

An Experimental Model for Peri-conceptual COVID-19 Pregnancy Loss and Proposed Interventions to Optimize Outcomes

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Reports appear to give reassurance that vertical transmission near term is unlikely, but risks of incidental SARS-CoV-2 infection during fertility treatments, at embryo implantation, or in the first trimester remain unknown. If early pregnancy sequela in the current COVID-19 pandemic are modeled from the 2004 Coronavirus outbreak data, then SARS-CoV-2 infection proximate to blastocyst nidation is likely to cause implantation failure or spontaneous abortion. Our model explains why this outcome is less attributable to virus-associated maternal pulmonary distress and instead derives from systemic inflammation and interference with trophoblast-endometrium molecular signaling required for implantation. COVID-19 is often accompanied by high levels of IL-6, IL-8, TNF-alpha and other cytokines, a process implicated in pulmonary collapse and systemic organ failure. Yet when regarded in an early reproductive context, this “cytokine storm” of COVID-19 triggers a pro-coagulative state hostile to normal *in utero* blastocyst/fetal development. Evidence from obstetrics is accumulating to show that mothers with SARS-CoV-2 deliver placentas with abnormal interstitial villi fibrin deposits, diffuse infarcts, and hemangiomatous changes. This model classifies such lesions as permissive at term but catastrophic near embryo implantation or early first trimester pregnancy. Clinical experience with recurrent pregnancy loss offers workable interventions to address this challenge, but success will depend on prompt and accurate SARS-CoV-2 diagnosis. Although no professional guidelines currently exist for SARS-CoV-2 in early pregnancy, this model would warrant a high-risk designation for such cases; these patients should receive priority access to screening and treatment resources.

Key words: SARS-CoV-2, hypercoagulation, inflammation, implantation

All coronaviruses are single-stranded, positive sense RNA viruses (Baltimore Class IV) within the family *Coronaviridae*, a viral type first described in the 1960's. Their ~30 Kb genome

places coronaviruses among the largest known RNA viruses. The initial population of asymptomatic SARS-CoV-2 carriers permitted explosive contagion, leading to a crushing

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challenge for clinics and hospitals. While public awareness programs emphasizing hand hygiene, reduced travel, and social distancing can diminish the rate of viral spread, testing and treatment demands continue to outpace provision of basic services in many jurisdictions.

To develop a phylogenetic tree from multiple reference sources, Benvenuto et al. (1) used an unconstrained Bayesian analysis to find both nucleocapsid and S-protein under positive pressure, with 2019-nCoV having substantial homology with a previously isolated (2015) bat SARS-like coronavirus sequence. These data suggest the “novel Coronavirus” is distinct from the virus responsible for SARS and was likely acquired from bats after a mutation which conferred the ability to infect humans (1). Li et al. (2) identified three phylogenetic and transmission clusters via network analysis, with only one cluster identified from banked genome sequences of 2019-nCoV strains. An estimated mean evolutionary rate for 2019-nCoV has been calculated, providing useful information for disease mapping and public health

guidance (2, 3). SARS-CoV-2 as a human pathogen is thought to be an independent emergence, distinct from the SARS-CoV infections of 2002-2003. Although SARS-CoV-2 is not a descendent of SARS-CoV, the two viruses are sufficiently similar that their shared evolutionary histories are informative (4). As efforts to develop an effective vaccine, treatment, and rapid screening for COVID-19 are redoubled, increased appreciation of underlying cytokine processes is gaining momentum. Perhaps the most intriguing of these involves a type of virus-induced hypofibrinolysis which brings the most relevance to the early reproductive sequence.

Early reproduction and coronavirus

While most COVID-19 research has focused on pulmonary angiotensin-converting enzyme 2 (ACE-2) receptor, *ACE-2* mRNA from endometrium localizes this receptor to the uterine interior as well (5). Our hypothesis incorporates data from endometrial samples assessed by immunoassay throughout the menstrual cycle (Figure 1). Such cells obtained by transcervical

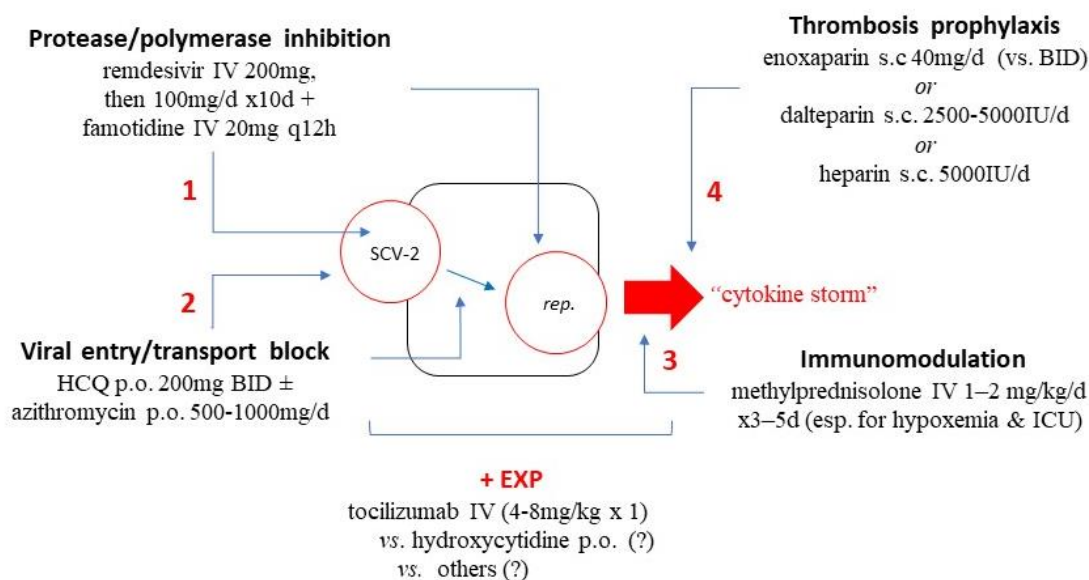


Fig. 1. Representative multifactorial treatment model for synergistic SARS CoV 2 (SCV 2) infection including antiviral, immunosuppressive, anticoagulant, and experimental (EXP) components. Some agents intervene by multiple mechanisms; exact dosing awaits multicenter clinical trial data. HCQ: hydroxychloroquine; rep: replication.

lavage identified the angiotensin receptor (Ang1-7) in glandular epithelium of mid- to late secretory phase (5). This finding is relevant to the current SARS-CoV-2 pandemic because the virus uses the ACE-2 receptor for cell access, setting the stage for immune hyperactivity. Crucially, where embryo implantation defects occur in experimental knock-out animal models, pro-inflammatory Th-1 cytokines are usually involved (6). Threshold levels exist for IF- γ , IL-2, and TNF- α , designated as “abortifacients” by prior research (7, 8).

The effects of anti-inflammatory mediators such as IL-4, IL-10, IL-13, and TGF- β are physiologically offset by proinflammatory cytokines like IFN- γ , IL-1 α , IL-1 β , IL-6, and IL-12. This balance is recalibrated in pregnancy as a putative fetal protective effect, although such permissive immunotolerance comes with increased maternal susceptibility to pathogens (9-11). COVID-19 is characterized by augmented IFN- γ and IL-1 β , IL-4, IL-10 and IFN- γ production (12). Preferential stimulation of Th1 immunity is often triggered in COVID-19 disease, yielding a sharp cytokine uptick for at least two weeks after initial infection (13). Interestingly, elevated IL-6 levels have been linked to increased mortality in COVID-19 (9, 14).

Xiong et al. (15) were among the first to report on a convalescing pregnant woman with COVID-19 in the third trimester. The uncomplicated vaginal delivery of a baby without SARS-CoV-2 infection suggested that intrauterine transmission of coronavirus is unlikely in late pregnancy. Italian experts (16) reviewed 13 studies, and found that vaginal delivery was performed in six cases and cesarean delivery (performed for worsening maternal status) in 31 cases (48.4%). Two newborns in this group were SARS-CoV-2 positive by RT-PCR assay, while IgG and IgM levels for SARS-CoV-2 were elevated with negative RT-PCR antigen for three offspring (16). Another summary identified 55 pregnant women infected with

COVID-19 and their 46 neonates showing no vertical transmission (9). A more recent analysis of COVID-19 in pregnancy (17) found a vertical transmission rate of 11%. While limited, the available evidence tends to show that maternal-fetal SARS-CoV-2 passage is low (16-18). Placental histology reviewed at delivery from new mothers with confirmed SARS-CoV-2 infection found fibrin deposition near and within villi, with occasional increased local syncytial nodularity. Notably, chorionic hemangioma and massive placental infarction were also identified (19).

Available research on COVID-19 (published after peer-review or posted on early preprint sharing platforms) provides almost no data on SARS-CoV-2 infection impact on early pregnancy. One large meta-analysis of 266 COVID-19 pregnancies illustrates the problem (20) as time of when SARS-CoV-2 infection initially occurred in pregnancy is not reported. Curiously, two studies did include early pregnancy data (21, 22) and these were the only samples where any miscarriage or abortion occurred. These events align with our “catastrophic error” model of COVID-19, such that infection with this virus soon after implantation or in early gestation is antagonistic with pregnancy viability.

Since embryo orientation, apposition, docking and invasion are all under cytokine control, any excess of pro- or anti-inflammatory signaling is detrimental to pregnancy outcome (23). Moreover, the local chemokine milieu prevailing at implantation and shortly thereafter determines the maternal macrophage phenotype (24). Normally, first trimester decidual cells rely on IL-1 β and TNF- α signaling via the NF- κ B pathway for macrophage colony-stimulating factor-1 production. This results in increasing maternal macrophage CD163 expression and reducing signal-regulatory protein- α expression (24). While numerous other pathways must also remain operational to facilitate implantation, interference with any of these by cytokine excess can be sufficient to disrupt decidual

Newer candidates? Mechanism and rationale

Hydroxychloroquine (HCQ) - FDA pregnancy category C

Pathogen attachment to exposed respiratory cells requires action by a spike (S) protein, and marks the first step in viral replication. The viral S-protein facilitates entry through ACE-2 receptors as well as by exploiting specific ganglioside markers present on cell surfaces. It appears that HCQ metabolites attach avidly to a highly conserved viral ganglioside binding domain, causing a mismatch between sialic acids and/or viral S-protein and host/target cell (31, 32). HCQ also serves as a weak base to raise interior pH of normally acidic lysosomes and endosomes, structures needed by SARS-CoV-2 for membrane fusion and cellular access (33). Because low pH is crucial for proper endosomal action, it has been surmised (34) that endosome maturation might be blocked after endocytosis, resulting in failure of further transport of virions for subsequent exteriorization. Researchers recently confirmed that HCQ effectively blocks both the initial insertion step and post-entry stages of SARS-CoV-2 across certain organelles, thus severely limiting viral transport and release functions (35-37). HCQ may also restore defective trophoblast function impaired secondary to antiphospholipid antibodies, but clinical data remains inconclusive and awaits appraisal through clinical trials (38, 39). HCQ has been shown not to adversely affect cell turnover, nutrient transport, cytokine release or overall placental explant function *in vitro*, and thus HCQ may have a protective anti-inflammatory effect (40).

Remdesivir - FDA pregnancy category: not assigned

An adenosine analog, remdesivir, causes early viral RNA truncation by interfering both with viral RNA polymerase and proofreading, resulting in dampened viral RNA production. Experience with remdesivir accumulated from its prior use in

treating Ebola and Marburg virus infections, but it is also effective against other single stranded RNA viruses including Lassa fever virus, Hendra virus, and several coronaviruses. Almost no reproductive system safety data exist on remdesivir, although its use in a neonate who contracted Ebola virus by vertical transmission was successful (41).

Tocilizumab - FDA pregnancy category: not assigned

Perhaps the most structurally complex of candidate drugs currently under study against COVID-19, tocilizumab is a humanized (genetically engineered) monoclonal antibody which targets the IL-6 receptor. IL-6 modulates immune response and is implicated in the pathogenesis of many diseases. Tocilizumab binds IL-6 receptors, blocking the pro-inflammatory action to attenuate the “cytokine storm” and associated pulmonary injury. Although reproductive health data are scant, preliminary reports agree on the safety of tocilizumab during pregnancy, and probably lactation. This medication likely has no negative effects on the embryo or fetus (42).

Hydroxycytidine - FDA pregnancy category: not assigned

Sheahan et al. (43) recently reported on ribonucleoside analog β -D-N4-hydroxycytidine (NHC, EIDD-1931), having potent antiviral activity in a murine model against SARS-CoV-2, MERS-CoV, SARS-CoV, and related zoonotic group 2b or 2c Bat-CoVs. Because this investigational agent retained antiviral effectiveness across a wide range of therapeutic targets, and particularly in the setting of remdesivir resistance (43), hydroxycytidine could be useful in refractory or atypical COVID-19 cases.

Conclusion

Why have so few successful outcomes been reported when COVID-19 occurs early in pregnancy? The “cytokine storm” of COVID-19 complicates early blastocyst- endometrial signaling required for embryo docking and amplifies microvascular thrombus formation at the

trophectoderm-endometrial interface, as discussed here. While both pathways are familiar in clinical reproductive medicine settings, there is no best standard to manage the problem. Our approach incorporates steroid immunomodulation and anticoagulation together with investigational antiviral therapy, and awaits findings from randomized controlled trial currently underway. The microvascular thrombosis in SARS-CoV-2 infection contrasts sharply with the consumptive coagulopathy encountered in Ebola and Lassa hemorrhagic fever (44, 45); available evidence supports our ‘cytokine-mediated inflammation and disruption of maternal coagulation’ model to explain the apparent adverse early reproductive outcomes with COVID-19.

Conflict of interest

The authors declare no conflict of interest.

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