 in the real-world [22], and suggests that mucins may be one culprit.

Despite dichotomous findings were clearly produced in real-life and laboratory conditions, there is still uncertainty on the presence of SARS-CoV-2 in the nosocomial and household settings, as some studies found extensive SARS-CoV-2 RNA contamination on inanimate surfaces in hospitals dedicated to patients with COVID-19 [24,25], while others did not [23]. Furthermore, a recent systematic review found that there is currently a lack of studies assessing SARS-CoV-2 viability on inanimate surfaces [26]. Therefore, until now, it was impossible to assess the real transmission potential of SARS-CoV-2 via surfaces.

Our study has been designed to precisely fill this gap of knowledge, comparing real-life with laboratory-based conditions. Indeed, some of the existing studies in the field are apparently at variance with our findings, providing direct evidence of SARS-CoV-2 viability on fomites for a length of time consistent with the possibility of onward transmission [27,28]. However, it must be emphasised that in those studies viable SARS-CoV-2 could be retrieved from fomites that were in continuous and long-lasting close contact with the patients, such endotracheal tubes [27]. Furthermore, hands, which in another study had shown a high infectious potential [29], were consistently negative for viable virus in our real-life study. Omicron's markedly different behaviour from previously isolated VOCs, may partially explain discrepancies with previous VOCs that were apparently characterised by a more prolonged viral persistence on fomites, using both laboratory adapted and infected patients' sputum-derived SARS-CoV-2 [3].

Although prudence suggests not to let our guard down, our findings provide evidence in support of somehow debunking the role of inanimate surfaces in SARS-CoV-2 transmission.

Conclusions, strengths and limitations

To our knowledge, this is the first study reporting a head-to-head comparison of *in vitro* propagated and *ex vivo* isolated SARS-CoV-2 survival in the environment under similar experimental conditions. With respect to the *ex vivo* experiments we aimed at reproducing real-life conditions using different inanimate materials and the patients' own hands. Moreover, the initial studies on SARS-CoV-2 contamination of inanimate surfaces were performed early in the course of the pandemic, before the appearance of VOCs. Thus, this study, which was designed at a time in which omicron subvariants prevailed, represents a significant new addition to the field. Finally, the study also focused on the patients' skin and not only on fomites, therefore mimicking a previously unexplored real-life setting.

It may be argued that the role of superspreaders has not been investigated. There are several factors that determine how someone transmits a virus. There are some people who, because of their own personal biology, they produce a higher percentage of aerosols *versus* respiratory droplets. The hypothesis is that some of these people are super-emitters, and they are responsible for superspreading events. We believe that in this setting, in which patients were asked to cough on surfaces under controlled conditions, data variability according to different virus-emitting potential is somehow attenuated.

Our main limitation is the small sample size, which is mostly attributable to low patients' compliance, and stringent Ct values cut-off. Moreover, we were unable to collect enough material from the sputum samples to perform both isolation of viable virus and

molecular quantitation of SARS-CoV-2 copies. However, the high reproducibility of the data significantly attenuates this relative limitation. We also lacked an *in vitro* replacement for the skin contamination setting, for obvious reasons. However, in consideration of the results obtained by Hirose R. et al. [3] and our own previous data [23], we see this as a minor limitation of the study.

Funding

This research was supported by EU funding within the Next Generation EU-MUR PNRR Extended Partnership initiative on Emerging Infectious Diseases (Project no. PE00000007, INF-ACT).

Conflict of interest

We have no conflict of interest to declare.

References

- [1] Jee Y. WHO international health regulations emergency committee for the COVID-19 outbreak. *Epidemiol Health* 2020;42:e2020013. <https://doi.org/10.4178/epih.e2020013>. Epub 2020 Mar 19. PMID: 32192278; PMCID: PMC7285442.
- [2] WHO; 2022. (<https://covid19.who.int/>) [Accessed on the 22th of December 2022].
- [3] Hirose R, Ikegaya H, Naito Y, Watanabe N, Yoshida T, Bandou R, et al. Survival of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) and influenza virus on human skin: importance of hand hygiene in coronavirus disease 2019 (COVID-19). *Clin Infect Dis* 2021;73(11):e4329–35. <https://doi.org/10.1093/cid/ciaa1517>
- [4] Lopez Bernal J, Andrews N, Gower C, Robertson C, Stowe J, Tessier E, et al. Effectiveness of the Pfizer-BioNTech and Oxford-AstraZeneca vaccines on covid-19 related symptoms, hospital admissions, and mortality in older adults in England: test negative case-control study. *BMJ* 2021;373:n1088. <https://doi.org/10.1136/bmj.n1088>
- [5] Singanayagam A, Hakki S, Dunning J, Madon KJ, Crone MA, Koycheva A, et al. ATACC Study Investigators. Community transmission and viral load kinetics of the SARS-CoV-2 delta (B.1.617.2) variant in vaccinated and unvaccinated individuals in the UK: a prospective, longitudinal, cohort study. *Lancet Infect Dis* 2022;22(2):183–95. [https://doi.org/10.1016/S1473-3099\(21\)00648-4](https://doi.org/10.1016/S1473-3099(21)00648-4)
- [6] Sharma A, Ahmad Farouk I, Lal SK. COVID-19: a review on the novel coronavirus disease evolution, transmission, detection, control and prevention. *Viruses* 2021;13(2):202. <https://doi.org/10.3390/v13020202>. PMID: 33572857; PMCID: PMC7911532.
- [7] Tharayil A, Rajakumari R, Mozetic M, Primc G, Thomas S. Contact transmission of SARS-CoV-2 on fomite surfaces: surface survival and risk reduction. *Interface Focus* 2021;12(1):20210042. <https://doi.org/10.1098/rsfs.2021.0042>. PMID: 34956610; PMCID: PMC8662391.
- [8] Virtanen J, Aaltonen K, Kivistö I, Sironen T. Survival of SARS-CoV-2 on clothing materials. *Adv Virol* 2021;2021:6623409. <https://doi.org/10.1155/2021/6623409>. PMID: 33927762; PMCID: PMC8049815.
- [9] Raiteux J, Eschlimann M, Marangon A, Rogée S, Dadvisard M, Taysse L, et al. Inactivation of SARS-CoV-2 by simulated sunlight on contaminated surfaces. *Microbiol Spectr* 2021;9(1):e0033321. <https://doi.org/10.1128/Spectrum.00333-21>
- [10] Gidari A, Sabbatini S, Bastianelli S, Pierucci S, Busti C, Bartolini D, et al. SARS-CoV-2 survival on surfaces and the effect of UV-C light. *Viruses* 2021;13(3):408. <https://doi.org/10.3390/v13030408>. PMID: 33807521; PMCID: PMC7998261.
- [11] Escalera A, Gonzalez-Reiche AS, Aslam S, Mena I, Laporte M, Pearl RL, et al. Mutations in SARS-CoV-2 variants of concern link to increased spike cleavage and virus transmission. *Cell Host Microbe* 2022;30(3):373–387.e7. <https://doi.org/10.1016/j.chom.2022.01.006>
- [12] Alteri C, Cento V, Piralla A, Costabile V, Tallarita M, Colagrossi L, et al. Genomic epidemiology of SARS-CoV-2 reveals multiple lineages and early spread of SARS-CoV-2 infections in Lombardy, Italy. *Nat Commun* 2021;12(1):434. <https://doi.org/10.1038/s41467-020-20688-x>. PMID: 33469026; PMCID: PMC7815831.
- [13] Reed IJ, Muench H. A simple method of estimating fifty per cent endpoints. *Am J Epidemiol* 1938;27:493–7. <https://doi.org/10.1093/oxfordjournals.aje.a118408>
- [14] Gaïarsa S, Giardina F, Batisti Biffignandi G, Ferrari G, Piazza A, Tallarita M, et al. Comparative analysis of SARS-CoV-2 quasispecies in the upper and lower respiratory tract shows an ongoing evolution in the spike cleavage site. *Virus Res* 2022;315:198786. <https://doi.org/10.1016/j.virusres.2022.198786>
- [15] Planas D, Saunders N, Maes P, Guivel-Benhassine F, Planchais C, Buchrieser J, et al. Considerable escape of SARS-CoV-2 Omicron to antibody neutralization. *Nature* 2022;602(7898):671–5. <https://doi.org/10.1038/s41586-021-04389-z>
- [16] Kozlov M. Omicron's feeble attack on the lungs could make it less dangerous. *Nature* 2022;601(7892):177. <https://doi.org/10.1038/d41586-022-00007-8>
- [17] Omicron: a shift in the biology of SARS-CoV-2. *Nat Microbiol*, 7(8); 2022 Aug. p. 1114–5. doi: [10.1038/s41564-022-01149-1](https://doi.org/10.1038/s41564-022-01149-1). PMID: 35821416.

- [18] Bálint G, Vörös-Horváth B, Széchenyi A. Omicron: increased transmissibility and decreased pathogenicity. *Signal Transduct Target Ther* 2022;7(1):151. <https://doi.org/10.1038/s41392-022-01009-8>. PMID: 35525870; PMCID: PMC9077027.
- [19] Fall A, Eldesouki RE, Sachithanandham J, Morris CP, Norton JM, Gaston DC, et al. The displacement of the SARS-CoV-2 variant Delta with Omicron: an investigation of hospital admissions and upper respiratory viral loads. *EBioMedicine* 2022;79:104008. <https://doi.org/10.1016/j.ebiom.2022.104008>. Epub 2022 Apr 20. PMID: 35460989; PMCID: PMC9020587.
- [20] Morris CP, Eldesouki RE, Sachithanandham J, Fall A, Norton JM, Abdullah O, et al. Omicron subvariants: Clinical, laboratory, and cell culture characterization. *medRxiv [Prepr]* 2022. <https://doi.org/10.1101/2022.09.20.22280154>
- [21] Wardzala CL, Wood AM, Belnap DM, Kramer JR. Mucins inhibit coronavirus infection in a glycan-dependent manner. *ACS Cent Sci* 2022;8:351–60. <https://doi.org/10.1021/acscentsci.1c01369>. PMID: 35345395. PMCID: PMC8864775.
- [22] Mondelli MU, Colaneri M, Seminari EM, Baldanti F, Bruno R. Low risk of SARS-CoV-2 transmission by fomites in real life conditions. *Lancet Infect Dis* 2021;21:e112.
- [23] Colaneri M, Seminari E, Novati S, Asperges E, Biscarini S, Piralla A, et al. COVID19 IRCCS San Matteo Pavia Task Force. Severe acute respiratory syndrome coronavirus 2 RNA contamination of inanimate surfaces and virus viability in a health care emergency unit. *Clin Microbiol Infect* 2020;26(8):1094.e1–5. <https://doi.org/10.1016/j.cmi.2020.05.009>. Epub 2020 May 22. PMID: 32450255; PMCID: PMC7243766.
- [24] Chia PY, Coleman KK, Tan YK, Ong SWX, Gum M, Lau SK, et al. Singapore 2019 novel coronavirus outbreak research team. Detection of air and surface contamination by SARS-CoV-2 in hospital rooms of infected patients. *Nat Commun* 2020;11(1):2800. <https://doi.org/10.1038/s41467-020-16670-2>. PMID: 32472043; PMCID: PMC7260225.
- [25] Kotwa JD, Jamal AJ, Mbareche H, Yip L, Aftanas P, Barati S, et al. Surface and air contamination with severe acute respiratory syndrome coronavirus 2 from hospitalized coronavirus disease 2019 patients in Toronto, Canada, March–May 2020. *J Infect Dis* 2022;225(5):768–76. <https://doi.org/10.1093/infdis/jiab578>. PMID: 34850051; PMCID: PMC8767887.
- [26] Gonçalves J, da Silva PG, Reis L, Nascimento MSJ, Koritnik T, Paragi M, et al. Surface contamination with SARS-CoV-2: a systematic review. *Sci Total Environ* 2021;798:149231. <https://doi.org/10.1016/j.scitotenv.2021.149231>. Epub 2021 Jul 24. PMID: 34325141; PMCID: PMC8302502.
- [27] Ahn JY, An S, Sohn Y, Cho Y, Hyun JH, Baek YJ, et al. Environmental contamination in the isolation rooms of COVID-19 patients with severe pneumonia requiring mechanical ventilation or high-flow oxygen therapy. *J Hosp Infect* 2020;106(3):570–6. <https://doi.org/10.1016/j.jhin.2020.08.014>. Epub 2020 Aug 21. PMID: 32828864; PMCID: PMC7441047.
- [28] Santarpia JL, Rivera DN, Herrera VL, Morwitzer MJ, Creager HM, Santarpia GW, et al. Aerosol and surface contamination of SARS-CoV-2 observed in quarantine and isolation care. *Sci Rep* 2020;10(1):12732. <https://doi.org/10.1038/s41598-020-69286-3>. Erratum in: *Sci Rep*. 2020 Aug 12;10(1):13892. PMID: 32728118; PMCID: PMC7391640.
- [29] Harbourt DE, Haddow AD, Piper AE, Bloomfield H, Kearney BJ, Fetterer D, et al. Modeling the stability of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) on skin, currency, and clothing. *PLoS Negl Trop Dis* 2020;14(11):e0008831. <https://doi.org/10.1371/journal.pntd.0008831>