

# Evaluating the impact of COVID-19 on male reproduction

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## Abstract

Invasion or damage of the male reproductive system is one of the reported outcomes of viral infection. Current studies have documented that SARS-CoV-2, which causes COVID-19, can damage the male reproductive system in large part by inflammatory damage caused by a cytokine storm. However, whether SARS-CoV-2 can infect the human testis directly and enter semen is controversial. Other adverse effects of SARS-CoV-2 on male reproduction are also of concern and require comprehensive evaluation. Here, we analyze the invasiveness of SARS-CoV-2 in the testis and examine reported mechanisms by which SARS-CoV-2 interferes with male reproduction. Long-term implications of SARS-CoV-2 infection on male reproduction are also discussed. It should be emphasized that although COVID-19 may induce testicular damage, a substantial decrease in male reproductive capacity awaits clinical evidence. We propose that there is an urgent need to track male COVID-19 patients during their recovery. The development of suitable experimental models, including human reproductive organoids, will be valuable to further investigate the viral impact on reproduction for current and future pandemics.

*Reproduction* (2021) **161** R37–R44

## Introduction

In the past months of combating COVID-19 (coronavirus disease 19), over 50 million people have been infected by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), and the impact of the disease may continue for several years. Moreover, the actual number of infected individuals worldwide has been estimated to be 12 times higher than official numbers as of July 2020 (Rahmandad *et al.* 2020). Although primarily a respiratory disease, this huge base amplifies the adverse effects of less severe aspects of COVID-19 and invites further evaluation. SARS-CoV-2 shares ~80% amino acid sequence with severe acute respiratory syndrome coronavirus 1 (SARS-CoV-1) (Gralinski & Menachery 2020) that caused more than 8000 infections in 2003. Both viruses use transmembrane serine protease 2 (TMPRSS2) and the receptor angiotensin-converting enzyme 2 (ACE2) to infect host cells. Besides lung, ACE2 is expressed in many tissues including the cardiovascular system, gastrointestinal tract and liver. Correspondingly, damage to those organs was observed in the COVID-19 patients (Xiao *et al.* 2020, Zhang *et al.* 2020b, Zheng *et al.* 2020). Notably, testis also was reported to express ACE2 in different cell types (Wang & Xu 2020) and it is worth noting that being male is a contributing factor in COVID-19 deaths (Williamson *et al.* 2020). In related studies, male patients showed higher plasma levels of innate immune cytokines than female patients (Takahashi *et al.* 2020) and a high proportion of male patients were of reproductive

age (Guan *et al.* 2020). Therefore, whether SARS-CoV-2 infection impairs the male reproductive system becomes an issue that deserves further investigation.

Current studies confirm that SARS-CoV-2 can affect the male reproductive system. Abnormal levels of sex hormones and declining sperm quality were observed in patients during and after recovery from COVID-19. Moreover, severe inflammation damage was detected in testes and the presence of virus in testicular tissue was reported at autopsy. Detrimental effects on male fertility have been reported during infections with Zika (ZIKV), mumps (MuV), and SARS-CoV-1 viruses, but the picture is less complete for COVID-19. A better understanding of the mechanisms of infection and pathophysiology will require long-term surveillance and investigation. However, previous studies of the aforementioned viruses, especially SARS-CoV-1, can supply meaningful guidance for analyzing the effect of COVID-19 on male reproduction. Here, based on the available evidence, we discuss tropism of SARS-CoV-2 for testicular cells, and analyze known and potential mechanisms by which SARS-CoV-2 affects male reproductive health with the aim of providing a reference for further investigations and treatment of COVID-19.

## The molecular basis of SARS-CoV-2 infecting the male reproductive system

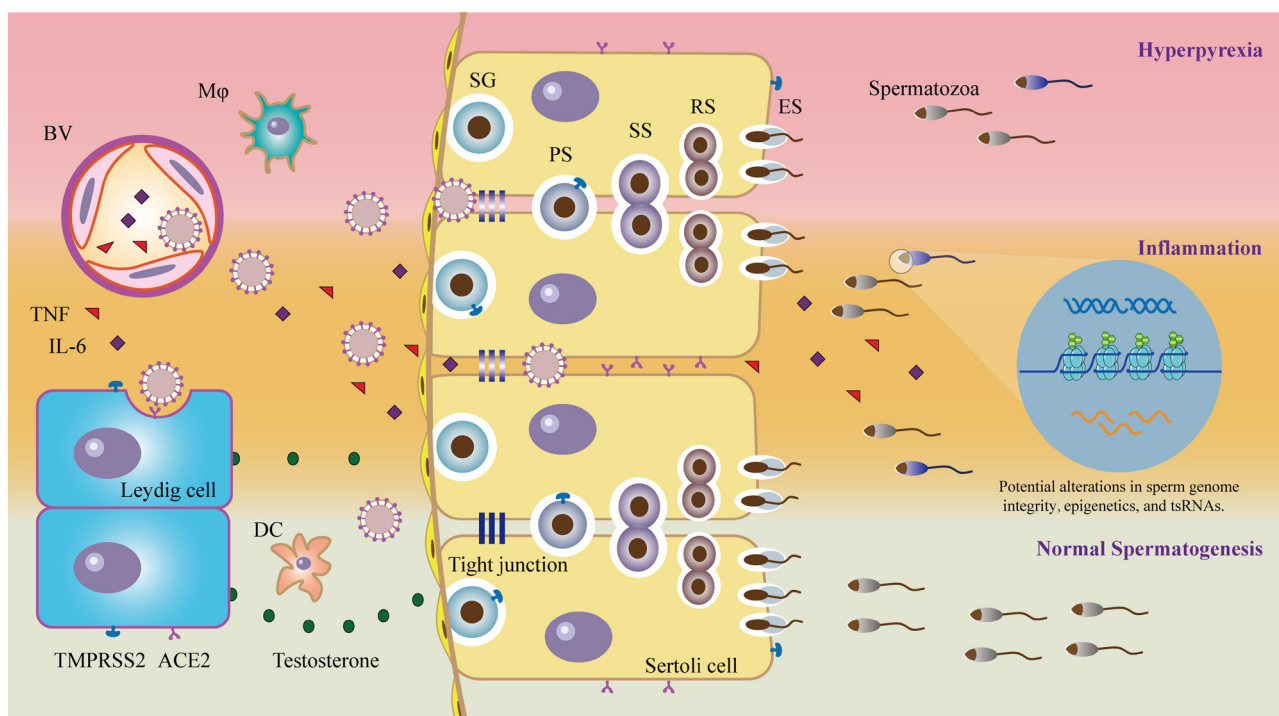
The testis consists of seminiferous tubules and interstitial cells that are separated by a basal lamina and are

important for spermatogenesis and hormone regulation, respectively. Within seminiferous tubules, germ cells mature centripetally between adjacent Sertoli cells. Meiotic and maturing spermatocytes are separated from more peripheral spermatogonia and pre-leptotene spermatocytes by cell–cell junctions between Sertoli cells that constitute a blood–testis barrier (BTB). This provides an immune-privileged microenvironment for maturing sperm by limiting the exchange of substances between interstitium and germ cells. Fully grown spermatids are released into the lumen of seminiferous tubules which drain into the rete testis and spermatozoa pass into the epididymis for maturation and storage (Suede *et al.* 2020). The interstitium of the testis is mainly composed of Leydig cells (LC), as well as smaller populations of immune, myoid and fibroblast cells. Testosterone secreted by LCs plays a crucial role in spermatogenesis and in maintaining secondary sex characteristics (Heinrich & DeFalco 2020). Overall, there is a highly integrated synergism among all testicular cells in ensuring male fertility and adverse effects on the testis risk male reproductive health (Fig. 1).

Previous bioinformatic analyses have documented the cellular distribution of the ACE2 receptor in testes

(Liu *et al.* 2020, Shen *et al.* 2020, Wang & Xu 2020). SARS-CoV-2 and SARS-CoV-1 share the common spike (S) protein. Endocytosis of SARS-CoV-2 virus into target cells depends on the recognition of the ACE2 receptor by the viral S protein (Hoffmann *et al.* 2020) and SARS-CoV-2 has a higher affinity for the receptor than SARS-CoV-1 (Wrapp *et al.* 2020). Within the testis, ACE2 is expressed mainly in SC, LC, and spermatogonia cells (Wang & Xu 2020). Expression of ACE2 in testes varies with age and men in their 30s had the highest expression of ACE2 (Liu *et al.* 2020, Shen *et al.* 2020). Thus, the testis is a potential target of SARS-CoV-2 and could adversely affect male fertility. Moreover, once the virus is present in semen, sperm that express ACE2 may also be infected by SARS-CoV-2. Theoretically, sperm could become vectors in the spread of COVID-19 which recently has been reviewed (Aitken 2020).

However, a viral infection of cells requires cofactors, particularly the TMPRSS2 protein which plays a vital role in enhancing SARS-CoV-2 entry by priming the S protein (Li *et al.* 2003, Matsuyama *et al.* 2010). Although spermatogonia and spermatozoon express TMPRSS2, high co-expression of the two proteins is not observed



**Figure 1** Potential routes of SARS-CoV-2 affecting male reproductive system. SARS-CoV-2 and high levels of cytokines may enter testicular tissue via hematogenous dissemination. LCs could be attacked by cytokines (or SARS-CoV-2 virus) which would influence testosterone secretion and spermatogenesis. Additionally, the BTB which normally prevents viral invasion is susceptible to the cytokine-mediated inflammation. The severe cytokine storm associated with SARS-CoV-2 infection could cause orchitis and epididymitis. Additional factors, including hyperpyrexia, inflammatory destruction of the BTB may increase the probability of virus crossing the BTB. Considering low co-expression of ACE2 and TMPRSS2 in SCs and germ cells, further investigations will be needed to determine the possibility of sexual transmission. Sperm DNA fragmentation, alterations in epigenome and sperm tsRNAs may be mediated indirectly by fever, inflammation, and psychological factors. Studies to investigate possible adverse effect on future pregnancies and progeny are also warranted. BV, blood vessel; Mφ, macrophage; DC, dendritic cell; SG, spermatogonia; PS, primary spermatocyte; SS, secondary spermatocyte; RS, round spermatid; ES, elongating spermatid.

in testicular cells (Pan *et al.* 2020, Stanley *et al.* 2020). For example, spermatogonia possess high levels of TMPRSS2, but low levels of ACE2 and SCs express high levels of ACE2, but low levels of TMPRSS2 (Liu *et al.* 2020, Pan *et al.* 2020, Wang & Xu 2020). Other cofactors, including endosomal cysteine proteases cathepsin L and cathepsin B facilitate SARS-CoV-1 entry, but are primarily expressed in spermatocytes and testicular macrophages, respectively (Liu *et al.* 2020).

Other potential cell receptors of the S protein include CD147, CD26, and Neuropilin-1 were reported recently, but need further validation. CD147 is mainly expressed in spermatocytes, but other testicular cells including SCs have a moderate expression (Liu *et al.* 2020, Wang *et al.* 2020). CD26 acts as a receptor of the middle east respiratory syndrome (MERS) coronavirus and its interaction with S protein has been documented (Vankadari & Wilce 2020). Meanwhile, the infection efficiency of SARS-CoV-2 mediated by Neuropilin-1 and TMPRSS2 reached half that mediated by ACE2 and TMPRSS2 (Cantuti-Castelvetri *et al.* 2020). Distribution of CD26 and neuropilin-1 in testis requires documentation, the role of these potential receptors warrant additional study. Overall, the expression of ACE2 and other potential receptors in the testes suggests a possible tropism of SARS-CoV-2 to the testis. However, based on current evidence it is not possible to conclude that SARS-CoV-2 infects testicular cells directly by interacting with ACE2 or other receptors. Moreover, given the low level of co-expression of ACE2 and other cofactors in testes, the probability of SARS-CoV-2 infecting cells of the testis via ACE2 remains low.

### Possible mechanisms of SARS-CoV-2 affecting male reproduction

The main mechanisms of viruses impacting male reproduction can be summarized as follows: (a) direct viral invasion of germ cells and spreading the viruses via sexual transmission; (b) virus affecting reproductive endocrinology necessary to maintain secondary sexual characteristics; (c) a secondary viral infection-induced inflammation response that severely affects the testes; and (d) viral infection-triggered fevers that interfere with the normal reproductive physiology. Of note, the above mechanisms often coexist and have a synergistic effect on mediating the impairment. Besides, the male reproductive system may also be impacted by drugs used to treat a viral infection, with the major reason to be gonadotoxic effect of drugs, such as glucocorticoids and interferons (Drobnis & Nangia 2017, Li *et al.* 2020c). Intriguingly, the binding of SARS-CoV-2 to ACE2 could cause up-regulation of angiotensin II, which may induce adverse effects on male fertility (Welter *et al.* 2014, Aitken 2020). Based on the current studies on SARS-CoV-2 and evidence that other viruses disrupted male reproduction, we analyzed known and potential mechanisms that SARS-CoV-2 affected male reproduction.

### Can SARS-CoV-2 directly infect the testis?

Evidence for SARS-CoV-2 infection of the testis is not compelling. In 2003-2004, several groups reported histopathological examination of testes from SARS-CoV-1 deaths. More reported negative (Zhang *et al.* 2003, Zhao *et al.* 2003b, Ding *et al.* 2004) than positive (Zhao *et al.* 2003a,c) results in evaluating the presence of the virus. Similarly, SARS-CoV-2 virus was not present in testis samples from a COVID-19 patient who died in the acute phase (Song *et al.* 2020). However, another study reported that the virus could be detected in multiple organs including testis, with varying degrees of germ cell reduction and damage (Bian *et al.* 2020). Recently, Yang *et al.* tested testis samples from 12 deceased COVID-19 patients. The virus was detected in only one patient who had a high viral load. The SARS-CoV-2 virus was not present in seminiferous tubules, but only in the interstitium, and it was unclear whether the virus came from blood rather than infection of the testis (Yang *et al.* 2020b).

However, whether the virus can be present in the testis does not entirely depend on the viral infection. There are other conditions where the virus might enter and disrupt testicular tissue, including high blood viral load, local inflammation, hyperpyrexia and imperfect BTB. Thus, the possibility of SARS-CoV-2 directly disrupting testis cannot be ruled out, but in the mild COVID-19 patients without risk factors, it seems unlikely.

### The possibility of sexual transmission of SARS-CoV-2

Current evidence does not support sexual transmission of the SARS-CoV-2 virus. Six small cohort studies (Guo *et al.* 2020, Holtmann *et al.* 2020, Li *et al.* 2020b, Ma *et al.* 2020, Pan *et al.* 2020, Song *et al.* 2020) from three counties and one case report (Paoli *et al.* 2020a) from Italy did not detect viral nucleic acids in semen samples from mildly affected and recover-stage male patients. Two independent studies also reported no SARS-CoV-2 virus in prostate fluid (Quan *et al.* 2020, Zhang *et al.* 2020a). However, one study showed that SARS-CoV-2 could be shed into semen. In this study, among the 23 clinical recovered and 15 acute phase patients, 2 (8.7%) and 4 (26.7%) patients, respectively, tested positive for virus components in their semen (Li *et al.* 2020a) which stands in sharp contrast to previous studies. But the detailed methodology of false-positives caused by sampling and detection errors was not provided (Paoli *et al.* 2020b). All these studies suffer from small sample sizes and the demographics of patients in convalescence after mild infection may have introduced selection bias. In addition, the abundance of viral RNA in semen was low compared to other tissues and procedures for the collection of ejaculates is vulnerable to contamination. Larger and multicenter trials are needed to conclude compelling results as to the presence or absence of



SARS-CoV-2 in semen, especially for patients early in the disease course or asymptomatic. However, according to available studies, the probability of virus entry into semen in mild or recover-stage patients is very low. Collectively, the absence of SARS-CoV-2 in the semen moots the possibility of sexual transmission of COVID-19.

### **Orchitis mediated by SARS-CoV-2-induced severe inflammation**

The BTB confers an isolated immune-privileged microenvironment for sperm in the testis. However, serious inflammation induced by infection can disrupt the BTB resulting in orchitis which adversely affects testicular function and male fertility. Notably, SARS-CoV-1 can cause severe orchitis with broadly damaged germ cells, few mature spermatozoa, thickened basement membranes and leukocyte infiltration. The absence of SARS-CoV-1 in the testis indicates that inflammation rather than the direct viral infection caused the histopathology and that substantial orchitis can occur in SARS patients (Xu *et al.* 2006).

Similar histopathological changes have been reported in COVID-19 patients. Testicular samples from 2 of 10 autopsies had orchitis of unknown origin, although few details were provided (Nunes Duarte-Neto *et al.* 2020). In two separate reports, histopathological changes consistent with orchitis were observed in 6 and 11 of 12 deceased COVID-19 patients (Li *et al.* 2020b, Yang *et al.* 2020b). These included inflammatory damages of seminiferous tubules with interstitial edema, congestion, inflammatory cell infiltrate and red blood cell exudation. Thinning of seminiferous tubules and desquamation of intratubular cells were also observed. There were decreased numbers of LCs in the interstitium. SCs were characterized by swelling, vacuolation and cytoplasmic rarefaction accompanied by marked shedding of cells from the basement membrane. In addition, eight patients had varying degrees of impaired spermatogenesis with hypospermatogenesis, maturation arrest and mild peritubular hyalinization (Yang *et al.* 2020b). Li *et al.* also detected the semen samples from COVID-19 inpatients, which showed that the sperm concentration declined with elevated levels of interleukin-6 (IL-6), tumor necrosis factor- $\alpha$  (TNF- $\alpha$ ), and monocyte chemoattractant protein-1 (MCP-1) in the semen compared with controls (Li *et al.* 2020b). Most importantly, in both studies, no SARS-CoV-2 virus was detected in those testis samples in which orchitis was confirmed, suggesting that inflammatory response rather than viral infection dominated these pathological changes.

Therefore, we speculate that a cytokine storm induced by SARS-CoV-2 plays a critical role in the observed orchitis. Patients with severe COVID-19 infection have high levels of plasma cytokines including IL-2, IL-6, IL-7, IL-10, TNF- $\alpha$ , and MCP-1 (Chen *et al.* 2020, Huang

*et al.* 2020). Male patients have a weaker T cell response and higher plasma level of innate immune cytokines than female patients, which is associated with the poor disease outcome (Takahashi *et al.* 2020). Unlike viral infections, cytokines such as IL-6 and TNF- $\alpha$  aggressively disrupt the integrity of the BTB (Li *et al.* 2006, Zhang *et al.* 2014). When hypercytokinaemia happens, it is likely that cytokines pass through the BTB and induce inflammation in seminiferous tubules. This is consistent with previous studies that implicated IL-6 disruption of the BTB in the pathogenesis of autoimmune orchitis (Rival *et al.* 2006, Perez *et al.* 2012). In adult infection with MuV (virus causing mumps), TNF- $\alpha$  can disrupt the BTB to cause orchitis and male infertility (Wu *et al.* 2019). Orchitis not only disrupts the immune balance of the testis microenvironment, but also impairs the seminiferous epithelium and spermatogonial stem cells. These detrimental effects are substantial and make complete recovery unlikely. Additionally, inflammation increases the probability of the virus entering the testis and it is not surprising that SARS-CoV-2 is detected in testes of patients with severe disease (Bian *et al.* 2020).

### **SARS-CoV-2 and dysfunction of hypothalamus–pituitary–gonad axis**

Androgens secreted from the testis are essential to maintain secondary male sex characteristics. Unlike spermatogenesis that occurs solely within the testis, the synthesis of male sex hormones involves a hypothalamus, pituitary, gonad (HPG) axis. Previous reports reported that male patients infected with SARS-CoV-1 had elevated luteinizing hormone (LH), follicle-stimulating hormone (FSH), prolactin (PRL) and decreased estradiol (E), progesterone (P), and testosterone (T) plasma levels (Wang *et al.* 2005, Wei *et al.* 2010). COVID-19 patients recapitulate these abnormalities. In a cohort study of 45 male COVID-19 patients, 68.6% and 48.6% of patients had low T and dihydrotestosterone levels, respectively (Schroeder *et al.* 2020). In comparison to patients in remission, 31 males in critical care (or deceased) had higher LH, lower total T and calculated free T which suggests hypogonadism in severely ill patients (Rastrelli *et al.* 2020). These and an additional study (Salciccia *et al.* 2020) indicate an inverse correlation between T levels and inflammatory factors. A retrospective single-center study involving 119 patients reported that male patients had no differences in serum levels of T compared with controls but had elevated LH levels and significantly decreased ratios of T/LH. There also was a negative correlation between T/LH ratios and inflammatory markers (Ma *et al.* 2020). Taken together, these findings suggest an impact of SARS-CoV-2 infection on the HPG axis and highlight the role of inflammation.

Impaired Leydig cells reduce androgen hormone secretion which feedbacks on the pituitary and could

account for decreased T and increased LH. However, previous studies indicated SARS-CoV-1 RNA was detected in the acidophilic cells of the pituitary and function of adenohypophyseal endocrine cells in infected patients was adversely affected (Zhang *et al.* 2003, Wei *et al.* 2010). Possible effects of COVID-19 on hypothalamus and pituitary have also been proposed (Ur & Verma 2020), although inadequately documented. Most importantly, the dysfunction of hypothalamus and pituitary not only affect the sex hormones but other hormones and adrenaline that help maintain normal sexual functions (Bhasin *et al.* 2007).

Therefore, abnormal sex hormone levels may be attributable to multiple targets in the HPG axis. Intriguingly, whether this result is caused by a virus affecting the pituitary or the testis, inflammation seems to play a vital role in it. Of course, other factors including corticosteroid treatment and mental stresses may be considerable contributors. Additionally, the long-term implications of altered sex hormones on male fertility needs to be recognized. There is a precedence in that the ZIKV induced testicular atrophy by decreasing testosterone (Uraki *et al.* 2017). More attention needs to be paid to the risk of SARS-CoV-2 on the HPG axis, and larger studies will be required to evaluate hypogonadism and sexual dysfunction caused by abnormal sex hormone levels in men with COVID-19.

### Febrility caused by SARS-CoV-2 may disrupt spermatogenesis

Decreased sperm number or increased sperm DNA fragmentation has been reported in males recovering from COVID-19 (Ma *et al.* 2020). Four patients with moderate infection and fever had impaired sperm quality (Holtmann *et al.* 2020), but whether this was

due to inflammation or fever remained unresolved. Fever associated with infections has a temporary adverse impact on spermatogenesis and sperm quality, but usually does not result in irreversible damage to male fertility. The adverse effect on sperm is characterized by decreased concentrations, alterations in morphology, diminished motility and increased DNA fragmentation (MacLeod 1951, Evenson *et al.* 2000). These defects can persist for months after the end of fever (Carlsen *et al.* 2003, Sergerie *et al.* 2007).

In patients infected with SARS-CoV-2, fever is one of the most common symptoms (Huang *et al.* 2020), but is often overshadowed by other manifestations of the disease. Increased sperm DNA fragmentation results in decreased fertility, poor embryo quality, and increased embryonic loss (Simon *et al.* 2011, Robinson *et al.* 2012). Of note, impaired sperm quality may persist and males recovering from COVID-19 should be monitored for gamete quality and delayed procreation may be required.

### Possible impact on reproduction of patients recovering from COVID-19

The effects of SARS-CoV-2 infection on male reproduction are generally consistent with previous results on SARS-CoV-1 (Table 1). Sexual transmission for patients recovering from COVID-19 seems unlikely, but the risk of impaired spermatogenesis has been observed in patients with moderate infection and in convalescents (Holtmann *et al.* 2020, Ma *et al.* 2020). The duration and severity of these abnormalities and their potential impact on progeny are not known. Thus, it will be important to carry out prospective and long-term studies for further clarification of adverse consequences of COVID-19. Establishing sperm quality as well as seeking clinical

**Table 1** Summary of known adverse effects of SARS-CoV and SARS-CoV-2 on male reproduction.

Adverse effects	SARS-CoV-1	SARS-CoV-2	References	
			SARS-CoV-1	SARS-CoV-2
Virus was detected in testicular tissues	+	+	Zhao <i>et al.</i> (2003c), Zhao <i>et al.</i> (2003a)	Bian <i>et al.</i> (2020), Yang <i>et al.</i> (2020b)
	–	–	Zhang <i>et al.</i> (2003), Zhao <i>et al.</i> (2003b), Ding <i>et al.</i> (2004)	Song <i>et al.</i> (2020)
Virus was detected in semen	ND	+		Li <i>et al.</i> (2020a)
		–		Guo <i>et al.</i> (2020), Holtmann <i>et al.</i> (2020), Li <i>et al.</i> (2020b), Ma <i>et al.</i> (2020), Pan <i>et al.</i> (2020), Song <i>et al.</i> (2020)
Semen parameters abnormalities	ND	+		Holtmann <i>et al.</i> (2020), Ma <i>et al.</i> (2020)
Orchitis	+	+	Xu <i>et al.</i> (2006)	Li <i>et al.</i> (2020b), Nunes Duarte-Neto <i>et al.</i> (2020), Yang <i>et al.</i> (2020b)
Sex hormone abnormalities	+	+	Wang <i>et al.</i> (2005), Wei <i>et al.</i> (2010)	Ma <i>et al.</i> (2020), Rastrelli <i>et al.</i> (2020), Salciccia <i>et al.</i> (2020), Schroeder <i>et al.</i> (2020)
Potential long-term effects	ND	ND		

+, indicates studies were performed and results were positive; –, indicates studies were performed and results were negative; ND, not determined indicates that assays were not performed.

examination and consultation are recommended before a planned pregnancy for convalescent patients.

Besides parameters that can be quantified, there may be some subtle alterations in spermatozoa. Environmental effects, including lifestyle, can be translated into heritable information stored in the epigenome and RNA of gametes that impact progeny (Zhang *et al.* 2019). For example, obesity is a phenotype that has a high probability of inheritance from overweight fathers. In a region associated with appetite control, a dramatic reprogramming of sperm DNA methylation is observed after bariatric surgery-induced weight loss (Donkin *et al.* 2016). Moreover, in mouse models of metabolic disorders, if male mice are given a high-fat diet (HFD), the expression profiles and modifications of the sperm tRNA-derived small RNAs (tsRNAs) is altered. Normal zygotes injected with tsRNAs from HFD mice also developed the metabolic disorder phenotype (Chen *et al.* 2016). These intriguing lines of evidence suggest that environmental inputs can have a profound effect on offspring. Given that the period of infection with SARS-CoV-2 is short and spermatogonial self-renewal, the possibility of a transgenerational effect may be low. Nevertheless, whether COVID-19-mediated adverse effects can act on sperm and induce abnormal infants will require long-term surveillance.

### Development of experimental models for future investigation

Considering the long-term threat to international public health posed by COVID-19, the effect of SARS-CoV-2 infection on male reproduction needs systematic investigation. However, human tissue from autopsies studies is difficult to obtain and often inadequate for investigations of molecular mechanisms. The establishment of human ACE2 transgenic mice (Bao *et al.* 2020) and Rhesus macaques (Shan *et al.* 2020) provide models but fail to mimic human reproduction physiology. Human organoids have played a role in studying susceptibility of liver and pancreas to SARS-CoV-2 (Yang *et al.* 2020a, Zhao *et al.* 2020), but the development of complete testis organoids is still in its infancy (Oliver & Stukenborg 2020). A simplified human testicular organoid to mimic the ZIKA virus infection has been reported, but lacks the integral structure of the BTB (Strange *et al.* 2018). Thus, further improvement of experimental models of the human reproductive system is vital to develop strategies to cope with the threat of current and future testis-tropic viruses.

### Conclusions and perspectives

The impact of SARS-CoV-2 on male reproduction has been confirmed in preliminary studies and summarized in Fig. 1. A virus-induced cytokine storm appears to be the major and most destructive effect of COVID-19

on male testes. Because of the low levels of ACE2 and TMPRSS2 expressed in the testis, the probability of SARS-CoV-2 infection in an ACE2-dependent manner remains relatively low. Instead, severe inflammation secondary to viral infection could disrupt the BTB and cause orchitis. The subsequent damage of germ cells and interstitium would adversely impact spermatogenesis and hormone production within the testes. Breaching the protective BTB could account for the seeding of the SARS-CoV-2 virus in the testis. If so, the adventitious presence of the virus in testes or seminal fluid should be low and it seems unlikely that SARS-CoV-2 is sexually transmitted.

While anti-inflammatory therapies will be helpful to treat acute orchitis, fertility evaluation and pre-pregnancy consultation are important for convalescent patients. Long-term follow-up studies should help validate the preliminary studies and provide guidance for the need for assisted reproductive technologies. Development of improved experimental models of human male reproduction will be valuable to determine the pathophysiology of SARS-CoV-2 viral infections. It will also be important to evaluate and monitor possible long-term transgenerational effects.

### Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of this review.

### Funding

This work was supported by National Key R&D Program of China (2018YFC1004502, 2018YFC1004001) and the National Natural Science Foundation of China (NSFC 31771661).

### Author contribution statement

Li-quan Zhou conceived the ideas and Yu Tian wrote the manuscript.

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Received 17 September 2020

First decision 12 October 2020

Revised Manuscript received 5 November 2020

Accepted 16 November 2020