



SARS-CoV-2 peritoneal positivity and emergency surgery: is there any putative predisposing factor?

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Dear Editor,

Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), causing coronavirus disease 2019 (COVID-19), has been declared a global pandemic. Belonging to beta-coronavirus species, infection causes a spectrum of potentially life-threatening symptoms, such as acute distress respiratory disease. Moreover, COVID-2019 patients have often to struggle with also severe gastrointestinal complications, including ischemia [1]. SARS-CoV-2 RNA has been detected in several infected tissues, including peritoneal fluid [2]. However, its clinical relevance as far as pathophysiological mechanisms are still unclear. Although viral tropism for angiotensin-converting enzyme 2 (ACE 2) has been clearly demonstrated, as a target domain for serine protease coupled cell entry mechanism, to this date, no clear evidences about ACE2 peritoneal expression are available. Since the SARS-COV-2 Spike (S) glycoprotein complex processed by the cellular protease TMPRSS2 represents a critical step in the suppression of the modulatory activity of ACE2, the subsequent predominance of the ACE/AngII/AT1R axis and reduction of the protective function of the Ang (1–7)–Mas complex could result into a cytokine–bradykinin inflammatory response, Fas-induced apoptosis and oxidative damage leading to mucosal disruption [3]. Furthermore, the imbalance of the Angiotensin II could cause a self-sustained inflammatory state due to synergic action on the ERK1/2

and p38 MAPK [4] with induction of proinflammatory cytokin gene expression as well as other soluble inflammatory mediators, such as the activation of the innate immune system tool-like-receptors (TLRs).

According to the need for ACE2-TMPRSS2 co-transport, the action of SARS-COV-2 is preferentially on tissues with high gene expressions. Target tissue, as reported by Zou et al. [5], in a scRNA-seq data analysis, are located specific cell lines into lung, heart, oesophagus, kidney, bladder and ileum; while it is not known whether a direct expression of ACE2 exists on peritoneal mesothelial cells. Regarding the gastrointestinal tract, the potential role of the gut in COVID-19 infection has been amply demonstrated [6] with a stool virionic RNA positivity ranging between 20 and 47% of cases. Surprisingly, SARS-COV-2 faecal shedding seems to be prolonged, persisting up to several weeks after respiratory swab negativisation [7]. It is still unclear whether a prolonged positivity at the faecal samples correlates with a prolonged infectivity, although long-standing immunoreaction has been described in paucisymptomatic young patients with uneventful courses. The expression of ACE2 and its transmembrane serine protease are not homogeneous in the gastrointestinal tract. In this setting, a putative theory that would justify peritoneal contamination from SARS-COV-2 could be the effects of a direct cellular damage as far as of an increased capillary permeability. The breakdown of the intestinal barrier could, therefore, be the *primum movens* of extraluminal translocation. However, the coexistence of an impaired host's responses remains a crucial aspect with disruption of enterocytic tight junctions and cryptic M cells. An inflammatory and ischemic microvascular induced dysbiosis could therefore justify the SARS-COV-2 contamination in the peritoneal fluid.

To this date, only case reports and a small series of patients are reported in the literature and are characterised by discrepancy of evidences. Emergency procedures as far as the heterogeneity of patients could justify the dissimilarity of results. Coccolini et al. [2], first, reported a case of

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peritoneal RT-PCR positivity in a surgically treated COVID-19 patient due an uncomplicated ileal volvulus. Adopting a nucleotide assay protocol based on the amplification of three viral targets (SARS-COV-2 RdRP, N, E), Authors identified the presence of a significantly higher peritoneal viral load compared to the upper airways. Similarly, Barberis et al. [8], in a 71-year-old woman who underwent subtotal intra-abdominal colectomy for severe colitis, confirmed the virionic presence in the peritoneal swab. Rimini et al. [9] showed an amplified RT-PCR positivity in one case of exploratory laparotomy and small bowel resection due to an incarcerated umbilical hernia with concomitant loop necrosis. On the other hand, both Negaserig et al. [10] and Vudayagiri et al. [11] did not find any intraperitoneal molecular positivity approaching cases of laparoscopic appendectomies in COVID-19 patients. In support of such evidence, a small case series of five patients, where the Authors [12] did not detect immunostaining in gene amplification assays. According to the scarce statistical representativeness of the available cases, exhaustive conclusions and evidences are far to be drawn. Despite this, some trends worthy of reflection emerge from the analysis of data. In fact, 50% (2/4) of patients with preoperative findings (radiological and endoscopic) of ischemia and/or indirect signs of microvascular injury (ulceration, bleeding) had positive peritoneal swabs. In contrast, only 16.67% (n.1/6) of patients without preoperative signs of ischemia showed RT-PCR SARS-CoV-2 abdominal positivity. The ischemic insult was associated with a threefold increased relative risk of peritoneal fluid positivity albeit in the absence of a relevant statistical correlation (RR 3.00 [95%CI 0.39–23.07; $p = 0.29$]). Although strong evidences are lacking, from a preliminary examination it would seem the peritoneal involvement in COVID-19 is mainly attributable to a secondary visceral outbreak rather than to a primitive serosal injury. The small cohort of patients, absolutely not representative, does not allow us to formulate further hypotheses that can refute or not what emerged. Surely, if the data were confirmed, COVID-19 surgical patients could be clustered into two scenarios: cases with preoperative or clinical findings suspected of visceral ischemia (at risk of peritoneal positivity) and patients with no findings of vascular organ insufficiency. In clinical practice this would entail significant implications as, for patients at risk, it would be necessary to strictly avoid minimally invasive surgical approaches and thus the risk of air-borne infections; while, for the second cohort, indications of a minimally invasive approach could be pursued by preserving the safety of healthcare professionals and the operating environment in general. Multicentric studies are urgently needed which can provide comprehensive answers in the immediate future.

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Declarations

Conflicts of interest The authors have no conflicts of interest to declare.

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