

SARS-CoV-2 Virions or Ubiquitous Cell Structures? Actual Dilemma in COVID-19 Era



To the Editor: Several reports have suggested ultra-structural evidence of direct infection of different types of kidney cells by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in postmortem

analysis and kidney biopsy specimens in patients with proven viral reverse-transcriptase polymerase chain reaction (RT-PCR) from nasopharyngeal smears. Detection of supposed viral particles by transmission electron microscopy (TEM) was used as sufficient evidence for viral invasion of renal tissue but data regarding detection of viral RNA or other valuable methods for viral detection of kidney specimens were missing.^{1,51} Likewise, an additional study proposed direct SARS-CoV-2 infection of endothelial cells in multiple organs of patients with coronavirus disease 2019 (COVID-19) on the basis of TEM solely.² The presented particles in the aforementioned studies

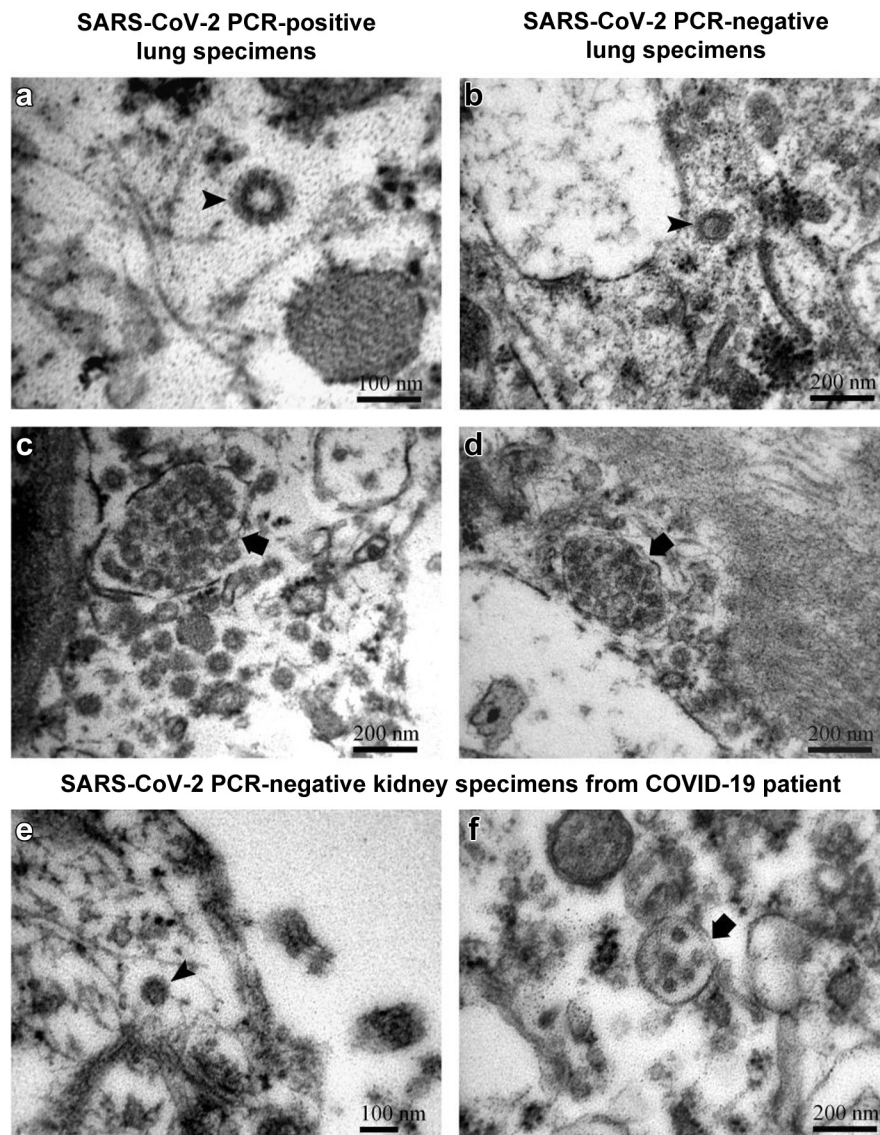


Figure 1. Individual vesicle with electron-dense coat (arrowhead) located freely in the cytosol of endothelial cell in lung with positive reverse-transcriptase polymerase chain reaction (RT-PCR) for severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) RNA (a) and in lung with negative RT-PCR for SARS-CoV-2 RNA (b). Note similar morphology of the 2 structures in images (a) and (b), which could be virus or coated vesicle. (Continued)

Table 1. Semiquantitative analysis of multivesicular bodies and clathrin-coated vesicles in 20 preimplantation donor kidney biopsies before (2018) and during COVID-19 era (2020)

No.	Year of biopsy	Multivesicular bodies			Clathrin-coated vesicles		
		Podocytes	Endothelial cells	Proximal tubular cells	Podocytes	Endothelial cells	Proximal tubular cells
1	2020	+	+	+	++	+	++
2	2020	+	-	-	++	++	+
3	2020	++	+	+	+	+	+
4	2020	+	+	+	+	+	++
5	2020	+	+	-	++	+	++
6	2020	+	-	+	+	+	++
7	2020	+	+	+	+	+	++
8	2020	+	+	-	+	+	++
9	2020	+	+	-	+	+	++
10	2020	+	+	-	+	+	++
11	2018	+	+	-	++	+	++
12	2018	+	+	-	+	+	+
13	2018	+	+	-	+	+	++
14	2018	+	+	-	++	+	+
15	2018	+	+	-	+	+	+
16	2018	+	-	+	+	+	++
17	2018	+	+	-	+	+	++
18	2018	+	+	-	+	+	++
19	2018	+	-	-	+	+	+
20	2018	+	+	+	+	+	+

-, no structures per cell; +, 1 to 5 structures per cell; ++, more than 5 structures per cell.

Specimens contained at least 1 glomerulus and 20 proximal tubules. Five coincidentally selected podocytes, 5 endothelial cells, and 5 proximal tubular cells were examined.

exhibit a diameter of 50 to 150 nm and crown-like electron-dense coat, so they may appear similar to Coronavirus but also similar to ubiquitous coated vesicles, such as clathrin-coated vesicles, or COPI- or COPII-coated vesicles. Moreover, clusters of viral particles inside the vacuole might resemble multivesicular bodies (MVBs), which are regular structures of the endocytic pathway.

Herein, to detect direct invasion of SARS-CoV-2 in the kidney, we performed RT-PCR on fresh post-mortem lung and kidney specimens of 4 patients with COVID-19. In all 4 patients, viral RNA was confirmed in all lung samples, but was negative in all kidney samples. However, ultrastructural examination revealed intracellular vesicular structures of similar size and morphology in lung with proven viral RNA and in kidney with no viral RNA. In lung specimens with proven viral RNA, we observed many structures that could be either viral particles with typical corona or coated vesicles with electron-dense protein coat (Figure 1a). In addition, in the same SARS-CoV-2-positive lung specimens, we found vacuoles that

could be either membrane-bound clusters of viral particles or MVBs with intraluminal vesicles inside (Figure 1c). On the other hand, ultrastructural examination of the lung specimens of 2 patients without SARS-CoV-2 (1 autopsy specimen of lung with negative RT-PCR for viral RNA and 1 biopsy specimen of lung before COVID era) revealed the same structures, resembling viral particles, coated vesicles, or MVBs, as in a specimen with positive SARS-CoV-2 (Figure 1b and d).

TEM analysis of postmortem kidney specimens of patients with COVID-19 revealed numerous individual vesicles and clusters of vesicles in different types of kidney cells, despite negative RT-PCR for SARS-CoV-2 RNA (Figure 1e and f). We additionally performed semiquantitative analysis of these intracellular structures in different types of renal cells from 20 preimplantation donor kidney biopsies, before and during the outbreak of SARS-CoV-2, with negative RT-PCR for SARS-CoV-2 RNA to demonstrate that these cell structures are numerous and ubiquitous in various types of cells (Table 1).

Figure 1. (Continued) In view of the RT-PCR results, the observed structures might be virus in image (a) but not in image (b). Vacuole with many small vesicles inside the limiting membrane (arrow) in the cytosol of endothelial cell in lung with positive RT-PCR for SARS-CoV-2 RNA (c) and in lung with negative RT-PCR for SARS-CoV-2 RNA (d). Note again similar morphology of the 2 structures in images (c) and (d), which could be a cluster of viral particles or multivesicular bodies (MVBs) with intraluminal vesicles inside. In view of the RT-PCR results, the observed structures might be a cluster of viral particles in (c) but not in (d). (e,f) Structures resembling virions, coated vesicles or MVBs were observed in the cytosol of kidney podocytes in a SARS-CoV-2-positive patient but with negative RT-PCR for SARS-CoV-2 RNA. In view of the RT-PCR results, the presented structures are not viruses but ubiquitous coated vesicles and MVBs.

There has been increasing evidence to indicate that coated vesicles and MVBs may mimic viral particles.^{3,S2,S3} However, it is known that the budding of enveloped viruses (to which SARS-CoV-2 belongs) from the plasma membrane, or the limiting membrane of the endosome, resembles the formation of intraluminal vesicles inside MVBs. Moreover, the 2 processes share some components of the same protein machinery.^{4,S4} Indeed, we detected intraluminal vesicles budding from discontinued limiting membrane of the vacuole in lung endothelial cells with positive SARS-CoV-2 RNA (Figure 1a). This finding strongly suggests that this structure is not MVBs but rather a cluster of viral particles with some of budding virions. Indisputable evidence of virions would thus be provided only by immunoelectron microscopy. In addition, coated vesicles might also faintly resemble viral particles, but it is necessary to be cautious about the intracellular location of coated vesicles. Specifically, coated vesicles are transient and are therefore mostly found in close vicinity to the membrane from which they bud, because they shed their coat within seconds after their formation.

To conclude, although TEM may serve as a useful diagnostic method for the detection of viral infection, caution should be exercised when confirmation of viral invasion relies only on TEM. Additional convincing methods, including immunoelectron microscopy, immunohistochemistry, and viral genetic material analysis, are needed for indisputable proof of viral invasion in organs.

SUPPLEMENTARY MATERIAL

Supplementary File (PDF)

Supplementary References.

1. Su H, Yang M, Wan C, et al. Renal histopathological analysis of 26 postmortem findings of patients with COVID-19 in China. *Kidney Int.* 2020;98:219–227.
2. Varga Z, Flammer AJ, Steiger P, et al. Endothelial cell infection and endotheliitis in COVID-19. *Lancet.* 2020;395:1417–1418.
3. Miller SE, Brealey JK. Visualization of putative coronavirus in kidney. *Kidney Int.* 2020;98:231–232.
4. Booth AM, Fang Y, Fallon JK, et al. Exosomes and HIV Gag bud from endosome-like domains of the T cell plasma membrane. *J Cell Biol.* 2006;172:923–935.

Maja Frelj¹, Andreja Erman²,
Karmen Wechtersbach¹, Jerica Pleško¹,
Tatjana Avšič-Županc³ and Nika Kojc¹

¹Institute of Pathology, Faculty of Medicine, University of Ljubljana, Ljubljana, Slovenia; ²Institute of Cell Biology, Faculty of Medicine, University of Ljubljana, Ljubljana, Slovenia; and

³Institute of Microbiology and Immunology, Faculty of Medicine, University of Ljubljana, Ljubljana, Slovenia

Correspondence: Maja Frelj, Institute of Pathology, Faculty of Medicine, University of Ljubljana, 1000 Ljubljana, Slovenia. E-mail: maja.frelj@mf.uni-lj.si

Received 2 July 2020; accepted 7 July 2020; published online 15 July 2020

Kidney Int Rep (2020) 5, 1608–1610; <https://doi.org/10.1016/j.ekir.2020.07.003>

© 2020 International Society of Nephrology. Published by Elsevier Inc. This is an open access article under the CC BY-NC-ND license (<http://creativecommons.org/licenses/by-nc-nd/4.0/>).

Transient Renal Tubular Syndromes Associated With Acute COVID-19 Disease



To the Editor: We report 2 transient renal tubular syndromes associated with coronavirus disease 2019 (COVID-19).

A 47-year-old patient in a neurorehabilitation unit was diagnosed with COVID-19 following onset of respiratory symptoms and pyrexia, confirmed by reverse transcriptase polymerase chain reaction. Ten days later, he developed hypernatremia with an acute kidney injury. He was exclusively fed by percutaneous endoscopic gastrostomy tube.

Investigations (Table 1) supported a diagnosis of nephrogenic diabetes insipidus. He was managed with increased enteral water intake via the percutaneous endoscopic gastrostomy and intravenous 5% dextrose over 24 hours. Biochemistry improved progressively, with serum sodium renal function returning to baseline by day 23.

A 52-year-old woman with diabetic nephropathy and a kidney-pancreas transplant was recovering from a below-knee amputation. She developed fever and a cough; reverse transcriptase polymerase chain reaction confirmed severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection. Three days later, she developed a severe metabolic acidosis associated with profound hypophosphatemia, hyperphosphaturia, and low molecular weight proteinuria, diagnostic of the renal Fanconi syndrome (Table 1). She required aggressive i.v. potassium, bicarbonate, and phosphate supplementation; she was weaned off all supplementation by day 18.

Kidney disease is widely recognized in COVID-19; there is evidence of direct viral invasion of the