The lectin-like protein 1 in *Lactobacillus rhamnosus* GR-1 mediates tissue-specific adherence to vaginal epithelium and inhibits urogenital pathogens

Mariya I. Petrova\(^1,2\), Elke Lievens\(^1,2\), Tine L. A. Verhoeven\(^3\), Jean M. Macklaim\(^3,4\), Gregory Gloor\(^4\), Dominique Schols\(^5\), Jos Vanderleyden\(^1\), Gregor Reid\(^3,4\) & Sarah Lebeer\(^4,2\)

The probiotic *Lactobacillus rhamnosus* GR-1 has been documented to survive implantation onto the vaginal epithelium and interfere with urogenital pathogens. However, the molecular mechanisms involved are largely unknown. Here, we report for the first time the construction of dedicated knock-out mutants in *L. rhamnosus* GR-1 to enable the study of gene functions. In a search for genes responsible for the adherence capacity of *L. rhamnosus* GR-1, a genomic region encoding a protein with homology to lectin-like proteins was identified. Phenotypic analyses of the knock-out mutant of *L. rhamnosus* GR-1 revealed a two-fold decreased adhesion to the vaginal and ectocervical epithelial cell lines compared to wild-type. In contrast, the adhesion to gastro-intestinal epithelial (Caco2) and endocervical cell lines (HeLa and End1/E6E7) was not drastically affected by the mutation, suggesting that the LGR-1_Llp1 lectins mediates tissue tropism. The purified LGR-1_Llp1 protein also inhibited biofilm formation and adhesion of uropathogenic *Escherichia coli*. For the first time, an important role for a novel lectin-like protein in the adhesion capacity and host cell-specific interaction of a vaginal probiotic *Lactobacillus* strain has been discovered, with an additional role in pathogen inhibition.

Probiotics are defined as "live microorganisms which, when administrated in adequate amounts, confer a health benefit on the host"\(^1\). *Lactobacillus rhamnosus* GR-1 is a well-known probiotic strain isolated from a healthy female urethra. This strain has been shown to adhere well to urogenital epithelial and vaginal cells in *vitro*\(^2\), and to temporarily colonize the human vagina\(^3\) and intestine following oral uptake\(^4,5\). The fact that oral application of *L. rhamnosus* GR-1 can result in vaginal colonization\(^3–5\) is of interest in view of the natural ascension of lactobacilli from the gastro-intestinal tract to the vagina. The ability of *L. rhamnosus* GR-1 to inhibit the growth and adhesion of urogenital pathogens is believed to be important in its probiotic activity. This activity is well documented *in vitro* for pathogens such as *Escherichia coli*\(^6,7\), *Enterococcus*, *Gardnerella vaginalis*, *Atopobium vaginae* and *Candida albicans*\(^8\). However, the molecular mechanisms by which this vaginal probiotic strain interacts with pathogenic bacteria and host cells are largely unknown, despite this strain being a model for vaginal probiotics.

Studies of gastro-intestinal probiotic lactobacilli indicate key roles for surface molecules in host interactions\(^11\). An interesting class of surface molecules are the lectins, i.e. proteins that bind carbohydrates with high specificity, but do not modify them. Lectins are well characterized in animals and plants\(^12,13\), while the information in bacteria is relatively poor. The best documented bacterial lectins are present in uropathogenic bacteria, such as the FimH adhesin located at the tip of type 1 pili of the uropathogenic *E. coli* (UPEC) where they play a role in attachment to urothelium by binding to mannosylated glycoreceptors\(^14\).

\(^1\)KU Leuven, Centre of Microbial and Plant Genetics, Leuven, Belgium. \(^2\)University of Antwerp, Department of Bioscience Engineering, Antwerp, Belgium. \(^3\)The Lawson Health Research Institute London, Canada Research and Development Centre for Probiotics, London, ON, Canada. \(^4\)University of Western Ontario, London, ON, Canada. \(^5\)KU Leuven, Rega Institute for Medical Research, Leuven, Belgium. Correspondence and requests for materials should be addressed to M.I.P. (email: mariya.petrova@biw.kuleuven.be) or S.L. (email: sarah.lebeer@uantwerpen.be)
In the present study, we aimed to better understand the molecular factors that contribute to \textit{L. rhamnosus} GR-1 vaginal adherence, immunomodulation and pathogen inhibition. Because of the presence of various glycans on the vaginal mucosa and surfaces of pathogenic microorganisms, we investigated whether lectin-like proteins could play a role in adhesion and immunomodulation capacity of \textit{L. rhamnosus} GR-1, and in its capacity to prevent uropathogenic \textit{E. coli} infections.

**Results**

**Identification and annotation of the \textit{LGR1} \_llp1 gene encoding the lectin-like protein 1.** To identify genes encoding putative lectin-like proteins, the draft genome sequence of \textit{L. rhamnosus} GR-1 was screened for the presence of open reading frames (ORFs) containing a lectin L-type domain (PF00139). A 4060 bp genomic region was identified (Fig. 1a) containing a 2040 bp sequence encoding a polypeptide of 680 amino acid residues, with an \textit{N}-terminal Legume-type (L-type) lectin domain (PF00139) and a \textit{C}-terminal WxL domain (PF13731) (Fig. 1b). We annotated this gene sequence as \textit{LGR1} \_llp1, encoding the putative lectin-like protein 1.

The L-type lectin domain represents approximately 250 amino acid residues in length and is found in several cell surface proteins of Gram-positive bacteria\(^{15}\). The \textit{C}-terminal WxL domain, detected also in proteins from several other Gram-positive bacteria, is suggested to be responsible for the non-covalently anchoring of proteins to the microbial surface, possibly by interaction with the peptidoglycan\(^{16}\).

**LGR1 \_llp1 mediates tissue-specific adhesion of \textit{L. rhamnosus} GR-1 to vaginal epithelial cells.** In order to develop a DNA transformation protocol for \textit{L. rhamnosus} GR-1, two different plasmids were used: the \textit{Lactobacillus} genome integrating pEM40 vector\(^{17}\) and the self-replicative \textit{Lactobacillus} pLAB1301 vector\(^{18}\). The electroporation protocol for \textit{L. rhamnosus} GG\(^{19}\) was used as a starting point. Both plasmids could be transformed with a similar efficiency as for \textit{L. rhamnosus} GG, with an electroporation efficiency of ca. $\approx 1.7 \times 10^6$ CFU/$\mu$g DNA for pLAB1301 and $\approx 1 \times 10^5$ CFU/$\mu$g DNA for pEM40. This is in agreement with the latter being an integrative plasmid in \textit{L. casei} and \textit{L. rhamnosus} strains\(^{17}\), thus requiring an integration step in the genome in addition to efficient transformation, resulting in lower efficiency.

By using the optimized electroporation protocol, a knock-out \textit{LGR1} \_llp1 mutant was constructed by double homologous recombination. The correct allelic replacement event in \textit{LGR1} \_llp1 mutant CMPG10744 with tetracycline resistant antibiotic cassette was confirmed by PCR and Southern hybridization. Subsequently we investigate the role of Llp1 protein in the adhesion capacity of \textit{L. rhamnosus} GR-1. The \textit{LGR1} \_llp1 mutant CMPG10744 showed a significant ($p = 0.0006$) ca. two-fold reduction in adhesion capacity to the vaginal epithelial cell line VK2/E6E7 compared to \textit{L. rhamnosus} GR-1 wild type (Fig. 2a). In addition, the \textit{LGR1} \_llp1 mutant CMPG10744 showed also a significant ($p = 0.04$) reduction in adhesion capacity with 26% to the ectocervical epithelial cells Ect/E6E7, which is also nonkeratinized stratified squamous epithelium (Fig. 2a). To confirm the genotype-phenotype relation for the \textit{LGR1} \_llp1 gene, mutant CMPG10744 was subsequently complemented by re-introducing the \textit{LGR1} \_llp1 gene. This complemented strain CMPG10746 showed complete restoration of the...
adhesion phenotype, reaching the same adhesion capacity levels as the wild type (Fig. 2a). We subsequently investigated whether the LGR1_Llp1 protein is also involved in the adhesion capacity to other, simple columnar epithelial cell lines such as the endocervical End1/E6E7, the model cervical cancer Hela cell line and the colon carcinoma Caco-2 cell line. Interestingly, the adhesion capacity of LGR1_Llp1 CMPG10744 was only slightly but not significantly reduced for the intestinal Caco2 cells line, the endocervical End1/E6E7 cell line and the Hela cell line as compared to the L. rhamnosus GR-1 wild type (Fig. 2a). This suggests that LGR1_Llp1 mediates tissue tropism with specificity towards vaginal epithelial and stratified squamous epithelial cells.

To confirm tissue specific adhesion by LGR1_Llp1, the lectin domain of LGR1_Llp1 was expressed in E. coli and purified. Subsequently, the binding of the FITC-labelled lectin domain of LGR1_Llp1 was explored by fluorescence microscopy for VK2/E6E7 and Caco-2 cells grown on cover slips. In contrast to Caco-2 cells, the lectin domain recognized and bound strongly to the VK2/E6E7 epithelial cells (Fig. 2b), suggesting indeed that the lectin plays a role in the well-documented adhesion capacity of L. rhamnosus GR-1.

Indication for a lectin-like role for Llp1 in adhesion of GR-1 to mannosylated vaginal epithelial cells. To validate that the adhesion role of LGR1_Llp1 is due to its sugar-binding capacity, we performed several complementary experiments. First, we designed a competition experiment with lectins with known specificity. After preincubation of VK2/E6E7 cells with the α(1,6)-α(1,3) mannos- specific lectin ConA, α(1,2) mannosse- specific AH and GRFT, but not with (GlcNAc)n specific UDA and Nictaba, a significant ca. two-fold reduction in the adhesion capacity of L. rhamnosus GR-1 wild type was observed (Fig. 2c). On the other hand, the adhesion capacity of the LGR1_Llp1 mutant CMPG10744 did not vary and was always ca. 40% of wild-type. These results indicate that the ConA, AH and GRFT block important mannosylated receptors on the cell surface of VK2/E6E7 and that LGR1_Llp1 possibly competes with these lectins for the same receptors on these vaginal epithelial cells.

Subsequently, agglutination assays with mannan-containing yeasts were performed to confirm the mannosse specificity of LGR1_Llp1. The FITC-labelled LGR1_Llp1 did not aggregate the yeast Saccharomyces cerevisiae BY4741 (Fig. 3a), but did lead to aggregation of the important opportunistic pathogen Candida albicans SC5314.
(Fig. 3b) as compared to the negative control. The affinity of the purified lectin domains towards various sugars was subsequently further examined by pull-down sugar-binding assays with beads coated with purified mannann, D-mannose (lane 3), D-glucose (lane 4), D-fucose (lanes 5) GluNAc (lane 6) or not coated with any sugar (lane 1, used as negative control). (d) Mammalian glycan array used to determine the carbohydrate binding specificity of the lectin domain of Llp1. Glycans to which the FITC labelled lectin domains show the strongest binding are depicted.

**Figure 3.** Carbohydrate specificity of the lectin domain of LGR1_Llp1. (a,b) Fluorescent images of the agglutination assay of *S. cerevisiae* BY4741 and *C. albicans* SC5314 in the presence of the FITC labelled lectin domain of LGR1_Llp1. (c) Proteins that bound to sugar-coated Sepharose beads were separated by SDS-PAGE. Sepharose beads were coated with mannan (lane 2), D-mannose (lane 3), D-glucose (lane 4), D-fucose (lanes 5) GluNAc (lane 6) or not coated with any sugar (lane 1, used as negative control). (d) Mammalian glycan array used to determine the carbohydrate binding specificity of the lectin domain of Llp1. Glycans to which the FITC labelled lectin domains show the strongest binding are depicted.

LGR1_Llp1 only moderately induces immune responses in vaginal epithelial cells. The host-cell signalling interaction capacity of LGR1_Llp1 in vaginal VK2/E6E7 cells was then investigated by cytokine arrays that explore the upregulation and downregulation of the mRNA expression of selected genes. Interestingly, the mRNA expression level of the majority of cytokines and chemokines tested was unchanged after incubation with the bacterial strains or with the purified lectin domain of LGR1_Llp1 (Fig. 4). However, the lectin domain of LGR1_Llp1 appeared to induce mRNA levels of the anti-inflammatory cytokines IL13 and IL10 and the pro-inflammatory IL27 by more than four-fold. Of note, IL13 mRNA expression was also upregulated by the wild
LGR1_Llp1 inhibits adhesion and biofilm formation of the key urogenital pathogen *E. coli* UTI89. Since pathogen inhibition is a key hallmark of probiotic bacteria, whether LGR1_Llp1 could block adhesion and biofilm formation of UPEC was investigated. When *E. coli* UTI89 was pre-incubated with the lectin domain of LGR1_Llp1 at concentration of 50 μg/ml, it caused a significant almost two-fold reduction in the adhesion capacity of *E. coli* UTI89 to VK2/E6E7 cells (Fig. 5a), indicating a role in pathogen exclusion from host cells. When the purified lectin domain of LGR1_Llp1 was added at the beginning of biofilm development at concentration 50 and 200 μg/ml, a reduction to 17% and 8% of the control situation was observed respectively (Fig. 5b). When the lectin domain of LGR1_Llp1 at 50 μg/ml was added after 1.5 or 24 h, a reduction to 18% and 43% respectively in the biofilm formation was still observed compared to the control (Fig. 5b). The absolute CFU counts of *E. coli* UTI89 in the formed biofilms were in agreement. The lectin domain of LGR1_Llp1 caused then a significant reduction of *E. coli* UTI89 biofilm to ca. 40% (for 50 μg/ml concentration) and 30% (for 200 μg/ml concentration) of the undisturbed pathogenic biofilms (Fig. 5c). Since the lectin domain is part of a full-length protein, we subsequently investigated if the full length LGR1_Llp1 (which is more difficult to express and purify in large amounts) also possesses an anti-biofilm activity. Indeed, when the full-length LGR1_Llp1 was added at the beginning of the biofilm formation at concentration of 50 μg/ml, also a ca. two-fold reduction was observed.
(Fig. 5d). Given the capacity of the lectin to prevent *E. coli* UTI89 biofilms, bioscreens were performed to investigate whether the lectin also shows an antimicrobial effect on growth in suspension. Interestingly, no inhibitory effect on planktonic growth was observed (Fig. 5e).

To explore how the lectin domain structurally interferes with *E. coli* UTI89 biofilm formation, microscopic analyses were performed. Addition of FITC-labeled lectin domain of LGR1_Llp1 at the onset of the biofilm at concentration 50 μg/ml resulted in the formation of loose biofilms with large holes (Fig. 5f), compared to the control without lectins, which formed dense biofilm (Fig. 5g). The lectin domain of LGR1_Llp1 was also shown to be clearly distributed across the biofilm (Fig. 5e).

**Figure 5.** Role of LGR1_Llp1 in inhibiting adhesion and biofilm formation of *E. coli* UTI89. (a) Effect of the lectin domain (LD) of LGR1_Llp1 on the adhesion of *E. coli* UTI89 after pre-incubating the bacterial cells with the lectin. (b) Effect of the lectin domain of *L. rhamnosus* GR-1 on *E. coli* UTI89 biofilms. The purified lectin domain (LD) of LGR1_Llp1 was added after 0, 1.5 and 24 hours to the biofilms. (c) Biofilm formation of *E. coli* UTI89 based on absolute cell counts. Biofilms were grown in 1/20 TSB medium without (control) or with 50 μg/ml and 200 μg/ml of LD of LGR1_Llp1. (d) Effect of full length (FL) lectin (50 μg/ml) on *E. coli* UTI89 biofilms added at zero-time point to the biofilms. (e) Growth of *E. coli* UTI89 in the presence of lectin domain of LGR1_Llp1 added at concentrations of 50 μg/ml and 200 μg/ml. The error bars represent standard deviations of three independent experiments. The dataset comparisons are considered significant (p < 0.05 indicated with one asterisk in the picture, p < 0.01 indicated with two asterisks or p < 0.001 indicated with three asterisks). (f) Biofilms of *E. coli* UTI89 grown with 50 μg/ml of FITC-labeled lectin domain of LGR1_Llp1 and (g) alone without adding lectin in 1/20 TSB medium (negative control). Holes in biofilms are indicated with arrows.
LGR1_Llp1 inhibits adhesion but not biofilm formation of Staphylococcus aureus. As for E. coli UTI89, the effect of the lectin domain of LGR1_Llp1 against the adhesion capacity of S. aureus Rosenbach and S. aureus SH1000 to VK2/E6E7 cells was investigated by pre-incubating the pathogenic bacterial cells with 50 μg/ml of the lectin domain. A two-fold reduction in the adhesion of S. aureus Rosenbach was observed compared to the control treatment (Fig. 6a). No significant differences were notable for LGR1_Llp1 on the biofilm capacity of S. aureus Rosenbach and S. aureus SH1000 as compared to the negative control (Fig. 6b).

LGR1_Llp1 increases biofilm formation of vaginal Lactobacillus species. Potential new anti-bacterial agents should not affect the beneficial Lactobacillus-dominated vaginal microbiota. Of interest, biofilm formation of the vaginal Lactobacillus strains tested here was shown to increase in the presence of LGR1_Llp1. Biofilm formation of L. reuteri RC-14 and L. gasseri ATCC 33323 increased significantly ca. two-fold and 2.5-fold respectively when LGR1_Llp1 was added in the initial steps of the biofilm formation (Fig. 6c).
Furthermore, a significant 1.6 to 1.7-fold increase in biofilm formation was observed for *L. jensenii* ATCC 25258, *L. crispatus* NCIMB 4505 and *L. plantarum* CMPG5300 respectively. Finally, the biofilm formation of *L. rhamnosus* GR-1 itself was also significantly increased with a 1.25-fold when LGR1_Llp1 was added (Fig. 6c).

**Discussion**

The vaginal ecosystem represents a special niche for the application of probiotic lactobacilli, because of the natural dominance of *Lactobacillus* species in healthy reproductive-age females and the link between *Lactobacillus* disappearance and disease phenotypes20,21. Therefore, in this study we biochemically and genetically identified and characterized a novel lectin-like protein LGR1_Llp1 from the vaginal probiotic strain *L. rhamnosus* GR-1. The lectin was shown to mediate vaginal niche-related functions for *L. rhamnosus* GR-1, including tissue tropism-mediated adhesion to vaginal and ectocervical epithelial cells and immunomodulation. In addition, the lectin domain showed to have a unique capacity to inhibit adhesion and biofilm formation of vaginally-associated pathogens, without affecting the normal *Lactobacillus*-dominated microbiota.

First, we were able to successfully optimize a protocol for electroporation for the construction of a dedicated knock-out mutant in *L. rhamnosus* GR-1. This report describes the development of genetic tools that permit mechanistic investigations of the probiotic strain *L. rhamnosus* GR-1. Phenotypic analyses of knock-out mutants have the advantage that the function of cell-surface molecules such as adhesions and lectins can be studied in situ on the surface of live bacteria in their physiological context. An isogenic LGR1_llp1 mutant was constructed in a gene encoding a putative lectin-like protein. This LGR1_llp1 mutant showed a significantly reduced adhesion capacity to vaginal and ectocervical epithelial cells, but not to endocervical, cervical carcinoma cells and colon carcinoma cells. Similar results were also observed when we investigated the binding capacity of the FITC-labelled LGR1_llp1. This suggests that the LGR1_llp1 protein provides tissue tropism to *L. rhamnosus* GR-1, likely determined by the presence of different glycosylated structures on the cell membrane of these epithelial cells. The fact that LGR_Llp1 preferentially binds to sugar residues on vaginal epithelial cells, and not intestinal cells, may promote the natural ascension and easier transit from the gastrointestinal tract to the vaginal epithelium by this vaginal strain. Of note, both Hela and Caco-2 cell lines were recently shown to have similar complex glycosylation profile22. The glycan structure of surface molecules of VK2/E6E7 cells is not yet well studied. However, previous studies suggest that glucose, mannose and glucosamine are important sugars in the vaginal niche23,24.

To substantiate the lectin-like role of LGR1_Llp1 in the adhesion capacity of *L. rhamnosus* GR-1, competition experiments with lectins with well-known specificity for adhesion to vaginal epithelial cells were performed. These experiments suggest that LGR1_Llp1 on the cell surface of *L. rhamnosus* GR-1 is involved in the recognition of similar glycosylated receptors on the surface of VK2/E6E7 cells as the tested α1,6 Man, α1,3 Man specific ConA and α1,2 Man specific AH and GRFT lectins. However, studying the exact sugar specificity of LGR1_llp1 is difficult in the context of the bacterial cells, because various other cell surface molecules may interfere. Therefore, we also determined the sugar specificity of the purified LGR1_llp1 lectin domain by using yeast agglutination assay, combined with Sepharose beads binding assays and dedicated mammalian glycan array screenings. All these experiments indicate specific binding of LGR1_Llp1 to complex N-glycan structures, in agreement with recent studies on plant lectins showing that their specificity is complex and cannot be merely described by single sugar monomers25. The glycan array especially revealed a strong binding of LGR1_Llp1 to agglutinate and saliva26,27. In addition, LGR1_Llp1 showed a capacity to bind to the glycosphingolipid GQ1 receptor expressed on various epithelial cells, such as gastrointestinal epithelium, vaginal epithelium, embryonic tissues and saliva26,27. In addition, LGR1_llp1 showed also a capacity to bind to the glycosphingolipid GQ1 receptor (Neu5Aco2-8Neu5Aco2-3Gal31-3GalNAc31-4(Fuc1-3)GlcNAc, as confirmed with glycan array29).

In addition to the surface glycan binding capacity, LGR1_Llp1 mediated the adhesion capacity of *L. rhamnosus* GR-1, as we also investigated its role in modulating cytokine responses in VK2/E6E7 cells. Probiotic strains have been reported to promote health by stimulating the host immune response30, but such stimulation of vaginal host cells is poorly documented31. We did not observe strong cytokine responses when VK2/E6E7 epithelial cells were treated with *Lactobacillus* strains or purified lectin protein. *L. rhamnosus* GR-1 only induced a modest 3 fold upregulation of IL13 and IL17. Of interest, IL17 has been previously suggested to control *C. albicans* vulvovaginal infections, by stimulating the production of antimicrobial peptides by vaginal epithelial cells31. The LGR1_Llp1 protein also moderately induced mRNA levels of the anti-inflammatory cytokines IL10 and showed a strong downregulation of IL7 mRNA. Of interest, IL-17 has been detected in the plasma of HIV-infected patients and possibly facilitates HIV-1 transmission32. Nevertheless, the exact cytokine signalling events in more physiological relevant conditions remain to be further substantiated.
In addition to adhesion and immunomodulating effects, we also explored the direct role of the LGR1_Llp1 lectin in pathogen exclusion, considering the fact that UPEC also shows tissue tropism for the vaginal epithelium via dedicated lectins such as FimH. The lectin domains of LGR1_Llp1 were found to have a major impact on the adhesion capacity and biofilm development of the model strain E. coli UTI89. The adhesion capacity of this UPEC was almost reduced by half when the lectins were added in 50 μg/ml concentration. The biofilm formation was even ca. 10-fold reduced, when the lectin domain was added at the onset of the biofilm, after 1.5 h or after 24 h of biofilm formation, indicating that the lectin domain of LGR1_Llp1 can both prevent and disrupt established biofilms of this important uropathogenic pathogen, by forming loose biofilm structures. This supports previous reports of L. rhamnosus GR-1 inhibiting the growth and adhesion of UPEC, but the exact molecular mechanism was not known till now. The localization of LGR_Llp1 within UPEC biofilms suggests that the lectin domain interact with components of the UPEC biofilm matrix. This matrix contains the polysaccharides cellulose (3-1, 4-D-glucose polymer) and colanic acid (heteropolysaccharide of glucose, galactose, fucose and glucuronic acid) in most E. coli strains. Of note, these sugars closely resemble the composition of the complex N-type glycans to which LGR1_Llp1 shows specificity. Therefore, LGR1_Llp1 might be able to bind to these exopolymeric substances and in this way destabilize the biofilm structure. This would explain the observed holes in the biofilms and the unstable biofilms formed by UPEC after adding the lectin. The clear biofilm-inhibiting effect is worth further exploration, given the prevalence of problems associated with biofilms and the increased resistance of various bacteria against antibiotics. Furthermore, LGR1_Llp1 reduced the adhesion capacity of S. aureus Rosenbach by half, albeit not its biofilm formation. Nevertheless, taken together, our results show the inhibitory capacity of a bacterial, and more specifically a Lactobacillus lectin, against urogenital pathogens. It is important to note that LGR1_Llp1 did not interfere with the biofilm formation capacity of several vaginal Lactobacillus isolates tested, which are of crucial importance to keep the homeostasis of the vaginal environment. On the contrary, LGR1_Llp1 was even found to increase their capacity to form biofilms, suggesting that LGR1_Llp1 could play a role in maintaining a normal Lactobacillus-dominated vaginal microbiota or supporting their re-establishment after infections.

In conclusion, the current report describes the optimization of genetic tools for the clinically well-documented vaginal probiotic strain L. rhamnosus GR-1, as well as the identification, annotation and functional analysis of LGR1_llp1 gene as the first described adaptation factor of L. rhamnosus GR-1 involved in vaginal adhesion and immunomodulation. The prominent inhibitory effects of LGR1_Llp1 on biofilm formation and adhesion of UPEC and S. aureus holds potential for further applications related to pathogen exclusion, either alone or, in combination with other antibacterials and in the context of the probiotic L. rhamnosus GR-1.

**Material and Methods**

**Bacterial strains, plasmids and growth conditions.** The bacterial strains and plasmids used in this study are listed in Table 1. L. rhamnosus GR-1 wild type and the corresponding mutants were routinely grown non-shaking in de Man Rogosa Sharpe (MRS) medium (Difco) at 37°C. Escherichia coli strains were grown in Luria Bertani (LB) medium (1% NaCl, 1% peptone, 0.5% yeast extract) with aeration at 37°C. Saccharomyces cerevisiae was grown in YPD medium (1% yeast extract, 2% peptone, 2% glucose) with aeration at 30°C. If required, antibiotics were used at following concentrations: 10 μg/ml tetracycline, 100 μg/ml ampicillin, 5 μg/ml (for L. rhamnosus GR-1) or 130 μg/ml (for E. coli) erythromycin. The antimicrobial effect of the lectin-like protein on E. coli UTI89 growth was assessed by using 100-well microtiter plates (Honeycomb, Oy Growth Curves Ab Ltd.) (Bioscreen) as previously described.

**DNA manipulations.** Routine molecular biology techniques were performed as described before. PCR primers used in this study were purchased from Integrated DNA Technologies (IDT) (Belgium) (Table 2). Enzymes for molecular biology were purchased from New England Biolabs and used according to the suppliers’ instructions. Plasmid DNA from E. coli was purified using QIAGEN miniprep kits. Chromosomal DNA from L. rhamnosus GR-1 was isolated as previously described.

**Electroporation in L. rhamnosus GR-1.** Electroporation of the replicating and integrating plasmids (Table 1) into L. rhamnosus GR-1 was performed as previously reported for L. rhamnosus GG with minor modifications. Briefly, serial dilutions (10^6 to 10^8 fold) were made from an overnight culture of L. rhamnosus GR-1 wild type into freshly prepared pre-warmed MRS medium supplemented with 2% glycine. L. rhamnosus GR-1 was then incubated without aeration at 37°C. After overnight growth, ~5 ml of the selected dilution in the exposure buffer (fresh MRS medium supplemented with 2 mM CaCl2 and 2 mM MgCl2) was added and resuspended in 1 ml of the same buffer. Electroporation was performed in the MicroPulser electroporator (BioRad) (2 mm cuvettes) under the following conditions: 2 kV, 25 μF, 200 Ω. Subsequently, 1 ml of the regeneration buffer (fresh MRS medium supplemented with 2 mM CaCl2 and 2 mM MgCl2) was added and the cells were incubated for 3 hours at 37°C without aeration. Finally, the cells were plated on MRS plates with appropriate antibiotics followed by incubation for 72 hours at 37°C.

**Identification and sequence analysis of the L. rhamnosus GR-1 LGR1_llp1 gene.** The draft genome sequence of L. rhamnosus GR-1 was mined for the presence of putative lectin-like proteins by BLAST using the lectin-like protein 1 (Llp1) of L. rhamnosus GG. This resulted in the identification of a genomic region encoding a putative lectin-like protein of which the putative gene sequence was designated LGR1_llp1. This genomic region of LGR1_llp1 and its flanking regions was submitted to Genbank under the ID accession number: KF295830.
the EcoRI site. Subsequently, HR2 was amplified by PCR using primers Pro7468 and Pro7469 and cloned into pApR, ampicillin resistance, KmR, kanamycin resistance.

as previously reported for L. rhamnosus the Table 1. L. rhamnosus Construction of the GR-1 plasmid is derived from pFAJ530140 by ligation of a tetracycline resistance gene from PCR. Primers were designed with restriction sites for the corresponding enzymes at the 5′ by PCR using primers Pro7466 and Pro7467 and subsequently cloned into pCMPG10205. The pCMPG10205 Cloning vector; pUC18 containing tetracycline resistant pCMPG10208 pLAB1301 derivative driven by promoter of the pET 28 a( pEM40 pCMPG10775 for secretion of the N-His6 tagged full domain with 22 amino acids extension of the Lectin-like domain of L. rhamnosus GR-1, KmR This study

Table 1. Strains and plasmids used in this study. TetR, tetracycline resistance; EmR, erythromycin resistance; ApR, ampicillin resistance, KmR kanamycin resistance.

Construction of the L. rhamnosus GR-1 LGR1_llp1 mutant pCMPG10744. To determine the role of the LGR1_llp1 gene, a corresponding knock-out mutation was constructed by double homologous recombination as previously reported for L. rhamnosus GG.39 Briefly, two regions of ~1000 bp upstream and downstream of llp1 gene, designated respectively as homologous region 1 (HR1) and homologous region 2 (HR2), were amplified by PCR. Primers were designed with restriction sites for the corresponding enzymes at the 5′ end. HR1 was amplified by PCR using primers Pro7466 and Pro7467 and subsequently cloned into pCMPG10205. The pCMPG10205 plasmid is derived from pFAJ530140 by ligation of a tetracycline resistance gene from L. plantarum MD5057 in the EcoRI site.41 Subsequently, HR2 was amplified by PCR using primers Pro7468 and Pro7469 and cloned into the plasmid containing already HR1, resulting in plasmid pCMPG10743. This suicide vector was transferred to highly competent L. rhamnosus GR-1 wild type by electroporation as described above. Putative knock-out mutants resulting from double homologous recombination were selected based on resistance to tetracycline and sensitivity to erythromycin. Confirmation of DNA recombination was performed by PCR using primers Pro8018

| Strain/plasmid | Relevant genotype/description | Reference or source |
|----------------|------------------------------|---------------------|
| E. coli strains |                             |                     |
| Top10F | F’ (lacP, Tn5) mcrA Δ(mrr-hsdRMS-mcrBC) F80LacZΔM15 ΔlacX74 deoR recA1 araD139 Δ (ara-leu)7697 galU galK rpsL(56) endA1 supG | Invitrogen |
| BL21 (DE3) | E. coli B F- dcm ampT haSl (rpsL-m15) gal λ (DE3) | Invitrogen |
| E. coli UT189 | Wild type, clinical isolate | 55 |
| C. MCMG10774 | E. coli BL21 (DE3) carrying the overexpression plasmid pCMPPG10747 for secretion of the N-His6 tagged Llp1 full protein of L. rhamnosus GR-1, KmR | This study |
| C. MCMG10775 | E. coli BL21 (DE3) carrying the overexpression plasmid pCMPPG10775 for secretion of the N-His6 tagged Llp1 lectin-like domain from L. rhamnosus GR-1, KmR | This study |
| L. rhamnosus GR-1 strains |                             |                     |
| Wild type | Human urethra isolate | ATCC 5582 |
| C. MCMG10744 | lllp1 knock-out mutant of L. rhamnosus GR-1; lllp1::tetR | This study |
| C. MCMG10746 | C. MCMG10744 mutant complemented by electroporation of pCMPPG10746 | This study |
| Lactobacillus strains |                             |                     |
| L. reuteri RC-14 ATCC 55845 | Wild-type, female urethra isolate | 56 |
| L. crispatus NCIMB 4505 | Wild-type, human vaginal isolate | 57 |
| L. jensenii ATCC 25258 | Wild-type, human vaginal isolate | 58 |
| L. gasseri ATCC 33323 | Wild-type, human vaginal isolate | 58 |
| L. plantarum CMPG5300 | Wild-type, human vaginal isolate | 59 |
| Other strains |                             |                     |
| S. cerevisiae BY 4741 | MATa; his3∆1; leu2∆0; met15∆0; ura3∆0 | 60 |
| C. albicans SC5314 (ATCC MYA-2876) | Wild type, human clinical isolate | 61 |
| S. aureus SH1000 | rsbU positive derivative of S. aureus 8325-4 | 62 |
| S. aureus Rosenbach (ATCC 33591) | Wild type, clinical isolate | ATCC |
| Plasmids |                             |                     |
| pLAB1301 | E. coli – Lactobacillus shuttle vector, ErmR, AmpR | 18 |
| pEM40 | pUC19E-derived integration vector (attB located at the 3′ end of the tRNA locus) containing a 1.6-kb int-attP cassette of phage A2; ErmR, AmpR | 17 |
| pET 28a (+) | KmR, T7 lac, N and C-terminal His Tag | Novagen |
| pCMPG10205 | Cloning vector; pUC18 containing tetracycline resistant cassette from pGR13 in the BspHI site | 39 |
| pCMPG10208 | pLAB1301 derivative driven by promoter of the dlt operon of L. rhamnosus GG, AmpR, ErmR | 63 |
| pCMPG10743 | pCMPPG10205 derivative used to inactivate the lllp1 gene by insertion of a tetR marker via double homologous recombination (for details, see text) | This study |
| pCMPG10746 | pCMPPG10208 derivative containing the lllp1 gene (2040 bp) in the Xmal/SacI site, AmpR, ErmR | This study |
| pCMPPG10774 | pET28 (a +) derivative carrying the N-His6 tagged lllp1 gene of L. rhamnosus GR-1 in the SalI/NotI site, KmR | This study |
| pCMPPG10775 | pET28 (a +) derivative carrying the N-His6 tagged lectin-like domain with 22 amino acids extension of the lllp1 gene of L. rhamnosus GR-1 in the SalI/NotI site, KmR | This study |
and Pro8019 and by Southern hybridization using primers Pro7467 and Pro8018. One colony showing the correct homologous recombination event was selected for further analysis and designated as CMPG10744.

CMPG10744 was complemented by electroporation of pCMPG10746, containing the lpl1 gene, amplified by PCR with primers Pro8675 and Pro8676, ligated downstream of the dlt promoter in the vector pCMPI0208, resulting in the complemented strain CMPG10746.

**Construction of overexpression constructs in E. coli BL21 (DE3).** For heterologous expression of LGR1_lpl1 in E. coli, a pET (a (+) system (Novagen) was used. The LGR1_lpl1 gene and the corresponding lectin domain from L. rhamnosus GR-1 were amplified by PCR (Table 2) and subsequently cloned into the pET-28 a (+) vector resulting in plasmids pCMPI0774 and pCMPI0775 respectively. Competent E. coli strain BL21 (DE3) cells were transformed with plasmid pCMPI0774 and pCMPI0775, resulting in strain CMPG10774 and CMPG10775 respectively. The recombinant E. coli strains for overexpression of putative lectin-like proteins were subjected to recombinant protein expression and subsequent detection by SDS-PAGE electrophoresis and Western blot.

**Production of recombinant lectins and lectin domains and their purification.** The recombinant E. coli BL21 (DE3) strains expressing the full length lectin or the corresponding lectin domain were grown overnight in LB with 50 μg/ml Kanamycin. Each culture was diluted 100-fold in LB with Kanamycin and grown for 2 to 3 hours at 37 °C under shaking conditions until an optical density (OD) (595 nm) between 0.3 and 0.4 was reached. Then the production of recombinant protein was induced with 1 mM isopropyl β-D-thiogalactopyranoside (IPTG) and the cultures were incubated at 25 °C under shaking until an OD of 0.8 to 1 was reached. Subsequently, the cells were pelleted and suspended in non-denaturing lysis buffer (NaH2PO4 50 mM, NaCl 300 mM, imidazole (IPTG) and the cultures were incubated at 25 °C under shaking until an OD of 0.8 to 1 was reached. Subsequently, the cells were pelleted and suspended in non-denaturing lysis buffer (NaH2PO4 50 mM, NaCl 300 mM, imidazole 20 mM). The full length lectin and the corresponding lectin domain were purified from the cell lysate using affinity chromatography. The lysate was run through a HisTrapTM HP column (GE Healthcare), which contains Nickel ions embedded in a matrix of sepharose. The lectin (domain) was further purified from the eluted sample using size exclusion chromatography. The sample was applied on a HighloadTM 16/60 column packed with a matrix of Superdex™ prep grade (GE Healthcare). Fractions containing the lectin (domain) were collected, analyzed using SDS-PAGE, pooled together and concentrated.

**SDS-PAGE and Western blot.** To verify the expression of recombinant proteins, as well as the presence of pure lectin (domain) after purification steps, each fraction was separated by SDS-PAGE in Bolt 12% Bis-Tris Plus gels (Life sciences). The gel was used for a Western blot or stained with Coomassie Brilliant Blue R-250 (Bio Rad) or Sypro® Ruby protein gel stain (Invitrogen).

For performing the Western blot, the proteins from the gel were transferred to a polyvinylidifluoride (PVDF) membrane by electroblotting at 500 mA and 30 V for 1 h. For protein detection, the membrane was incubated with 0.2 μg/ml primary mouse monoclonal anti-His6 antibodies (serial no. 11922416001, Roche) in 20 ml 0.3% bovine serum albumin (BSA) (Sigma-Aldrich). Subsequently after washing, the membrane was incubated with 1:10000 diluted secondary anti-mouse antibodies conjugated with alkaline phosphatase (A3562-25ML, Sigma-Aldrich).

Finally, protein detection was based on color reaction adding nitro blue tetrazolium and bromo-chloro indolyl phosphate as substrate. The reaction was stopped using 1x PBS with 25 mM ethylenediaminetetraacetic acid (EDTA).

**S. cerevisiae and C. albicans agglutination assays.** To determine the lectin-binding capacity of the lectin-like domain of LGR1_lpl1, S. cerevisiae and C. albicans agglutination assays were performed as previously described with minor modifications. Briefly, overnight-grown cultures of S. cerevisiae BY4741 or C. albicans SC5314 cells were washed and suspended in PBS to final concentration of a 1% w/v cell suspension. 50 μl of these cell suspensions was added to the wells of 96-well U-bottomed plates (Cellstar® 650180, Greiner bio-one) and subjected to recombinant protein expression and subsequent detection by SDS-PAGE electrophoresis and Western blot.

**Pull-down carbohydrate binding assay using Sepharose beads.** Sepharoses® 6B beads (Sigma-Aldrich) were coated with 20% D-glucose, GlcNac, D-mannose, D-fucose, mannose of S. cerevisiae as previously described with little modification. For the sugar-binding assay, 25 l of each functionalized bead was washed with binding buffer (25 mM MES, 25 mM NaCl and 1% polyvinylalcohol) as previously described. Briefly, 1 ml of binding buffer containing 100 μg/ml of the purified lectin domain was added to each bead. The mixture was then incubated at 4 °C for 2 h. The beads were washed twice with 1 ml of wash buffer and bound lectin domains were eluted by boiling the beads in SDS-PAGE loading buffer (Fermentas, Life Sciences) for 10 min at 95 °C. The bound lectin domains were resolved by SDS-PAGE through 12% polyacrylamide gels (Life Sciences), which were stained with Sypro® Ruby protein gel stain (Invitrogen) and scanned by using the Typhoon scanner (GE Healthcare Life Sciences).

**Glycan array analysis.** The mammalian-glycan array version 5.2 was used to profile the sugar specificity of the lectin domain of LGR1_lpl1. The array consists of 609 glycan targets of natural and synthetic mammalian glycans with amino linkers and it is printed onto N-hydroxysuccinimide (NHS)-activated glass microscope slides (SCHOTT Nexterion), forming covalent amide linkages. The purified lectin domain of LGR1_lpl1 was labelled
Table 2. List of primers used in the study.

| Primer | Primer sequence (5'–3') | Restriction site | Remarks |
|--------|-------------------------|-----------------|---------|
| Pro564 | AGCAGGACGGGAAACGAATGAATG | / | Forward primer to check pEM40 integration in Lactobacillus genome |
| Pro565 | GCCGGGTTGGGGGAGATTGGACAG | / | Reverse primer to check pEM40 integration |
| Pro7129 | ATGTTCAATGTAATCTCTCTATTAC | / | Reverse primer to check insertion in multiple cloning site pLAB1301 |
| Pro7130 | ATAGGGCTCAGAAAAGGAGCCTTATAATTGTA | / | Forward primer to check insertion in multiple cloning site pLAB1301 |
| Pro7466 | ATCCCCGGGCAAATACATCCTG | XmnI | Forward primer HR1 llp1 gene LGR-1 |
| Pro7467 | ATCCCCGGGCAATGATCGTCAAATTCA | XmnI | Reverse primer HR1 llp1 gene LGR-1. Used for Southern hybridisation |
| Pro7468 | ATGGCGGCGCGCTGAGTCGTCATCGTCAGG | NotI | Forward primer HR2 llp1 gene LGR-1 |
| Pro7469 | ATGGCGGCGCGCTGAGTCGTCATCGTCAGG | NotI | Reverse primer HR2 llp1 gene LGR-1 |
| Pro8018 | TTACACTCCGACTTCTAACCCTG | / | Forward primer to check llp1 replacement with TcR cassette |
| Pro8019 | CTAACAGGCATGCCTTAGTCG | / | Reverse primer to check llp1 replacement with TcR cassette |
| Pro8675 | ATCCCCGGGATGAAGAATGGCCGCTA | XmnI | Forward primer complementation llp1 gene |
| Pro8676 | ATGGAGCCGCTGATCGTCACTCCTTCTTGA | ScaI | Reverse primer complementation llp1 gene |
| Pro8880 | CACCGGCAACGGAAGAAGATATCA | SalI | Forward primer for full length llp1 gene for pET28 a(+) |
| Pro8881 | ATGGCGGCGCGCTGATGGAAGGCTAGGAGTAG | NotI | Reverse primer for full length llp1 gene for pET28 a(+) |
| S&P-00517 | ATGGCGGCGCGCTGATGGAAGGCTAGGAGTAG | SalI | Forward primer upstream of lectin-like domain of llp1 gene |
| S&P-00518 | ATGGCGGCGCGCTGATGGAAGGCTAGGAGTAG | NotI | Reverse primer upstream of lectin-like domain of llp1 gene |
| S&P-0044 | TGGCAGCAGCGGCAACTCGGCTT | / | Reverse primer for MCS of pET28 a(+) |
| S&P-0045 | TATAGGCGGCGGCAACTCGGCTT | / | Forward primer for MCS of pET28 a(+) |

with FITC using FluoReporter® FITC Protein Labeling Kit (Life Technologies) according to the producer’s manual. The experiment was performed by the Consortium for Functional Glycomics (CFG, www.functionalglycomics.org).

**In vitro adhesion assay to a human epithelial cell lines.** Adhesion assays using the VK2/E6E7 (ATCC CRL-2616™), Ect/E6E7 (ATCC CRL-2614™), End1/E6E7 (ATCC CRL-2615™), Hela (ATCC CCL-2™) and Caco-2 (ATCC HTB-37™) cell lines were performed as previously described45,46. The adhesion ratio was calculated by comparing the number of adherent cells (CFUs) to the determined CFUs of the initially added bacterial suspension. Adhesion of *L. rhamnosus* GR-1 wild type, CMPG10744 (mutant) and CMPG10746 (complemented mutant) was tested in triplicate in three independent experiments. Alternatively, a fluorescence assay was performed as previously described47 with minor modifications. The FITC labelled lectin domain of LGR1_Llp1 protein was suspended in the DMEM medium in the absence of serum and antibiotics, and incubated for 1 h with the monolayer of VK2/E6E7 and Caco-2 cells grown on 13-mm coverslips. After incubation, the cells were sequentially washed three times with PBS, and fixed with 4% paraformaldehyde for 10 min. Slides were examined with a Zeiss Axio Imager Z1 microscope with an EC Plan Neofluar (X40 magnification/0.3 numerical aperture) objective (excitation 488 nm, emission 511 nm). Pictures were acquired with an Axioscam MRm and the AxioVision software.

**Induction of cytokine gene expression in VK2/E6E7 epithelial cells.** To determine the production of different pro-inflammatory and anti-inflammatory cytokines upon co-incubation with the *L. rhamnosus* GR-1 wild type, its *LGR1* _llp1_ mutant and purified LGR1_Llp1 lectin domain, the vaginal VK2/E6E7 cell line was used. Cytokine expression was monitored by quantitative reverse transcription-PCR (qRT-PCR) as described previously39, with minor modifications. VK2/E6E7 cells growing in 12-well tissue culture plates were deprived of FCS one day prior the mRNA induction experiments. Strains were grown overnight in MRS medium and subsequently centrifuged at 2,000 × g at 4 °C for 10 min. After washing once with 1x PBS, cells were resuspended in DMEM without FCS and adjusted to a final concentration of 1 × 10^7 CFU/ml. A 1.5-ml volume of the bacterial cell suspension was then added to the VK2/E6E7 epithelial cells for 1.5 h. Afterwards, the epithelial cells were rinsed twice with prewarmed 1x PBS. RNA was extracted from the VK2/E6E7 cells by using the high pure RNA isolation kit (Roche) following the manufacturer’s protocol. Cytokine gene expression was measured by qRT-PCR. In addition, the RT2 Profiler PCR Array Human Cytokines and Chemokines (Qiagen, PAHS-150Z) was used. This RT2 PCR array profiles the expression of 84 key secreted proteins central to the immune response.
In vitro biofilm assay with selected pathogens and Lactobacillus species. The effect of LGR1_Llp1 on the biofilm formation capacity of E. coli UTI98, S. aureus SH1000 and S. aureus Rosenbach and various Lactobacillus strains was investigated as previously described.38,39 The strains were grown on polystyrene pegs in the presence of the purified lectin domain of GR1_Llp1 at a final concentration of 50 or 200 μg/ml. The experiment was performed at least three times with eight technical repeats. The total cell count of the biofilms for E. coli UTI89 was also determined. E. coli UTI89 was allowed to form a biofilm on the bottom of polystyrene wells of 12-well culture plates (Cellstar®) and the lectin domain was added at 50 μg/ml. The biofilm was detached from the bottom of the wells using scrapers (Greiner bio-one) and pushed through a needle (25 G, 0.5 × 16 mm, Terumo) to dissolve cellular aggregates. The dissolved biofilms were serially diluted in PBS and plated on LB. For each strain, the experiments were performed at least three times with threetechnical repeats.

For the visualization of E. coli UTI89 biofilms, the wild-type E. coli UTI89 and FITC-labeled lectin domains were used. The lectin domain was added at the onset of the biofilm formation at 50 μg/ml and the biofilms were grown for 48 h. Microscopic epifluorescence imaging was performed using a Zeiss Axio Imager Z1 microscope with an EC Plan Neofluar (X40 magnification/0.3 numerical aperture) objective (excitation 488 nm, emission 511 nm). Pictures were acquired with an AxioCam MRm and the AxioVision software.

Statistical analysis. To determine significant differences, the unequal variance t-test was applied. A P-value below 0.05 was considered as statistically significant.

References

1. WHO/FAO. Evaluation of health and nutritional properties of powder milk and live lactic acid bacteria. Food and Agriculture Organization of the United Nations and World Health Organization Expert Consultation Report. (2001).
2. Reid, G. & Bruce, A. W. Selection of lactobacillus strains for urogenital probiotic applications. J. Infect Dis., 183, S77–S80 (2001).
3. Gardiner, G. E., Heinemann, C., Bruce, A. W., Beuerman, D. & Reid, G. Persistence of Lactobacillus fermentum RC-14 and Lactobacillus rhamnosus GR-1 but not L. rhamnosus GG in the human vagina as demonstrated by randomly amplified polymorphic DNA. Clin. Diagn. Lab. Immunol., 9, 92–96 (2002).
4. Reid, G. et al. Oral use of Lactobacillus rhamnosus GR-1 and L. fermentum RC-14 significantly alters vaginal flora: randomized, placebo-controlled trial in 64 healthy women. FEMS Immunol. Med. Microbiol., 20(35), 131–134 (2003).
5. Reid, G. et al. Oral probiotics can resolve urogenital infections. FEMS Immunol. Med. Microbiol., 30, 49–52 (2001).
6. McGroarty, J. A. & Reid, G. Detection of a Lactobacillus substance that inhibits Escherichia coli. Can. J. Microbiol., 34, 974–978 (1988).
7. Reid, G., McGroarty, J. A., Angotti, R. & Cook, R. L. Lactobacillus inhibitor production against Escherichia coli and coaggregation ability with uropathogens. Can. J. Microbiol., 34, 344–351 (1988).
8. McGroarty, J. A. & Reid, G. Inhibition of Enterococcus by Lactobacillus species in vitro. Microbial Ecology in Health and Disease, 1, 215–219 (1988).
9. McMillan, A. et al. Disruption of urogenital biofilms by lactobacilli. Colloids Surf. B Biointerfaces, 86, 58–64 (2011).
10. Kohler, G. A., Assafa, S. & Reid, G. Probiotic interference of Lactobacillus rhamnosus GG and Lactobacillus reuteri RC-14 with the opportunistic fungal pathogen Candida albicans. Infect. Dis Obstet. Gynecol., 2012, doi:10.1155/2012/636474 (2012).
11. Lebeer, S., Vanderleyden, J. & De Keersmaecker, S. C. Host interactions of probiotic bacterial surface molecules: comparison with commensals and pathogens. Nat. Rev. Microbiol., 8, 171–184 (2010).
12. Van Damme, E. J., Broekaert, W. F. & Peumans, W. J. The Urтика dioica agglutinin is a complex mixture of lectins. Plant Physiol., 86, 598–601 (1988).
13. Van Damme, E. J. M., Allen, A. K. & Peumans, W. J. Related mannose-specific lectins from different species of the family Amaryllidaceae. Plant Physiol., 73, 52–57 (1988).
14. Wellens, A. et al. The tyrosine gate as a potential entropic lever in the receptor-binding site of the bacterial adhesin FimH. Biochemistry, 49(11), 4790–4799 (2012).
15. Clarke, S. R., Harris, L. G., Richards, R. G. & Foster, S. J. Analysis of Ebb, a 1.1-megadaton cell wall-associated fibronectin-binding protein of Staphylococcus aureus. Infect. Immun., 70, 6680–6687 (2002).
16. Marraffini, L. A., Dedent, A. C. & Schneewind, O. Sortases and the art of anchoring proteins to the envelopes of gram-positive bacteria. Microbiol. Mol. Biol. Rev., 70, 192–220 (2006).
17. Alvarez, M. A., Herrero, M. & Suarez, J. E. The site-specific recombination system of the Lactobacillus species bacteriophage A2 integrates in gram-positive and gram-negative bacteria. Virology, 250, 185–193 (1998).
18. Josson, K. et al. Characterization of a gram-positive broad-host-range plasmid isolated from Lactobacillus hilgardii. Plasmid, 21, 9–20 (1989).
19. De Keersmaecker, S. C. et al. Flow cytometric testing of green fluorescent protein-tagged Lactobacillus rhamnosus GG for response to defensins. Appl. Environ. Microbiol., 72, 4923–4930 (2006).
20. Petrova, M. I., van den, B. M., Balzarini, J., Vanderleyden, J. & Lebeer, S. Vaginal microbiota and its role in HIV transmission and infection. FEMS Microbiol. Rev., 37, 762–792 (2013).
21. Petrova, M. I., Lievens, E., Malik, S., Imholz, N. & Lebeer, S. Lactobacillus species as biomarkers and agents that can promote various aspects of vaginal health. Front. Physiol., 6, 81 (2015).
22. Arndt, N. X., Tiralongo, J., Madge, P. D., van, I. M. & Day, C. J. Differential carbohydrate binding and cell surface glycosylation of human cancer cell lines. J. Cell Biochem., 112, 2230–2240 (2011).
23. Rajan, N. et al. Roles of glycoproteins and oligosaccharides found in human vaginal fluid in bacterial adherence. Infect Immun., 67, 5027–5032 (1999).
24. Venegas, M. F. et al. Binding of type 1-piliated Escherichia coli to vaginal mucus. Infect Immun., 63, 416–422 (1995).
25. Van Damme, E. J., Smith, D. F., Cummings, R. C. & Peumans, W. J. Glycan arrays to decipher the specificity of plant lectins. Adv. Exp. Med. Biol., 705, 757–767 (2011).
26. Navas, E. L. et al. Blood group antigen expression on vaginal cells and mucus in women with and without a history of urinary tract infections. J. Urol., 152, 345–349 (1994).
27. Corfield, A. P. & Berry, M. Glycan variation and evolution in the eukaryotes. Trends Biochem. Sci., 40, 351–359 (2015).
28. Syytgat, H. L. & Lebeer, S. The sweet tooth of bacteria: common themes in bacterial glycoconjugates. Microbiol. Mol. Biol. Rev., 78, 372–417 (2014).
29. Topin, J. et al. Deciphering the glycan preference of bacterial lectins by glycan array and molecular docking with validation by microcalorimetry and crystallography. PLoS. One, 8, e71149 (2013).
30. Lebeer, S., Vanderleyden, J. & De Keersmaecker, S. C. Genes and molecules of lactobacilli supporting probiotic action. Microbiol. Mol. Biol. Rev., 72, 728–764 (2008).
Author Contributions

interpretation, or the decision to submit the work for publication. Sarah Lebeer). We thank the Protein-glycan Interaction Resource of the CFG and the supporting grant R24 program financing from the KU Leuven (PF/10/018, Jan Balzarini and Dominique Schols, Jos Vanderleyden, grant KaN 28960. Elke Lievens holds an IWT-SB doctoral grant. Work at KU Leuven was supported by BOF funding from the CMPG, KU Leuven, Belgium for their technical assistance with the fluorescence microscope. We also thank David De Coster and Stefanie Roberfroid for their help with sugar-binding assays and biofilm formation of bacterial species. We also thank Nicole Imholz and Karolien Bijnens (Master students of the KU Leuven) for their help with PCR-mediated gene disruption and other applications. Yeat 14, 115–132 (1998).

Gillum, A. M., Tsay, E. Y. & Kirsch, D. R. Isolation of the gene for orotidine-5′-phosphate decarboxylase by complementation of S. cerevisiae ura3 and E. coli pyrF mutations. Mol. Gen. Genet. 198, 179–182 (1984).

Horsburgh, M. J. et al. Sigma8 modulates virulence determinant expression and stress resistance: characterization of a functional rnb strain derived from Staphylococcus aureus 8325–4. J. Bacteriol. 184, 5457–5467 (2002).

Malik S. et al. The strong auto-aggregative and adhesive phenotype of a vaginal Lactobacillus plantarum strain CMPG 5300 is sortase-dependent. Appl. Environ. Microbiol., doi:10.1128/AEM.00926-13 (2013).

Acknowledgements

We thank Nicole Imholz and Karolien Bijnen (Master students of the KU Leuven) for their help with sugar-binding assays and biofilm formation of bacterial species. We also thank David De Coster and Stefanie Roberfroid from the CMPG, KU Leuven, Belgium for their technical assistance with the fluorescence microscope. We thank Professor Giudo Van Ham for kindly providing VK2/E6E7, Ect/E6E7 and End1/E6E7 cell lines used in this study. Mariya Petrova holds a postdoctoral grant from the Fund for Scientific Research (FWO Vlaanderen). Sarah Lebeer was supported from the FWO Vlaanderen with a previous postdoctoral grant and a research grant KαN 28960. Elke Lievens holds an IWT-SB doctoral grant. Work at KU Leuven was supported by BOF program financing from the KU Leuven (PF/10/018, Jan Balzarini and Dominique Schols, Jos Vanderleyden, Sarah Lebeer). We thank the Protein-glycan Interaction Resource of the CFP and the supporting grant R24 GM098791 for conducting the glycan array analysis. The funders had no role in study design, data collection and interpretation, or the decision to submit the work for publication.

Author Contributions

Mariya Petrova and Sarah Lebeer designed the experiments and wrote the manuscript. Mariya Petrova, Elke Lievens and Tine Verhoeven performed the experimental work. Gregor Reid, Greg Gloor and Jean Macklaim
sequenced and annotated the genome of *L. rhamnosus* GR-1. Gregor Reid, Greg Gloor, Jean Macklaim, Dominique Schols and Jos Vanderleyden reviewed the manuscript.

**Additional Information**

**Competing financial interests:** The authors declare no competing financial interests.

**How to cite this article:** Petrova, M. I. *et al.* The lectin-like protein 1 in *Lactobacillus rhamnosus* GR-1 mediates tissue-specific adherence to vaginal epithelium and inhibits urogenital pathogens. *Sci. Rep.* 6, 37437; doi: 10.1038/srep37437 (2016).

**Publisher’s note:** Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.

This work is licensed under a Creative Commons Attribution 4.0 International License. The images or other third party material in this article are included in the article’s Creative Commons license, unless indicated otherwise in the credit line; if the material is not included under the Creative Commons license, users will need to obtain permission from the license holder to reproduce the material. To view a copy of this license, visit http://creativecommons.org/licenses/by/4.0/

© The Author(s) 2016