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Article

Candida Vaginal Colonization in Pregnancy and Its Perinatal Impact: Cause, Effect or Coincidence?

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Abstract: In the vaginal ecosystem there is a symbiotic relationship between the microbiota and the host, particularly *Candida* spp., which are considered commensals of the vaginal mucosa. During pregnancy there is an increase in hormonal secretion that alters the vaginal microbiota and favors the development of *Candida* infection. Several studies have shown the increased prevalence of *Candida* during pregnancy, as well as its association with perinatal complications and current clinical practice guidelines recommend fluconazole as the treatment of choice during pregnancy in case of infection by *Candida albicans* and non-*albicans* *Candida*. However, in the Mexican population, this association has not been studied. So in this study, an intentional search for colonization by *Candida* spp. in pregnant women who attended the obstetric triage of the Hospital Regional de Alta Especialidad de Ixtapaluca during the period from May to October 2019, and they were followed up until the end of the pregnancy. Data on age, number of pregnancies, gestational age, body mass index, and perinatal outcomes were analyzed. Of 83 patients included in the study, yeasts corresponding to *C. albicans* (12) and *C. glabrata* (4) were isolated in 16. In the group of patients with *Candida*, the maternal age range was from 15 to 36 years old, of which 7 were adolescents and 2 with advanced maternal age, with an average of 2 gestations, the range of body mass index was from 19.3 to 42.4, of which 4 were overweight and 7 obese. Perinatal complications were observed in three patients, one presented chorioamnionitis and in two the neonates were small for gestational age. No association was found between colonization by *Candida* spp. and the number of pregnancies, gestational age, body mass index or perinatal results. However, it was found that age (adolescence) may be a risk factor for the development of vulvovaginitis during pregnancy.

Keywords: *Candida albicans*; colonization; vulvovaginal candidiasis; preterm delivery; premature rupture of membranes; small for gestational age; perinatal mortality

1. Introduction

The vagina provides a humid, nutritious and warm habitat for microbiota, which is composed of a large number of protective microorganisms and others that turn out to be opportunistic pathogens [1–4]. The vaginal microbiota is responsible for the production of antimicrobial and anti-inflammatory factors, that provide a protection mechanism against non-native microorganisms, as well as a balance between the microorganisms that make up the microbiota [2,4–8]. This balance can be altered by both internal factors (hormonal status, age, and the immune system) and external factors (use of antibiotics, infections, and microbial exposure to the environment), which favors the development of dysbiosis [2,4–8]. So far, five different microbial communities have been identified

that maintain a symbiotic relationship with the host and, in particular, *Candida* spp., are considered commensals of the vaginal mucosa and coexist as part of the complex vaginal ecosystem [1,2,9–14].

During pregnancy, physiological changes in the immune system, increased estrogen and progesterone levels, glycogen storage, low vaginal pH, and decreased cell-mediated immunity favor the development of *Candida* infection [9–14]. Vulvovaginal candidiasis during pregnancy has a high prevalence, which presents a wide geographical variation (Table 1), e.g, in Selangor in Malaysia they report 17.2%, Burkina Faso 22.71%, and Argentina 24.78%, are the countries with the lowest prevalence and, on the other hand those with the highest prevalence are the state of Enugu in Nigeria with 62.2%, Taif in Saudi Arabia 70.2% and Kenya 90.38%. In Latin America, a prevalence between 24.8 and 44.8% has been reported (Argentina and Brazil); as for Mexico there are no representative and recent studies on the prevalence of *Candida* in the general population, the latest report on its study in pregnancy dates from 2014 by Solís-Árias et al. where we found a prevalence of 12.6%. Vulvovaginal candidiasis, or *Candida* vaginitis, is caused primarily by *Candida albicans* in 85-95% of cases and the remainder by non-*albicans* *Candida* species (*C. glabrata*, *C. parapsilosis*, and *C. tropicalis*) [7,9–12,19,20].

Table 1. Prevalence of vulvovaginal candidiasis and/or positive vaginal candidiasis (from highest to lowest).

City, country	Prevalence (%)
Kenya	90.38
Taif, Saudi Arabia	70.2
Enugu State, Nigeria	62.2
Ibb, Yemen	61.5
Northwest Nigeria	60.8
Maroua, Cameroon	55.4
Sana'a, Yemen	51.6
Lebanon	44.8
Mato Grosso, Brazil	44.8
Natal, Brazil	44.8
South Libya	43.8
South Karnatka, India	42.37
Sarajevo, Bosnia and Herzegovina	40.9
Lebanon	39
Adana, Turkey	37.4
Middle Betl of Ghana	36.5
Southwestern Nigeria	36
Ardabil, Iran	35
Janakpur, Nepal	35
Tunisia	32.87
Ghana	30.7
Anambra State, Nigeria	30
Kathmandu, Nepal	29.5
Northwest Ethiopia	25
Argentina	24.8
Burkina Faso	22.71
Selangor, Malaysia	7.2

Vulvovaginal candidiasis can be classified as uncomplicated or complicated depending on factors including the severity of the infection, the yeast species, and the integrity of the immune system. Complicated vulvovaginal candidiasis is generally difficult to treat, making it necessary to administer more aggressive treatment regimens to cure it. For uncomplicated vaginal candidiasis, azole antifungals are the mainstay of treatment and are adequate to resolve *Candida* infections in most

cases. These medications come in a variety of formulations and can be administered orally or topically as vaginal creams, ointments, or suppositories. [7]

Azole antifungals (Table 2) are a group of medicines that contain an azole ring and inhibit the growth of a wide range of fungi. They are classified into two groups: those with two nitrogens in the azole ring (the imidazoles; examples include clotrimazole, econazole, ketoconazole, miconazole, and tioconazole) and those with three nitrogens in the azole ring (the triazoles; examples include fluconazole, itraconazole, posaconazole, and voriconazole). Azole antifungals work by inhibiting the cytochrome P450 dependent enzyme lanosterol 14- α -demethylase, which converts lanosterol to ergosterol, the main sterol in the fungal cell membrane. Depletion of ergosterol damages the cell membrane resulting in cell death.[21]

Rotem et al. evaluate the risk for major malformations following first-trimester exposure to vaginal azoles was not associated with either major or specific malformations according to organ systems.[22]

Table 2. Azole antifungals for Vulvovaginal candidiasis.

Drug	US FDA pregnancy category
Clotrimazole	B
Econazole	C
Ketoconazole	C
Miconazole	C
Tioconazole	Not Assigned
Fluconazole	D
Itraconazole	Not Assigned
Posaconazole	Not Assigned
Voriconazole	Not Assigned
Boric acid	Not Assigned
Terconazole	C
Nystatin	C
Oteseconazole	Not Assigned

US – United States; FDA - Food and Drug Administration.

For recurrent vulvovaginal candidiasis (three or more symptomatic episodes in a 12-month period), treatment options include fluconazole, itraconazole, miconazole, clotrimazole, terconazole, and nystatin, with the recent addition of oteseconazole and a vaccine directed at a hyphal virulence factor of *Candida albicans*. When vulvovaginal candidiasis is caused by non-*albicans Candida*, the recommended treatment options are fluconazole or boric acid. The greatest experience in the treatment of vulvovaginal candidiasis is with fluconazole, and a systematic review on congenital malformations and first trimester use of fluconazole found a potential association between general malformations (odds ratio [OR] 1.10, confidence interval of 95% [CI] 0.98–1.25), cardiac defects (OR 1.29, 95% CI 1.05-1.58), craniofacial defects (OR 1.25, 95% CI 0.88-1 .77) and abortion, although the rate of general malformations and craniofacial defects were not significant. Despite these findings, current guidelines state that only topical azole therapy should be used to treat vulvovaginal candidiasis during pregnancy.[7]

Several studies have shown the increase in the prevalence of *Candida* during pregnancy, which can alter the vaginal microbiota, causing an increase in proinflammatory cytokines with induction of the inflammatory cascade and generate adverse pregnancy outcomes such as preterm labor, premature rupture of membranes, preeclampsia, spontaneous abortion, restriction in the fetal growth, low birth weight, fetal death and neonatal sepsis. On the other hand, a positive culture result not associated with symptoms is considered a sign of colonization and not of infection, and therefore does not generate any consequences. The aim of this study is to contribute to identifying whether there is an association between colonization by *Candida* and adverse perinatal outcomes. [3,8–11,16,23–27].

2. Materials and Methods

2.1. Study design

Descriptive, observational, prospective study that included pregnant patients without signs or symptoms of vaginal infection, and without antimicrobial treatment, who attended the Hospital Regional de Alta Especialidad de Ixtapaluca (HRAEI) obstetric triage service for a period of six months (May to October 2019), agreed to participate in the study and were followed up until the end of pregnancy.

2.2. Cervicovaginal sampling

Samples of cervicovaginal exudate (posterior cul-de-sac and cervix) were taken with a sterile swab, after speculum placement. Vaginal samples were immediately inoculated into Petri dishes with Sabouraud agar and incubated at 37°C for 24-72h.

From the primary isolations, a colony was taken, reseeded on Sabouraud agar plates and incubated at 37°C for 24h, for subsequent identification.

2.3. Phenotypic identification of *Candida* spp.

Initially, the yeasts were identified by germ tube production, through the inoculation of the yeasts in 500 µL of human serum and incubation at 37 °C for 2 h. Subsequently, 20 µL of the serum inoculated with *Candida* were placed between slides and coverslips and microscopic observation (40X) was performed to search for the germ tube. The yeasts that presented a germ tube were considered *C. albicans*, and those that did not develop a germ tube were considered *C. non-albicans*.

2.4. Genotypic identification of *Candida* spp.

DNA was extracted from each of the vaginal isolates using the commercial Yeast DNA Preparation kit (Jena Bioscience, GE), following the instructions by the company. The DNA was amplified with the oligonucleotides CandF (5'-AGCTTGCGTTGATTACGTCCCTGCCC-3') and CandR (5'-TTCACCTCGCCGCTACTAAGGCAATCCC-3'), which identify, based on the size of the amplicon, *C. albicans* (850 bp), *C. glabrata* (1000 bp), *C. tropicalis* (790 bp), *C. parapsilosis* (731 bp), *C. krusei* (800 bp), *C. guilliermondii* (1100 bp), *C. lusitaniae* (590 bp) and *C. dubliniensis* (810 bp)[28]. As positive controls, DNAs from the reference strains of *C. albicans* ATCC 10231 and *C. glabrata* ATCC 90030 were used. [27]

2.5. Perinatal results

The digital record of the patients with positive results for *Candida* was reviewed to see the maternal, fetal, and neonatal results at the end of the pregnancy (preterm delivery, premature rupture of membranes, small for gestational age, and perinatal mortality).

2.6. Statistic analysis

The possible associations between vaginal colonization by *Candida* and perinatal complications were determined using Fisher's test (95% CI, p<0.05) using the GraphPad Prism 8.0 statistics software.

3. Results

During the study period, 981 pregnant patients were attended to in the HRAEI obstetric triage; 385 patients were admitted to the hospital, 112 for control of some maternal or fetal complication and 273 for termination of pregnancy, 15 patients were sent to another less complex unit for termination of pregnancy, of the remaining 581 patients, 221 attended for symptoms of vaginal infection, 58 for urinary tract infection, and 302 for non-urgent care, of which only 83 patients out of them met the inclusion criteria and agreed to participate in the study. The age of the patients was between 15 and

45 years old, with a gestational age of 13.2 to 42 weeks. The number of gestation was from 1 to 5. The BMI of the patients ranged from 18.5 to 44.8 Kg/m² (Table 3).

Table 3. Data of the asymptomatic pregnant patients who attended the obstetric triage at the Hospital Regional de Alta Especialidad de Ixtapaluca in a period of six months.

Item No.	Age (years)	Pregnancy	Gestational age (weeks)	BMI	Culture
1	18	2	40	27.2	bacteria
2	19	1	33.6	26.1	bacteria
3	16	1	22.6	22.8	bacteria
4	26	4	25.4	21.6	bacteria
5	32	1	39.1	29.7	bacteria
6	18	1	40.5	31.4	ND
7	15	1	39	27.5	yeast
8	24	1	32	34.9	bacteria
9	26	4	25.6	21.6	bacteria
10	25	2	33.5	22.8	ND
11	45	3	41.2	25.9	ND
12	17	1	38.3	27.4	yeast
13	22	2	18.6	20.9	bacteria
14	35	3	33.5	22.2	bacteria
15	16	1	39.5	27	bacteria
16	15	2	35.3	23.8	bacteria
17	31	1	40.3	31.7	bacteria
18	20	2	30.3	21.4	bacteria
19	18	3	35.1	25.6	ND
20	24	3	40.3	24.3	bacteria
21	17	2	38.2	30.4	yeast
22	22	1	22	38.2	bacteria
23	29	3	40.4	32.1	yeast
24	25	3	33	33	ND
25	30	3	38.6	29.5	ND
26	27	2	34.3	26	yeast
27	23	1	34.2	30	bacteria
28	32	2	24.4	25.7	ND
29	19	1	38.3	30	yeast
30	17	1	35.4	26.6	bacteria
31	41	4	31.6	38.4	bacteria
32	31	3	30.2	44.8	bacteria
33	27	2	41.5	33.3	ND
34	21	1	36.6	23.2	ND
35	34	3	38.6	34.1	yeast
36	36	5	18.6	32.6	bacteria
37	37	3	20	26.6	bacteria
38	18	1	19	37.2	yeast
39	22	2	39.5	34.3	ND
40	22	3	37.6	25.5	bacteria
41	28	2	16.1	26.2	bacteria
42	27	3	25.1	19	ND
43	36	2	36.3	34.1	yeast
44	35	3	29.6	29.4	bacteria
45	29	3	36.4	28.7	yeast

46	21	1	38.4	23.3	ND
47	23	2	22.6	21.3	bacteria
48	35	1	37.2	27.8	bacteria
49	24	3	25.2	27.2	ND
50	22	1	33.5	31	ND
51	35	4	26.4	21.9	bacteria
52	29	2	40.6	34.2	bacteria
53	18	1	15.2	26.6	bacteria
54	34	4	36.6	42.4	yeast
55	21	1	33	26.4	ND
56	41	5	39	32	ND
57	28	2	38.2	32.1	ND
58	30	4	42	33.3	ND
59	32	3	33.3	32.5	bacteria
60	32	5	39.5	30	bacteria
61	25	3	38.2	31.8	yeast
62	32	3	38.6	23.6	bacteria
63	21	3	39.6	28.1	bacteria
64	23	2	35.4	26.2	bacteria
65	20	1	41	25.7	bacteria
66	34	4	25.2	28.9	bacteria
67	21	2	38.1	25	yeast
68	21	1	39.5	29.6	bacteria
69	18	1	13.2	18.5	bacteria
70	18	2	34.4	24.1	yeast
71	16	1	37.1	27.3	bacteria
72	17	1	26.3	19.3	yeast
73	31	3	39.3	33.8	bacteria
74	35	3	37.6	25	yeast
75	31	2	29	37.1	bacteria
76	33	4	39.5	36.8	ND
77	28	2	39.4	32.4	bacteria
78	31	4	38.1	21.1	ND
79	24	2	36.4	30.8	ND
80	24	2	39.4	26.3	ND
81	33	4	20	22.2	bacteria
82	23	2	38.6	34.5	bacteria
83	29	4	37.5	22.2	bacteria

ND – No development.

Out of the cultures of the 83 cervicovaginal samples, bacterial colonies were observed in 45 (54.2%), yeast-like colonies developed in 16 (19.3%) (7, 12, 21, 23, 26, 29, 35, 38, 43, 45, 54, 61, 67, 70, 72, 74), and in 22 (26.5%) no development of microorganisms was observed (Table 3). Regarding the 16 yeast isolates, 12 (75.0%) developed germ tubes (7, 12, 21, 23, 26, 29, 35, 43, 45, 61, 72, 74), suggestive of the species *C. albicans*, while 4 (25%) corresponded to *Candida non-albicans* (Figure 1a). Similarly, amplification with *Cand* oligonucleotides generated an 850 bp amplicon in 12 samples, while an amplicon of 1000 bp was obtained in the four remaining samples, which corresponds to the identification of the species *C. albicans* and *C. glabrata*, respectively (Figure 1b).

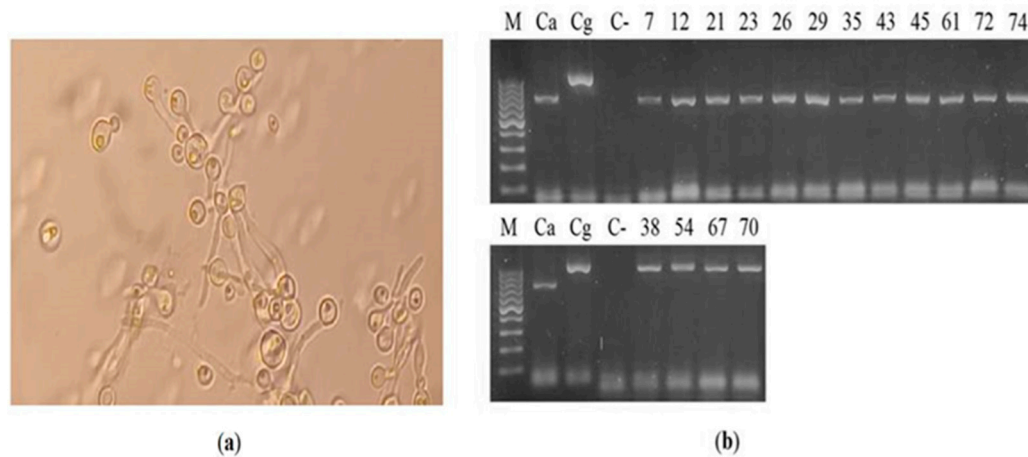


Figure 1. (a) Positive germ tube observed under a microscope (40X), corresponding to the yeast isolated from sample 7. (b) Electrophoresis gel revealed under UV light, where the amplification, by PCR, of the DNA of the yeast strains is shown. reference Ca (*Candida albicans* ATCC 10231) and Cg (*Candida glabrata* ATCC 90030), M: 100 bp molecular size marker; C-: negative control.

Regarding antimicrobial susceptibility (Table 4), eleven of the samples positive for *C. albicans* are sensitive to fluconazole and only one was resistant; all the samples positive for *C. glabrata* were resistant to fluconazole.

According to the antecedents in the clinical history, all the patients presented uncomplicated vulvovaginal candidiasis, there were no recurrent or complicated ones.

Table 4. Data of the asymptomatic pregnant patients with yeast-like colonies developed and antimicrobial susceptibility.

Item No.	Genotypic identification	Antimicrobial susceptibility (US FDA pregnancy category)					
		Caspofungin (NA)	Micafungin (NA)	Amphotericin b (B)	Voriconazole (NA)	Flucytosine (NA)	Fluconazole (D)
7	<i>C. albicans</i>	S	S	S	S	S	S
12	<i>C. albicans</i>	S	S	S	S	S	S
21	<i>C. glabrata</i>	S	S	S	S	R	R
23	<i>C. albicans</i>	S	S	S	S	S	S
26	<i>C. albicans</i>	S	S	R	S	S	R
29	<i>C. glabrata</i>	S	S	S	S	S	R
35	<i>C. albicans</i>	S	S	S	S	R	S
38	<i>C. albicans</i>	S	S	S	S	S	S
43	<i>C. albicans</i>	S	S	S	S	S	S
45	<i>C. glabrata</i>	S	S	S	S	S	R
54	<i>C. albicans</i>	S	S	S	S	S	S
61	<i>C. glabrata</i>	S	S	S	S	S	R
67	<i>C. albicans</i>	S	S	S	S	S	S
70	<i>C. albicans</i>	S	S	S	S	S	S
72	<i>C. albicans</i>	S	S	S	S	S	S
74	<i>C. albicans</i>	S	S	S	S	S	S

NA – not assigned; S – Sensitive; R – Resistant.

The perinatal results of the positive group for *Candida* and of the patients who had no development in the cultures for preterm delivery, premature rupture of membranes, small for gestational age, and perinatal mortality are described in Tables 5 and 6.

Table 5. Characteristics of the patients with positive culture for *Candida* vs. without development.

	With <i>Candida</i>	No development	<i>p</i>
No.	16	22	>0.9999
Average age (years)	24	27	0.0067
< 19 years	7	2	0.0052
> 35 years	2	2	0.2809
Average pregnancy	2	2	>0.9999
Initial average gestational age (weeks)	36	36	>0.9999
Final average gestational age(weeks)	39	40	>0.9999
Average BMI	30	28	0.2431
Overweight	4	9	0.2031
Obesity	7	10	0.4110

Table 6. Characteristics of the patients with positive culture for *Candida* vs. without development.

	With <i>Candida</i>	No development	<i>p</i>
No.	16	22	
Premature rupture of membranes	0	1	>0.9999
Corioamnionitis	1	0	0.4211
Preterm	0	0	>0.9999
Small for gestational age	2	2	>0.9999
Perinatal mortality	0	0	>0.9999

In the group of patients with *Candida*, the maternal age range was from 15 to 36 years old, of which 7 were adolescents and 2 with advanced maternal age, with an average of 2 gestations, the range of body mass index was from 19.3 to 42.4, of which 4 were overweight and 7 obese. Regarding perinatal complications, one presented chorioamnionitis and 2 neonates were small for gestational age. The patients who had small neonates had a history with epilepsy and the other with anemia.

In the patients that did not develop in the cultures, the age range was from 18 to 45 years old (2 adolescents and 2 with advanced maternal age), with 2 pregnancies on average, the range of body mass index was from 19 to 36.8 (7 overweight and 10 obese), of the perinatal complications, one presented premature rupture of membranes and two small neonates for gestational age. No data were found in the medical record of the patients with a small-for-gestational-age fetus that could be associated with this outcome.

4. Discussion

Epidemiological reviews have shown that at least 75% of all women will present a picture of vulvovaginal candidiasis (VVC) during their lives and that it will be associated with pregnancy between 9 and 55%; in this study the frequency of colonization was 19.3 % [6,24].

Regarding the etiology, the species *C. albicans* is the most frequent; however, other emerging species such as *C. glabrata*, *C. parapsilosis* and *C. tropicalis* may be present, in this work 75% were *C. albicans* and 25% *C. non-albicans*. [6,24]

Traditionally, it was considered that the fetoplacental unit was free of germs and that the newborn's first exposure to microorganisms occurred during delivery, and that any microbial growth in the amniotic fluid or placenta originated in the lower genital tract, leading to poor perinatal outcomes (preterm labor, congenital cutaneous candidiasis, neonatal candidiasis, chorioamnionitis, and premature delivery), although it has recently been shown that the bacteria break through the intact

maternal-fetal membranes without causing any consequences. When analyzing adverse perinatal outcomes in this study, a higher frequency was not identified in patients with VVC, compared to healthy pregnant women, which is consistent with the systematic review and meta-analysis by Schuster et al. and may be explained by the fact that complex changes in the maternal immune system protect the pairing (mother-fetus) from infections by favoring the development of fetal immunity, avoiding rejection of the fetus and its attachments by the mother. In addition, the microbiome of the maternal genital tract regulates this immunitary behavior, generating a greater tolerance to microorganisms, through an increase in anti-inflammatory cytokines, the initiation of tolerance to endotoxins, and the suppression of autophagy, which leads to a downward modulation of the immune response or as referred by Farr et al. and Sust et al. that a positive result in the culture is a sign of colonization but not of infection and therefore it is asymptomatic (He et al.), so that it is important to differentiate between colonization and infection. Colonization has been described as the presence of *Candida* in the intact vaginal tract of immunocompetent individuals that are generally unrelated to symptoms and when colonization progresses to infection frequently manifests by vaginal itching, burning, pain, dysuria, flushing, dyspareunia, and vaginal discharge (Chatzivasileiou and Vyzantiadis). On the other hand, it is known that any alteration in the vaginal microbiota can lead to an increase in proinflammatory cytokines with induction of the inflammatory cascade and adverse pregnancy outcomes such as preterm labor, premature rupture of membranes, preeclampsia, spontaneous abortion, restriction in the fetal growth, low birth weight, fetal death and neonatal sepsis [6,10,12,13,18,19,23,24,28,29].

As a result of the reduced estrogenization of the vagina at the extremes of life (before menarche and postmenopause, in the absence of hormone replacement therapy), the probability of colonization by *Candida* is lower. In this study, it is striking that the frequency of VVC in adolescents (43.7%) versus healthy patients (9%), probably due to high estrogen levels that generate an increase in glycogen content in vaginal fluid and contrary to what was reported by Chatzivasileiou - Vyzantiadis, Sasani et al. and Bender et al., where adolescence is associated with a high resistance to vaginal *Candida* infection despite the fact that sexual activity in this age group predisposes to a higher vaginal fungal load and that the age range with the highest number of infected is between 25 and 34 years of age. [10,12,13,18,19,23,24].

Even though the Center for Disease Control (CDC) recommendations for first-line treatment for vulvovaginal candidiasis is fluconazole, in our study, *C. glabrata* species were not sensitive to this antifungal.[7]

Finally, as a secondary analysis, the number of patients who tested positive for bacteria (54.2%) draws attention; this will be the subject of another study, considering the same perinatal complications, since it is described that the presence of *C. albicans* in the vagina can generate dysbiosis due to the presence of virulence factors, as well as the production of proteolytic enzymes that alter vaginal immunity [1,18,23,29].

The limitation of this study for the recruitment of a greater number of patients is related to the fact that the hospital, due to its level of complexity, can only monitor high-risk pregnant women.

5. Conclusions

No association was found between colonization by *Candida* spp. and the number of pregnancies, gestational age, body mass index or perinatal results. However, it was found that age (adolescence) may be a risk factor for the development of vulvovaginitis during pregnancy. Fluconazole is a good therapeutic option for vulvovaginal candidiasis caused by *C. albicans* and not for those caused by *C. glabrata*, in our study population.

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