The Effect of Microbes Isolated from The Female Reproductive Tract of Pregnant Women on Fetuses

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ABSTRACT

Sections Info

Article history: Submitted: May 15, 2025 Final Revised: May 31, 2025 Accepted: June 14, 2025 Published: July 01, 2025

Keywords:

Maternal microbiome
Fetal development
Pregnancy complications
Reproductive tract infections
Lactobacillus
Dysbiosis
Immune regulation
Miscarriage
Preterm birth

Objective: This study explores the role of microorganisms isolated from the female reproductive tract during pregnancy and their potential impact on fetal development and pregnancy complications. Method: A descriptive, literature-based approach was employed to analyze microbial populations across various anatomical sites - vaginal, cervical, endometrial, tubal, and ovarian - focusing on their interactions with the maternal immune system and reproductive outcomes. Results: The findings reveal that microbial imbalance, particularly the reduction of Lactobacillus species, disrupts immune regulation and compromises epithelial integrity, leading to adverse outcomes such as premature labor, miscarriage, infections, and congenital abnormalities. Pathogens including Candida, Trichomonas, Cytomegalovirus, and Group B Streptococcus were identified as major contributors to these complications. Novelty: Unlike prior studies that often examine isolated reproductive sites, this research integrates microbiome data from both upper and lower reproductive tracts, offering a comprehensive view of organ-specific dysbiosis and its influence on maternal-fetal health. These insights suggest new directions for preventive strategies, targeted therapies, and the development of biomarkers aimed at restoring microbial equilibrium to safeguard pregnancy outcomes.

DOI: https://doi.org/10.61796/jmgcb.v2i7.1368

INTRODUCTION

The balance and diversity of microorganisms during pregnancy play a vital role in the health of the mother and fetus, as they have a direct impact on regulating pregnancy hormones and inflammation, as well as immune responses and maintaining the integrity of the female reproductive system through their direct interaction with the body, any disruption in the work of these organisms causes several problems or various complications (such as infection, premature birth, and recurrent miscarriages), when a change occurs in the natural microbe, it leads to the growth of organisms that cause inflammation in the urinary tract and vaginal infections. Therefore, any disruption in the microbial balance may lead to premature birth or miscarriage by hindering the process of fetal development [1][2]. In the context of the widespread prevalence of various vaginal infections during pregnancy or its early stages, the need for new strategies to diagnose and treat these infections emerges to avoid any diseases that lead to the loss of the fetus at any stage of its development [3][4].

The female reproductive tract is one of the most important mucosal environments, colonized by diverse microbes that naturally interact with the host to form a delicate and complex relationship that can be termed an ecosystem [5][6]. Most research has shown

that the reproductive tract microbiome plays a fundamental role in this microenvironment, interacting precisely with anatomical structures and immune responses, and holding significant potential for maintaining maternal health [7][8]. One of the major influences on human physiological and pathological functions is the existence of natural, symbiotic relationships between organisms and their hosts throughout the healthy body, including the female reproductive system. Each part has its own microbe, representing 9% of the total bacteria in the female body, such as the ovaries, fallopian tubes, endometrium, cervix, and vagina [9][10]. These parts have their own microbiome[11]. Lactobacillus bacteria are responsible for vaginal health in women. The upper genital tract (UGT) contains a less dense but highly diverse bacterial population [12][13].

MATERIAL AND METHODS

The methodology of this study was based on an integrative and descriptive analysis of microbial populations within the female reproductive tract, aiming to understand their influence on fetal development and pregnancy outcomes. The researchers adopted a literature-based, cross-sectional approach supported by observational data on microbial communities isolated from various anatomical regions, including the vagina, cervix, endometrium, fallopian tubes, and ovaries. Through a combination of microbiological characterization and analysis of clinical case findings, the study systematically identified dominant microbial taxa particularly Lactobacillus species and contrasted their presence in healthy versus dysbiotic states. Sources of microbial dysbiosis were categorized into bacterial, fungal, viral, and parasitic agents, with emphasis on clinical correlations such as premature birth, miscarriage, and congenital abnormalities. Data interpretation involved comparative analysis between microbial profiles under normal and pathological conditions, examining shifts in immune responses, epithelial barrier integrity, and cytokine activity. In addition, findings from peer-reviewed studies were synthesized to evaluate microbial-host interactions and their implications for maternal and fetal immune regulation. The method integrated tables and figures to illustrate differences in microbial composition and immune activation across reproductive tract compartments. By aligning microbial patterns with clinical indicators, the methodology not only enabled the identification of pathogenic mechanisms but also informed the development of preventive and therapeutic strategies. The approach relied on secondary data from microbiome research, immunological assays, and infection outcomes, enabling the formulation of hypotheses about microbiota-driven complications in pregnancy. This comprehensive methodology served as the basis for linking microbial dysbiosis to specific reproductive and fetal health risks.

RESULT AND DISCUSSION

Result

Pregnancy

Pregnancy occurs inside a woman's uterus, either with one fetus or multiple fetuses, as in the case of twins. Pregnancy can occur as a result of several processes, which may occur naturally after vaginal intercourse or through assisted reproductive techniques [14].

Components of the Female Reproductive System

The female reproductive system consists of two parts: the lower and the upper. The lower system includes the vulva and vagina (The birth canal), while the upper system consists of the ovaries (A pair of organs that produce eggs and female hormones such as estrogen), fallopian tubes, uterus (An internal organ where the fetus develops and develops before birth), and cervix (The round, thick end of the uterus, with an opening through which menstrual blood and the baby exit during birth) [15][16].

Vaginitis

When vaginitis occurs, the fetus is at risk of premature birth due to membrane rupture and microbial imbalance. It is one of the most important causes of congenital malformations caused by organisms present in the reproductive tract, such as chlamydia, mycoplasma, and ureaplasma. This can also lead to congenital malformations in fetuses or stillbirth [17].

Discussion

Symptoms of genital tract infections:

Vaginal itching, burning, pain, or irritation. Internal itching or irritation. Pain during intercourse. Pain during urination [18].

Causes of miscarriage

During pregnancy, many physiological changes occur. Pregnancy is a complex process that requires many changes to contain and nourish the fetus with essential nutrients such as fatty acids, amino acids, vitamins, and minerals. The female reproductive tract (FGT) is also exposed to numerous sexually and non-sexually transmitted diseases, which can lead to infections [19][20].

1. Bacterial cause

One of the most common bacteria found in the female reproductive tract Group B streptococcus (GBS), can cause infections in newborns, such as meningitis and pneumonia. Therefore, the mother should be given antibiotics before delivery to prevent transmission (Table 1).

Table 1. Shows the parts of the female reproductive system and some of the microbes that can infect it.

	1 _Vaginal	The vagina is considered the entrance to	(Chen et
1-lower	microbiome	the female reproductive tract and	al., 2017;
reproductive		contains the highest bacterial biomass.	Ravel et
tract		Lactobacillus is found in a higher	al., 2011;
		relative abundance (over 89%), while	Integrative,

		the presence of Staphylococcus,	2019;
		Synthetia, Veillonella, Streptococcus, Prevotella, and others remains	Santella et al., 2022).
	• • • •	controversial.	(01
	2 _Cervical	It was previously believed that the	(Chen et
	microbiome	cervical microbiome was the same as	al., 2017).
		the vaginal microbiome; however,	(0
		recent studies have confirmed	(Onywera
		differences between the two phyla .	et al.,
		Firmicutes was found to be the most	2019a;
		abundant phylum among the cervical	Onywera
		microbiome, and Lactobacillus was the dominant genus within this phylum . L.	et al., 2019b).
		crispatus in the cervix was found to	
		produce lactic acid and antimicrobial compounds, thus preventing many infections and acting as a shield against many diseases. Bacteroidetes is the second most abundant phylum, with Prevotella bacteria being the dominant.	genus (Keburiya et al., 2022).
2-upper	1_Endometrial	Some studies have shown that	(Chen B. et
reproductive	microbiome	Acinetobacter, Lactobacillus,	al., 2021).
tract	microbiome	Methylobacterium, Sphingobium, and Streptococcus dominate the endometrium. Several previous studies have also revealed that Lactobacillus is dominant in the endometrial microbiome (Lactobacillus >90%, other bacteria <10%), and that clinical pregnancy rates and live birth rates (i.e., reproductive success) are higher. However, a decrease in these bacteria (Lactobacillus <90%, other bacteria >10%) leads to microbial imbalance, which increases pregnancy complications, such as preterm birth, recurrent miscarriage, recurrent implantation failure, and biochemical pregnancies .Currently, the presence of an endometrial microbiome is considered a positive predictor of	(Moreno et al., 2016). (Bonzon-Jimenez and Labarta, 2021).

	reproductive success, which may	
	provide new insights and research	
	directions for the prevention and	
	treatment of pregnancy complications.	
2 _Tubal	The entire reproductive tract of women	(Walther-
microbiome	who had undergone total hysterectomy	Antonio et
	was studied and the results showed the	al., 2016).
	presence of Shigella bacteria in the	
	fallopian tubes	
3 _Ovarian	Many studies have described the gut	(Miles et
microbiome	microbiome as rare in healthy	al., 2017;
	individuals but found in malignant	Zhou et al.,
	diseases	2019;
		Banerjee et
		al., 2017).

The female reproductive system consists of organs that are continuously and sequentially involved in the birth of new life and are predisposed to resist many diseases that cause prenatal miscarriage ,Figure (1) illustrates the most important interactions between the microbiome and the female reproductive system in both normal and pathological conditions [21][22]. The Lactobacillus microbiome and its main metabolites, especially lactic acid, create a healthy environment for the female reproductive system, enhancing the integrity of the epithelial barriers and mucosal membranes. It also stabilizes local immune defenses, creating an efficient immunological environment [23][24].

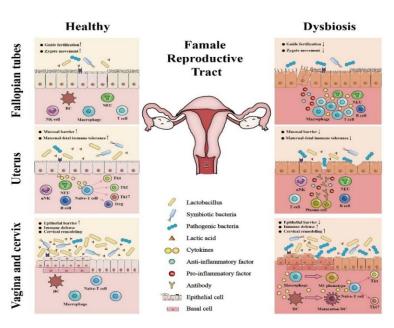


Figure 1. The figure shows the reproductive system in a state of health and dysbiosis.

The table 2 compares the vaginal and cervical microenvironment in healthy and dysbiotic states. In health, *Lactobacillus* dominance, intact epithelial barriers, and anti-inflammatory responses maintain immune balance. In contrast, dysbiosis features increased pathogenic bacteria, disrupted barriers, elevated pro-inflammatory cytokines, and macrophage-mediated inflammatory responses, compromising reproductive tract immunity and function [25][26].

Table 2. Vagina and Cervix

No.	Healthy	Dysbiosis
1	Dominated by Lactobacillus (yellow	Increase in pathogenic bacteria (pink
	rods).	and purple).
2	Lactic acid and anti-inflammatory	Disrupted epithelial barrier.
	cytokines are present.	
3	Epithelial barrier is intact.	More pro-inflammatory cytokines .
4	Macrophages and T-cells maintain	Macrophages release pro-
	immune homeostasis.	inflammatory mediators.

The table 3 highlights microbial and immunological differences between healthy and dysbiotic vaginal and cervical environments. Healthy states are marked by *Lactobacillus* dominance, intact epithelial barriers, and anti-inflammatory conditions. Dysbiosis involves increased pathogenic bacteria, epithelial disruption, elevated pro-inflammatory cytokines, and macrophage activation, contributing to compromised mucosal immunity and increased risk of reproductive complications [27][28].

Table 3. Uterus

No.	Healthy	Dysbiosis
1	Immune tolerance is maintained by	Immune tolerance is impaired.
	anti-inflammatory cytokines and	
	regulatory T-cells.	
2	Balanced microbiota supports a	Elevated inflammatory cytokines.
	favorable environment for implantation	
	and pregnancy.	
3		Increased recruitment of neutrophils
		(NE) and reduced regulatory T-cell
		activity.

The table 4 presents the impact of microbial balance in the fallopian tubes. A healthy state supports fertilization and early zygote movement through balanced microbiota and cytokines. In contrast, dysbiosis disrupts the fertilization environment, where inflammation and microbial imbalance hinder sperm-egg interaction or zygote transport, potentially impairing reproductive success [29][30].

Table 4. Fallopian Tubes

N.	Healthy	Dysbiosis
1	Support for fertilization and early	Disruption of fertilization
	zygote movement.	environment.
2	Balanced microbiota and cytokines.	Inflammation and bacterial imbalance
		can interfere with sperm-egg
		interaction or zygote transport.

2. Fungal cause

Fungal diseases are considered one of the most important causes of clinical diseases [31]. Hippocrates and Galen identified the oldest types of infection, fungal infections, which can be chronic, superficial, or deep [32]. Vaginal infections caused by yeast-like fungi of the genus Candida are the main infection during pregnancy, Fungi were identified in the 19th century as infections of the reproductive system during pregnancy and newborns, and how they interact with each other [33][34]. The infection can be transmitted to the fetus via ascending (from the vagina) or through the bloodstream in exceptional cases, even when the placenta and fetal membranes are protective against infection. They can cross the barrier without damaging the mucous membranes ,As a result of congenital candidiasis and Candida fungi invading the membranes, a serious infection occurs in newborns in the twenty-third week of pregnancy, leading to intrauterine infection with Candida fungi,This causes elevated levels of inflammation in the mother's blood (leukocytes, procalcitonin, C-reactive protein), which are also detected in the baby's blood after birth. Therefore, it is preferable to terminate the pregnancy early [35][36].

3. Parasitic cause

Trichomoniasis is a sexually transmitted parasitic infection that affects the reproductive system. During pregnancy, trichomoniasis may increase the risk of premature birth and low birth weight [37].

4. Viral Cause

Viral infections are among the most important factors that increase the risk of miscarriage, stillbirth, fetal brain injury, hearing loss, intrauterine growth retardation, cataracts, and other fetal abnormalities [38][39]. The placenta also acts as an immune shield to protect the fetus from pathogens during maternal viral infections. However, most viruses have evolved mechanisms to match the pathogenesis of the disease. In response to infection, the mother's immune system increases cytokine activation, which negatively affects the fetus. Cytomegalovirus (CMV) is one of the most important viral infections and the most common cause of birth defects in fetuses [40]. It is a viral infection that can infect the body at any age. Once infected, the virus remains in the body for life, and the disease often does not produce any signs or symptoms and is not detected until after diagnosis. However, its symptoms may appear in some patients, especially those with weakened immunity, as it is transmitted to cells through direct contact with the body fluids of an infected person, such as urine, sweat, blood, and semen. It may also be

transmitted from mother to child during pregnancy or breastfeeding. It is called cytomegalovirus because it enlarges the size of the infected cell, slowly replicating within it [41].

Maternal immune system

The effect of the microbiome on the fetal immune system and its development, The diversity of the maternal microbiome contributes to the development of the immune system through exposure to multiple antigens and may lead to the maturation of the immune system, especially microbes that help in the production of short-chain fatty acids (SCFAs), These acids have anti-inflammatory properties, which can reduce the risk of diseases in children [42].

The placenta and its role

The placenta was previously believed to be a completely sterile environment, but research has proven otherwise. It contains a unique community of organisms that directly influence the fetus, particularly its growth and the development of the maternal immune system. It also guides fertilization and promotes zygote transfer. However, any disruption of the microbial balance and the invasion of pathogenic bacteria negatively impact reproductive function through several factors, as shown in Table (5) ,it is important to highlight these interactions and the essential role of the microbiome in regulating physiological processes within the female reproductive tract, including fertilization, embryo implantation, fetal development, childbirth, and protection against infection.

The table 5 outlines key pathological effects of microbial imbalance in the female reproductive tract. Dysbiosis leads to reduced barrier integrity, heightened immune activation, premature cervical remodeling, impaired maternal-fetal immune tolerance, and decreased ciliary function. These disruptions collectively compromise reproductive health and increase the risk of adverse pregnancy outcomes.

Table 5. Factors Contributing to Microbial Dysbiosis-Induced Reproductive Complications

No.	Factors	Low/High
1	Damage to epithelial and mucosal barriers	↓ barrier integrity
2	Overstimulation of immune responses	↑ immune
		activation
3	Causing premature cervical remodeling	↑ cervical changes
4	Impairment of maternal and fetal immune tolerance	↓ tolerance
5	Causing edema, necrosis, and loss of fallopian tube cilia	↓ ciliary function
	function	

Sexually Transmitted Diseases (STIs)

One of the direct causes of early miscarriage is sexually transmitted infections (STIs). Most research focuses on the fact that changes in the composition and stability of the reproductive tract microbiome may also predispose individuals to a greater risk of STIs. Immune responses, through microbial variations in the vagina, affect the integrity

of the mucosal barrier. Any changes in the vaginal microbiome can exacerbate pregnancy complications caused by STIs, even during pregnancy, which can exacerbate pregnancy complications.

Prevention and treatment

Vaginal infections during pregnancy are very common and can be prevented and treated if the causative agent is identified, which may be bacterial, fungal (such as a yeast infection), or parasitic (such as trichomoniasis), Topical treatments (such as metronidazole and clindamycin) can be used to prevent them through good personal hygiene, changing wet clothes, avoiding irritants, wearing cotton underwear, and practicing safe sex.

CONCLUSION

Fundamental Finding : This study establishes a strong link between maternal reproductive tract microbiota composition and fetal health, revealing that microbial dysbiosis—especially the depletion of beneficial *Lactobacillus* species—can disrupt immune regulation, weaken epithelial barriers, and elevate the risk of adverse outcomes such as miscarriage, preterm birth, and congenital infections. **Implication :** These findings underscore the clinical potential of microbiome monitoring and modulation as a preventative and therapeutic strategy in maternal-fetal medicine, offering new avenues for early diagnosis and personalized interventions to improve pregnancy outcomes. **Limitation :** However, the study is primarily based on secondary data and lacks empirical validation through direct clinical or experimental trials, limiting the generalizability of its conclusions. **Future Research :** Subsequent studies should focus on longitudinal, multi-omics analyses of maternal microbiomes, including mechanistic investigations of host-microbiota interactions, to better understand causal pathways and support the development of microbiome-targeted therapies in obstetric care.

REFERENCES

- [1] C. Abbe and C. M. Mitchell, "Bacterial vaginosis: a review of approaches to treatment and prevention," Frontiers in Reproductive Health, vol. 5, p. 1100029, 2023.
- [2] J. M. Baker, D. M. Chase, and M. M. Herbst-Kralovetz, "Uterine microbiota: residents, tourists, or invaders?," Frontiers in Immunology, vol. 9, p. 208, 2018.
- [3] S. Banerjee et al., "The ovarian cancer oncobiome," Oncotarget, vol. 8, no. 22, pp. 36225, 2017.
- [4] A. Bjelica, N. Cetkovic, A. Trninic-Pjevic, and L. Mladenovic-Segedi, "The phenomenon of pregnancy A psychological view," Ginekologia Polska, vol. 89, no. 2, pp. 102–106, 2018.
- [5] C. Chen et al., "The microbiota continuum along the female reproductive tract and its relation to uterine-related diseases," Nature Communications, vol. 8, p. 875, 2017.

- [6] P. Chen, P. Chen, Y. Guo, C. Fang, and T. Li, "Interaction between chronic endometritis caused endometrial microbiota disorder and endometrial immune environment change in recurrent implantation failure," Frontiers in Immunology, vol. 12, p. 748447, 2021.
- [7] J. Eleutério Jr, A. B. Campaner, and N. S. De Carvalho, "Diagnosis and treatment of infectious vaginitis: Proposal for a new algorithm," Frontiers in Medicine, vol. 10, p. 1040072, 2023.
- [8] J. M. Franasiak and R. T. Scott Jr, "Reproductive tract microbiome in assisted reproductive technologies," Fertility and Sterility, vol. 104, no. 6, pp. 1364–1371, 2015.
- [9] H. Gao et al., "Deciphering the role of female reproductive tract microbiome in reproductive health: A review," Frontiers in Cellular and Infection Microbiology, vol. 14, p. 1351540, 2024.
- [10] S. Giakoumelou et al., "The role of infection in miscarriage," Human Reproduction Update, vol. 22, no. 1, pp. 116–133, 2016.
- [11] B. L. Hainer and M. V. Gibson, "Vaginitis: diagnosis and treatment," American Family Physician, vol. 83, no. 7, pp. 807–815, 2011.
- [12] H. K. Hussein, "Candida albicans and Abortion," Advances in Candida albicans, p. 101, 2021
- [13] Integrative Human Microbiome Project, "The Integrative Human Microbiome Project," Nature, vol. 569, no. 7758, pp. 641–648, 2019.
- [14] M. Ito et al., "A role for IL-17 in induction of an inflammation at the fetomaternal interface in preterm labour," Journal of Reproductive Immunology, vol. 84, no. 1, pp. 75–85, 2010.
- [15] L. K. Keburiya et al., "Does the uterine microbiota affect the reproductive outcomes in women with recurrent implantation failures?," BMC Women's Health, vol. 22, no. 1, p. 168, 2022
- [16] O. P. Lebedeva et al., "Female reproductive tract microbiome and early miscarriages," APMIS, vol. 131, no. 2, pp. 61–76, 2023.
- [17] C. Leeper and A. Lutzkanin, "Infections during pregnancy," Primary Care: Clinics in Office Practice, vol. 45, no. 3, pp. 567–586, 2018.
- [18] W. Mendling et al., "Vulvovaginal candidosis (excluding chronic mucocutaneous candidosis)," Geburtshilfe und Frauenheilkunde, vol. 75, no. 4, pp. 342–354, 2015.
- [19] S. M. Miles, B. L. Hardy, and D. S. Merrell, "Investigation of the microbiota of the reproductive tract in women undergoing a total hysterectomy and bilateral salpingo-oopherectomy," Fertility and Sterility, vol. 107, no. 3, pp. 813–820, 2017
- [20] C. M. Mitchell et al., "Colonization of the upper genital tract by vaginal bacterial species in nonpregnant women," American Journal of Obstetrics and Gynecology, vol. 212, no. 5, p. 611.e1, 2015.
- [21] S. S. Morelli et al., "The maternal immune system during pregnancy and its influence on fetal development," Research and Reports in Biology, pp. 171–189, 2015.

- [22] I. Moreno and J. M. Franasiak, "Endometrial microbiota—new player in town," Fertility and Sterility, vol. 108, no. 1, pp. 32–39, 2017.
- [23] I. Moreno and C. Simon, "Deciphering the effect of reproductive tract microbiota on human reproduction," Reproductive Medicine and Biology, vol. 18, no. 1, pp. 40–50, 2019.
- [24] I. Moreno et al., "Evidence that the endometrial microbiota has an effect on implantation success or failure," American Journal of Obstetrics and Gynecology, vol. 215, no. 6, pp. 684–703, 2016.
- [25] C. A. Muzny, P. Łaniewski, J. R. Schwebke, and M. M. Herbst-Kralovetz, "Host-vaginal microbiota interactions in the pathogenesis of bacterial vaginosis," Current Opinion in Infectious Diseases, vol. 33, no. 1, pp. 59–65, 2020.
- [26] N. Neu, J. Duchon, and P. Zachariah, "TORCH infections," Clinics in Perinatology, vol. 42, no. 1, pp. 77–103, 2015.
- [27] H. Onywera et al., "The cervical microbiota in reproductive-age South African women with and without human papillomavirus infection," Papillomavirus Research, vol. 7, pp. 154–163, 2019.
- [28] H. Onywera et al., "Factors associated with the composition and diversity of the cervical microbiota of reproductive-age Black South African women: a retrospective cross-sectional study," PeerJ, vol. 7, p. e7488, 2019.
- [29] M. Pal et al., "Growing role of fungi in the reproductive disorders of animals," Journal of Advances in Microbiology Research, vol. 5, no. 1, pp. 37–42, 2024.
- [30] N. Parveen et al., "Frequency of vaginal candidiasis in pregnant women attending routine antenatal clinic," Journal of the College of Physicians and Surgeons Pakistan, vol. 18, no. 3, pp. 154–157, 2008.
- [31] P. Punzón-Jiménez and E. Labarta, "The impact of the female genital tract microbiome in women health and reproduction: a review," Journal of Assisted Reproduction and Genetics, vol. 38, no. 10, pp. 2519–2541, 2021.
- [32] S. D. Rathod et al., "Epidemiologic Features of Vulvovaginal Candidiasis among Reproductive-Age Women in India," Infectious Diseases in Obstetrics and Gynecology, vol. 2012, p. 859071, 2012.
- [33] J. Ravel et al., "Vaginal microbiome of reproductive-age women," Proceedings of the National Academy of Sciences, vol. 108, suppl. 1, pp. 4680–4687, 2011.
- [34] B. Santella et al., "Microbiota and HPV: The role of viral infection on vaginal microbiota," Journal of Medical Virology, vol. 94, no. 9, pp. 4478–4484, 2022.
- [35] A. Stejskalova et al., "In vitro modelling of the physiological and diseased female reproductive system," Acta Biomaterialia, vol. 132, pp. 288–312, 2021.
- [36] E. L. Sweeney et al., "Group B Streptococcus serotypes Ia and V induce differential vaginal immune responses that may contribute to long term colonization of the female reproductive tract," American Journal of Reproductive Immunology, vol. 83, no. 1, p. e13199, 2020.

- [37] H. Verstraelen et al., "Characterisation of the human uterine microbiome in non-pregnant women through deep sequencing of the V1-2 region of the 16S rRNA gene," PeerJ, vol. 4, p. e1602, 2016.
- [38] K. M. A. Waldorf and R. M. McAdams, "Influence of infection during pregnancy on fetal development," Reproduction, vol. 146, no. 5, pp. R151–R162, 2013.
- [39] M. R. Walther-António et al., "Potential contribution of the uterine microbiome in the development of endometrial cancer," Genome Medicine, vol. 8, pp. 1–15, 2016.
- [40] B. Zhou et al., "The biodiversity composition of microbiome in ovarian carcinoma patients," Scientific Reports, vol. 9, no. 1, p. 1691, 2019.
- [41] N. Zhu et al., "Iron triangle of regulating the uterine microecology: Endometrial microbiota, immunity and endometrium," Frontiers in Immunology, vol. 13, p. 928475, 2022.
- [42] A. Chudnovets et al., "Role of inflammation in virus pathogenesis during pregnancy," Journal of Virology, vol. 95, no. 2, p. 10–1128, 2020.

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