



SARS-CoV-2 and male infertility: from short- to long-term impacts

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Abstract

Purpose The coronavirus 2019 (COVID-19) pandemic—caused by a new type of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)—has posed severe impacts on public health worldwide and has resulted in a total of > 6 million deaths. Notably, male patients developed more complications and had mortality rates ~ 77% higher than those of female patients. The extensive expression of the SARS-CoV-2 receptor and related proteins in the male reproductive tract and the association of serum testosterone levels with viral entry and infection have brought attention to COVID-19's effects on male fertility.

Methods The peer-reviewed articles and reviews were obtained by searching for the keywords SARS-CoV-2, COVID-19, endocrine, spermatogenesis, epididymis, prostate, and vaccine in the databases of PubMed, Web of Science and Google Scholar from 2020–2022.

Results This review summarizes the effects of COVID-19 on the male reproductive system and investigates the impact of various types of SARS-CoV-2 vaccines on male reproductive health. We also present the underlying mechanisms by which SARS-CoV-2 affects male reproduction and discuss the potentially harmful effects of asymptomatic infections, as well as the long-term impact of COVID-19 on male reproductive health.

Conclusion COVID-19 disrupted the HPG axis, which had negative impacts on spermatogenesis and the epididymis, albeit further investigations need to be performed. The development of vaccines against various SARS-CoV-2 variations is important to lower infection rates and long-term COVID risks.

Keywords SARS-CoV-2 · Male infertility · Testis · Epididymis

Introduction

The coronavirus disease 2019 (COVID-19) outbreak occurred in December 2019, and rapidly spread worldwide [1]. To date, more than 600 million COVID-19 cases have been reported, and the total death toll exceeds 6.4 million (as of September 5, 2022; <https://coronavirus.jhu.edu/map.html>). COVID-19 is caused by the novel single-stranded RNA virus severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) of the *Coronaviridae* family [2, 3]. More comorbidities and higher mortality rates were observed in male than in female patients [4]. Testosterone has been found to affect the SARS-CoV-2 entry and priming in male hosts, and was correlated with a weaker immune response, higher infection rates, and greater predisposition to thromboembolism [5]. A meta-analysis indicated that diabetes is a key factors associated with high mortality rates in men and women diagnosed with COVID-19 [6]. SARS-CoV-2 encodes four main proteins: a spike (S) protein that mediates virus entry into host cells, N protein that regulates

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nucleocapsid development, and envelope (E) and membrane (M) proteins that mediate viral assembly [7].

The potential impact and molecular mechanisms of SARS-CoV-2 on the reproductive system, particularly in male patients, have been reported in many studies [8–13], though often with conflicting results. In this review, we describe the endocrine status associated with COVID-19 infection, relationship between SARS-CoV-2 and spermatogenesis, effect of the virus on the epididymis and prostate, and potential impact of COVID-19 vaccines on male fertility.

Methods

The current review was conducted using related English literature from electronic databases, including the Web of Science, PubMed, and Google Scholar. The literature search considered publications from 2020 to 2022 with the terms COVID-19, SARS-CoV-2, endocrine, hypophysis, hypothalamus, testosterone, sperm, spermatogenesis, testis, epididymitis, prostate, fever, inflammation, diabetes, and asymptomatic. The studies were further classified by topic. The validity of each publication was assessed by reviewing the title, abstract, and conclusion. The reference lists of valid papers were checked for other relevant studies. All original research or review papers were assessed by two independent investigators and the data of each study were used to summarize their conclusions and perform a meaningful classification.

Receptors for SARS-CoV-2 invasion in the male reproductive system

To infect or enter a cell, the SARS-CoV-2 S protein binds to angiotensin I-converting enzyme 2 (ACE2) expressed on a cell-surface receptor. The host transmembrane protease serine 2 (TMPRSS2) assists in this process by further activating and cleaving the S protein [14]. ACE2 occurs in many tissues and organs, including the lungs, liver, kidneys, and testes [15, 16]. According to the GETx, FAMTOM5, and Human Protein Atlas databases, the testis is among the tissues expressing the highest RNA level of ACE2 [17]. Both ACE2 and TMPRSS2 protein levels were found to be enriched in the testes [18]. Thus, the male reproductive system is highly susceptible to SARS-CoV-2 [19, 20].

Single-cell RNA sequencing (scRNA-seq) suggested that ACE2 was enriched in various cells of the testis, including Sertoli cells, Leydig cells, spermatogonia, and somatic cells [21–23]. Compared to females, symptom severity and mortality rates are higher in males [24, 25]. The transcription of TMPRSS2 could be promoted by an androgen response element [26] and further activate its androgen [27]. Shastri et al.

demonstrated a delay in SARS-CoV-2 clearance in male compared with female subjects, likely associated with the higher level of ACE2 present in the testes, which creates a viral reservoir and promotes viral persistence [17]. Indeed, SARS-CoV-2 antigens have been detected in fibroblasts, spermatogonia, Sertoli cells, and Leydig cells, while the viral particles reside in the cytoplasm of multiple cell types, such as Sertoli cells, Leydig cells, spermatids, endothelium cells, fibroblasts, and epithelial cells in the rete testis [28]. However, Rastrelli et al. concluded that poor prognosis and mortality were associated with lower total testosterone (T), based on the levels of free T [29]. Meta-analyses revealed that COVID-19 negatively impacted T production in the short term, which was related to an increased risk of death or admission to an intensive care unit (ICU) [11]. Although the contradictory effects of androgen on COVID-19 progression need further clarification, at least it proves that the dysregulation of T occurs after SARS-CoV-2 invasion.

Apart from ACE2, the transmembrane glycoprotein CD147 was shown to mediate SARS-CoV-2 pseudovirus entry into host cells via ADP-ribosylation factor 6 (Arf6)-regulated endocytosis [30, 31]. Our previous studies showed that CD147 was expressed at various stages of spermatogenesis, and CD147 null mutants resulted in infertility [32]. Furthermore, CD147 regulated the migration of spermatogonia and spermatocytes via metalloproteinases-2 (MMP-2) signals [33]. CD147 also repressed the extrinsic apoptosis of spermatocytes via nuclear factor κ B (NF- κ B) signaling [34, 35]. Moreover, CD147 played an indispensable role in sperm motility and acrosome reactions, suggesting that CD147 acts as a therapeutic target against asthenozoospermia [36]. Our previous investigations implied that SARS-CoV-2 might affect spermatogenesis via CD147 regulation. However, whether CD147 is involved in the entry of SARS-CoV-2 into the testes remains unclear.

Apart from ACE2, TMPRSS2, and CD147, other recently identified factors involved in SARS-CoV-2 cell invasion may contribute to the impairment of spermatogenesis by acting as receptors (e.g., receptor tyrosine kinase [AXL], kringle containing transmembrane protein 1 [KREMEN1], and asparaginase and isoaspartyl peptidase 1 [ASGL1]), co-receptors (e.g., N-acetylneuraminase pyruvate lyase [NPL], C-type lectin domain family 4 member D [CLEC4D], neuropilin-1 [NRP1], and CD4), and cofactors (ectonucleotide pyrophosphatase/phosphodiesterase 1 [ENPP1], serine protease 1 [PRSS1], TMPRSS6, cathepsin F [CTSF], and paired basic amino acid cleaving enzyme [FURIN]; Fig. 1).

SARS-CoV-2 impairs male fertility by interfering with the hypothalamus–pituitary–gonadal (HPG) axis

Gonadotropins and various steroids serve as a molecular bridge between the brain and testes by regulating HPG axis

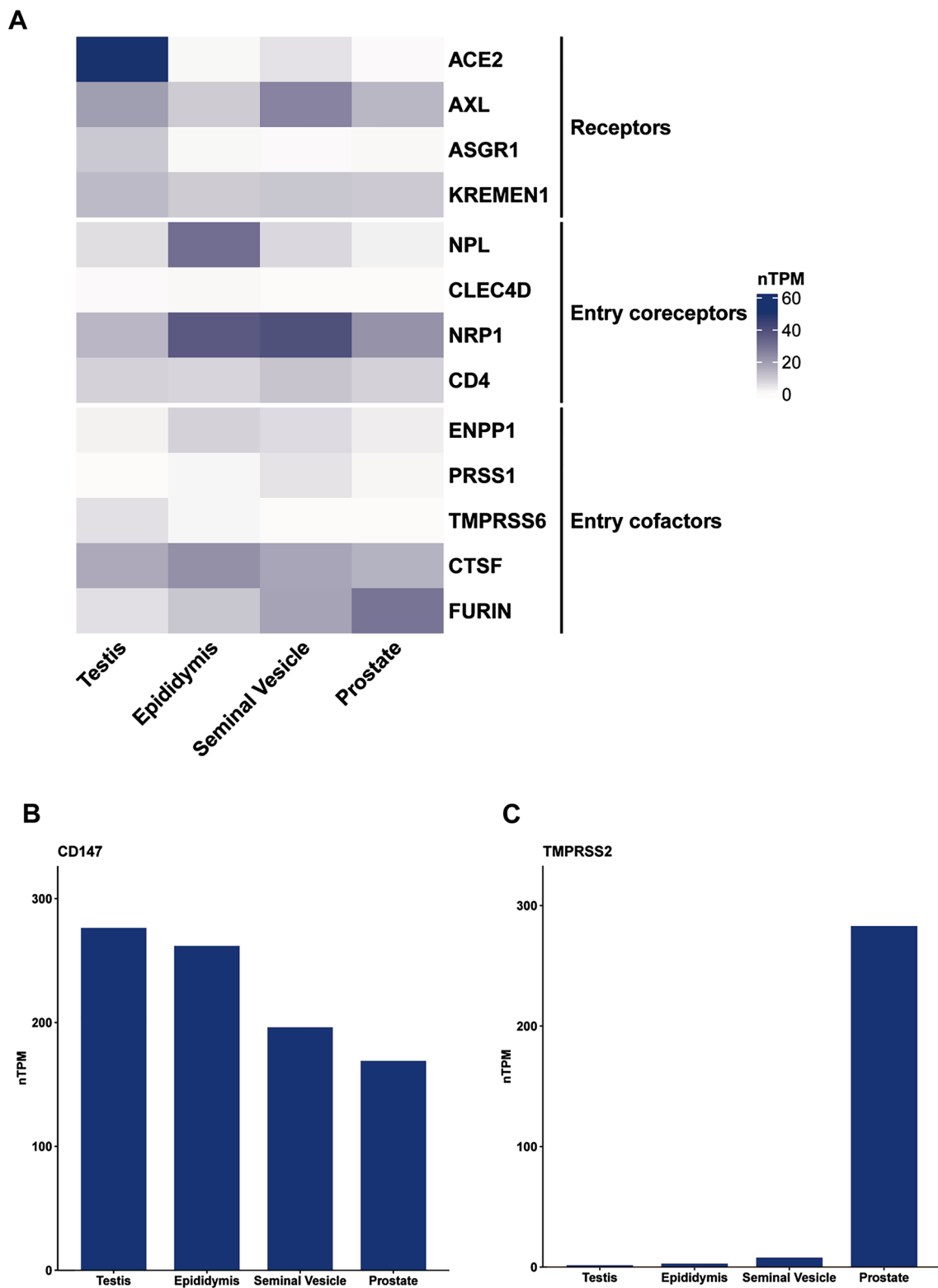


Fig. 1 Expression profile of SARS-CoV-2 invasion-related factors in male reproductive system. **A** RNA expression of receptors, coreceptors, and cofactors related to SARS-CoV-2 cell entry. Gene expression data were acquired from the Human Protein Atlas [162]. ACE2

is abundant in the testis but rarely expressed in the epididymis, seminal vesicle, and prostate. **B**, **C** Differential expression of TMPRSS2 and CD147 in different tissues and organs. The expression values were clarified as normalized transcript per million (nTPM)

activity. The activity of the HPG axis is initiated by the secretion of gonadotropin-releasing hormone (GnRH) from specialized hypothalamic neurons. Gonadotrophs, including follicle-stimulating hormone (FSH) and luteinizing hormone (LH), are secreted by GnRH-dependent adenohypophysis. FSH primarily acts on Sertoli cells to regulate spermatogenesis and keep the seminiferous tubule intact, whereas LH plays a crucial role in stimulating Leydig cells to produce T [37]. Notably, the production of androgens by steroidogenesis is restricted to Leydig cells. Androgens play a vital role in spermatogenesis and regulate gonadotropin levels via a negative feedback mechanism [38].

Male infertility is closely associated with pathological changes in HPG regulation [39, 40]. Previous studies have shown that male patients or animal models infected with SARS-CoV-2 exhibited aberrant hormone levels. Ma et al., in the first report of the effects of SARS-CoV-2 infection on male gonadal function, showed that infected males had considerably increased serum-luteinizing hormone (LH) levels and lower levels of T and FSH. The concentration of C-reactive protein (CRP, a marker of viral infection) was strongly correlated with the T:LH ratio after COVID-19 infection [41]. Coincidentally, serum T, FSH, and LH levels decreased acutely in infected patients, particularly in the case of viral pneumonia [42]. Furthermore, a series of retrospective cohort studies noted that decreased concentrations of T were mostly found in severe COVID-19 cases that required admission to the ICU and prolonged hospitalization [43–45]. In hospitalized men, this low T level was associated with an overactive immune response and inflammatory storm manifested by elevated levels of D-dimer, CRP, interleukin 6 (IL-6), and procalcitonin [46–48]. Lower levels of total T and free T were associated with poor prognosis and increased mortality rates in patients of the respiratory ICU [29]. Interestingly, estradiol levels were substantially increased in critically ill male COVID-19 patients, and their interferon γ (IFN- γ) levels were positively linked to serum estradiol levels, thereby increasing the requirement for extracorporeal membrane oxygenation (ECMO) treatment and risk of mortality [47, 49].

In a cohort of men recovering from COVID-19, a 7-month follow-up study revealed that T levels were further decreased in 10% of the patients, suggesting persistent hypogonadism, especially in cases with a higher burden of comorbid conditions. Moreover, the LH and 17 β -estradiol levels in men with restored T levels decreased significantly [50]. In contrast, Apaydin et al. showed that hypogonadism persisted in 48.2% of men with lower T concentrations over a 6-month follow-up post-recovery [48]. In summary, these studies revealed that the male endocrine system was both directly and indirectly disrupted by COVID-19 infection, and reproductive health was negatively affected overall. The possible adverse effects of SARS-CoV-2 on the HPG axis

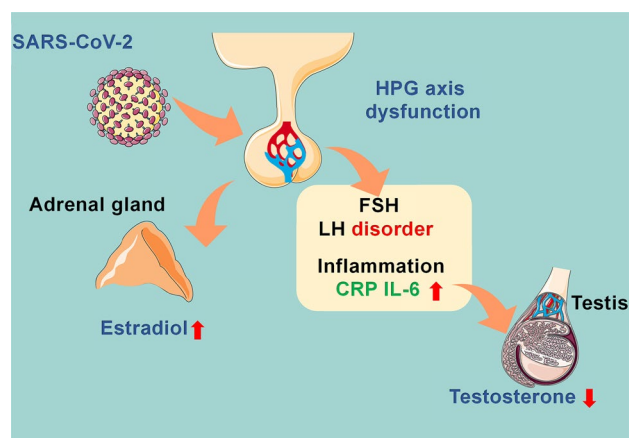


Fig. 2 SARS-CoV-2 invasion caused the dysfunction of the HPG axis. Abnormal FSH and LH serum levels observed during SARS-CoV-2 infection, along with systemic inflammation, which eventually led to a lower concentration of T. Serum estradiol content was elevated by SARS-CoV-2 infection

are summarized in Fig. 2. The characteristics of the cited studies are listed in Table 1.

COVID-19 and spermatogenesis

Spermatogenesis can be divided into three consecutive phases. First, spermatogonia differentiate through mitosis into a B-type, which acts as the precursor of tetraploid primary spermatocytes. Second, diploid secondary spermatocytes are generated from primary spermatocytes via meiotic I division, and haploid spermatids are produced during meiosis II. Third, spermatids undergo spermiogenesis—nuclear elongation and condensation and acrosome biogenesis—to form spermatozoa. During spermatogenesis, different types of germ cells attach to Sertoli cells through specialized cell junctions to enable cell migration from the basement membrane to the abluminal compartment. Additionally, the blood–testis barrier (BTB), constituted by cell–cell junctions among Sertoli cells, limits mature sperm penetration into the circulatory system [51].

Evidence suggests that SARS-CoV-2 infection downregulates the expression of spermatogenesis-related genes [21]. The higher levels of ACE2 expression in the testes of infertile men suggests that COVID-19-mediated reproductive disorders likely depend on ACE2 activation, or that men with reproductive abnormalities may be more susceptible to viral infection [18, 22]. Indeed, SARS-CoV-2 mRNA has been detected in semen [52, 53] and COVID-19 has been associated with decreased numbers of Leydig cells in the testes [54]. Several studies have shown the effects of COVID-19 on sperm quality and spermatogenesis during infection and recovery. For instance, semen volume and total sperm number were lower in COVID-19 patients than those in control

Table 1 The effect of COVID-19 on the HPG axis

Reference number	COVID-19 severity/stage	Tissue assayed	Study design	Age range/ median age	Main conclusion
[41]	Mild, moderate, severe, critical	Serum	Case report	20–54/38	Levels of LH, FSH, and T were influenced
[42]	Severe	Serum	Case report	18–50	T level decreased
[43]		Serum	Prospective cohort study	19–88	T level decreased
[44]	Mild, moderate, severe	Serum	Prospective cohort study	25–91	Lower total T level in serum predicted poor prognosis
[45]	Mild, moderate, severe	Serum	Prospective cohort study	20–65	Increased LH and prolactin, and declined total T level
[46]	Mild, moderate, severe	Serum	Case control report		Lower T level was related to severe clinical outcomes
[47]		Serum	Case report		Decreased T level linked with overactivated immunity
[48]		Serum, semen			Lower T level at baseline with higher inflammatory marker levels
[49]	Severe	Serum	Retrospective cohort study		Lower T and higher estradiol levels may be related to disease severity
[50]	Recovery	Serum	Cohort study	49–65/57	Male hypogonadism persisted 7 months post-recovery

subjects (non-infected men) [55–59]. Sperm viability, motility, and progressive motility were decreased [56–59], and the sperm DNA fragmentation index (DFI) was positively correlated with COVID-19 [57, 59–61]. The deterioration of sperm function and T levels have also been associated with the dysregulation of serum FSH and LH levels [56, 59, 62, 63]. Nonetheless, a number of studies reported unaffected testis or epididymis function, but abnormal sperm parameters. Scropo et al. showed that the levels of T, gonadotropins, and inflammatory factors were unaffected in young males with mild or moderate COVID-19 infections, in spite of abnormal seminal values [64]. Guo et al. showed that the semen of 23 COVID-19-infected male patients had no detectable SARS-CoV-2 RNA, and their sperm counts, and morphology were within normal ranges [65]. Holtmann et al. found that mild COVID-19 infection did not affect testis and epididymis function, but semen parameters were impaired in moderate infection cases [66]. Emerging evidence suggests that the detrimental impact was imposed on spermatogenesis under SARS-CoV-2 invasion, although few studies indicating no influence were reported on sperm quality, the reason for which was mostly due to the males infected with COVID-19 being asymptomatic or with very mild conditions.

Oxidative stress (OS)—induced by an imbalance between reactive oxygen species (ROS) production and clearance by antioxidants—is an important driver of male infertility and is increased under COVID-19 infection [67]. Testicular dysfunctions induced by OS include impaired sperm quality and endocrine function, with oxidative damage in sperm corresponding predominantly to increasing DFI [68].

Total antioxidant capacity was negatively correlated with COVID-19, along with a higher DFI [59]. Similarly, ROS and DFI scores were found to be markedly higher at 14 d after COVID-19 diagnosis compared to those at 120 d [57]. A case report revealed that SARS-CoV-2 invasion into male germ cells may occur prior to the onset of symptoms and disrupt spermatogenesis, as evidenced by high levels of oxidative DNA damage in sperm [61].

In addition to OS, inflammation associated with COVID-19-induced testicular lesions may adversely affect spermatogenesis. Although an unusual presentation, testicular pain can occur with COVID-19 infection, suggesting a link to orchitis [69–71]. Duarte-Neto et al. observed testis congestion, testicular basilemma thickening, and interstitial edema in COVID-19-infected men, which are comparable to the symptoms of interstitial orchitis [28]. Moreover, the number of Sertoli and Leydig cells decreased [28, 72, 73], and testicular blood vessels showed strong expression of vascular cell adhesion molecule (VCAM) in infected men. Fibroblasts, spermatogonia, Sertoli, and Leydig cells all tested positive for the SARS-CoV-2 antigen [28]. COVID-19-infected men also had higher levels of pro-inflammatory cytokines, such as tumor necrosis factor α (TNF- α), IFN- γ , IL-8, IL-10, IL-1 β , and IL-6, in seminal plasma [74, 75]. With the upregulation of these pro-inflammatory cytokines, the expression of junctional proteins involved in the BTB, such as connexin-43, claudin-11, and occludin, is disrupted [76]. The expression of genes involved in apoptosis, such as BAX, caspase-3, caspase-8, and caspase-9, was markedly increased in testicular specimens or sperms of infected men [77]. In deceased individuals, researchers observed a

marked loss of germ cells and increased levels of apoptotic cells, CD3⁺ (mature T lymphocytes), CD68⁺ (macrophage-derived immune cells), CD20⁺ (B cell-derived immune cells), and IgG in the testis/epididymis [18, 74]. Together, these observations suggest that SARS-CoV-2 infection disrupts reproductive microenvironments by triggering OS and inflammation, thereby inhibiting spermatogenesis (Fig. 3).

Males with COVID-19 mostly suffered long-term repercussions and their reproductive functions were compromised even after recovery. Total sperm number at 37 d after recovery was still lower than that of age-matched viral-negative men [78]. Similarly, Guo et al. found that sperm number, concentration, and motility were lower in recovered men after 29 d than those in control subjects, while sperm parameters, including sperm morphology, improved considerably at 56 d after recovery [79]. In contrast, another report found that the concentration and total motility of sperm at 80 d after recovery remained poor [80]. Another study revealed that sperm quality only returned to normal after half a year [81]. A meta-analysis of seven studies corresponding to 934 subjects (median age of 37.34 ± 10.5 years) revealed deteriorating sperm quality and increased LH and prolactin levels during recovery. Therefore, whether deteriorating sperm quality is regulated only by the testes remains unclear [82]. Adamyan et al. investigated transcriptional

alterations in semen from COVID-19-recovered men and found that genes in sperm mitochondria involved in mitochondrial oxidative phosphorylation and toll-like receptor signaling were inhibited. In fact, all protein-coding genes of the mitochondrial genome were dramatically downregulated. This could potentially explain how sperm motility is compromised in convalescent males after viral infection [83]. Moreover, semen proteomics revealed that the proteins linked to male fertility, including prosaposin and semenogelin 1, were markedly downregulated after recovery. The signaling pathways associated with reproductive functions, such as sperm motility, adhesion regulation, endopeptidase activity, oocyte–sperm recognition, and T reaction, were also repressed during recovery [84]. Gacci et al. showed that IL-8 was expressed at pathological levels in 75% of men (33 individuals) during recovery [85]. Negative correlations were found between sperm number and IL-1 β and TNF α levels in semen after recovery [86]. Overall, current findings suggest that it is necessary to conduct accurate follow-ups focusing on the fertility status of patients in convalescence. Additionally, some individuals recovering from COVID-19 showed better recovery of spermatogenetic capacity than others. Gharagozloo et al. found that males with moderate COVID-19 infections along with azoospermia could recover rapidly as the infection waned [61]. Paoli et al. demonstrated

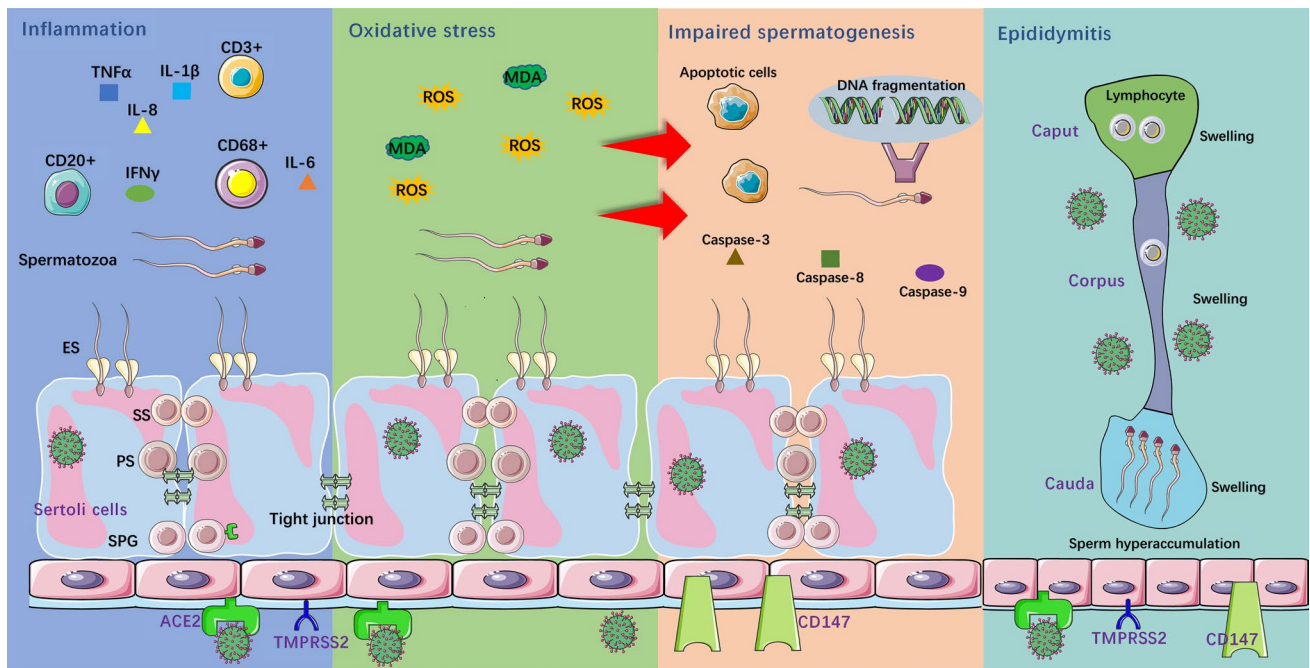


Fig. 3 Possible mechanisms of SARS-CoV-2-mediated impairment of spermatogenesis and sperm function. Spermatogenesis was directly affected by SARS-CoV-2 infection of the testis. CD3⁺, CD68⁺, or CD20⁺-positive inflammatory cells infiltrated the seminiferous tubule upon SARS-CoV-2 infection, which elevated the levels of inflammatory factors, including TNF- α and IL-1 β . Oxidative stress (OS)-

mediated disruptions of male fertility occurred after virus entry, further causing germ cell apoptosis and sperm DNA fragmentation. Epididymitis was observed and mainly characterized by lymphocyte infiltration, sperm hyperaccumulation in the cauda, and a swollen lumen. SPG spermatogonia, PS primary spermatocyte, SS secondary spermatocyte, ES elongated sperm

that testicular function was not damaged directly, and that indirect damage was transient during recovery [10]. Hormone disorders and semen abnormalities gradually disappeared as COVID-19 was cleared [87]. The characteristics of the cited studies are listed in Table 2.

Effect of SARS-CoV-2 on the epididymis

The epididymis is a complex highly coiled duct that connects the vas deferens to the efferent ducts through four anatomically distinct parts, namely the caput, corpus, cauda, and initial segments. A highly specialized microenvironment exists along the lumen that secretes proteins, ions, and cytokines and reabsorbs rete testis fluid. Sperm generated in the testis are matured in the epididymis and acquire their full motility and fertility. The epididymis also acts as an environment to concentrate, store, and transport sperm before ejaculation [88]. Although ACE2 has limited expression in the epididymis [89], other cofactors and receptors, including NPL, NRP1, and CD147, are highly expressed in the epididymis (Fig. 1), suggesting potential for impairment by SARS-CoV-2. Furthermore, SARS-CoV-2 can attach to epididymal sperm via the spike protein [90]. Histological analysis of the epididymis in deceased patients with COVID-19 demonstrated that a large number of sperm and immature spermatocytes accumulated in the cauda [91]. In a pediatric case diagnosed concurrently with COVID-19 and orchiepididymitis, Gagliardi et al. confirmed inhomogeneous testis swelling and epididymal inflammation with reactive hydrocele [92]. Similarly, Marca et al. reported that COVID-19 induced slight swelling and gentle accentuation of vascularization in the epididymis, which were symptoms of epididymitis [69]. Epididymitis caused by COVID-19 can present as reactional hydrocele with nonuniform echo or microcyst dissemination, both of which appear as caput augmentation (> 1.2 cm) and scrotum incassation [93]. In an extensive cohort study (142 patients hospitalized with COVID-19) conducted by Chen et al. 32 patients were diagnosed with acute orchitis, epididymitis, or orchiepididymitis. Acute scrotal infection was found to increase with age, with an incidence rate of 53.3% in men > 80 years of age. Notably, compared with mild infections, severe infections were associated with a higher risk of developing orchiepididymitis [94]. In contrast, Holtmann et al. found that mild infection had no effect on the function of the epididymis and testis [66]. Histopathological assessment of the epididymis after COVID-19 invasion indicated noticeable alterations compared with the control group. Interstitial hyperemia and edema, as well as exudation of red blood cells, were observed in the epididymis. In addition, a few T-lymphocytes appeared in the epididymal duct and infiltrated the testicular blood vessels [8]. Collectively, these results suggest that SARS-CoV-2 infection affects the

epididymal microenvironment and interferes with sperm maturation. Therefore, the risk of epididymitis, especially in young aspiring parents, requires greater clinical attention. The key mechanisms associated with epididymal injury in SARS-CoV-2 are shown in Fig. 3.

Effect of SARS-CoV-2 on the prostate

The prostate, an essential accessory gland of the male reproductive system, consists of the epithelium and stroma. The epithelial compartment secretes prostatic fluid, accounting for approximately one-fifth to one-third of the ejaculate volume. A large number of factors in the prostatic fluid regulate ejaculation and mediate sperm motility, semen liquefaction, and the clotting cycle [95]. Regarding the expression of ACE2 and TMPRSS2 in prostate tissues, single-cell RNA sequencing revealed that epithelial cells expressing ACE2 and TMPRSS2 accounted for 0.32% and 18.65% of the total cells, respectively. Notably, 0.61% of the cells co-expressed ACE2 and TMPRSS2 [96]. Regarding the pathological characteristics of the prostate following SARS-CoV-2 infection, Zhang et al. demonstrated for the first time that infected males had no detectable SARS-CoV-2 RNA in the expressed prostatic secretion (EPS), despite elevated levels of CRP, erythrocyte sedimentation, and IL-6 [97]. SARS-CoV-2 RNA was also absent 80 d after complete clearance of the virus (based on RT-PCR) in mild, moderate, and severe pneumonia [80]. The hospitalized patients did not show signs of prostate inflammation, and the mean serum level of prostate-specific antigen was normal (1.13 ng/mL) [98]. The characteristics of the cited studies are listed in Table 3.

Indirect effects of COVID-19 on male reproduction

Indirect injury to the HPG axis

Salonia et al. reported that 85% of COVID-19 patients showed secondary hypogonadism with lower T levels, which predicted severe clinical outcomes [46]. A follow-up study suggested that a slower recovery rate of T levels was associated with a higher burden of comorbid conditions. Different high-dose corticosteroid formulations for treating COVID-19 have drawn widespread attention [99, 100]. It is well established that the pituitary–gonadal axis can be disturbed by hypercortisolism [101]. Decreased serum T levels, perturbed spermatogenesis, and Sertoli cell dysfunction were associated with hypercortisolism in animal models [102–104]. Excessive treatment with corticosteroids for congenital adrenal hyperplasia in male patients appeared to decrease sperm quality and promote hypogonadism [105]. Compared with long-term corticosteroid treatment, high-dose therapy for critical COVID-19 infections is the safer

Table 2 Effect of COVID-19 on spermatogenesis

References number	COVID-19 severity/stage	Tissue assayed	Study design	Age range/median-age	Main conclusion
[54]		Testis	Case report	42–87/65	Decreased Leydig cell count, seminiferous tubular damage
[55]		Semen	Case report		Decreased semen volume, sperm number, FSH, and LH levels
[56]		Semen, serum	Case report		Sperm viability, motility, and progressive motility decreased
[57]	Moderate	Semen	Case report	20–50	Deterioration of sperm parameters associated with OS
[58]	Mild, moderate	Semen	Multicenter study	20–45	Sperm motility and vitality decreased
[59]		Semen, serum		20–45	LH and DFI levels increased, while T, sperm motility, viability, and total antioxidant capacity decreased
[61]	Moderate	Semen	Case report	55	Sperm count and motility reduced, DNA damage observed. Sperm parameters recovered rapidly in patient recovery
[62]	Mild	Semen, serum	Cross-sectional pilot study	18–60	Abnormal sperm morphology. FSH, LH, and T levels decreased
[63]	Mild, moderate, severe, critical	Semen, serum	Case report	20–49	Poor sperm motility and morphology, low T/LH ratio
[64]	Mild, moderate	Semen, serum	Prospective cohort study	18–50	No effect on spermatogenesis or T and gonadotropin levels
[65]	Mild, moderate, severe, critical, and recovery	Semen	Case report	20–62	Sperm count, motility, and morphology were within normal limits
[66]	Mild, moderate	Semen	Pilot cohort study	42.7 ± 10.4 (mild); 40.8 ± 10.7 (moderate)	Normal testis and epididymis function in mild case. Impaired semen parameters in moderate case
[70]		Questionnaire	Case report	18–75	Testicular pain observed more frequently
[71]			Case report	37	Testicular pain
[73]		Testis (autopsy samples)	Case report		Sertoli and Leydig cell count decreased
[76]		Lung, testis	Case report	45–58	Pro-inflammatory cytokine levels increased; junctional protein levels declined
[77]		Testis	Case report		Apoptotic cell number increased, OS observed

Table 2 (continued)

References number	COVID-19 severity/stage	Tissue assayed	Study design	Age range/median-age	Main conclusion
[78]	Severe, recovery	Semen	Prospective observational study	18–70	Sperm count decreased
[79]	Recovery	Semen, serum	Cohort study	18–45	Sperm count, motility, and concentration decreased at first sampling around 56 d (median)
[80]	Recovery	Semen, urine, expressed prostatic secretions	Cohort study	31	Sperm quality declined slightly, with normal hormone levels after 80 d
[81]	Recovery	Semen	Cohort study	31.75 ± 5.77 (recovered participants); 31.49 ± 3.10 (Control group)	Recovery time of sperm quality was about half a year
[83]	Recovery	Semen	Retrospective observational cohort study	18–65	Levels of energy metabolism-related genes in mitochondria and protein-coding genes in mitochondrial genome downregulated
[84]	Recovery	Semen	Cohort study		Major pathways related to male reproductive functions were inhibited
[85]	Recovery	Saliva, pre-ejaculation urine, semen, and post-ejaculation urine	Prospective cross-sectional study	18–65	Semen impairment, SARS-CoV-2 negative
[86]	Recovery	Semen	Prospective cohort study	18–65	IL-1β and TNF-α were negatively associated with sperm concentration

Table 3 Effect of COVID-19 on epididymis and prostate parameters

References number	COVID-19 severity/stage	Tissue assayed	Study design	Age range/median age	Main conclusion
[90]	Critical	Testis, epididymis	Cohort autopsy study	53–88	SARS-CoV-2 present in sperm, testis, blood–testis barrier, epididymis
[91]		Testis, epididymis, seminiferous duct	Cohort autopsy study	22–83/49.5	Samples negative for SARS-CoV-2. Testis and seminiferous tubule injury, sperm and immature spermatocytes accumulated in epididymis
[92]		Testis	Case report	14	Orchepididymitis-associated with COVID-19
[93]	Mild, moderate	Testis, epididymis	Cross-sectional study	18–55	No orchitis detected, 42.3% of males had epididymitis
[94]		Testis, epididymis	Single-center-based study		Acute epididymitis or orchitis or orchiepididymitis was found in 22.5% patients
[97]		Expressed prostatic secretion (EPS)	Case report	57.5	SARS-CoV-2 not detected in EPS
[98]		Serum	Case report	57.1	No prostate inflammation, serum prostate-specific antigen level was normal

approach, given the potential risks of cortisol drugs on testosterone production and semen quality.

Effects of COVID-19-mediated fever and inflammation on male reproduction

Higher scrotal temperatures caused by fever has an adverse impact on germ cell development as well as sperm DNA integrity, quality, and viability, and can transiently induce sperm apoptosis [106]. Fever was found to occur in 80% of infected patients, and was likely to decrease sperm quality, even in the absence of SARS-CoV-2 viral particles [107, 108]. During recovery, patients presenting with fever had a decreased sperm volume, motility, concentration, and total number compared with those in patients without fever [66]. In addition, the secondary inflammatory response in the male reproductive tract is frequently associated with subfertility [109]. The “cytokine storm” caused in severe and critical SARS-CoV-2 infections presents as the dysregulation of several pro-inflammatory factors, including IL-6, IL-1 β , IL-8, and TNF- α [110], which disrupts the integrity and permeability of the BTB and potentially decreases semen quality and fertilization ability. Studies have shown that, even without the expression of SARS-CoV-2 RNA in semen plasma during infection or recovery, sperm count, concentration, and motility were downregulated and frequently accompanied by alterations in the expression of

oxidation and apoptosis markers [85, 111]. In general, the impact of COVID-19-mediated fever and inflammation on male fertility is unclear, especially as studies rarely regard the contribution of comorbidities and other pathologies. Further studies are needed to clarify the relationship between reproductive injury and the myriad symptoms of COVID-19.

Diabetes: a potential cofactor affecting male reproduction in COVID-19 patients

SARS-CoV-2 causes acute hyperglycemia and insulin resistance in non-diabetic patients and exacerbates diabetes in pre-diabetic individuals [112–114], which is likely mediated by the increased expression of ACE2 in the exocrine glands and islets of the pancreas. SARS-CoV-2 infiltration with concomitant pancreatic injury has been observed in infected patients, which may cause acute β -cell dysfunction [115]. In addition, virus-induced ROS and pro-inflammatory cytokine production promote insulin resistance [116]. Markers of insulin resistance, such as the glucose index and triglyceride content, have been related to COVID-19 severity and mortality rates [117]. Higher blood glucose levels were found to promote COVID-19 progression by facilitating cytokine production and viral replication in diabetic patients [118]. Mounting evidence indicates a relationship between diabetes and testicular injury in murine and human diabetic models [119–121]. Indeed, serum testosterone, FSH, LH, and

antioxidant levels along with sperm number and viability were increased after diabetes resistance treatment [122]. The apelin (APLN) peptide and its receptor are overactivated in the testes of diabetic patients, which promotes the dysfunction of the BTB and spermatogenesis. ML221, an antagonist of the APLN receptor, relieved BTB damage and enhanced spermatogenic ability in cultured human testicular cells [123]. In summary, current evidence shows that diabetes exacerbates the effect of COVID-19 on male reproduction.

SARS-CoV-2 vaccine and male fertility

A vaccine can provide effective protection for susceptible populations against the more serious symptoms of an infectious disease [124]. At the end of 2020, several COVID-19 vaccines were granted Emergency Use Authorization after third-phase clinical trials, which included attenuated amplicons from China and two mRNA vaccines from Moderna and Pfizer [125, 126]. Numerous studies have investigated the safety of these vaccines in the short to medium term. Common side effects included local pain, swelling, redness at the injection site, fatigue, chills, and fever. Serious side effects have also been documented in rare cases, such as Bell's palsy, right leg paresthesia, and paroxysmal ventricular arrhythmia [127, 128]. In addition, the vaccines had a potential deteriorative effect on spermatogenesis in the recovery phase of males with persistent hypothyroidism [78], resulting in elevated ROS content [57] and an inflammation storm in the testis [8, 129], which is similar to the physiological response that could theoretically occur via inoculation of COVID-19 [130]. These side effects can promote vaccine hesitancy (20.9% in males and 79.1% in females of the USA), along with fears that fertility could be affected [131]. Common concerns have also emerged regarding sperm quality among couples receiving assisted reproductive technology (ART) treatment and sperm donors [132, 133]. A small-cohort short follow-up (1–3 months) study of healthy males indicated that there were no noticeable changes in sperm parameters after COVID-19 mRNA vaccines administration, including mRNA-1273 (Moderna) and BNT162b2 (Pfizer-BioNTech) [134–137]. A detailed analysis revealed that motility, DNA fragmentation, and the ultrastructure of sperm remained unchanged, while sperm concentration increased compared to that before three doses of mRNA vaccination [138]. A retrospective cohort study determined that there were no marked differences in sperm parameters after administration of the inactivated vaccine for SARS-CoV-2 after two doses [139]. Furthermore, vaccination with viral vectors or mRNA vaccines did not exacerbate poor fertility rates or sperm motility and concentration in patients with male-factor infertility [136, 140]. A meta-analysis showed that sperm quality was not affected by any of the mRNA COVID-19 vaccines [11]. A larger analysis involving

subjects from seven countries concluded that there was no association between subfertility and COVID-19 vaccines, including BNT162b2 and mRNA-1273, in women or men [141]. mRNA vaccines also showed no relationship with the risk of developing orchitis and/or epididymitis [142]. Overall, COVID-19 vaccination has not affected sperm quality or testis and epididymis function. However, the long-term effect (> 10 years) of vaccination on male fertility requires a thorough follow-up investigation.

Conclusions and perspectives

It is clear that COVID-19 can cause dysfunction of the HPG axis [43–45]. Deviations in T levels reflected the pathological status of steroidogenesis in the testis, which was related to the dysregulated levels of LH and FSH in COVID-19-affected subjects. Moreover, the lowered T levels could cause erectile dysfunction and altered spermatogenesis, which promote subfertility [143]. Higher concentrations of LH and FSH reflect testicular damage and other pathological outcomes [144]. Dysregulation of the HPG axis could result in not only hypothyroidism but also neurodegenerative senescence, liver cirrhosis, and chronic kidney disease [145]. The multiple possible mechanisms by which SARS-CoV-2 affects spermatogenesis are as follows: direct damage to testicular tissue or sperm; exaggeration of immune response, OS, and apoptosis mediated by SARS-CoV-2 infection; dysregulation of hormone levels. However, SARS-CoV-2 may have longer-lasting effects on the HPG axis and spermatogenesis. First, several studies ignored the importance of simultaneously detecting hormone levels and semen quality and did not consider the existence of SARS-CoV-2 in testicular tissue. Second, it is plausible to conclude that male reproduction is disturbed during the recovery phase [83]. Therefore, greater attention should be paid to the molecular mechanisms of male reproductive disorders and the long-term impact of COVID-19 to distinguish transient and irreversible injuries to the HPG axis and spermatogenesis. Furthermore, medical interventions, such as newly developed oral medications, should be considered during recovery to improve the transient subfertility state and avoid further deterioration of the reproductive system. Third, unlike for moderate and mild disease states, data are lacking for reproductive outcomes in critical and severe cases. In addition, the relationship between disease severity, hormone levels, and semen parameters has not yet been reported. Finally, because of the high concentration of prolactin, which inhibits the signaling of the HPG axis [146], prognosis and clinical management of COVID-19 could be improved by measuring the baseline level of prolactin in male patients.

Although epididymal damage has been verified according to the clinical pathologic status, there is no evidence that the virus is present in the epididymis, and the pathologic changes observed in biopsy and autopsy are also limited. The analysis of sperm parameters and hormone levels is recommended in the evaluation of males with COVID-19 diagnosed with epididymo-orchitis. The pathophysiological changes observed in the epididymis are likely due to an exaggerated immune response or cytokine storm in cases of prolonged illness, though this requires further consideration. Furthermore, research indicated that the prostate may not be targeted by SARS-CoV-2; however, the relationship between EPS and semen parameters remains unclear. Regarding the potential influence of SARS-CoV-2 vaccines, follow-ups should be continued to accurately determine the effective recovery of spermatogenesis and reproductive capacity. Although sperm viability has not been influenced by any of the COVID-19 vaccines [134–136, 138, 139], hormone levels and clinical characteristics should not be neglected in follow-up studies, especially given the known side effects. An increasing number of vaccines have been approved for clinical use, and even if the features and side effects are similar among distinct vaccines, it is unlikely that results can be generalized to other vaccines for emergency use.

To date, five primary variants of SARS-CoV-2 have been reported, including alpha (B.1.1.7), beta (B.1.351), gamma (P.1), delta (B.1.617.2), and omicron (B.1.1.529). Although these variants are associated with lower morbidity rates, they have shown higher transmission rates along with increased resistance to previous antiviral, antibody, and immune plasma treatments, which promotes the reinfection of recovered or vaccinated individuals, especially in the case of the omicron variant [147, 148]. Owing to its highly contagious nature, the omicron variant has aroused concerns of a new worldwide wave of COVID-19. In addition, findings also indicated a higher rate of asymptomatic cases associated with omicron infections, based on clinical analysis and lung computed tomography [149, 150]. A delicate balance between inflammation and antiviral action occurs in asymptomatic patients, depending on the pathogen's capacity to clear virus-specific T cells [151]. Moreover, higher functional cellular immune responses are triggered in asymptomatic than in symptomatic individuals [151]. Higher levels of IL-2 and IFN- γ secretion were found in asymptomatic patients (including omicron-infected individuals), which were related to the increased production of IL-10, IL-6, IL-1 β , and TNF- α triggered by virus-specific T cells [149, 152]. Therefore, asymptomatic individuals have a more robust cellular immune response than those with weak antiviral immunity.

In general, the innate immune response in the testicular microenvironment is counteracted by the immunosuppressive system [153]. However, severe systemic inflammation

can negatively affect the male reproductive system through blood-borne transmission or secondary inflammation [109, 154]. Thus, the potential damage to the male reproductive system caused by SARS-CoV-2 cannot be neglected in asymptomatic individuals. While SARS-CoV-2 infection in young males is mostly mildly symptomatic or asymptomatic, its prevalence has been underestimated because children are often excluded from screening tests [155]. In fact, serological analysis revealed that the rate of infection among children under 18 years was 68% as compared with the reported 33% from December 2021 to February 2022 [156]. Previous studies have shown that the integrity of the BTB can be compromised by SARS-CoV-2 infestation [76], which is linked to the production of nitric oxide, IL-6, and TNF- α and increased macrophage infiltration [76]. Since the BTB is only functional at the time of puberty [157], children could be at a higher risk of long-term COVID-19 infections with direct or indirect damage caused by SARS-CoV-2-related multisystem inflammation [155]. Therefore, the potential impact of SARS-CoV-2 infection on the development of the reproductive system of preadolescent or pubertal males and their fecundity should not be overlooked.

“COVID-19 illness” is used to describe prolonged symptomatic infections or the persistence of symptoms after recovery [158]. Studies have indicated that “long COVID-19” occurs in approximately 30% of infected individuals [159]. Davis et al. characterized COVID-19 through an online international survey conducted for 7 months. The common symptoms after 6 months were cognitive impairment, fatigue, and post-exertional malaise. The mean prevalence of symptoms related to endocrine, reproductive, and genitourinary was 62.25% [160]. According to the largest cohort study, vaccination only 15% decreased the long-term risk of COVID-19 [161]. The high prevalence of long COVID symptoms associated with the male reproductive system and the limited protection offered by vaccination emphasize the need for further research on the impact of the various SARS-CoV-2 variants on male reproduction. Furthermore, more effective vaccines against various variants of SARS-CoV-2 need to be developed to reduce infection rates and long-COVID risks. Future studies should focus on the early diagnosis of reproductive abnormalities caused by COVID-19 to promote early treatment.

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Data availability All data generated or analysed during this study are included in this published article.

Declarations

Conflict of interest The authors declare no conflicts of interest.

Ethical approval The present article does not contain any studies with human participants or animals performed by any of the authors.

Informed consent The formal consent is not required by this type of study.

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