



Vaginal Infections in Pregnant Women at the Bè Hospital in Lomé (Togo) from 2008 to 2013

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Abstract

Sexually transmitted infections remain a public health problem almost all around the world. Women of childbearing age frequently have vulvovaginitis. The aim of this work is to evaluate the prevalence of the main germs responsible for vaginal infections in pregnant women who have had their vaginal swabs taken at Bè hospital. We conducted a retrospective study between May 2008 and December 2013, on 126 pregnant women in whom a vaginal swab was performed. We vaginally sampled 126 pregnant women, of whom 90 or 71.43% had vaginal infections. *Candida albicans* was the most frequently isolated species (47.97%) followed by *Gardnerella vaginalis* (38.21%). The most affected age group was between 25 and 30 years old. *Candida albicans* were respectively resistant to miconazole, clotrimazole and econazole. Our study showed that pregnant women are highly exposed to vaginal infections with a predominance of *Candida albicans* and *Gardnerella vaginalis*. This shows that it is necessary to do at least one culture of vaginal swabs during pregnancy for each woman.

Subject Areas

Microbiology

Keywords

Vaginal Infection, Bacterial Vaginosis, Vaginitis Infectious, Pregnancy, Togo

1. Introduction

The vaginal flora of healthy women is made up of a wide variety of anaerobic

and aerobic bacteria. The most represented species is the kind *Lactobacillus* (Döderlein Bacillus) [1]. Lactobacilli are involved in maintaining the natural balance of the vaginal flora, and thus protect this flora. This role is especially important during pregnancy, because of vaginal abnormalities such as bacterial vaginosis and vaginitis infectious [2]. These abnormalities have been described as mechanisms responsible for premature births and perinatal complications [3].

We talk about bacterial vaginosis when the normal balance of bacteria in the vagina is disrupted and replaced by a proliferation of pathogenic bacteria. Lactobacilli are replaced by anaerobic microorganisms such as *Gardnerella vaginalis* and *Mycoplasma hominis* [4] [5]. Bacterial vaginosis is known to be a significant risk factor for adverse pregnancy outcomes [5].

Infectious vaginitis is defined as an infection related to a pathogen; it is also an abnormal manifestation of a bacterium, a parasite or a fungus that usually occurs in minute quantities. These are infections with *Gardnerella* [4], *Neisseria gonorrhoeae*, *Trichomonas vaginalis* [6] [7] [8] but also fungal infections. The fungi responsible for vaginitis are mainly caused by yeasts of the genus *Candida*; in this case we are talking about vaginal candidiasis.

We have superficial or local candidiasis and deep or visceral candidiasis. Superficial candidiasis comes when we have the passage of yeasts from the commensal state to the pathological state; deep candidiasis is defined by the involvement of at least one deep organ.

In Togo, a similar study was conducted in Sokodé in the central region in 2011 [9].

The main objective of our study was to determine the prevalence of vaginal infections in pregnant women in a secondary hospital in Lomé.

2. Material and Methods

This is a retrospective descriptive study carried out from the registers of the gynecological services and the registers of the microbiology laboratory of the Bè hospital. It covers a period of 5 years from May 2008 to December 2013. This hospital includes the services of gynecology-obstetrics, pediatrics, general medicine, ophthalmology, stomatology, radiology, social service, and laboratory of biology, service of maintenance, hygiene and vaccination. The inclusion criteria are all registrations of pregnant women who came to the laboratory during the study period and the non-inclusion criteria are all incomplete registrations during the same period. 126 pregnant women were selected for this study.

To perform the vaginal samples, the patient goes to the laboratory without personal hygiene, suspend any antibiotic 48 hours rather, as well as sex 24 hours instead.

In the laboratory, the sample is made with three swabs; a swab is used to take the sample from the endocervix and two other swabs are used to take the sample from the vagina. One of the vaginal swabs is used to make fresh condition for looking for clues-cells, *Trichomonas vaginalis*, red and white blood cells, and

yeasts.

Gram stain is made from the other vaginal swab and the endocard swab. The Gram stain makes it possible to look for Gram diplococci on the one hand intracellular and extracellular (gonococcal) negatives, and intracellular and extracellular Gram-variable coccobacilli (*Gardnerella vaginalis*), Gram-negative intracellular and extracellular Gram-negative bacilli (*Mobiluncus* spp.), gram-positive bacilli of varying length (Bacillus Doderlein), Gram-positive, Corynebacterium-type bacilli, other bacteria (Gram-negative bacilli and Gram-positive cocci), and the presence of yeast and mycelial pseudo-filaments [10].

Smears were classified and interpreted according to the Nugent score [11] as shown in **Table 1** and **Table 2** respectively [12].

Seeding is systematic on Thayer Martin (TM) agar and Sabouraud chloramphenicol agar. Gram directs the choice of bromo-cresol-purple agar (BCP), nalidixic acid agar and Eosine Methylene Blue agar (EMB).

The media were incubated for 24 hours at 37°C and under CO₂ for Thayer-Martin agar. On each medium, suspicious colonies were identified and their identification was carried out according to the classic bacteriological characters (catalase, appearance of colonies with a color shift of the media, metallic reflection) and by biochemical characters using mini-galleries (Kligler-Hajna, Simmons citrate, mannitol/mobility and urea/indole). The antibiotic susceptibility of the different strains isolated was determined by the standard antibiogram antimicrobial diffusion method on agar medium, according to the recommendations of the antibiogram committee of the French Society of Microbiology [13] [14].

Table 1. Nugent's scoring of vaginal swabs (number based on average of 10 fields).

| Morphotype | Number of organisms per oil immersion field | | | | |
|---|---|----|-------|--------|-----|
| | None | <1 | 1 - 4 | 5 - 30 | >30 |
| <i>Lactobacillus</i> species | 4 | 3 | 2 | 1 | 0 |
| <i>Gardnerella</i> & anaerobic Gram-negative bacilli | 0 | 1 | 2 | 3 | 4 |
| Curved Gram-negative bacilli (<i>Mobiluncus</i> species) | 0 | 1 | 1 | 2 | 2 |

Table 2. Interpretation of nugent's score.

| Nugent's score | And | Interpretation |
|----------------|-----------------------------|---|
| 0 - 3 | No clue cells | Normal vaginal flora |
| 4 - 6 | No clue cells | Intermediate or Not consistent with Bacterial vaginosis |
| 4 - 6 | Clue cell present | Indicative of bacterial vaginosis |
| ≥7 | Clue cell present or absent | Indicative of bacterial vaginosis |

Statistical analyzes.

The statistical software SPSS version 22 for Windows and MedCal 13 were used for data processing and comparison of proportions. Chi-square was used and the statistical threshold was set at $P < 0.05$.

3. Result

The retrospective analysis of the data identified a total of 126 pregnant women, who performed their consultation at the Department of Gynecology at Bè Hospital and their vaginal swabs in the microbiology laboratory of the same hospital.

The age group was 12 to 41 years old with an average age of 28.10 years. Their age distribution is shown in **Table 3**.

We recorded 62 women (49.21%) with scanty leucorrhoea (**Table 4**). Of the women recruited, 90 (71.42%) had an infection (**Figure 1**). *Candida albicans* was the most isolated with 47.97% followed by *Gardnerella vaginalis* (47.97%); *Trichomonas vaginalis* was poorly recovered (4.06%) (**Table 5**). The distribution of isolated organisms by age is shown in **Table 6**. *Candida albicans* is isolated in 6.5% of women aged 35 to 40, followed by *Gardnerella vaginalis* 4% and *Trichomonas vaginalis* 2%. The 25 to 30 age group has the highest rate of isolation germ and is respectively 16%, 14%, 4% and 1% for *Candida albicans*, *Gardnerella vaginalis*, *Mobiluncus* spp. and *Trichomonas vaginalis*.

Strains of *Candida albicans* were 100% resistant to the following antifungals: miconazol, clotrimazol and econazol.

Table 3. Distribution of pregnant women by age.

| Age (years) | Effectif | Percentage |
|--------------|------------|-------------|
| [10 - 15] | 1 | 0.79% |
| [15 - 20] | 5 | 3.97% |
| [20 - 25] | 25 | 19.84% |
| [25 - 30] | 46 | 36.51% |
| [30 - 35] | 31 | 24.60% |
| [35 - 40] | 16 | 12.70% |
| [40 - 45] | 2 | 1.59% |
| Total | 126 | 100% |

Table 4. Classification of leucorrhoea abundance.

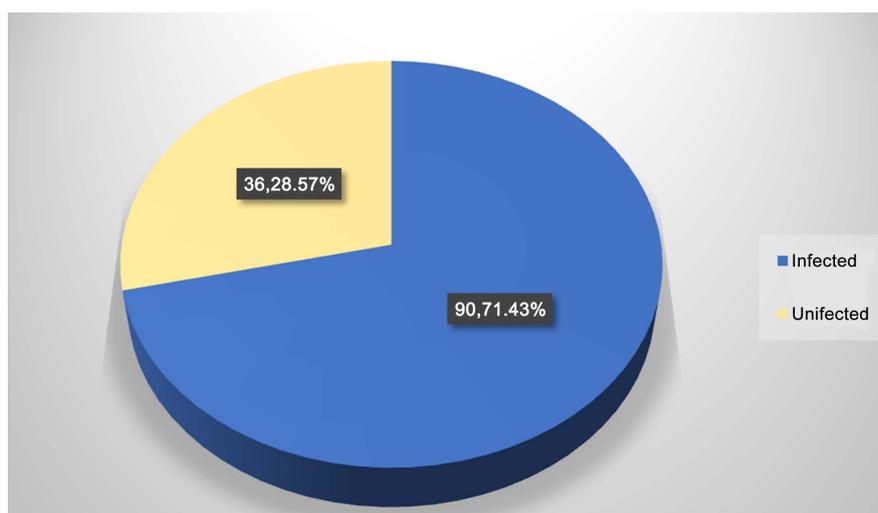
| Leucorrhoea abundance | Effectif | Percentage |
|-----------------------|------------|-------------|
| Scanty | 62 | 49.21% |
| Abundant | 47 | 37.30% |
| Very abundant | 17 | 13.49% |
| Total | 126 | 100% |

Table 5. Distribution of microorganisms found.

| Microorganismes | Effectif | Percentage |
|---------------------------------|------------|-------------|
| <i>Trichomonas intestinalis</i> | 5 | 4.06% |
| <i>Candida</i> spp. | 6 | 4.88% |
| <i>Mobiluncus</i> spp. | 6 | 4.88% |
| <i>Gardnerella vaginalis</i> | 47 | 38.21% |
| <i>Candida albicans</i> | 59 | 47.97% |
| Total | 123 | 100% |

Table 6. Distribution of microorganisms by age group.

| Age (years) | [10 - 15] | [15 - 20] | [20 - 25] | [25 - 30] | [30 - 35] | [35 - 40] | [40 - 45] | Total |
|------------------------------|-----------|-----------|-------------|-------------|-------------|-------------|-----------|-------------|
| <i>Candida albicans</i> | 0 | 2 (1.63%) | 13 (10.57%) | 20 (16.26%) | 16 (13.00%) | 8 (6.50%) | 0 | 59 (47.96%) |
| <i>Gardnerella vaginalis</i> | 0 | 0 | 10 (8.13%) | 18 (14.63%) | 14 (11.38%) | 5 (4.07%) | 0 | 47 (38.21%) |
| <i>Mobiluncus</i> spp. | 0 | 0 | 1 (0.81%) | 5 (4.07%) | 0 | 0 | 0 | 6 (4.88%) |
| <i>Candida</i> spp. | 0 | 0 | 1 (0.81%) | 2 (1.63%) | 1 (0.81%) | 2 (1.63%) | 0 | 6 (4.88%) |
| <i>Trichomonas vaginalis</i> | 0 | 0 | 0 | 2 (1.63%) | 0 | 3 (2.44%) | 0 | 5 (4.07%) |
| Total | 0 | 2 (1.63%) | 25 (20.32%) | 47 (38.22%) | 31 (25.19%) | 18 (14.64%) | 0 | 123 (100%) |

**Figure 1.** Prevalence of vaginal infection.

4. Discussion

In this study conducted between May 2008 and December 2013, we recorded 126 vaginal specimens in the Bè hospital, in pregnant women who met the criteria for inclusion and non-inclusion. Nadembega found 118 cases [15], which is in agreement with our data; however, other studies were performed with a larger sample size [9] [16] [17] [18] [19]. A large sample size would reflect the true proportions of the population conditions.

The age of the study population varies between 12 and 41 years, an average of

28.10 years; this is an average close to that obtained in a similar study conducted by Tchelougou in the central region of Togo in 2011 [9].

The most exposed age group is between 20 and 35 years old; it is the age of procreation and during which the sexual activity is intense; this would explain this high rate. This is the most represented age group. Unlike us, Benchellal in 2011 found an age range of 25 to 53 years in Morocco [20] while Anane found 20 to 39 years in 2010 in Tunisia [21] which is close to our results.

Leucorrhoea is often abundant among women in our study in the maritime region of Togo; this result is similar to that of Tchelougou (45.37%) in the central region of the country [9]; we noticed a significant difference depending on the character “scanty” (6.95%) and “very abundant” (45.37%). This difference can be explained by the social and cultural diversities and behaviors of populations [22] [23] [24] [25] that could affect their physiology.

The percentage of vaginal contamination observed in the study population is 71.42%. These results confirm those obtained by Tchelougou [9] in 2011 in Sokodé (73.18%).

Gardnerella vaginalis and *Candida albicans* account for most laboratory-identified germs (86.18%) during the study period; this result is similar to that of Tchelougou [9] in 2011 which is 86.08%.

There is a high prevalence of *Candida albicans* (47.97%) among pregnant women compared to the works of Tchelougou in 2011 in Sokodé [9]. *Candida albicans* can increase premature delivery rates and decrease neonatal birth weight [18], hence the need to follow women during pregnancy.

According to Benchellal, the risk factors involved in the occurrence of vulvovaginal candidiasis are: pregnancy, poor hygiene habits and frequent intimate toilet [20]. In fact, there is a marshy area, so that the water used by the population of Bè could be wet; This could also explain the high rate of vaginal infection in the area.

We found that *Mobiluncusspp* had a rate of 4.88%; this result is not similar to that found by Tchelougou in 2011 [9]. *Candida* spp. represented 4.88%; this result is similar to that of Cravello [26] in 2001, which reports that in mycotic vulvovaginitis, *Candida albicans* occupies 9 out of 10 cases, compared to 1 out of 10 cases for *Candida* spp. *Trichomonas intestinalis* (4.06%) has a similar rate to that found in 2011 by Tchelougou (3.66%) in the central region [9], unlike us Nadembega did not find *Trichomonas intestinalis* [15].

In this study, we noted that *Candida albicans* was resistant to miconazol, clotrimazol and econazol. This resistance could be explained by the use of unknown products for intimate toilets. This resistance is at the origin of the therapeutic failure. Nadembega gets results different from ours [15].

5. Conclusion

From the analysis of the retrospective study carried out between May 2008 and December 2013, it appears that the rate of vaginal infection is 71.42%; *Candida*

albicans and *Gardnerella vaginalis* were the most isolated respectively with a frequency of 47.97% and 38.21%. The most exposed age group was 20 to 35 years old. All strains of *Candida albicans* isolated are resistant to Miconazol, Clotrimazol and Econazol. These preliminary results provide information on the prevalence of vaginal infections and on the susceptibility profile of isolated *Candida albicans* strains. However, it would be desirable to continue the studies over a much larger period and on a larger sample size for a better appreciation of the frequency of isolated organisms.

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