



An update on the role of *Atopobium vaginae* in bacterial vaginosis: what to consider when choosing a treatment? A mini review

Werner Mendling¹ · Ana Palmeira-de-Oliveira^{2,3} · Stephan Biber⁴ · Valdas Prasauskas⁴

Received: 23 January 2019 / Accepted: 26 March 2019 / Published online: 5 April 2019
© The Author(s) 2019

Abstract

Introduction *Bacterial vaginosis* (BV) is the most common vaginal disorder in reproductive-age women. The condition is characterised by the replacement of a healthy, lactobacilli-dominated vaginal microbiota by anaerobic and facultative anaerobic bacteria. BV increases the risk of acquisition of STIs and is associated with pregnancy complications. Although the composition of the bacteria in BV varies between individuals, there are some species such as *Gardnerella*, *Atopobium*, *Mycoplasma*, *Sneathia*, *Megasphaera*, *Dialister*, etc., that are found most frequently.

Material and Methods Literature research to the importance of *Atopobium vaginae* in BV and treatment options.

Results *Atopobium* (*A.*) *vaginae* is an important component of the complex abnormal vaginal flora in BV; even though *A. vaginae*, like *Gardnerella vaginalis*, has also been detected in the normal flora, it is much more common in BV patients. *A. vaginae* has been shown to play an important role in the pathophysiology of BV and is thought to be at least a partial cause of the known negative sequelae. The presence of *A. vaginae* in the BV-associated biofilms and its resistance to some antimicrobial substances has been described - this seems to have a major impact on treatment outcome.

Conclusion Current scientific data demonstrate that dequalinium chloride (Fluomycin[®]) is one of the valid therapeutic options for BV treatment, since it displays a broad antimicrobial spectrum against relevant vaginal pathogens, especially against *G. vaginalis* and *A. vaginae*, without having safety concerns.

Keywords Bacterial vaginosis · Bacterial biofilm · *Atopobium vaginae* · Metronidazole · Clindamycin · Dequalinium chloride · Microbial resistance

Introduction

Bacterial vaginosis (BV) is the most common vaginal disorder in reproductive-age women [1]. The condition is characterised by the replacement of a lactobacilli-dominated vaginal microbiota by anaerobic and facultative anaerobic bacteria. It is still unknown whether the loss of lactobacilli precedes or follows the upheaval of flora in BV [2].

However, it seems to be quite certain that it is a preferential proliferation of the BV-related bacteria, rather than an exogenous acquisition [2].

Only some women with BV have symptoms, displaying a malodorous, watery, grey discharge. Because of the absence of inflammation, BV is not called vaginitis—there is no pain, no itching, no dyspareunia, no redness of the vulva or vagina, and no toxic leucocytes—only a microbial shift to anaerobic pathogens [3]. However, BV increases the risk of acquisition and transmission of STIs [4] and is associated with adverse obstetric and gynaecologic outcomes including miscarriage, premature labour, preterm birth, preterm prelabour rupture of membranes, chorioamnionitis, intrauterine infection, post-caesarean endometritis, upper genital tract infections, and pelvic inflammatory disease [5, 6].

Although the composition of the bacteria in BV varies between individuals, *Gardnerella*, *Atopobium*, *Mycoplasma*, *Prevotella*, *Bifidobacterium*, *Megasphaera*, *Lep-totrichia*, *Sneathia*, *Dialister*, *Clostridium*, and Bacterial

✉ Werner Mendling
w.mendling@t-online.de

¹ German Center for Infections in Obstetrics and Gynaecology, Wuppertal, Germany

² Labfit-HPRD: Health Products Research and Development, Lda, Covilhã, Portugal

³ CICS-UBI: Health Sciences Research Centre, University of Beira Interior, Covilhã, Portugal

⁴ Scientific and Medical Department, Medinova AG, Zurich, Switzerland

Vaginosis-Associated Bacterium (BVAB)-1, -2 and -3 species are found most frequently [7, 8]. The association of *G. vaginalis* with BV was originally described by Gardner and Dukes already in 1955 [9]. *G. vaginalis* and *Prevotella spp.* are found in the disturbed vaginal microbiome, but they are also present in lower loads in healthy women [7, 10]. However, the involvement of *A. vaginae* in BV, although it rarely occurs in the absence of *G. vaginalis*, has only been established in recent years [11–13]. Increasing evidence on its involvement in BV biofilm formation, as well as on specific resistances of *A. vaginae* against standard antibiotics may explain therapeutic failures and recurrences of BV [14].

Despite the current knowledge on BV many questions regarding treatment remain unanswered—all current therapies have disadvantages and gaps [14]. This mini review describes the current scientific knowledge regarding the relevance of *A. vaginae* in the pathogenesis of BV and what should be considered when choosing a treatment.

Role of *A. vaginae* in BV

Atopobium vaginae is a newly discovered bacterium frequently found in women with BV [15]. The name *Atopobium*, meaning “strange living thing” in Greek, was proposed in 1992 [16] to reclassify three bacterial species formerly designated *Lactobacillus minutus*, *Lactobacillus rimae* and *Streptococcus parvulus*. Genus *Atopobium* belongs to the *Coriobacteriaceae* family and *A. minutum*, *A. rimae*, *A. parvulum*, and, later described *A. deltae* and *A. fossor*, can be distinguished [15]. In 1999 Rodriguez et al. [17] first described *A. vaginae* isolated from the vagina of a healthy women in Sweden. These are Gram-positive, elliptical or rod-shaped cocci, nonmotile and non-spore-forming organisms, and occur alone, in pairs, in clumps or in short chains (Fig. 1). They produce major amounts of lactic acid next to acetic and formic acids and are strictly anaerobic.

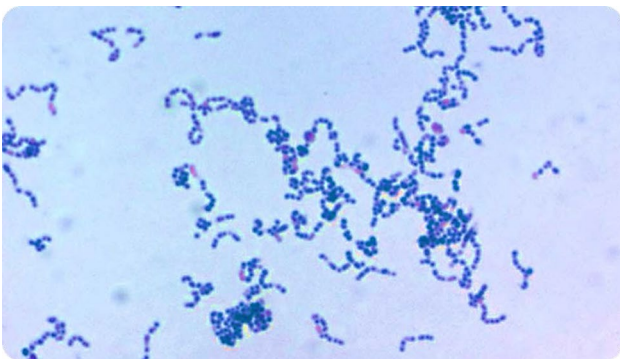


Fig. 1 *Atopobium vaginae* culture, Gram stain, magnification 100 × (courtesy of M. Vaneechoutte, Belgium)

Only recently, the association of *A. vaginae* with BV was reported [11, 12] and many subsequent studies have confirmed the bacterium to be an important component of the complex abnormal vaginal flora in BV [11, 12, 18–21]. Even though *A. vaginae*, as *G. vaginalis*, has also been detected in the normal vaginal flora (8% [11] up to 25% [19]), it is found much more commonly in BV patients (50% [12], 55% [11], 83% [18], 96% [19]). Also, the differences between African and Caucasian women, referring the prevalence of *A. vaginae* in the first ones have been described [22]. Loads of *G. vaginalis*, *A. vaginae*, and other typical BV-pathogens are significantly higher in the BV-positive group than in healthy controls [23]. Additionally, *A. vaginae* has been positively associated with BV typical vaginal discharge, an elevated pH and the presence of clue cells [18, 24]. It was also described that high vaginal loads of *A. vaginae* in combination with *G. vaginalis* are associated with late miscarriage and prematurity [6, 25]. Additionally, it was demonstrated using an in vitro model that *A. vaginae* stimulates an innate immune response from epithelial cells, leading to localised IL-6 and IL-8 and an antimicrobial β -defensin peptide production occurring after triggering the toll-like receptor 2, and this possibly contributes to the pathogenesis of BV [26]. Hence, in BV *G. vaginalis* and *A. vaginae* still belong to ‘the main suspects’, a possible synergism between the two organisms has been considered, and because of this several authors using molecular-based techniques have examined the possibility of combining loads of *A. vaginae* and *G. vaginalis* as a means of diagnosing BV [15, 27, 28].

Involvement of *A. vaginae* in biofilm formation

Bacteria rarely exist as single-species planktonic forms but thrive in complex polymicrobial adhering communities enveloped by extracellular matrices, so-called biofilms. The bacteria account for less than 10% of biofilm mass, whereas the biofilm matrix usually accounts for more than 90% and provides the best living conditions for the bacteria [29]. Costerton et al. in 1999 [30] have described the association of a bacterial biofilm with various chronic infections. Regardless of the location in the human body, biofilm infections share similar clinical characteristics. They grow slowly, and bacterial communities are rarely fully destroyed by the host-defence mechanisms. Bacteria in biofilm release antigens resulting in an increase in antibody production. However, due to the biofilm structure, the produced antibodies are not capable of killing the biofilm bacteria [30].

The biofilm formation in BV is a virulence mechanism that enhances pathogenicity [31]. The polymicrobial BV-biofilm can be seen with the Gram-stain method in the form of clue cells, which are vaginal epithelial cells covered by

layers of adherent Gram-negative and/or -variable cells. Using fluorescence in situ hybridisation (FISH) method, the structure and composition of the biofilm can be studied in more detail [32], especially considering the combined presence of *G. vaginalis* and *A. vaginae*. Hardy et al. [33], similar to that previously described by Swidsinski et al. [31], have demonstrated that adherent *A. vaginae* and *G. vaginalis* were visualised in, respectively, 54% and 82% of samples with bacterial biofilm in BV. It was detected that *G. vaginalis* accounted for 60% or more and *A. vaginae* accounted for 40% or less of the film bacterial composition. It is assumed that *G. vaginalis* acts as an initial coloniser to establish early biofilm structures to which secondary colonisers, such as *A. vaginae* can attach [33, 34]. The fact that *G. vaginalis* is capable of displacing protective lactobacilli on pre-coated vaginal epithelial cells, is probably related to its ability to promote biofilm formation. In contrast, the other anaerobes, including *A. vaginae*, are easily outcompeted by *L. crispatus* [35]. Hardy et al. [33] also demonstrated the important role of *A. vaginae* together with *G. vaginalis* in BV-associated biofilm. Interactions among these species within a biofilm are synergistic: these include co-aggregation, metabolic cooperation and increased resistance to antibiotics or host immune responses and have important clinical implications [36]. The presence of a biofilm—due to increased resistance to treatment—is thought to be one of the possible reasons for the BV recurrence [37, 38].

Considerations on conventional treatments

Concerning the treatment options, the mainstay of BV treatment in many countries remains either oral or vaginal metronidazole once a day for 5–7 days or vaginal clindamycin as first-line treatments. The efficacy of the treatment with

metronidazole is comparable to topical clindamycin [39]. Cure rates, following intravaginal treatment with metronidazole or clindamycin, account for 70–90% at the end of treatment and 1 month after the end of therapy [3, 4, 40]. However, as it was first and best described by Larsson and Forsum in 2005 [39], 3 months after the treatment the recurrence rate can exceed 30% [41].

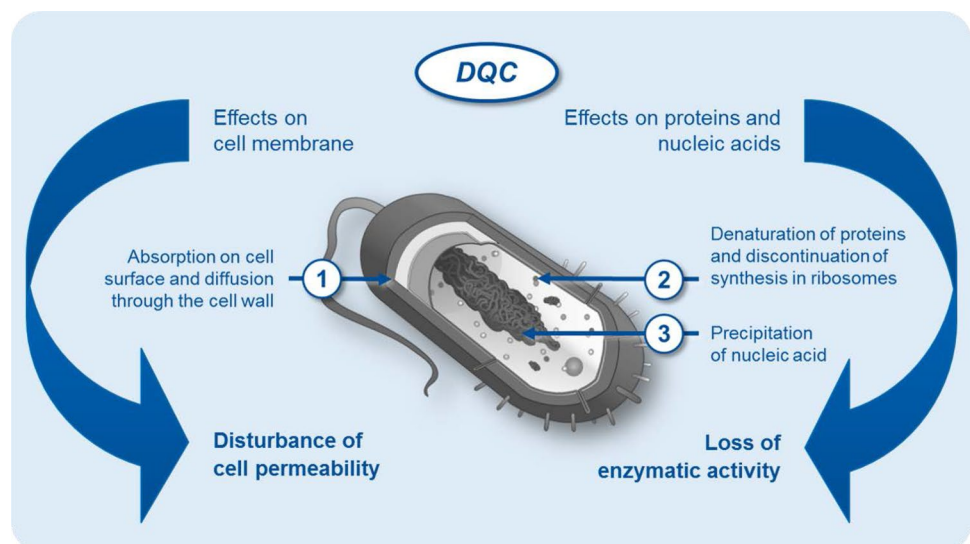
De Backer et al. [42] demonstrated that susceptibility to metronidazole varied significantly across various *A. vaginae* strains in vitro. Some of the investigated clinical isolates of *A. vaginae* were shown to be highly resistant to nitroimidazoles such as metronidazole and secnidazole [11, 43] and it was suggested that this could play a role in treatment failure [11, 42]. Susceptibility testing for metronidazole of additional *A. vaginae* isolates demonstrated that the minimum inhibition concentration (MIC) is variable, ranging from 2 µg/mL (sensitive) to more than 256 µg/mL (resistant) [12]. So far more than half of the tested isolates were resistant.

Beigi and colleagues described a significant increase of clindamycin-resistant anaerobic bacteria after treatment [44]. It is unclear if this is true for *A. vaginae* specifically. Even though these resistance findings seem to be interesting, it is questionable whether it influences clinical efficacy in patients with recurrent BV formerly treated with clindamycin. In vitro data suggest that clindamycin is effective against *A. vaginae* already at low concentrations [45].

Exploring alternative approaches

Alternatives to current antibiotic treatments against BV are increasingly being explored: antiseptics, probiotics, plant-derived compounds, vaginal acidifying and buffering agents, as well as different combination therapies are increasingly used [46–48]. A big interest, due to beneficial

Fig. 2 Mode of action of dequalinium chloride [47]. Dequalinium chloride (DQC) acts as a microbicidal against all main vaginal pathogens and due to the multiple modes of action, the development of resistances is unlikely



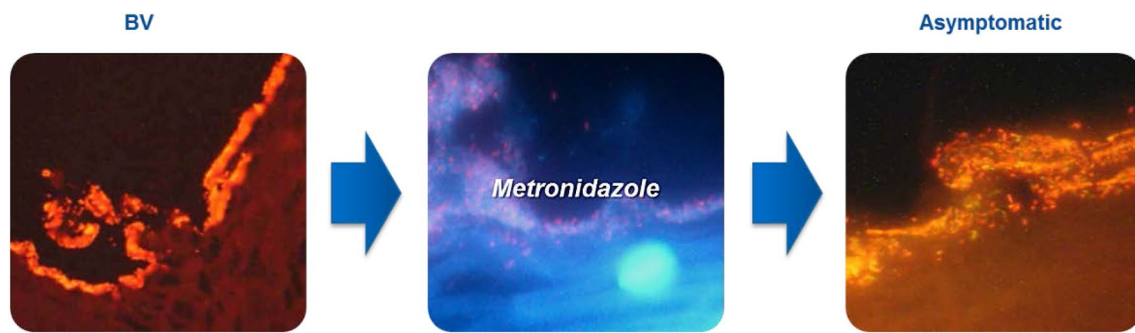


Fig. 3 BV-biofilm [31, 38]. A bacterial biofilm, an adhering microbial community enveloped by extracellular matrices, is considered one of the possible reasons for the BV recurrence

characteristics, has been seen in a group of antimicrobial substances belonging to antiseptics, such as dequalinium chloride (DQC) (Fig. 2) [47]. DQC was recently listed in an international guideline as an alternative treatment for BV [49]. Some authors [50] have suggested that DQC, as an antiseptic substance, could be preferable to a repeated course of antibiotics for patients with frequently recurrent BV. The antimicrobial activity of DQC has been investigated and demonstrated over the past decades by several investigators [51–55]. Della Casa et al. [51] has demonstrated the in vitro antimicrobial activity of this substance against different pathogens that are relevant for vaginal infections, including anaerobic bacteria (*G. vaginalis*, etc.), aerobic bacteria (staphylococci, streptococci, etc.), and *Candida* species. Additionally, the non-inferiority of DQC vaginal tablets to clindamycin vaginal cream for the treatment of BV has been demonstrated in a clinical study by Weissenbacher et al. [48]. Based on Amsel's criteria, the clinical cure rates 4 weeks after the end of treatment with DQC were 79.5% and 77.6% with clindamycin, respectively.

It has been shown that a high concentration of *A. vaginae* before treatment was associated with complete or partial failure of treatment for BV [11, 42]. Additionally, the presence of *A. vaginae* in the BV-associated biofilm has been described [31] and it seems that this knowledge has a major impact on treatment [33].

A recent study by Lopes dos Santos Santiago et al. [45] has investigated the in vitro susceptibility of *A. vaginae* to DQC in comparison to established substances (metronidazole and clindamycin). The MIC (minimal inhibition concentration) and MBC (minimal bactericidal concentration) range of DQC for 28 strains, belonging to 4 species of the genus *Atopobium*, i.e., *A. minutum*, *A. rimae*, *A. parvulum*, and *A. vaginae*, have been determined. The MIC and MBC for *Atopobium* spp. to DQC ranged between < 0.0625 and 2 µg/mL with an MIC₉₀-value of 2 µg/mL. The MIC₉₀-value for *A. vaginae* was demonstrated to be 0.5 µg/mL, i.e., *A. vaginae* was more sensitive than the other species tested. Not only was growth of *A. vaginae* inhibited at the MIC-levels,

but the bacterial cells were also killed. The MICs of clindamycin and DQC for *A. vaginae* in this study were similar (sensitive), whereas the MIC of metronidazole was much higher (resistant) [45].

Biofilm disruption for treatment success

Persistence of an adherent bacterial biofilm, containing mostly *G. vaginalis* and *A. vaginae*, can be considered as the major reason for failure of BV treatment (Fig. 3) [31, 38]. Bacteria in biofilms are less susceptible to antibiotics compared to planktonic cells and have a higher tolerance towards antimicrobial treatment [56]. It was found, that although all patients recovered after oral metronidazole treatment, a large reservoir of *A. vaginae* (together with *G. vaginalis*) was persisting as a bacterial biofilm [38]. Additionally, an in vivo study with topical metronidazole gel by Bradshaw et al. [57] found that rates of recurrence of BV were higher when *A. vaginae* was present in addition to *G. vaginalis*. Interestingly, an in vitro study has demonstrated the ability of beneficial *Lactobacillus* spp. to disrupt the biofilm consisting of *A. vaginae* and *G. vaginalis* [58].

In the course of recent in vitro investigations, Gottschick and colleagues have screened various compounds for vaginal Biofilms (consisting of *G. vaginalis*): The antibiotics, such as metronidazole and tobramycin were effective in preventing biofilm formation, but had no effect on an established biofilm, while some antiseptic substances led to the disintegration of existing biofilms [59]. Recent not yet published in vitro data are suggesting that DQC could be effective in disrupting the BV-biofilm under experimental conditions [60].

Conclusions

A. vaginae is an important component of the complex abnormal vaginal flora in BV. Resistance of *A. vaginae* to metronidazole, one of the current first-line treatments, as well

as the presence of *A. vaginae* in the BV-associated biofilm have been described. Although more clinical data would be needed on this subject, this seems to have a major impact on BV treatment outcomes. Nevertheless, recently available scientific data confirms that DQC (Fluomizin®) is one of the valid therapeutic options for BV treatment, as it displays a broad antimicrobial spectrum against relevant vaginal pathogens, especially against *G. vaginalis* and *A. vaginae*, without having safety concerns.

Acknowledgements The authors would like to thank Gabriela Zwysig and Dr Philipp Grob, Medinova AG, Switzerland for their comments on the manuscript.

Author contributions The author WM received a honorary for the manuscript. All authors contributed to the manuscript.

Funding This review and online free access of the publication was sponsored by Medinova AG, Switzerland.

Compliance with ethical standards

Conflict of interest Dr Valdas Prasauskas and Stephan Biber are employees of Medinova AG, Switzerland.

Open Access This article is distributed under the terms of the Creative Commons Attribution 4.0 International License (<http://creativecommons.org/licenses/by/4.0/>), which permits unrestricted use, distribution, and reproduction in any medium, provided you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons license, and indicate if changes were made.

References

- Kenyon C, Colebunders R, Crucitti T (2013) The global epidemiology of bacterial vaginosis: a systematic review. *Am J Obstet Gynecol*
- Srinivasan S, Fredricks DN (2008) The human vaginal bacterial biota and bacterial vaginosis. *Interdiscip Perspect Infect Dis* 2008:750479
- Donders G (2010) Diagnosis and management of bacterial vaginosis and other types of abnormal vaginal bacterial flora: a review. *Obstet Gynecol Surv* 65:462–473
- Nasioudis D, Linhares IM, Ledger WJ, Witkin SS (2017) Bacterial vaginosis: a critical analysis of current knowledge. *BJOG* 124:61–69
- Witkin S (2014) The vaginal microbiome, vaginal anti-microbial defence mechanisms and the clinical challenge of reducing infection-related preterm birth. *BJOG*
- Menard JP, Mazouni C, Salem-Cherif I, Fenollar F, Raoult D, Boubli L, Gamorre M, Bretelle F (2010) High vaginal concentrations of *Atopobium vaginae* and *Gardnerella vaginalis* in women undergoing preterm labor. *Obstet Gynecol* 115:134–140
- Srinivasan S, Hoffman NG, Morgan MT, Matsen FA, Fiedler TL, Hall RW, Ross FJ, McCoy CO, Bumgarner R, Marrazzo JM, Fredricks DN (2012) Bacterial communities in women with bacterial vaginosis: high resolution phylogenetic analyses reveal relationships of microbiota to clinical criteria. *PLoS ONE* 7:e37818
- Cox C, Watt AP, McKenna JP, Coyle PV (2016) *Mycoplasma hominis* and *Gardnerella vaginalis* display a significant synergistic relationship in bacterial vaginosis. *Eur J Clin Microbiol Infect Dis* 35:481–487
- Gardner HL, Dukes CD (1955) Haemophilus vaginalis vaginitis: a newly defined specific infection previously classified "nonspecific" vaginitis. *Am J Obstet Gynecol* 69:962–976
- Fredricks DN, Fiedler TL, Thomas KK, Oakley BB, Marrazzo JM (2007) Targeted PCR for detection of vaginal bacteria associated with bacterial vaginosis. *J Clin Microbiol* 45:3270–3276
- Ferris MJ, Masztal A, Aldridge KE, Fortenberry JD, Fidel PL Jr, Martin DH (2004) Association of *Atopobium vaginae*, a recently described metronidazole resistant anaerobe, with bacterial vaginosis. *BMC Infect Dis* 4:5
- Burton JP, Devillard E, Cadieux PA, Hammond JA, Reid G (2004) Detection of *Atopobium vaginae* in postmenopausal women by cultivation-independent methods warrants further investigation. *J Clin Microbiol* 42:1829–1831
- Verstraelen H, Verhelst R, Claeys G, Temmerman M, Vaneechoutte M (2004) Culture-independent analysis of vaginal microflora: the unrecognized association of *Atopobium vaginae* with bacterial vaginosis. *Am J Obstet Gynecol* 191:1130–1132
- Donders GG, Zozzika J, Rezeberga D (2014) Treatment of bacterial vaginosis: what we have and what we miss. *Expert Opin Pharmacother* 15:645–657
- Lamont R, Sobel J, Akins R, Hassan S, Chaiworapongsa T, Kusanovic J, Romero R (2011) The vaginal microbiome: new information about genital tract flora using molecular based techniques. *BJOG* 118:533–549
- Collins MD, Wallbanks S (1992) Comparative sequence analyses of the 16S rRNA genes of *Lactobacillus minutus*, *Lactobacillus rimae* and *Streptococcus parvulus*: proposal for the creation of a new genus *Atopobium*. *FEMS Microbiol Lett* 74:235–240
- Rodriguez Jovita M, Collins MD, Sjoden B, Falsen E (1999) Characterization of a novel *Atopobium* isolate from the human vagina: description of *Atopobium vaginae* sp. nov. *Int J Syst Bacteriol* 49(Pt 4):1573–1576
- De Backer E, Verhelst R, Verstraelen H, Alqumber MA, Burton JP, Tagg JR, Temmerman M, Vaneechoutte M (2007) Quantitative determination by real-time PCR of four vaginal *Lactobacillus* species, *Gardnerella vaginalis* and *Atopobium vaginae* indicates an inverse relationship between *L. gasseri* and *L. iners*. *BMC Microbiol* 7:115
- Fredricks DN, Fiedler TL, Marrazzo JM (2005) Molecular identification of bacteria associated with bacterial vaginosis. *N Engl J Med* 353:1899–1911
- Marconi C, Cruciani F, Vitali B, Donders GG (2012) Correlation of *Atopobium vaginae* amount with bacterial vaginosis markers. *J Low Genit Tract Dis* 16:127–132
- Verhelst R, Verstraelen H, Claeys G, Verschraegen G, Delanghe J, Van Simaey L, De Ganck C, Temmerman M, Vaneechoutte M (2004) Cloning of 16S rRNA genes amplified from normal and disturbed vaginal microflora suggests a strong association between *Atopobium vaginae*, *Gardnerella vaginalis* and bacterial vaginosis. *BMC Microbiol* 4:16
- Zhou X, Brown CJ, Abdo Z, Davis CC, Hansmann MA, Joyce P, Foster JA, Forney LJ (2007) Differences in the composition of vaginal microbial communities found in healthy Caucasian and black women. *ISME J* 1:121–133
- Wang KD, Su JR (2014) Quantification of *Atopobium vaginae* loads may be a new method for the diagnosis of bacterial vaginosis. *Clin Lab* 60:1501–1508
- Silva D, Henriques A, Cereija T, Martinez-de-Oliveira J, Miranda M, Cerca N (2014) Prevalence of *Gardnerella vaginalis* and *Atopobium vaginae* in Portuguese women and association

- with risk factors for bacterial vaginosis. *Int J Gynaecol Obstet* 124:178–179
25. Bretelle F, Rozenberg P, Pascal A, Favre R, Bohec C, Loundou A, Senat MV, Aissi G, Lesavre N, Brunet J et al (2015) High *Atopobium vaginae* and *Gardnerella vaginalis* vaginal loads are associated with preterm birth. *Clin Infect Dis* 60:860–867
 26. Libby EK, Pascal KE, Mordechai E, Adelson ME, Trama JP (2008) *Atopobium vaginae* triggers an innate immune response in an in vitro model of bacterial vaginosis. *Microbes Infect* 10:439–446
 27. Menard JP, Fenollar F, Henry M, Bretelle F, Raoult D (2008) Molecular quantification of *Gardnerella vaginalis* and *Atopobium vaginae* loads to predict bacterial vaginosis. *Clin Infect Dis* 47:33–43
 28. Henriques A, Cereija T, Machado A, Cerca N (2012) In silico vs in vitro analysis of primer specificity for the detection of *Gardnerella vaginalis*, *Atopobium vaginae* and *Lactobacillus* spp. *BMC Res Notes* 5:637
 29. Flemming HC, Wingender J (2010) The biofilm matrix. *Nat Rev Microbiol* 8:623–633
 30. Costerton JW, Stewart PS, Greenberg EP (1999) Bacterial biofilms: a common cause of persistent infections. *Science* 284:1318–1322
 31. Swidsinski A, Mendling W, Loening-Baucke V, Ladhoff A, Swidsinski S, Hale LP, Lochs H (2005) Adherent biofilms in bacterial vaginosis. *Obstet Gynecol* 106:1013–1023
 32. Hardy L, Jespers V, Dahchour N, Mwambarangwe L, Musengamana V, Vaneechoutte M, Crucitti T (2015) Unravelling the bacterial vaginosis-associated biofilm: a multiplex *Gardnerella vaginalis* and *Atopobium vaginae* fluorescence in situ hybridization assay using peptide nucleic acid probes. *PLoS ONE* 10:e0136658
 33. Hardy L, Jespers V, Abdellati S, De Baetselier I, Mwambarangwe L, Musengamana V, van de Wijgert J, Vaneechoutte M, Crucitti T (2016) A fruitful alliance: the synergy between *Atopobium vaginae* and *Gardnerella vaginalis* in bacterial vaginosis-associated biofilm. *Sex Transm Infect*
 34. Machado A, Cerca N (2015) Influence of biofilm formation by *Gardnerella vaginalis* and other anaerobes on bacterial Vaginosis. *J Infect Dis*
 35. Machado A, Salgueiro D, Harwich M, Jefferson KK, Cerca N (2013) Quantitative analysis of initial adhesion of bacterial vaginosis-associated anaerobes to ME-180 cells. *Anaerobe* 23:1–4
 36. Elias S, Banin E (2012) Multi-species biofilms: living with friendly neighbors. *FEMS Microbiol Rev* 36:990–1004
 37. Swidsinski A, Dorffel Y, Loening-Baucke V, Mendling W, Verstraelen H, Dieterle S, Schilling J (2010) Desquamated epithelial cells covered with a polymicrobial biofilm typical for bacterial vaginosis are present in randomly selected cryopreserved donor semen. *FEMS Immunol Med Microbiol* 59:399–404
 38. Swidsinski A, Mendling W, Loening-Baucke V, Swidsinski S, Dorffel Y, Scholze J, Lochs H, Verstraelen H (2008) An adherent *Gardnerella vaginalis* biofilm persists on the vaginal epithelium after standard therapy with oral metronidazole. *Am J Obstet Gynecol* 198:96–97
 39. Larsson PG, Forsum U (2005) Bacterial vaginosis—a disturbed bacterial flora and treatment enigma. *APMIS* 113:305–316
 40. Livengood CH (2009) Bacterial vaginosis: an overview for 2009. *Rev Obstet Gynecol* 2:28–37
 41. Bradshaw CS, Morton AN, Hocking J, Garland SM, Morris MB, Moss LM, Horvath LB, Kuzevska I, Fairley CK (2006) High recurrence rates of bacterial vaginosis over the course of 12 months after oral metronidazole therapy and factors associated with recurrence. *J Infect Dis* 193:1478–1486
 42. De Backer E, Verhelst R, Verstraelen H, Claeys G, Verschraegen G, Temmerman M, Vaneechoutte M (2006) Antibiotic susceptibility of *Atopobium vaginae*. *BMC Infect Dis* 6:51
 43. De Backer E, Dubreuil L, Brauman M, Acar J, Vaneechoutte M (2010) In vitro activity of secnidazole against *Atopobium vaginae*, an anaerobic pathogen involved in bacterial vaginosis. *Clin Microbiol Infect* 16:470–472
 44. Beigi RH, Austin MN, Meyn LA, Krohn MA, Hillier SL (2004) Antimicrobial resistance associated with the treatment of bacterial vaginosis. *Am J Obstet Gynecol* 191:1124–1129
 45. Lopes dos Santos Santiago G, Grob P, Verstraelen H, Waser F, Vaneechoutte M (2012) Susceptibility testing of *Atopobium vaginae* for dequalinium chloride. *BMC Res Notes* 5:151
 46. Polatti F (2012) Bacterial vaginosis, *Atopobium vaginae* and nifuratel. *Curr Clin Pharmacol* 7:36–40
 47. Mendling W, Weissenbacher ER, Gerber S, Prasauskas V, Grob P (2016) Use of locally delivered dequalinium chloride in the treatment of vaginal infections: a review. *Arch Gynecol Obstet* 293:469–484
 48. Weissenbacher ER, Donders G, Unzeitig V, de Martinez TB, Gerber S, Halaska M, Spacek J (2012) A comparison of dequalinium chloride vaginal tablets (Fluomizin(R)) and clindamycin vaginal cream in the treatment of bacterial vaginosis: a single-blind, randomized clinical trial of efficacy and safety. *Gynecol Obstet Invest* 73:8–15
 49. Sherrard J, Wilson J, Donders G, Mendling W, Jensen JS (2018) European (IUSTI/WHO) International Union against sexually transmitted infections (IUSTI) World Health Organisation (WHO) guideline on the management of vaginal discharge. *Int J STD AIDS* 2018:956462418785451
 50. Hay P (2017) Bacterial vaginosis. *F1000Res* 6:1761.
 51. Della Casa V, Noll H, Gonser S, Grob P, Graf F, Pohlig G (2002) Antimicrobial activity of dequalinium chloride against leading germs of vaginal infections. *Arzneimittelforschung* 52:699–705
 52. D'Auria FD, Simonetti G, Strippoli V (1989) Caratteristiche antimicrobiche di una tintura al dequalinio cloruro. *Ann Ig* 1:1227–1241
 53. Cox WA (1965) Site of action of certain antibacterial heterocyclic quaternary ammonium compounds. *Appl Microbiol* 13:956–966
 54. Babbs M, Collier HOJ, Austin WC, Potter MD, Taylor EP (1956) Salts of decamethylene-bis-4-aminoquinaldinium (Dequadin), a new antimicrobial agent. *J Pharm Pharmacol* 8:110–119
 55. Cella JA, Eggenberger DN, Harriman LA, Harwood HJ (1952) The relation of structure and critical concentration to the bactericidal activity of quaternary ammonium salts. *J Am Chem Soc* 74:2061–2062
 56. Anderl JN, Franklin MJ, Stewart PS (2000) Role of antibiotic penetration limitation in *Klebsiella pneumoniae* biofilm resistance to ampicillin and ciprofloxacin. *Antimicrob Agents Chemother* 44:1818–1824
 57. Bradshaw CS, Tabrizi SN, Fairley CK, Morton AN, Rudland E, Garland SM (2006) The association of *Atopobium vaginae* and *Gardnerella vaginalis* with bacterial vaginosis and recurrence after oral metronidazole therapy. *J Infect Dis* 194:828–836
 58. McMillan A, Dell M, Zellar MP, Cribby S, Martz S, Hong E, Fu J, Abbas A, Dang T, Miller W, Reid G (2011) Disruption of urogenital biofilms by lactobacilli. *Colloids Surf B Biointerfaces* 86:58–64
 59. Gottschick C, Szafranski SP, Kunze B, Sztajer H, Masur C, Abels C, Wagner-Dobler I (2016) Screening of compounds against *Gardnerella vaginalis* biofilms. *PLoS ONE* 11:e0154086
 60. Palmeira-de-Oliveira A, Personal communication, 2018.

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.