




# *Lactobacillus rhamnosus* Lcr35 Stimulates Epithelial Vaginal Defenses upon *Gardnerella vaginalis* Infection

 Sylvie Miquel,<sup>a</sup> Julien Verlaquet,<sup>a</sup> Sophie Garcin,<sup>b</sup> Thomas Bertran,<sup>b</sup> Bertrand Evrard,<sup>b,c</sup>  Christiane Forestier,<sup>a</sup> Marjolaine Vareille-Delarbre<sup>a</sup>

<sup>a</sup>Université Clermont Auvergne, UMR CNRS 6023 LMGE, Laboratoire de Bactériologie, Clermont-Ferrand, France

<sup>b</sup>Université Clermont Auvergne, INRA, UNH, Unité de Nutrition Humaine, Laboratoire d'Immunologie, Clermont-Ferrand, France

<sup>c</sup>CHU Clermont-Ferrand, Service d'Immunologie, Clermont-Ferrand, France

**ABSTRACT** Dysbiosis of the vaginal microbiome as a result of overgrowth of anaerobic bacteria, such as *Gardnerella vaginalis*, and low levels of “healthy” lactobacilli leads to bacterial vaginosis (BV), usually associated with a low-grade inflammatory process. Despite appropriate antibiotic treatment, *G. vaginalis*-associated BV is characterized by significant recurrence. The use of probiotics could be an interesting alternative therapy due to their ability to rebalance vaginal microbiota. In this study, we investigated the effects of a well-characterized probiotic strain, *Lactobacillus rhamnosus* Lcr35, on epithelial vaginal and dendritic cell (DC) immune responses after *G. vaginalis* infection. In an *in vitro* coculture model with human monocyte-derived dendritic cells and a vaginal epithelial cell (VEC) monolayer, the Lcr35 strain induced DC activation, as evidenced by the induction of maturation and synthesis of interleukin-8 (IL-8) and CCL-20 chemokines upon apical challenge of the VECs by *G. vaginalis*. Analysis of the vaginal epithelial response showed that the presence of Lcr35 significantly increased the production of the proinflammatory cytokines IL-8 and IL-1 $\beta$  and human  $\beta$ -defensin 2 (HBD-2), whereas the concentration of secretory leukocyte protease inhibitor (SLPI) was decreased in *G. vaginalis*-infected vaginal epithelial cells. Treatment with recombinant SLPI was associated with upregulation of Lcr35-stimulated IL-8 and HBD-2 production. These results suggest that inhibition of SLPI by Lcr35 in vaginal epithelial cells contributes to the host defense response against *G. vaginalis* infection.

**KEYWORDS** *Gardnerella vaginalis*, *Lactobacillus*, antimicrobial peptides, cytokines, dendritic cells, epithelial cells, vaginosis

The human vaginal microbiome is a complex bacterial community that interacts closely with vaginal epithelial cells impacting the mucosal immune response and its responses to pathogenic bacteria. A healthy vaginal microbiome is dominated by *Lactobacillus*, including *Lactobacillus crispatus*, *Lactobacillus jensenii*, *Lactobacillus gasseri*, and *Lactobacillus vaginalis*, resulting in a dynamic balance that inhibits overgrowth of potential anaerobic pathogens, such as *Gardnerella vaginalis*, *Atopobium* spp., and *Prevotella* spp. (1–3).

The female reproductive tract (FRT) can be divided into two immunological regions, the upper (endocervix, uterus, and oviduct) and lower (vagina and ectocervix) FRT. In the lower FRT, interactions between vaginal cells, lactobacilli, and their metabolic products create a physical and immunological barrier against pathogens that contributes to the prevention of invasion by exogenous microbes and regulation of the inflammatory response (4–6).

Despite these local host defense mechanisms, as many as 75% of women may experience an occasional vaginal infection, and some 5% to 10% suffer from recurrent bacterial vaginosis (BV) (7, 8).

BV is characterized by a polymicrobial imbalance, or dysbiosis, of the natural microflora of the cervicovaginal (CV) space. *G. vaginalis* is found in up to 94% of all cases of BV and has recently been suggested as the main “early colonizer species” that displaces lactobacilli and

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Address correspondence to Christiane Forestier, christiane.forestier@uca.fr.

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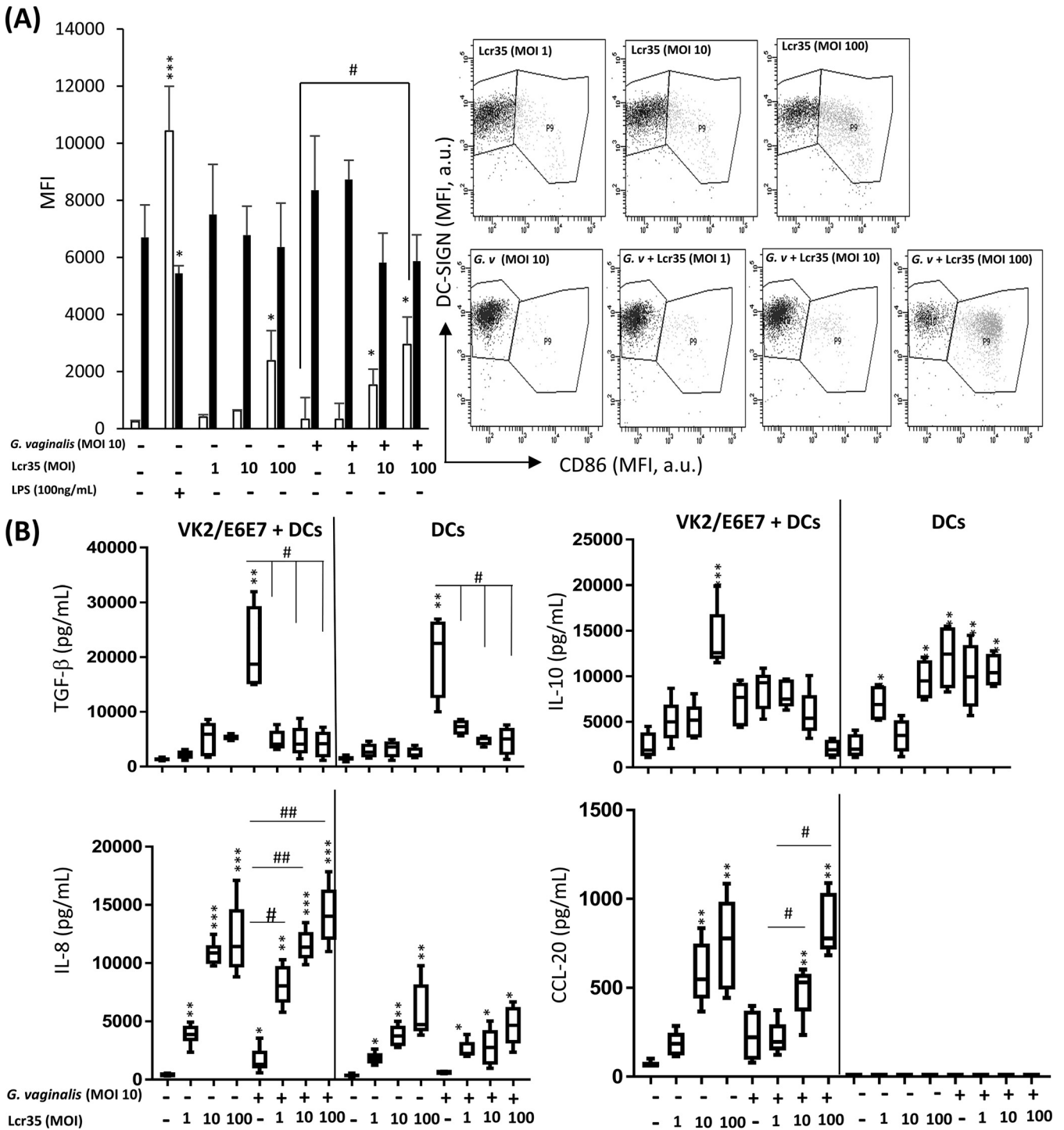
forms a biofilm, thereby creating a favorable environment for colonization by BV-associated bacteria, such as *Prevotella bivia*, *Atopobium vaginae*, and *Megasphaera* type 1 (9). BV can also produce a state of local immunosuppression that increases susceptibility to HIV and other sexually transmitted diseases (8, 10). Unlike vulvovaginal candidiasis and vaginitis, BV is typically distinguished by a lack of inflammation: IL-8 concentrations and leukocyte counts are not significantly greater in BV-positive women than in healthy control subjects (11–13). A deficiency in antimicrobial peptides, such as human  $\beta$ -defensin-1 and -2, has also been reported in vaginal fluid from women with BV (14). In contrast, vaginal interleukin-1 $\beta$  (IL-1 $\beta$ ) levels are largely increased (13, 15), suggesting that BV-associated *G. vaginalis* interacts with the host vaginal immunity specifically to dampen IL-8 production. BV has also been associated with a significant reduction in vaginal levels of the secretory leukocyte protease inhibitor (SLPI). SLPI is produced by both epithelial and immune cells (16) and is bactericidal for pathogens such as *Neisseria gonorrhoeae* (17); it has also been shown to participate in the mucosal defense by reducing inflammation (18).

The currently recommended treatment for BV is antibiotics (metronidazole and clindamycin), but the rates of recurrence following antibiotic treatment are extremely high and >50% of women have recurrent episodes within 6 to 12 months (19). In this context, probiotics appear as an interesting alternative strategy due to their role in maintaining a healthy vaginal flora (3, 20). Their beneficial action could be related to the competitive exclusion of pathogenic bacteria, competition for nutrients, production of antimicrobial substances, and/or activation of the immune system (21, 22).

*Lactobacillus rhamnosus* Lcr35, recently renamed *Lactocaseibacillus rhamnosus* Lcr35, is a well-characterized probiotic strain that has been used in clinical practice for more than 50 years, in particular to restore vaginal health (GynOphilus). The antimicrobial effects of this probiotic strain have been largely documented over the past decade (23, 24), but little is known about its interactions with the vaginal immune system. In the present study, we assessed the immunomodulatory effect of Lcr35 in innate vaginal cells infected by *G. vaginalis*. We show that Lcr35 has an immunostimulatory effect on vaginal cells infected by *G. vaginalis* by activating both epithelial and dendritic cell (DC) responses. These effects are associated with modulation of SLPI production, offering a novel mechanism for the regulation of innate response by lactobacilli during *G. vaginalis* infection.

## RESULTS

***Lactocaseibacillus rhamnosus* Lcr35 stimulates the cross talk between epithelial and dendritic cells when cocultured with *Gardnerella vaginalis*.** In a coculture model composed of human monocyte-derived dendritic cells in the lower part and vaginal epithelial cell monolayer (VK2/E6E7) in the upper part (see Fig. S1 in the supplemental material), the addition of *G. vaginalis* in the upper part did not induce any change in the DC surface phenotype compared to that of immature DCs as measured by determination of CD86 and dendritic cell-specific intercellular adhesion molecule-3-grabbing nonintegrin (DC-SIGN) expression (Fig. 1A). In contrast, the addition of Lcr35 in the upper part of the same coculture system induced a dose-dependent higher surface expression of CD86 and a lower surface expression of DC-SIGN than noninfected cells (Fig. 1A). Analysis of other activation markers such as CD40, CD206, CD83, and HLA-DR confirmed that incubation with Lcr35 in the presence or absence of *G. vaginalis* induced maturation of DCs (see Fig. S2B and C in the supplemental material). In addition, increased expression of CD86 and decreased expression of DC-SIGN were also observed when both Lcr35 and *G. vaginalis* were added directly onto DCs (see Fig. S3 in the supplemental material). For controls, an immature DC profile was observed with low levels of CD86 and high levels of DC-SIGN expression, whereas addition of lipopolysaccharide (LPS) at the apical surface of the epithelial monolayer induced a phenotype characteristic of fully mature DCs with increased levels of CD86 combined with decreased levels of DC-SIGN (Fig. 1A). Determination of the concentrations of immune attractive chemokines (CCL-20 and IL-8) and immunosuppressive cytokines (transforming growth factor  $\beta$  [TGF- $\beta$ ] and IL-10) in the lower compartment of the cell coculture model (corresponding to the basolateral surface of the epithelial monolayer) showed an increase in both TGF- $\beta$  and IL-8 production when the cells were incubated with *G. vaginalis* alone



**FIG 1** Effect of Lcr35 on *G. vaginalis*-infected dendritic cells in a coculture model. Human monocyte-derived dendritic cells were exposed indirectly through the vaginal epithelial cell monolayer to UV-inactivated *G. vaginalis* (MOI, 10) alone or UV-inactivated Lcr35 (MOI, 1 to 100) for 48 h. (A) The effects of DC functional maturation were determined by measuring the DC-surface expression of DC-SIGN (black bars) and CD86 (white bars) by flow cytometry. The dot plots and histograms show MFI values of gated DCs. DC-SIGN/CD86 dot plots gated on human DCs. (B) The secretions of cytokines TGF-β, IL-10, IL-8, and CCL-20 were measured in the supernatant of coculture VK2/E6E7 + DC (lower part) or DC alone by ELISA. Values are the means ± SEM; n = 4 to 5; \*, P < 0.05; \*\*, P < 0.01 compared with noninfected DCs. #, P < 0.05; ##, P < 0.01 compared with DCs infected with *G. vaginalis*.

(multiplicity of infection [MOI], 10). Incubation of the same cells with Lcr35 induced dose-dependent increased secretion of IL-8, IL-10, and CCL-20 but not TGF-β. Infection of DCs with both bacteria, *G. vaginalis* and Lcr35, resulted in phenotypes similar to those observed when cells were incubated with Lcr35 alone (Fig. 1B). Direct contact of Lcr35 with DCs

**TABLE 1** Production of innate vaginal components by VK2/E6E7 vaginal epithelial cells after incubation with Lcr35<sup>a</sup>

Mediator	Production of innate vaginal components (pg/mL)						
	NI	Lcr35			<i>G. vaginalis</i>		
		MOI, 1	MOI, 10	MOI, 100	MOI, 1	MOI, 10	MOI, 100
Cytokine							
IL-8	16.8 ± 6.1	814.5 ± 133.3**	1,757.9 ± 135.9***	2,391.2 ± 322.7***	192.2 ± 8.6	216.1 ± 28.1	317.3 ± 34.5
IL-1β	5.2 ± 1.8	7.9 ± 1.2	12.6 ± 1.9	81.8 ± 20.6*	3.1 ± 1.3	4.7 ± 2.1	18.2 ± 3.6
IL-6	ND	ND	ND	ND	ND	ND	ND
IL-10	ND	ND	ND	ND	ND	ND	ND
Antimicrobial peptide							
HBD-2	182.0 ± 7.2	421.6 ± 30.2*	550.3 ± 41.8*	648.9 ± 34.4**	350.3 ± 20.8*	335.4 ± 11.2*	318.9 ± 10.4*

<sup>a</sup>VK2/E6E7 cells were incubated with Lcr35 or *G. vaginalis* (MOI, 1 to 100) for 6 h. Subsequently, cytokines or host defense peptides concentration were measured in the culture supernatants by ELISA. Each value is the mean ± SEM of *n* = 3 to 8 independent experiments. ND, not detected. \*, *P* < 0.05; \*\*, *P* < 0.01; \*\*\*, *P* < 0.001 compared with noninfected cells (NI).

induced an increase of IL-8 secretion and a decrease of TGF-β and IL-10 levels compared to those of noninfected cells in the presence or absence of *G. vaginalis* (Fig. 1B). No production of CCL-20 was detected when DCs were directly incubated with bacteria (Fig. 1B).

**Lcr35 with and without *Gardnerella vaginalis* induces human epithelial vaginal cell response in a NF-κB-dependent manner.** To determine whether Lcr35 influences the immune response of VK2/E6E7 cells to *G. vaginalis*, the production of inflammatory cytokines by cells incubated separately with Lcr35 or *G. vaginalis* was first assessed. After incubation with Lcr35, the cells specifically induced the production of proinflammatory cytokine IL-8 and IL-1β and antimicrobial human β-defensin 2 (HBD-2) in a dose-dependent manner, whereas no significant change in the production of IL-6 and IL-10 was observed (Table 1). Incubation with different doses of *G. vaginalis* (MOI from 1 to 100) induced a low vaginal epithelial response, as shown by the absence of significant change in all innate cell markers except IL-8 (Table 1).

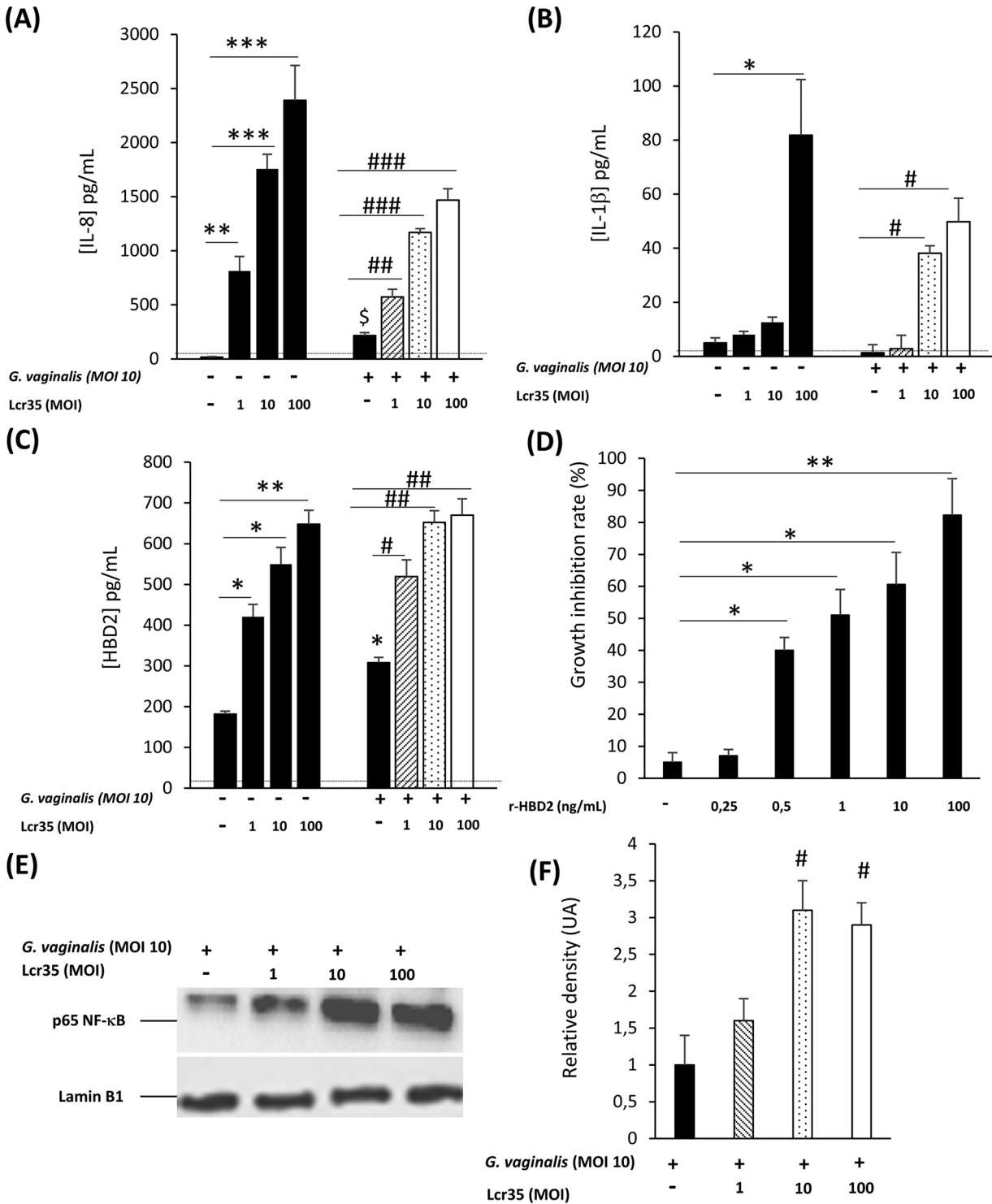
The cells were then treated with both Lcr35 and *G. vaginalis*. Increased production of IL-8 and IL-1β in an Lcr35 MOI-dependent manner was observed compared to that of cells infected with *G. vaginalis* alone (Fig. 2A and B). In all cases, no cytotoxic effect was observed (see Fig. S4 in the supplemental material), and no significant change in the production of IL-6 and IL-10 was observed (data not shown). Similar assays performed with other *Lactobacillus* species (*L. vaginalis*, *L. gasseri*, *Lactobacillus plantarum*, and *L. crispatus*) gave rise to similar results (see Fig. S5 in the supplemental material).

Likewise, the production of antimicrobial peptide HBD-2 increased significantly in vaginal epithelial cells treated by both Lcr35 and *G. vaginalis* compared to that of cells infected with *G. vaginalis* alone (Fig. 2C). Incubation of purified HBD-2 in the range of 0.5 to 100 ng/mL impaired *G. vaginalis* growth (Fig. 2D) but had no impact on Lcr35 growth (data not shown). Moreover, SLPI at the range of 0.2, 0.5, 1, 5 and 10 ng/mL did not show antimicrobial effect with Lcr35 or *G. vaginalis* (data not shown).

The level of p65 NF-κB in VK2/E6E7-infected cells was then measured to determine if the NF-κB pathway was involved in response to the presence of *G. vaginalis* and Lcr35. Infection with both *G. vaginalis* and Lcr35 resulted in larger amounts of p65 NF-κB protein than cells infected with the pathogen alone, as shown by Western blotting assays (Fig. 2E).

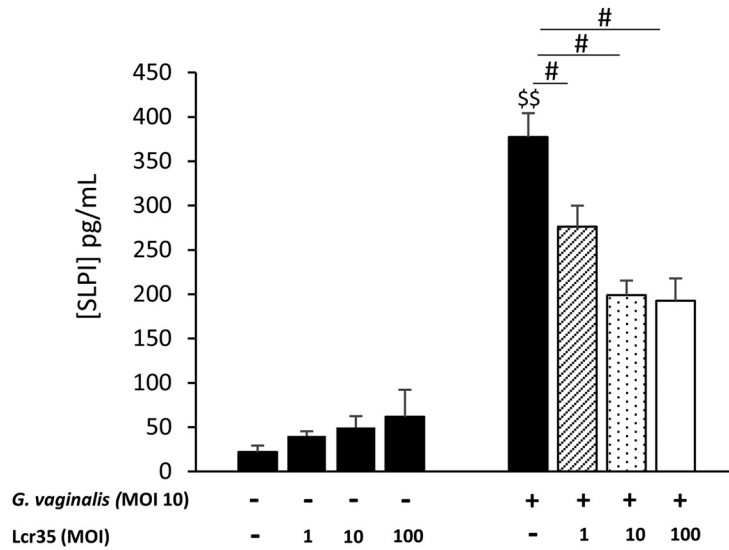
**Modulation of innate immune factor production by Lcr35 under the control of SLPI production in *G. vaginalis*-infected cells.** To further characterize the vaginal epithelial innate immune response to Lcr35 and/or *G. vaginalis*, we monitored SLPI host defense peptide production in vaginal cells. Incubation with Lcr35 did not elicit secretion of SLPI whatever the bacterial concentration tested, whereas a strong induction was observed in *G. vaginalis*-infected vaginal cells (Fig. 3A). Coincubation of the two bacterial strains, Lcr35 and *G. vaginalis*, with VK2/E6E7 cells resulted in lower SLPI levels in the cell supernatants than in cells infected with *G. vaginalis* alone (Fig. 3A).

Because SLPI has immunomodulatory properties via inhibition of the NF-κB pathway (25), we examined the effect of purified peptide on *G. vaginalis* growth and proinflammatory

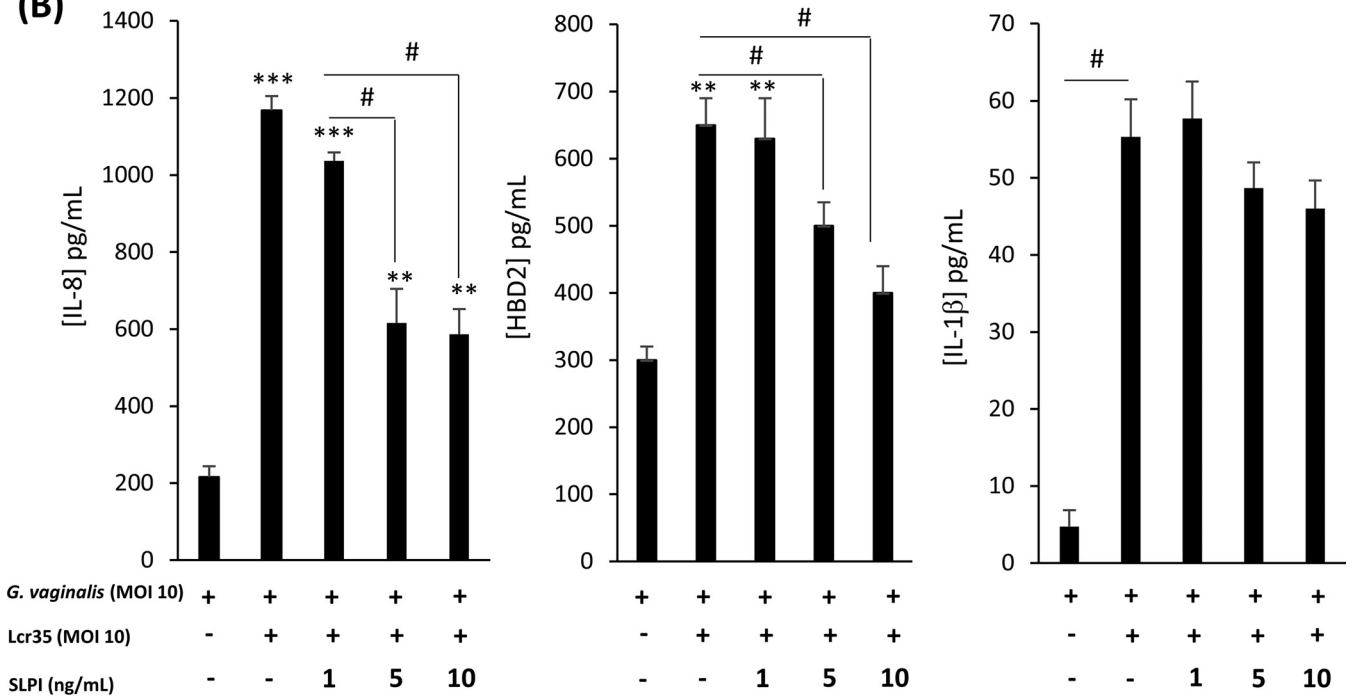


**FIG 2** Effect of Lcr35 on innate immune response in *G. vaginalis*-infected vaginal epithelial cells. VK2/E6E7 cells were infected with *G. vaginalis* (MOI, 10) alone or with Lcr35 at different MOIs (MOI, 1 to 100). IL-8 (A), IL-1β (B), and HBD-2 (C) concentrations were analyzed 6 h postinfection by ELISA. The detection limits of IL-8 and IL-1β were 312 and 39 pg/mL, respectively. (D) Growth inhibition rates of *G. vaginalis* after treatment with various concentrations of HBD-2 for 4 h. (E) The presence of the p65 NF-κB and Lamin B1 proteins were detected by Western blotting in cellular extracts of VK2/E6E7 cells stimulated for 3 h with *G. vaginalis* alone or with Lcr35. (F) Densitometric analysis of the data in Fig. 2E by using Image Lab 2.0 software ( $n = 3$ ). Representative data of 3 to 8 independent experiments. Values are the means  $\pm$  SEM; \*,  $P < 0.05$ ; \*\*,  $P < 0.01$  compared with uninfected cells. #,  $P < 0.05$ ; ##,  $P < 0.01$  compared with *G. vaginalis*-infected cells.

(A)



(B)



**FIG 3** Modulation of innate immune factor production by Lcr35 under the control of SLPI production in *G. vaginalis*-infected cells. VK2/E6E7 cells were infected with *G. vaginalis* (MOI, 10) alone or with Lcr35 at different MOIs (MOI, 1 to 100). (A) SLPI concentration was analyzed 6 h postinfection by ELISA. Recombinant SLPI protein was added to VK2/E6E7 cells incubated with *G. vaginalis* (MOI, 10) and Lcr35 (MOI, 10). (B) IL-8, HBD-2, and IL-1β concentrations were detected by ELISA in supernatants after 6 h of incubation. Values are the means ± SEM; n = 3 to 4; \$\$, P < 0.05 compared with uninfected cells; \*, P < 0.05; \*\*, P < 0.01 compared with *G. vaginalis*-infected cells. #, P < 0.05; #, P < 0.01 compared with vaginal cells infected with *G. vaginalis* and Lcr35.

cytokine production in *G. vaginalis*-infected cells. The addition of recombinant SLPI significantly decreased IL-8 and HBD-2 production compared to that observed with *G. vaginalis* and Lcr35 (MOI, 10) in vaginal epithelial cells (Fig. 3B). No significant decrease in IL-1β concentration was observed (Fig. 3B).

**DISCUSSION**

Several studies have previously demonstrated the efficacy of the probiotic strain Lcr35 in the prevention of recurrent vulvovaginal candidiasis and bacterial vaginosis (23, 24, 26, 27). Clinical studies performed with GynOphilus, the commercial vaginal form of Lcr35, reported



a positive effect on the recurrence of infections and patient quality of life (26, 28). In the present study, we showed that Lcr35 modified the host defense response of innate immune cells exposed to the pathogen *G. vaginalis*, the key player in the pathogenesis of BV.

Using a coculture model mimicking the vaginal epithelium with both vaginal epithelial cells (VK2/E6E7 cell line) and adjacent DCs, we showed that apical challenge of the epithelial cells with Lcr35 induces a dose-dependent maturation of DCs, whereas incubation with *G. vaginalis* had no effect. When the two bacteria were cocultured on the cells, the immunostimulatory effect of Lcr35 was not modified. Analysis of cytokine and chemokine concentrations in the basolateral compartment demonstrated an induction of chemokine secretion (IL-8 and CCL20) and a decrease in the TGF- $\beta$  immunosuppressive cytokine when challenged with Lcr35 regardless of the presence of *G. vaginalis*. Altogether these data suggest that (i) Lcr35 is able to activate DCs across the epithelial barrier and (ii) vaginal epithelial cells have an active role during *G. vaginalis* infection and are able to “sense” bacteria. Similar results were obtained in a murine coculture model (intestinal epithelial cells and bone marrow-derived DCs) mimicking the intestinal barrier by Grangette et al., who observed differential responses in intestinal epithelial activation and DC maturation between *Escherichia coli* and lactobacilli strains (29). These findings suggest that vaginal epithelial cells, like intestinal epithelial cells, have an “immunological filtering” role that predicts DC activation and consequently modulates adaptive response.

To characterize this signal, we first assessed the effect of Lcr35 on VK2/E6E7 cell response by measuring the release of several cytokines. We showed that Lcr35 stimulates production by VK2/E6E7 cells of the proinflammatory cytokines IL-8 and IL-1 $\beta$  and the antimicrobial peptide HBD-2 via the NF- $\kappa$ B pathway. Although the immunomodulatory properties of lactobacilli are strain-specific, we observed that this specific pathway was also induced by four other lactobacilli species: *L. vaginalis*, *L. gasseri*, *L. crispatus*, and *L. plantarum*. In the vagina, lactobacilli have been described as noninducers of cytokine production (5, 30, 31). Nevertheless, vaginal epithelial cells possess Toll-like receptor (TLR), and some *in vitro* studies propose that they respond to both commensal bacteria and invading pathogens by producing cytokines and immune mediators (32, 33). In addition, incubation with *Lactobacillus iners* has been associated with higher production of IL-6 and IL-8 by VK2 cells (34), colonization of vaginal epithelial cells by *L. jensenii* led to increased IL-6 and tumor necrosis factor alpha (TNF- $\alpha$ ) synthesis (35), and *L. rhamnosus* GG, *L. rhamnosus* LC705, and *L. rhamnosus* GR-1 were able to induce IL-8, IL-6, and TNF- $\alpha$  production in VK2 cells (36). The effect of lactobacilli on vaginal epithelial cells, in the absence of another immune stimulus, is therefore still a controversial issue. However, the Lcr35-induced inflammatory effect observed in this study is moderate compared with that induced by the pathogen *Candida albicans* with the level of IL-8 reaching  $2,635 \pm 213 \mu\text{g/mL}$  (our unpublished data). Tolerance of Lcr35 (GynOphilus) was previously evaluated in clinical studies, and no inflammatory signs (tingling, dryness, burning, itching, and pelvic pain) were reported (37). The same clinical study reported high basal levels of some cytokines in healthy women (12, 13, 38) in accordance with our *in vitro* results. All of these data suggest that a “basal moderate inflammatory state” is naturally induced in the vaginal environment to activate vaginal innate and adaptive immunity against pathogens.

*In vitro*, infection by *G. vaginalis* has been associated with divergent effects on immune cell populations and the cytokine network in the vaginal mucosa. Clinically, BV is mainly differentiated from vaginitis by the absence of an inflammatory response and is characterized by the absence of the chemotactic effect of IL-8 correlated the lack of leukocyte recruitment and increased vaginal antigen-specific immune levels (39–41). Our results show that *G. vaginalis* induces a low inflammatory response in vaginal epithelial cells characterized by low levels of IL-8, IL-6, and IL-1 $\beta$ . When Lcr35 was added together with *G. vaginalis* on the upper part of the cell monolayer, the response was exacerbated compared to that observed with *G. vaginalis* alone in terms of proinflammatory and antimicrobial signals involving the NF- $\kappa$ B pathway. Most studies describe an anti-inflammatory effect of *Lactobacillus* strains when they are associated with proinflammatory vaginal pathogens like *C. albicans* or *Atopobium vaginae* or used in pretreatment before inflammatory stimuli (5, 30, 42, 43). In our study, the effects of Lcr35 were assessed on epithelial cells infected with *G. vaginalis*, a noninflammatory pathogen. In

this tripartite relation, Lcr35—and other Lactobacilli species—induced an immunostimulatory effect. We thus hypothesized that lactobacilli are able to permit immune homeostasis through an interaction with epithelial cells. In a low inflammatory environment, such as the one encountered in BV, *Lactobacillus* could induce an “alert” immunostimulatory signal. However, when the vaginal cavity is contaminated by highly proinflammatory pathogens, lactobacilli could decrease the production of cytokines to control the inflammation and therefore avoid deleterious hyperinflammation. Although the precise mechanism underlying the Lcr35 immunomodulatory action remains to be elucidated, exogenous addition of Lcr35 is also likely to restore the balance of microbiota and thus contribute to an effective vaginal immune defense against pathogens. Specific induction of HBD-2 by Lcr35 could also be an effective mechanism to increase antimicrobial defense against both *G. vaginalis* and bacterial vaginosis-associated bacteria (44, 45). In our study, *G. vaginalis* alone strongly induced SLPI production by vaginal epithelial cells, whereas decreased levels of SLPI were observed *in vivo* in vaginal fluids of BV-positive women (46). This multifunctional antimicrobial protein expressed at mucosal surfaces plays a key role in infectious and inflammatory processes and suppresses inflammatory responses and proinflammatory cytokines (e.g., IL-8 and HBD-2) by controlling the activity of NF- $\kappa$ B in various immune cells (25, 47–49). Our findings showed that the induction of IL-8 and HBD-2 by Lcr35 in *G. vaginalis*-infected cells involves the inhibition of SLPI, which suggests a novel role of lactobacilli in the modulation of the epithelial immune response. The direct or indirect action of Lcr35 on the SLPI response still remains to be elucidated in the immune escape mechanism during *G. vaginalis* infection. The bactericidal activity of Lcr35 against *G. vaginalis* (27) could in part explain the anti-SLPI mechanisms. However, Lcr35 could also directly interact with epithelial cells as demonstrated by its highly dynamic adhesion abilities to vaginal ectocervical and endocervical cell lines (27). The mechanism(s) underlying the process by which SLPI is regulated by Lcr35 is likely multifaceted.

In summary, this study shows that the probiotic Lcr35 induces host responses against *G. vaginalis* by modulating the synthesis of both proinflammatory cytokines and host defense peptides in vaginal cells and also by enhancing the dialogue between epithelial and dendritic cells (Fig. 4). These immunostimulatory properties could restore a beneficial local immune response during *G. vaginalis* infections in line with the positive effects observed in clinical studies.

## MATERIALS AND METHODS

**Bacterial strains and culture conditions.** *Lactocaseibacillus rhamnosus* 35 (Lcr35), *Lactobacillus crispatus* BLL2005, *Lactobacillus vaginalis* CIP 105932, *Lactobacillus gasseri* CIP 102991-43, and *Lactobacillus plantarum* CIRM653 (24, 50) were grown without agitation in De Man, Rogosa, Sharpe (MRS) medium (Becton, Dickinson, Franklin Lakes, NJ, USA) at 37°C overnight. *G. vaginalis* ATCC 14018 was grown in brain heart infusion (BHI) medium (Becton, Dickinson) supplemented with maltose (0.1%), glucose (0.1%), yeast extract (1%), and horse serum (10%) in 5% CO<sub>2</sub> at 37°C for 72 h.

Bacterial cells were harvested by centrifugation (11,000 × *g* for 10 min), and the pellet was resuspended in appropriate cell culture medium. Optical density (OD) was measured at 620 nm to adjust the final concentration of the bacterial suspension, and the exact number of CFU was determined by plating serial dilutions of the inoculum on MRS or BHI plates (Becton, Dickinson).

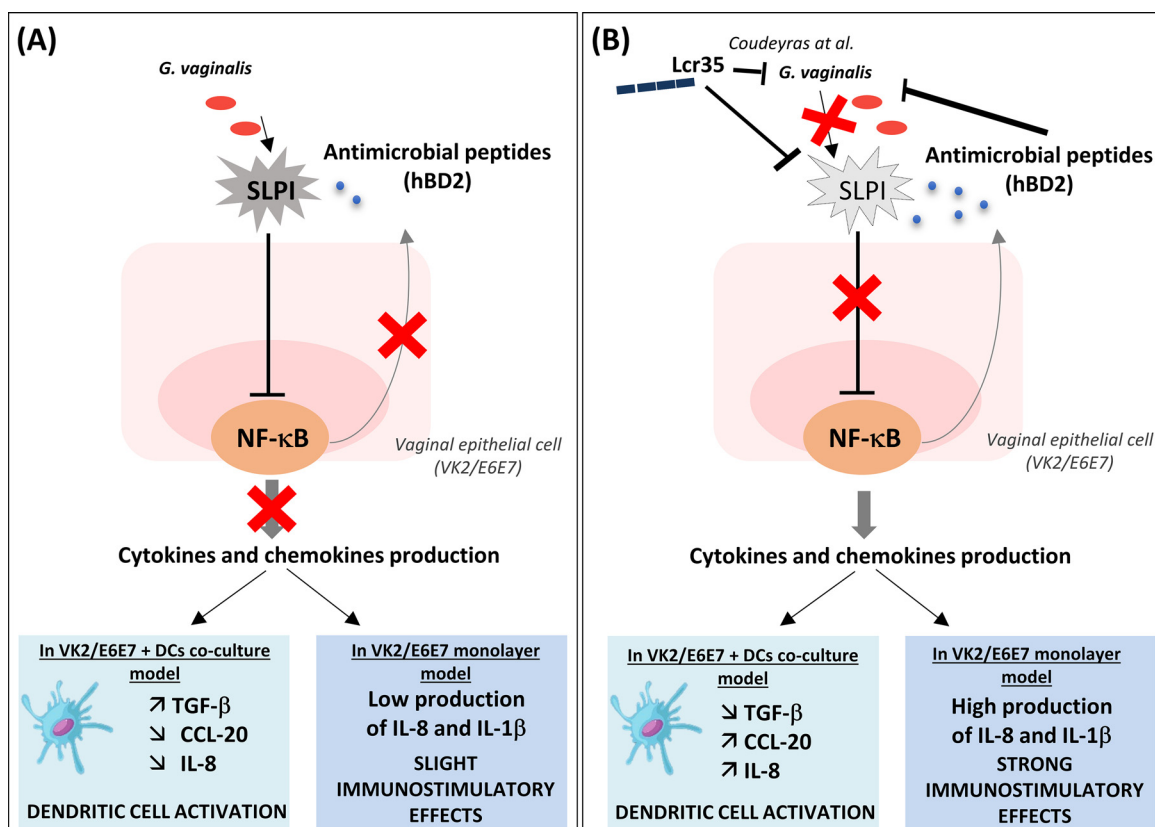
For experiments with DCs, the bacterial cells were inactivated by exposure to UV for 1 h. Successful inactivation of bacteria was assessed by plating the final suspension on agar plates.

**Cultures of epithelial cells and infections.** Human vaginal (VK2/E6E7, ATCC CRL-2616) epithelial cells were grown to confluence in keratinocyte serum-free medium (Thermo Fisher Scientific) supplemented with 0.1 ng/mL human recombinant epidermal growth factor (EGF), 0.05 mg/mL bovine pituitary extract, and additional calcium chloride at 44.1 mg/L (final concentration, 0.4 mM) (all from Sigma-Aldrich). For bacterial infection, cells were seeded in 24- or 6-well tissue culture plates (Thermo Fisher Scientific) and incubated with *G. vaginalis* and/or Lcr35 at a multiplicity of infection of 1 to 100 in complete medium in the presence or absence of 10 to 100 ng/mL recombinant human SLPI (R&D Systems, Minneapolis, MN). Following infection, cells were washed, and (i) proteins were extracted from lysed cells (3 h postinfection), and (ii) supernatants were collected (6 h postinfection) to determine cytokine secretion in the supernatants.

**In vitro differentiation of monocyte-derived dendritic cells.** DCs were generated from peripheral blood mononuclear cells (PBMC) obtained from the local French blood agency (Etablissement Français du Sang [EFS], Saint-Etienne). Blood donation requires the systematic information of the volunteers (article R.1221-5 of the Public Health Code, 01/12/2009 and 06/11/2006 decrees) and written informed consents were obtained by EFS from all donors involved in our study.

Briefly, PBMCs were isolated from the buffy coat of healthy volunteers by Ficoll-Histopaque (Sigma, Saint-Quentin Fallavier, France) density gradient centrifugation. For phenotypic assays, the PBMCs were





**FIG 4** Model for immunostimulatory effect of Lcr35 on *G. vaginalis*-infected vaginal epithelial cells. *G. vaginalis* induces the secretion of SLPI, which interacts with the NF- $\kappa$ B pathway and can inhibit immune cell recruitment (A). Lcr35 counteracts these immunosuppressive effects by SLPI inhibition, which leads to secretion of immunostimulatory cytokines and defensins and an increase in immune cell (e.g., dendritic cells) recruitment (B).

washed twice with RPMI 1640 (Cambrex Bio Science, Verviers, Belgium), and the monocytes were then isolated by adherence (2 h). The surface of the plates (75 cm<sup>2</sup> flasks; BD Falcon, Le Pont de Claix, France) was precoated with 1  $\mu$ g/mL poly-L-lysine (PLL) (Sigma) for 2 h at 4°C. Conversely, for transcriptional assays, the PBMCs were resuspended in PBS supplemented with 2% fetal calf serum (FCS) and 1 mM EDTA at a final concentration of 5.10<sup>7</sup> cells/mL, and the monocytes were purified by negative selection using the EasySep human monocyte enrichment kit as recommended by the manufacturer (StemCell Technologies, Grenoble, France). In both cases, the monocytes were then cultured for 5 days in RPMI 1640 supplemented with 1% L-glutamine (Sigma), 10% FCS (Biowest-Abcys, Paris, France), and 0.5% penicillin-streptomycin (Sigma) and containing 500 U/mL IL-4 (R&D systems, Lille, France) and 800 U/mL granulocyte-macrophage colony-stimulating factor (GM-CSF) (from R&D systems). After 3 days of incubation, one-half volume of fresh medium containing 2 $\times$  doses of IL-4 and GM-CSF was added to each well. The purity of the DCs was assessed by flow cytometry analysis using a marker highly specific of the DC lineage, the DC-SIGN, and was always above 90% (see Fig. S2A in the supplemental material).

**Vaginal epithelial and dendritic cell coculture.** VK2/E6E7 cells were seeded ( $2.5 \times 10^5$  cells/well) on a 6-well format cell culture Transwell insert (3- $\mu$ M nucleopore size; Costar, Corning Inc., USA). The medium was changed every 2 days, and cells were cultured until confluence. For coculture experiments, inserts containing confluent vaginal epithelial cell monolayers were transferred to six-well plates containing DCs ( $2 \times 10^6$  cells/well in 3 mL complete RPMI) (see Fig. S1 in the supplemental material). The apical surface of VK2/E6E7 monolayers was challenged by addition of UV-treated bacteria in the upper chamber of each well. For direct interaction, DCs ( $2 \times 10^6$  cells/well in 3 mL complete RPMI) were directly challenged by addition of the bacteria at  $1 \times 10^7$  CFU/well or LPS (10 ng/mL). All plates were incubated at 37°C in a 5% CO<sub>2</sub>/95% air atmosphere for 48 h. After incubation, cells were collected, centrifuged, and resuspended in phosphate-buffered saline (PBS) with 1% bovine serum albumin (BSA) (Sigma). Cell surfaces were stained with the appropriate fluorescence-labeled murine antibodies as follows: APC-Cy7-conjugated anti-CD14 (specific molecule of the monocytes, coreceptor of the LPS), PE-conjugated anti-CD86 (costimulatory molecule, activation marker), fluorescein isothiocyanate (FITC)-conjugated anti-CD83 (specific marker of mature DCs whose biological functions are not yet clear), PE-conjugated anti-HLA-DR, PerCP-Cy5.5-conjugated anti-DC-SIGN (CD209, member of the CLR family, specific marker of immature DCs), Alexa488-conjugated anti-MR (Mannose Receptor or CD206, member of the CLR family, specific marker of immature DCs and macrophages), and streptavidin APC-conjugated anti-TLR4 (biotin antibody, a LPS receptor with activation functions). Antibodies were obtained from BD Biosciences except anti-MR from BioLegend and anti-HLA-DR from Beckman Coulter. The cells were analyzed using BD-LSRII with FACSDiva Software (BD Biosciences) from the Centre d'Imagerie Cellulaire Santé (CICS), Université Clermont

Auvergne. Gates were set on living DCs based on forward/side scatter properties (Fig. S2A). The analysis was based on a count of 3,000 DCs. The level of staining was expressed as the mean fluorescence intensity (MFI). The culture supernatant was collected and stored at  $-20^{\circ}\text{C}$  until cytokine analysis.

**ELISA.** The human IL-1 $\beta$ , IL-8, HBD-2, SLPI, TGF- $\beta$ , IL-10, CCL20, and IL-6 cytokines were assayed in culture supernatants with enzyme-linked immunosorbent assay (ELISA) kits (BioLegend, San Diego, CA, USA) according to the manufacturers' instructions.

**Protein extraction and Western blotting.** Cells were lysed with the NE-PER nuclear protein extraction kit (Thermo Fisher Scientific) containing a protease inhibitor cocktail (Sigma-Aldrich) and phosphatase inhibitor cocktail (Sigma-Aldrich). Protein concentrations were determined by the BCA protein assay (Thermo Fisher Scientific). Western blotting was performed with 20  $\mu\text{g}$  of protein per lane. Membranes were probed with a rabbit anti-p65NF- $\kappa$ B polyclonal antibody (BioLegend; 1:1,000), or a rabbit anti-lamin B1 polyclonal antibody (BioLegend; 1:5,000) overnight at  $4^{\circ}\text{C}$ . After washes, the membranes were incubated for 1 h at room temperature with anti-rabbit horseradish peroxidase-conjugated IgG (Sigma-Aldrich). Signals were visualized with Clarity Western enhanced chemiluminescence (ECL) substrate (Bio-Rad). Quantification was performed with Image Lab 2.0 software (Bio-Rad). Most representative full, uncropped blots used for Fig. 1E are available in Fig. S6 in the supplemental material.

**Antimicrobial activity.** The antibacterial activity of HBD-2 (PeproTech) and SLPI (R&D systems) was evaluated by incubating *G. vaginalis* ( $1.10^8$  CFU/mL) or Lcr35 inoculum ( $1.10^7$  CFU/mL) with different concentrations of HBD-2 (0.25 to 100 ng/mL) or SLPI (0.2 to 10 ng/mL) for 4 h. After incubation, serial decimal dilutions of cultures were prepared in PBS and plated onto BHI or MRS agar and incubated for 48 h to 72 h. The number of CFU was assessed after incubation. The results of antibacterial activity were expressed as the percentage of growth inhibition.

**Cytotoxicity assays.** The impact of bacteria on the viability of the cultured VK2 cells was determined with the 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT) assay. After the 6 h exposure, 5  $\mu\text{g}$ /mL of 3-(4,5-dimethyl-2-thiazolyl)-2,5-diphenyl-2H-tetrazolium bromide (MTT reagent, Life Technologies) was added to each well, and the plates were incubated at  $37^{\circ}\text{C}$  for 2 h in the dark. Subsequently, the media was thoroughly removed, about 400  $\mu\text{L}$  of dimethyl sulfoxide (DMSO) was added to each well and the plate was incubated with shaking (150 rpm) at room temperature for 15 min to allow color development. The OD was measured at 540 nm.

**Statistical analysis.** One-way analysis of variance (ANOVA) (Kruskal-Wallis with Dunn's multiple comparison test) tests were performed with GraphPad Prism 6 software. A *P* value of less than 0.05 was considered statistically significant. \*, *P* < 0.05; \*\*, *P* < 0.01; \*\*\*, *P* < 0.001. Error bars depict mean  $\pm$  standard error of the mean (SEM).

## SUPPLEMENTAL MATERIAL

Supplemental material is available online only.

**SUPPLEMENTAL FILE 1**, PDF file, 1.5 MB.

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We declare that no conflict of interest exists.

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