Immune responses at the maternal-fetal interface

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Pregnancy poses an immunological challenge because a genetically distinct (nonself) fetus must be supported within the pregnant female for the required gestational period. Placentation, or the establishment of the fatally derived placenta, is a common strategy used by eutherian mammals to protect the fetus and promote its growth. However, the substantial morphological differences of the placental architecture among species suggest that the process of placentation results from convergent evolution. Although there are considerable similarities in placental function across placental mammals, there are important differences that arise owing to species-specific immunological (and other biological) constraints. This Review focuses on the immunological similarities and differences that occur at the maternal-fetal interface in the context of human and mouse pregnancies. We discuss how the decidua and placenta of these different species form key immunological barriers that sustain maternal tolerance yet generate innate immune responses that prevent microbial infections.

INTRODUCTION

Placentation is a common strategy used across eutherian mammals to protect and promote fetal growth. Although mice are commonly used to study the maternal-fetal interface within the immunological context of pregnancy, differences exist in placental architecture, gestational period, and mechanisms of maternal tolerance from humans. Throughout this Review, we focus on the immunological similarities and differences during human and mouse pregnancies. We define the fetal and maternal components, their interactions, and mechanisms of mediating maternal tolerance in both species. We also examine the role of the placenta as a barrier to maternally transmitted pathogens and conclude with a discussion on the strengths and weaknesses of commonly used models of the human placenta.

THE MATERNAL-FETAL INTERFACE IN HUMANS AND MICE

The maternal-fetal interface is composed of the maternal-derived decidua and the fetally derived placenta (Fig. 1). In both mice and humans, the placenta develops from the trophectoderm of the blastocyst. During implantation, invading trophoblasts anchor the blastocyst to the specialized uterine epithelium (the decidua), on which placentation ensues. Over the course of pregnancy, the placenta is the sole site for all gas, nutrient, and waste exchange between the fetus and mother.

Decidua formation

The placenta is embedded within the decidua, the maternal component of the maternal-fetal interface. The decidua only exists during pregnancy and originates from the endometrial lining of the uterus (the endometrium). At the conclusion of pregnancy (parturition), the decidua is shed, to be rebuilt only upon subsequent pregnancy. However, signs of predecidualization can be observed within the nonpregnant human endometrium halfway through the luteal phase (around days 23 to 25 of the menstrual cycle), including increased prominence of the spiral arterioles, differentiation of endometrial fibroblast-like cells into enlarged and granulated decidual stromal cells, and an influx of leukocytes (1). Timing is critical for pregnancy to occur. Implantation must take place before this predecidualization because this thickening of the endometrium is not amenable for implantation.

During decidualization, there is both fetally and maternally mediated remodeling of the spiral arteries so that the placenta becomes bathed in maternal blood, which facilitates exchange of nutrients, gases, and waste. After implantation, the endothelial lining of the spiral arteries is eroded (as well as the local decidual stromal cells), creating a fibrinoid wall embedded with invasive fetal placental trophoblasts (2). Maternal leukocytes, such as natural killer (NK) cells and macrophages, have been implicated in this remodeling process. These concordant efforts of fetal trophoblasts and maternal leukocytes result in the dilation of the spiral arteries, which decreases the force and maximizes the volume of the maternal blood bathing the placenta (2).

Placenta development

In humans, the definitive structure of the placenta is composed of villous trees and is established by the third week of gestation (Figs. 1 and 2). The structure of the human placenta is composed of both floating and anchoring villi. A single layer of contiguous multinucleated syncytiotrophoblasts (SYNs) lines the outermost surface of the human placenta villous trees and acts as the major cellular barrier between the fetal compartment and maternal blood. Underlying the SYN layer are the undifferentiated, mononucleated cytotrophoblasts (CTBs). CTBs are progenitor trophoblast cells and can fuse to replenish the SYN layer or differentiate into multinucleated extravillous trophoblasts (EVTs), which are located at the tips of the anchoring villi. During the first trimester, the human placenta is hemochorial, with two layers of trophoblasts separating the fetal and maternal bloodstreams (the SYN and CTBs). As the placenta grows, the underlying CTB layer thins and becomes dispersed; thus, the human placenta of the second and third trimesters is essentially hemomonochorial, with only a single layer of SYNs.
The SYN facilitates the transport of nutrients, gases, and waste across the maternal-fetal interface. The SYN also functions as the main endocrine cell of the placenta, producing human chorionic gonadotropin (hCG) and progesterone, vital hormones that support pregnancy (3). During the first 8 weeks of gestation, the SYN secretion of hCG is required to induce progesterone production by the corpus luteum (4). Afterward, the placenta itself becomes the major producer of progesterone (4). Although the mouse also requires progesterone during the course of gestation, its placenta does not synthesize progesterone and instead continuously throughout pregnancy relies on the corpus luteum for progesterone (3).

EVTs physically anchor the human placenta to the decidua. The invasive EVT is also important for remodeling the spiral arteries in the outer third of the myometrium. In the first trimester, EVTs act as a plug for the spiral arteries, thus creating a hypoxic environment by excluding the oxygen-rich maternal blood. During the transition from the first to the second trimester, the EVT plug is eroded, and the intervillous space (IVS) becomes flooded with maternal blood (Fig. 2). Later during gestation, the IVS can fill with as much as 150 ml of maternal blood. The presence of maternal blood in direct contact with the placenta allows for efficient gas, nutrient, and waste exchange, which is maintained throughout the rest of pregnancy. Because the direct contact of maternal blood with the placenta does not occur until the end of the first trimester, this event distinguishes the early (first trimester) and later (second and third trimesters) stages of pregnancy.

Fig. 1. Comparison of human and mouse placentation. (Left) Before placentation, the blastocysts of humans and mice are similar. (Middle and right) However, upon implantation, placental development progresses differently. (Top middle) After blastocyst implantation, the human SYN layer burrows into the maternal decidua. By the third week of gestation, the definitive human placenta is formed and is composed of villous trees. However, at this stage of human pregnancy, the fetal-derived placenta does not directly contact maternal blood. (Top right) EVTs anchor the villi to the decidua and are involved in the remodeling of the spiral arteries to flood the IVS with maternal blood toward the end of the first trimester of pregnancy. The surface of the villi is covered by the SYN layer, which directly contacts the maternal blood and facilitates the transport of nutrients, gases, and waste across the placental barrier. Underlying the SYN layer are mononucleated CTBs that can either fuse to replenish the syncytial layer or differentiate into EVTs. By contrast, mouse placental development and organization are different from those in humans. (Bottom middle) Upon implantation, the trophectoderm differentiates, and trophoblast giant cells encapsulate the developing mouse embryo. Halfway through gestation, the definitive mouse placenta is fully formed and functional, where (bottom right) the folded villi form a labyrinth structure that becomes perfused with maternal blood. The trophoblast giant cells channel maternal blood from the decidua through the spongiotrophoblast layer (a structure not present in the human placenta) toward the labyrinth zone. In the labyrinth zone, the maternal blood makes contact with the CTBs that overlay two separate layers of SYNs.
In the early stage of pregnancy, the IVS is filled with a clear fluid containing uterine gland secretions, which are phagocytosed by the SYN and serve as a nutrient source for the developing fetus (5). Uterine glands originate from invaginations of the endometrium and are required for establishing pregnancy. Among their varied secretions are growth factors that regulate placental development, including epidermal growth factor, vascular endothelial growth factor (VEGF), transforming growth factor–β (TGF-β), and leukemia inhibitory factor (6).

Decidualization and placenta in the mouse

Key differences exist between mice and humans during the stages of pregnancy. In particular, the time frame and establishment of the maternal-fetal interface in each species are distinct. Human gestation takes place over a period of 40 weeks, whereas in mice, it is about 3 weeks (Fig. 2). The timing of decidual formation and placenta also varies. In humans, the uterus is primed for decidualization, independent of fertilization, around menstrual cycle day 23 when the stromal cells near the (now prominent) spiral arteries begin to differentiate into large predecidual cells (1). However, in mice, spiral artery outgrowth and decidualization do not begin until fertilization and blastocyst attachment to the uterus, respectively (2). Likewise, the placenta does not have a definitive structure in mice until the midpoint of gestation (around days 10.5 to 11.5), whereas the definitive placenta in humans forms far earlier in relative development (around day 21) (3). Thus, timing is critical to experimental design and interpretation when using the mouse (or any other animal) to model human pregnancy. A timeline highlighting the differences between the human and mouse placenta and the key events that occur throughout pregnancy is shown in Fig. 2.

Although the hemochorial mouse placenta shares features with the human placenta, several differences exist that affect physiology, immunity, and development (Fig. 1). Whereas the human placenta is structured as villous trees bathed in maternal blood (after the first trimester), the mouse placenta has a labyrinth structure perfused by maternal blood. In the mouse, the maternal blood is directed through trophoblast giant cell–lined channels in the spongiotrophoblast layer (a cell type not present in the human placenta) to the chorionic plate and back through the labyrinth zone containing the fetal vasculature (7). Unlike the anchoring chorionic villi of humans, the mouse chorionic projections are highly interconnected, presenting a maze-like structure through which the maternal blood must pass to leave the placenta. This labyrinth chorionic structure is lined by three layers of trophoblasts: two layers of SYNs overlaid with CTBs. In further contrast to the human placenta, the mouse CTBs directly contact the maternal blood. The trophoblast giant cells (large polyploid cells) anchor the mouse placenta to the decidua. Unlike human EVT’s, mouse giant cells are minimally invasive and do not remodel the maternal spiral arteries. These differences in physiology may affect the placental barrier functionally and the types of strategies used to protect the fetus from activation of the maternal immune system, as well as affect the pathways used by circulating pathogens to access the fetus.

**LEUKOCYTES AT THE MATERNAL-FETAL INTERFACE**

In addition to stromal cells, a remarkably large portion (~40%) of the decidua is composed of maternal leukocytes (Fig. 3). In the first-trimester decidua basalis (the site of implantation and trophoblast invasion/remodeling), decidual NK (dNK) cells comprise the majority (~70%) of immune cells, followed by decidual macrophages (20 to 25%) and T cells (3 to 10%) (8, 9). Maternal leukocytes are present in the decidua throughout pregnancy, although the population...
dNK cells

dNK cells are the largest population of maternal leukocytes that accumulate at the maternal-fetal interface, where they contribute to decidualization and implantation. Unlike their circulating counterparts, dNK cells produce a vast array of growth factors, angiogenic factors, and cytokines (15). Through these secretions, they help to remodel the decidua and spiral arteries, promote trophoblast invasion, and increase the availability of maternal blood at the implantation site (15–19). In the mouse decidua, dilation of the spiral arteries is induced by dNK-secreted type II interferon (IFN-γ) (20). dNK production of interleukin-8 (IL-8) and CXCL10 has been implicated in promoting EVT migration into the placental bed (15). An absence of dNK cells in mouse pregnancies is associated with decreased fetus viability and abnormal formation of the decidual structure and spiral arteries at the implantation site (21, 22). Analogously, in human patients with unexplained infertility, endometrial biopsies have found substantially fewer NK cells than in fertile counterparts (23). Interestingly, a subpopulation of dNK cells that exhibit enhanced production of IFN-γ and VEGFα has been identified recently as highly enriched in multigravid women (24). These pregnancy trained dNK (PTdNK) cells are transcriptionally distinct from other dNK cells, with higher expression of genes related to NK cell activation, growth factors, and immuno-modulatory proteins (24). With single-cell transcriptomics, another recent study identified three subsets of dNK cells in the first-trimester decidua, including a highly active subset of dNK cells with characteristics similar to the previously described PTdNK cells (25). Because improved placentation is seen upon subsequent pregnancies (26), it is interesting to speculate that this subset of dNK cells may become enriched upon subsequent pregnancies and boost decidua receptivity.

The dNK cells of pregnancy are phenotypically distinct from the peripherally circulating NK cells, and their specific origins remain enigmatic and may differ between humans and mice (8). These cells are highly granulated and distinguished as CD56++ CD16− (human) or CD122+ CD3− (mouse) (10, 11). Maternal leukocytes are recruited by chemokine gradients produced by decidual stromal cells and trophoblasts (12, 13) and are typically distinct from their peripherally circulating counterparts in phenotype and function. Most of our understanding of maternal leukocytes at the maternal-fetal interface has been determined from mouse studies and correlated to observations in human patients. Whereas the majority of the decidual leukocytes are dNK cells and decidual macrophages, T cell subsets also have key functions. More detailed information about leukocytes present at the maternal-fetal interface are available at (8, 9, 14).

Decidual macrophages

Decidual macrophages are the primary antigen-presenting cells (APCs) at the maternal-fetal interface in early pregnancy (9). Like uterine NK cells, levels of uterine macrophages rise and fall with the
menstrual cycle and then increase upon fertilization (32, 33). Phenotypically, decidual macrophages are believed to exist as regulatory/homeostatic, anti-inflammatory cells of an M2-like phenotype (9). The phenotype of decidual macrophages is believed to be influenced by trophoblasts, which secrete macrophage colony-stimulating factor (M-CSF) and IL-10 (34). Human decidual macrophages are CD163+CD206+DC-SIGN+ and predominantly express IL-10, CCL2, and CCL18 (35–40).

Decidual macrophages have many functions during pregnancy. Like dNK cells, they aid in remodeling of the spiral arteries and trophoblast invasion (41, 42) and localize to sites of disruption near the spiral arteries (43). Decidual macrophages in vitro produce VEGF and matrix metalloproteinase 9 (MMP9), which may promote angiogenesis and tissue remodeling (38, 44, 45). Decidual macrophages are proposed to perform “cleanup” functions by phagocytosing apoptotic trophoblasts, which prevents activation of pro-inflammatory pathways in the decidua (46–49). These cells also produce indoleamine 2,3-dioxygenase (IDO), which catabolizes tryptophan and hinders T cell activation (50, 51). Decidual macrophages may also have a more canonical antimicrobial role in protecting the fetus against infections, as suggested by the surface expression of pattern recognition receptors CD163 (hemoglobin scavenger receptor), CD206 (mannose receptor), and CD209 (DC-SIGN) (52).

**Regulatory T cells**

The importance of regulatory T (Treg) cells in pregnancy has become increasingly apparent. Observational experiments with human samples demonstrate the presence of Treg cells in the human decidua, and cases of human infertility, recurrent spontaneous abortions, and other pregnancy complications have been inversely correlated with Treg cell frequencies or functionality (53–56). In mice, fetal-specific Treg cells are recruited to and induced at the maternal-fetal interface, where they confer tolerance to fetal antigens and help maintain a homeostatic environment conducive to fetal survival. Fetal-specific Treg cells are capable of persisting beyond parturition while maintaining their functionality (57). Upon subsequent pregnancy with the same paternal background, the expansion of these cells correlates with decreased fetal resorption (57). Intriguingly, the origins of these fetal-specific Treg cells may be linked to the in utero exposure of noninherited maternal antigens (NIMAs). A multigenerational mouse study found pregnancies to be more successful when the sire has overlapping allogenicity with the maternal grandmother (58). Thus, in utero exposure to NIMA may expand a female’s Treg cell repertoire and explain the presence of maternal Treg cells specific for fetal (nonself) antigen (58).

**MATERNAL TOLERANCE**

Maternal tolerance, which permits a mother to carry the fetus to term despite the presence of foreign fetal antigen, is a poorly understood phenomenon that seems to defy some of the basic tenants of immunology. For a successful pregnancy, maternal tolerance must be established, and failure of maternal tolerance is correlated with preeclampsia and miscarriage (59–61). In general, tolerance is mediated by the restriction and modulation of leukocytes that permeate the maternal-fetal interface. Although there is an abundance of NK cells in the decidua, the numbers of dendritic cells (DCs) and effector T cells are low. As has been demonstrated in the mouse decidua, this may be due to the absence of local lymphatic vasculature in the endometrium (62, 63) and epigenetic silencing of T cell chemoattractants in decidual stromal cells (64). As for leukocytes that do gain access to the maternal-fetal interface, intercellular communication between the resident decidual leukocytes, stromal cells, and trophoblasts can alter the functional profile of leukocytes and promote regulatory phenotypes. For example, first-trimester human placental explants produce granulocyte colony-stimulating factor (G-CSF), IL-10, and TGF-β, which are known to promote differentiation of peripheral circulating monocytes and T cells into M2 MØ and Treg cells, respectively (34). Apoptosis is also used to mediate immune privilege. The SYN secretes exosomes that express TNF-related apoptosis-inducing ligand (TRAIL) and Fas ligand on their surface, which are capable of binding to their cognate death receptors on leukocytes to trigger apoptosis (65).

Maternal tolerance may also occur through species-specific mechanisms. In humans, placental EVTs express human leukocyte antigen–G (HLA–G), a nonclassical major histocompatibility complex (MHC) molecule, for which there is no homolog in the mouse genome. Unlike the canonical class I MHC molecules, of which thousands of allelic variations that serve to distinguish self from nonself exist, there are only 16 protein variants of HLA–G (66). Solely expressed by EVTs, HLA–G binds to dNK inhibitory receptors KIR2DL4 (67) and LILRB (68) to protect the trophoblasts from NK-mediated cytolysis (69). Likewise, the membrane-localized regulator of complement, Crry, is an example of a rodent-specific mechanism that protects the mouse placenta from the deposition and activation of circulating maternal complement, and its expression is required for fetal survival (70). Because Crry is rodent specific, it remains to be determined whether the human placenta also expresses such complement regulatory proteins that inhibit complement deposition and activation. Such disparities in mediating maternal tolerance between mice and man may be due to differences in the degree of placental invasiveness and length of gestational period. Likewise, different mechanisms could have arisen independently to handle the same problem of maternal immune responses that antagonize fetal viability.

**INTRINSIC IMMUNE RESPONSES OF THE PLACENTA**

Once the IVS fills with maternal blood, the placenta is continuously exposed to any and all pathogens circulating systemically in the maternal circulation. Hence, the placenta has several intrinsic defenses to protect the fetus from infection (Fig. 4).

**Structure and location**

The architecture of the human placenta allows it to present its strongest cellular defense, the SYN layer, on its outermost surface. The SYN forms a single continuous cell and thus lacks cellular junctions that can be exploited by pathogens or modulated by inflammatory signals. This contrasts with the mouse, which is arranged with the two layers of SYN buried beneath the CTBs contacting the maternal bloodstream. Another physical property that confers microbial resistance to the SYN is the dense cytoskeletal network that creates a dense brush border formed at the apical surface. This brush border not only provides a vast surface area for nutrient and gas exchange between the maternal and fetal compartments but also protects from direct microbial invasion, in part because of the dense underlying actin network. For example, SYNs are highly resistant to infection by *Listeria monocytogenes* (71) but become more permissive upon pharmacological disruption of the actin cytoskeleton (72). In addition,
SYNs restrict *Toxoplasma gondii* entry, most likely via a particular plasma membrane composition not amenable to parasite attachment (73, 74).

**Secrected antiviral factors**

The placenta secretes antiviral molecules that broadly function to protect the fetus from congenital infections of all types, including the expression of pattern recognition receptors such as Toll-like receptors (TLRs), the constitutive expression of type III IFNs (IFN-λ), and the release of antimicrobial peptides. Inoculation of placent al trophoblasts with the parasite *T. gondii* induces the secretion of chemokines, including the potent T helper 2 and Treg cell chemoattractant CCL22, suggesting that parasite infection alters or signals to maternal-derived immune cells. Furthermore, SYN expression of the FcRn also suggests a protective role for maternal IgG within the fetal compartment through the development of passive immunity.

**Intracellular defenses**

In addition to these processes, the placenta can directly initiate innate defenses aimed at suppressing microbial infections and/or alerting the maternal immune system to infection. Placental trophoblasts recognize pathogens via Toll-like receptors and RIG-I–like receptors, which trigger the induction of antimicrobial signaling pathways (82, 83). Trophoblasts also exhibit high rates of basal autophagy, which can serve as a pan-antimicrobial strategy to restrict the replication of diverse intracellular pathogens (75, 84).

**MODELING THE MATERNAL-FETAL INTERFACE FOR STUDIES ON CONGENITAL INFECTION**

Despite the formidable barrier presented by the placenta, some pathogens are capable of overcoming these placental defenses and induce devastating consequences to the developing fetus. These pathogens are collectively referred to as TORCH pathogens with the acronym referring to *Toxoplasma*, *other* [ZIKV, *L. monocytogenes*, *Treponema pallidum*, varicella zoster virus (VZV), HIV, and others], rubella virus, cytomegalovirus (HCMV), and herpes simplex virus. To understand how these pathogens cause fetal disease, it is important to consider routes of entry to the fetal compartment and methods of evading the intrinsic defenses and barriers of the maternal-fetal interface. Several different models have been used to study this evasion, including immortalized trophoblast cell lines, primary trophoblast cultures, human tissue explants, and in vivo models. Each model has advantages and limitations, which must be understood to interpret and extrapolate the conclusions to human pregnancy.

**In vitro models to study human placental functions**

BeWo, JEG-3, and JAR cells are commonly used tractable trophoblast cell lines derived from choriocarcinomas. Although these cell lines all express trophoblast markers, they do not spontaneously fuse to form syncytia and thus are more suited to model either CTBs or EVTs and do not recapitulate the biology of the SYN layer. Accordingly, these cell lines do not recapitulate the microbial resistance phenotypes observed in primary trophoblasts or placental explants (73, 75, 85). BeWo cells can be compelled to syncytialize by treatment with agents that increase adenosine 3’5’-monophosphate (cAMP) levels to enhance the expression of endogenous retrovirus fusion proteins responsible for CTB fusion (86); however, the elevation of
intracellular cAMP levels can produce other phenotypes, and these cells still remain susceptible to microbial infection (73). In comparison, JEG-3 cells grown in a three-dimensional bioreactor-based system cocultured with human endothelial cells spontaneously fuse to form SYNs and are able to recapitulate resistance to T. gondii and viral infections, as well as the constitutive release of type III IFN-λ (85, 87). Primary human trophoblasts are excellent cellular models of CTBs and SYNs. However, the lack of efficient means to genetically manipulate these cells, coupled with their limited (usually 3 to 5 days) life span after isolation, limits their utility. Ex vivo placental explants maintain the distinct morphological structure of the placenta as well as the multicellular complexity and can be isolated at all stages of pregnancy (88). However, procurement of placentas from early gestation to midgestation is complicated by increasingly restrictive government regulations. Several countries and states within the United States have illegitimized medical research that uses fetal-derived tissue, which includes the placenta, and other localities have limited access to tissues obtained from elective terminations.

In vitro models to study human decidual functions
Although the abovementioned models can provide valuable information regarding specific trophoblast-pathogen interactions and trophoblast-intrinsic immunity, they lack the maternal component. As with trophoblast models, there are a variety of different endometrial epithelial cell lines available for studies, with particular lines better suited as models for particular regions of the endometrium (such as glandular versus luminal models) (89). Primary stromal cells can be obtained and are capable of decidualizing in culture. Cocultures of endometrial stromal cells and trophoblasts are used to model implantation (90, 91). These models have limitations, including the exclusion of maternal immune cells that also compose up to 40% of the decidua. Studies using ex vivo decidual explants are able to model the multicellular composition (including dNK cells and decidual macrophages) and three-dimensional structure better, and recent congenital transmission studies using this model have provided insight into the decidual innate immune response and mechanisms of viral transmission (92, 93).

Animal models
Animal models are necessary to understand the dynamic immunological complexities of maternal-fetal tolerance, inflammation at the maternal-fetal interface, and the disruption of tolerance associated with congenital infections. Although a number of studies on placenta biology have come from experiments in mice, other animal models have also provided insights. Commonly used in vivo models include nonhuman primates (NHPs), sheep, and rodents (94). As might be expected, NHPs are good models for human pregnancy because there are many common characteristics, including a hemochorial placenta, singleton pregnancies, and a long gestation period comparable with that of human pregnancy (94). However, these models are ethically challenging, may be difficult to access or generate for some researchers, and are costly. Sheep are also commonly used to study placental vasculature because their villous trees are shaped similarly to those of humans (94). However, placenta is different in sheep; in particular, the depth of implantation is minimal (with no trophoblast invasion through the endometrial epithelium), and there is a greater degree of separation between the fetal and maternal vasculature (epitheliochorial placenta). Rodents, and specifically mice, are the most commonly used animal models. Among rodents, the guinea pig shares more similarities with human pregnancy than the mouse, including the source and levels of progesterone produced, deep trophoblast invasion, and long gestation (around three times the length of mouse gestation).

Nonetheless, the most commonly used animal model for studying congenital transmission is the mouse because it facilitates the use of many valuable immunological tools and techniques; has a short gestation period and large litter size, which enables a robust sample size; and is relatively inexpensive. Mouse models are particularly advantageous in the context of genetic deficiencies (on the maternal or fetal side of the interface), and results from these animal studies can be complemented with data from human models. The use of the mouse model for congenital infections has elucidated some of the mechanisms by which pathogens cause congenital disease. Experiments on Listeria-induced fetal wastage have demonstrated the importance of maintaining maternal tolerance toward the fetus, where promoting the accumulation of fetal-specific CD8+ T cells in the decidua causes fetal resorption, and most of the damage to the fetus is likely the result of a loss of maternal tolerance rather than the maternal response necessary to control the bacterial infection (95). A similar phenomenon has been described in Salmonella-induced placental inflammation, in which the host response upsets the balance of maternal tolerance and leads to fetal loss (96).

However, when studying congenital infections, it is important to consider that many TORCH pathogens are species specific, and mice may lack susceptibility. Each model must be interpreted carefully, keeping in mind its limitations. One example is the use of a mouse pathogen analogous to the human TORCH pathogen, such as the use of the mouse cytomegalovirus (MCMV) as a substitute for HCMV. However, MCMV cannot cross the placental barrier (unlike HCMV) (97). Another approach to overcome the barrier of host specificity is the use of immunocompromised mice, such as studies on ZIKV that use mice lacking the receptor to type I IFN (77, 98, 99). Likewise, there also exists variability in both susceptibility and immune response between inbred strains of laboratory mice, as has been shown with Listeria (100). Notwithstanding these issues, mice are still a highly useful tool and have provided much insight on the complexity of the maternal immune response during congenital transmission.

CONCLUSIONS
Placenta is a common strategy used by eutherian mammals to generate a conduit and barrier between the maternal and fetal environments. The variety of strategies used to support placentation across the breadth of placental mammals is particularly interesting. Humans and mice both rely upon hemochorial placentas, but the structure and tissue organizations are distinct. Although the composition of the decidua is similar between mouse and human, the timing and mediators of decidualization are disparate. Through millions of years of evolution, each species has optimized the complex immunological equilibrium that is required to sustain a healthy pregnancy. Therefore, both the advantages and limitations of various models should be considered when extrapolating data to human congenital transmission and disease. Although no model is perfect, biological insights can be obtained through the continued use and development of in vitro, ex vivo, and in vivo models. Collectively, these systems will provide new paradigms for one of the most distinct aspects of human biology and, potentially, strategies to protect the developing fetus from potentially devastating congenital disease.
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