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## Does exposure to different menstrual products affect the vaginal environment?

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#### Abstract

The vaginal ecosystem is a key component of women's health. It also represents 37 an ideal system for ecologists to investigate the consequence of perturbations on 38 species diversity and emerging properties between organisational levels. Here, we 39 study how exposure to different types of menstrual products is linked to microbial, 40 immunological, demographic, and behavioural measurements in a cohort of young 41 adult women who reported using more often tampons (n = 107) or menstrual cups 42 (n = 31). We first found that cup users were older and smoked less than tampon 43 users. When analysing health indicators, we detected potential associations between 44 cups use reporting and fungal genital infection. A multivariate analysis confirmed 45 that, in our cohort, reporting using cups over tampons was associated with the higher 46 odds ratio to report a fungal genital infection diagnosis by a medical doctor within 47 the last 3 months. We did not detect significant differences between groups in terms 48 of their bacterial vaginal microbiota composition and found marginal differences in 49 the level of expression of 20 cytokines. However, a multivariate analysis of these 50 biological data identified some level of clustering based on the type of menstrual 51 product type preferred (cups or tampons). These results suggest that exposure 52 to different types of menstrual products could influence menstrual health. Larger 53 studies and studies with a more powered setting are needed to assess the robustness 54 of these associations and identify causal mechanisms. 55

Keywords: vaginal microbiota; fungal infections; immunity; epidemiology; women's
 health

#### 58 Introduction

There is a long history of considering organisms as ecosystems, and *vice versa* [1]. This parallel has been particularly exploited to study within-host parasite dynamics [2]. With the advent of metagenomics, there is a renewed interest in applying ecological theory to understand microbiota dynamics [3, 4]. Beyond parasites or even the microbiota, the importance of encompassing the diversity of these ecosystems, e.g. including immune cells, is increasingly acknowledged, especially since the border between self and non-self is tenuous in this context [5].

Some general questions pertaining to ecosystem functioning can be studied by considering an individual host as an ecosystem. For instance, it is challenging to clearly define properties that emerge at a higher level of organisation in large-scale ecological systems. However, the definition and measurement of individual health outcomes (i.e., systems-level outcomes) is routine in clinical research. In contrast to free-living ecosystems, individual hosts are also smaller, more numerous, and readily replicable, making empirical ecosystem ecology research more accessible and reproducible [6].

The vaginal ecosystem is an ideal study system to understand how within-host inter-73 actions shape host health. First, the vaginal microbiota is a key component of women's 74 health throughout their lives [7], modulating the risk of diseases such as fungal genital 75 infections, urinary tract infections, and sexually transmitted infections [8], such as hu-76 man immunodeficiency virus [9]. Contrary to its gut counterpart, the vaginal microbiota 77 is characterised by relatively few, clearly-defined, community state types (CST), most 78 of which are dominated by one species of Lactobacillus bacteria [10]. Second, the mi-79 crobes in the vaginal environment, including bacteria, fungi, and viruses, are involved in 80 a tight cross-talk with the local innate and adaptive immune response, the components 81 of which are known to be adapted to this highly variable environment [11]. In partic-82 ular, the interaction between the immune response and sex hormones throughout the 83 menstrual cycle is well documented [12]. Finally, the vaginal ecosystem is perturbed by 84 factors including menses, sexual activities, and antibiotic treatments [13]. Longitudinal 85 follow-ups demonstrate that such perturbation can lead to shifts from one CST to an-86 other [14]. Perturbations that jeopardise the balance between immunity, hormones, and 87

the vaginal microbiota could also hinder the efficiency of the immune system to control
bacteria, fungal, and viral pathogens [13].

Here, we explore whether exposure to a particular type of menstrual product, namely menstrual cups or tampons, is associated with differences in the vaginal environment. To this end, we analyse the vaginal microbiota, immune response (i.e., cytokines and antibodies), and, more generally, the emerging property of this ecosystem, which revolves around menstrual health [15].

Among the variety of products used during menses, menstrual cups are perceived as 95 a safe, practical, economical, and ecologically friendly alternative to tampons and san-96 itary pads [16]. The majority of women using them report wanting to continue to use 97 them, both in high and low-income settings, showing a good level of acceptability [17, 98 16]. Nevertheless, there have been some reported cases of toxic shock syndrome, renal gq colic, and allergies linked with menstrual cup usage [16]. For example, higher levels of 100 Staphylococcus aureus growth have been reported in menstrual cups compared to tam-101 pons [18]. In 2019, a systematic review identified 12 clinical trials and 23 observational 102 studies comparing this type of menstrual product to others [16]. In the vast majority of 103 existing studies, the outcome studied was the practicality of the cups implementation, 104 e.g. acceptability or risk of leakage. Therefore, we know little about the potential mi-105 crobiological and immunological implications of cup usage compared to other menstrual 106 hygiene products. 107

In the present study, we analysed biological, demographic, and behavioural longitudinal data from 138 women. We compared two populations of women defined by the type of menstrual product they reported using most often (tampons or menstrual cups) and analysed microbiological, immunological, and clinical data while statistically accounting for demographic and behavioural differences. Using statistical modelling, we identified profile differences depending on the type of menstrual product used.

#### <sup>114</sup> Materials and methods

#### <sup>115</sup> Cohort description and data curation

Women included in the present study were enrolled in the PAPCLEAR longitudinal clinical study [19], which followed 149 women between 2016 and 2020 to study human papillomavirus (HPV) infections [19]. The women enrolled were between 18 and 25 years old, lived in the area of Montpellier (France), and reported at least one new partner within the last 12 months. Their status for HPV and other genital infections, immunological responses (antibodies and cytokines), and behaviours were followed for up to two years (see [20] for details about the study protocol).

We selected participants who reported using tampons or cups as menstrual products 123 and for whom detailed cytokine profiles, microbiota metabarcoding data, and antibody 124 data at the inclusion visit were available. We assigned tampon or menstrual cup cate-125 gories when a participant reported using either type of menstrual product over 75% of 126 the time over the whole duration of the study (see Figure S1 for details on how the as-127 signment was performed). Women using menstrual cups more frequently than tampons 128 are denoted 'menstrual cups users' for convenience throughout the manuscript. There 129 was no difference in follow-up duration between women using mostly cups or mostly tam-130 pons (Table 1). Other analyses of demographic, behavioural, and biological data were 131 performed on the first visit of the participant (inclusion visit). Overall, our analysis 132 includes data from 138 women. 133

Since the PAPCLEAR study was designed to understand the ecology of HPVs and their interaction with immune responses and vaginal microbiota, the data from this cross-sectional study is not perfectly balanced in terms of exposure to each type of menstrual product. However, given the current lack of studies, it greatly improves our understanding of women's menstrual health.

#### 139 Biological analyses

HPV infections were assessed from cervical smears using the LiPA<sub>25</sub> genotyping assay, which discriminates between 25 genotypes [21]. Further details on HPV detection and prevalence in this cohort can be found elsewhere [20]. A 'focal' infection was defined as an infection by the same HPV genotype at the first and second visits.

The microbiota metabarcoding was performed on 200 µL of vaginal swabs specimen stored at -80° in Amies medium. DNA extraction was performed using the MagAttract PowerMicrobiome DNA/RNA Kit (Qiagen). Next-generation sequencing of the V3-V4 region of the 16S gene [22] was performed on an Illumina HiSeq 4000 platform (150 bp paired-end mode) at the Genomic Resource Center at the University of Maryland School of Medicine. Taxonomic assignment was performed using the internal software package SpeciateIT (https://github.com/Ravel-Laboratory/speciateIT) and
the community state type (CST) was determined using the VALENCIA software package
[23].

Antibodies were analysed using a multiplex Luminex assay [24], as already described in the context of HPV infections [20]. Cytokine data were obtained using MesoScale discovery technology from vaginal secretions collected using ophthalmic sponges placed directly on the cervical os for approximately one minute, as described in a previous study which analyses this data in the context of HPV infections [25]. We used the same protocol to obtain values that were normalised per total protein concentration in the sample.

#### 160 Statistical analyses

To study the difference between the main characteristics of our two populations of interest (Table 1 and Table 2), we used  $\chi^2$ -test or Kruskal-Wallis rank sum tests when applicable.

We then performed multivariate analyses using generalised linear models (GLMs) with a binomial distribution for the response variable for the models shown in Figure 1A, Table 3, and Supplementary Tables S4 to S8. In the first series of models, the response variable was the type of menstrual product (cups or tampons). In the second series, the response variable was being diagnosed or not with a fungal genital infection within the last three months by a medical doctor.

We identified 15 covariates of interest (Table S1). Given the exploratory nature of the study, we built models with all the possible combinations of covariates as explanatory variables. We then selected the best models using the Akaike Information Criterion corrected for small sample sizes (AICc). We used a lowest AICc +5 interval to identify the most probable best models [26]. For each model, we computed the odds ratios associated with each predictor along with a 95% confidence interval.

For clustering, factor analysis of mixed data was used when combining both factor and numeric data (Figure 3B and S2), and multiple correspondence analysis was used when analysing categorical data (Figures S2 and S4) [27]. Non-overlapping 95% confidence ellipses indicate a statistically significant difference between cups and tampons users. The covariates used for the socio-economic analysis using a multiple correspondence analysis are listed in Table S2.

Statistical analyses were performed in R 4.1.3 [28] with the with the kruskal.test 182 function for Kruskal-Wallis rank sum tests (in Tables 1 and 2 for numeric data), chisq.test 183 for  $\chi^2$  test (in Table 1 and Figure 2B for proportional data), wilcox.test function 184 for Mann-Whitney test (Figure 2C) and the glmulti function for binomial regressions 185 [29] (in Figure 1A and Table 3). The factor analysis of mixed data and the multiple 186 correspondence analysis (in Figures 1B, 3B, S2, S3, and S4) were processed and repre-187 sented using the FactorMineR and the FactoExtra (https://github.com/kassambara/ 188 factoextra) packages [27]. 189

#### 190 **Results**

The 138 women included in the analysis were aged from 18 to 25 years old and primarily 191 university students (119/138, 86%). We stratified the population according to the most 192 frequent type of menstrual product used, *i.e.* either tampons, n = 107, or menstrual cups, 193 n = 31. Analysis of the main demographic characteristics (shown in Table 1) using 194 a generalised linear model (GLM) selection approach identified significant differences 195 between these two populations in terms of age and smoking status: cups users were 196 older and reported smoking less than tampon users (Figure 1A and Table S3). On the 197 other hand, a multiple correspondence analysis on 12 covariates associated with socio-198 economic status and listed in Table S2 did not show any difference between the two 199 populations (Figure 1B). 200

We then explored associations between the type of menstrual product used and 201 microbiological covariates. First, we analysed vaginal microbiota diversity using 16S 202 metabarcoding data (Figure 2A). We found no significant difference in community state 203 types (CST) composition (Figure 2B), although there were some qualitative differences. 204 For instance, among our menstrual cup users, we found no occurrence of CST II, which 205 is dominated by L. gasseri, but slightly more CST I and V, which are dominated by 206 L. crispatus and L. jensenii [10]. We also did not find any significant differences in micro-207 biota diversity, assessed using Shannon diversity index, between women using menstrual 208 cups or tampons (Figure 2C). 209

We then analysed six health-related covariates, most of which correspond to infections 210 (Table 2). The only significant difference was that women using cups also reported more 211 often having been diagnosed with fungal genital infections within the last 3 months. To 212 further investigate this association, we used our GLM selection method to identify the 213 covariates associated with this fungal genital infection. We found that reporting using 214 cups is the only predictor with a significant odds ratio (OR of 4.73, 95% CI: 1.44-16.1) 215 associated with the risk of reporting a fungal genital infection (the number of lifetime 216 partners is also present in the model, but not significant). Similar models for the other 217 five health-related covariates are shown in Appendix (Tables S4 to S8). 218

To investigate the potential effect of menstrual cups on the local immune response,

we analysed cytokine and chemokine relative concentrations in cervical samples. Among the 20 analytes measured, IL-10, MIP-1 $\alpha$  and TNF- $\alpha$  appeared to be significantly lower in women using cups (IL-10: p = 0.012, MIP-1 $\alpha$ : p = 0.049 and TNF- $\alpha$ : p = 0.036), although these associations did not withstand correction for multiple testing comparisons (p = 0.24, p = 0.32, and p = 0.32 respectively) (Figure 3A).

Finally, we performed a profile analysis using a factor analysis of mixed data approach 225 involving CST data, Shannon diversity index, and cytokine and chemokine relative con-226 centrations. Women who use tampons and women using menstrual cups fall into two 227 largely non-overlapping clusters (Figure 3B and Supplementary Figure S2), suggest-228 ing that the type of menstrual product used could be associated with a different im-229 munological and microbial environment. Conversely, a similar multiple correspondence 230 analysis approach using blood seropositivity status for immunoglobulins G (IgG) and 231 immunoglobulins M (IgMs) of several sexually transmitted infections, including HPVs, 232 detected no clustering (Supplementary Figure S3), hinting that the women in these two 233 groups have similar exposure risks to sexually transmitted infections. 234

#### 235 Discussion

The vaginal ecosystem is an essential part of women's health and several 'perturbations' such as menses, sexual intercourse, or drug treatment can cause pronounced shifts [7]. In this study, we analysed whether exposure to different types of menstrual products, namely menstrual cups or tampons, could be associated with differences in health indicators or, more generally, in the vaginal environment.

Menstrual cups are gaining in popularity as an environmentally sustainable and af-241 fordable type of menstrual product, but the data surrounding their use are limited, 242 especially outside low-income countries [16]. Therefore, we first analysed the profiles 243 of the two populations of women defined based on the type of menstrual product they 244 reported using most. In terms of demographic covariates, we found that the main factors 245 that were associated with preferential use of cups over tampons were age (cup users were 246 older) and smoking status (tampon users smoked more). In terms of socio-economic sta-247 tus, a multiple correspondence analysis did not suggest any differences between the two 248 populations, which could be due to the fact that the majority of the cohort member was 249

university students. A similar multiple correspondence analysis based on serological data
for a variety of sexually transmitted infections also did not find any difference, further
suggesting that these two cohorts are comparable in terms of their general lifestyle.

On the microbiological side, we did not find a significant difference between the 253 bacterial community compositions of the vaginal environment depending on the type of 254 menstrual product used. A recent preprint [30] studied vaginal microbiota in a large 255 cohort of 3,345 women aged 18 to 98 with a median age of 32 years old. Their results 256 show that women who reported using menstrual cups are more often associated with a 257 Lactobacillus crispatus cluster (which also includes L. jensenii, and could be described 258 as encompassing both CST-I and CST-V); which is consistent with the trend detected 259 in our dataset. 260

Moving to health-related factors, we detected a strong association between reporting 261 using menstrual cups and a recent diagnosis of a fungal genital infection. None of 262 the other five health-related variables that we analysed, such as reporting urinary tract 263 infections or the diagnosis of chlamydia infections, were significantly different. To further 264 investigate this trend, we used a generalised linear model selection approach. In almost 265 all the best models, reporting using cups more than tampons was significantly associated 266 with a higher risk of reporting fungal genital infections. It is noteworthy that the number 267 of partners was present in most of the best models (98.75%) of the time), although never 268 significant. 269

Finally, we investigated the host component of the vaginal environment by analysing local cytokine and chemokine concentrations. Three cytokines or chemokines were found to exhibit potentially lower concentrations in women using cups (IL-10, MIP-1 $\alpha$ , and TNF- $\alpha$ ), although significance did not withstand correction by multiple hypothesis testing. Furthermore, the joint analysis of microbiota and immunological data suggests that women tend to segregate into two clusters based on the type of menstrual product they use most suggesting general profile differences in terms of their vaginal environment.

Our study has several limitations, the strongest being the relatively small sample size of the cohort used (n = 138), which was originally designed to study HPV infections. This may hinder our ability to detect moderate or subtle changes induced by menstrual cups in the vaginal environment. For example, we included the use of lubricants in the

analysis, but more detailed studies could also include details on contraception methods or 281 probiotic use. Another limitation of the study lies in its cross-sectional nature. Further 282 longitudinal analyses would be helpful to establish long-term potential impacts of the 283 use of menstrual cups on the local environment, for example, on the vaginal microbiota 284 composition, which can be highly variable over time for some women [31]. In particular, 285 prospective or retrospective cohort studies, where participants are stratified based on 286 the type of menstrual products used, could be a powerful means to study the occurrence 287 of a particular negative outcome, i.e. here fungal infections. 288

Regarding these infections, we cannot provide any further details regarding the fungal species or their abundance with the current data. A quantitative polymerase chain reaction approach targeting the ITS locus coupled with a metagenomics approach could provide a more accurate picture, especially thanks to existing gene catalogues for the vaginal microbiota, such as VIRGO which already includes more than 10,000 fungal genes [32].

The existing literature on the usage of menstrual cups focuses on availability and ac-295 ceptability, yet its health implications remain understudied. A recent systematic review 296 reported nine cases of urinary tract complaints [16], which we did not detect as being 297 significantly more frequent among menstrual cups users than for tampons users. Con-298 versely, the primary adverse health outcome associated with menstrual cup use in our 299 study, i.e. fungal infections, was not included in the systematic review. Our results are 300 consistent with another recent study, which does not show adverse effects of menstrual 301 cups on the vaginal microbiota, and even potentially an increased proportion of some 302 Lactobacillus-dominated communities, which are considered beneficial for maintaining 303 the vaginal health [30]. These findings can help inform public health policies regarding 304 the use of menstrual cups and underline the need for larger, balanced cohort studies. 305

Health is partly the result of dynamic interactions between hosts and their microbes that have a long coevolutionary history. The human vaginal microbiota is an ideal study system for studying these questions for several reasons. First, its composition is a known health moderator — *Lactobacillus*-dominated communities tend to decrease the risk of acquisition of sexually transmitted infections [10]. Second, the composition of the vaginal microbiota can vary over time as a result of menstrual cycles or age, as well

as a variety of 'perturbations' including sexual intercourse and chemical interventions 312 [14]. These two points are not unique to the vaginal microbiota but contrarily to other 313 microbiota, especially in the gut, the diversity is limited (both between and within 314 women) and the variability better understood. From an evolutionary point of view, 315 human vaginal mircobiota are unique for the high prevalence of *Lactobacillus*-dominance: 316 Lactobacilli are rarely dominant in other mammals, including non-human primates [33]. 317 This human specificity obviously complicates experimental studies using animal model 318 systems. However, three-dimensional cell culture models now offer new opportunities to 319 manipulative experiments, for instance to test how the microbiota composition affects the 320 interaction between a parasite and host cells [34]. As illustrated by our study, integrating 321 knowledge across scales, i.e. going from the microbiological and immunological variables 322 to individual health is data-intensive and challenging as it requires multidisciplinary 323 knowledge from biological to public health. Nonetheless, it is a necessary challenge to 324 bridge the gap between molecular mechanisms and health outcomes. 325

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#### 329 Disclosure of Interests

The authors declare the following financial interests/personal relationships which may be considered as potential competing interests: TW serves on advisory boards for MesoScale discovery (Merck) Sharp & Dohme.

#### 333 Contributions

NT, CLM, NBo, and SA designed the study. NT, CLM, BE, BR, TK, IGB, and SA
designed the experiments. CB, VB, SGro, MR, and NBe performed experiments. CH,
JRa, and TW contributed reagents, materials, and analysis tools. CB, VB, SGra, SGro,
MR, MB, CG, VT, VF, CLM, JRe, IGB, MSe, NBo, and SA contributed to study design,
patient recruitment, and clinical data acquisition. NT, IBU, BE, BR, CS, TK, CLM,

TK and SA performed data analyses. NT, IBU, IGB, CLM, TK and SA wrote the initial
version of the manuscript. All authors approved the final version of the manuscript.

#### 341 Ethics

The PAPCLEAR trial was promoted by the Centre Hospitalier Universitaire (CHU) de 342 Montpellier and has been approved by the Comité de Protection des Personnes (CPP) 343 Sud Méditerranée I on 11 May 2016 (CPP number 16 42, reference number ID RCB 2016-344 A00712-49); by the Comité Consultatif sur le Traitement de l'Information en matière de 345 Recherche dans le domaine de la Santé on 12 July 2016 (reference number 16.504); and 346 by the Commission Nationale Informatique et Libertés on 16 December 2016 (reference 347 number MMS/ABD/AR1612278, decision number DR-2016-488). This trial was autho-348 rised by the Agence Nationale de Sécurité du Médicament et des Produits de Santé on 20 349 July 2016 (reference 20160072000007). The Clinical Trials.gov identifier is NCT02946346. 350 All participants provided written informed consent. 351

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#### 360 Data availability

Table 1, 2, 3, 4, Figures 1, 2, 3, Supplementary Tables S1, S2, S3, S4, S5, S6, S7, S8 and supplementary figures S1, S2, S3 have associated raw data. The data that support the findings of this study are available from the corresponding author upon request, and data are available in the Zenodo public repository (10.5281/zenodo.6913875).

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Table 1: Key characteristics of participants included in the study. n indicates the number of individuals, p-value refers to the outcome of a Kruskal-Wallis Rank Sum Test (kruskal.test function in R), IQR is the interquartile range, SD the standard deviation.

	Tampon	Menstrual cup	р
n	107	31	
PAPCLEAR follow-up duration (days, median [IQR])	220.00 [116.50, 374.50]	270.00 [74.00, 437.50]	0.740
Lifetime number of partners (median [IQR])	8.00 [4.00, 15.00]	11.00 [4. 00, 16.50]	0.396
Age (years, median [IQR])	21.00 [20.00, 23.00]	23.00 [22.00, 24.00]	0.021
Age at menarchy (years, median [IQR])	13.00 [12.00, 14.00]	13.00 [12.00, 14.00]	0.695
Age at sexual debut (years, median [IQR])	17 [15.00, 18.00]	16 [15.00, 17.00]	0.338
Body Mass Index (BMI, median [IQR])	21.15 [19.78, 23.40]	21.62 [20.56, 23.22]	0.483
Smoking (%)			0.129
No	64(59.8)	24(77.4)	
Occasionally	15(14.0)	4(12.9)	
Often	28(26.2)	3(9.7)	
Antibiotics (last two weeks) = Yes $(\%)$	7 (6.5)	3 ( 9.7)	0.842
Lubricant (last two weeks) (%)	14(13.1)	5(16.1)	0.891
Intercourse with a regular partner (last two weeks) (%)	63(58.9)	18(58.1)	1.000
Intercourse with an occasional partner (last two weeks) (%)	15 (14.0)	4(12.9)	1.000
Stress level (past week) $(\%)$			0.168
0 (Min)	20 (18.7)	5(16.1)	
1	45 (42.1)	13(41.9)	
2	30(28.0)	13(41.9)	
3 (Max)	12 (11.2)	0 ( 0.0)	
Menses (last two weeks) (%) (Yes)	58(54.2)	20(64.5)	0.416
HPV vaccinated (%)	55(51.4)	19(61.3)	0.443
HPV positive (focal) (%)	49 (45.8)	10(32.3)	0.256
HPV positive (multiple HPVs) (%)	33 (30.8)	9 (29.0)	1.000

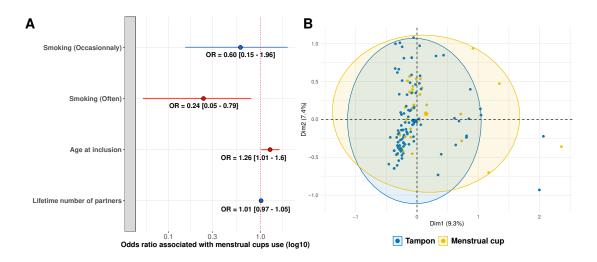


Figure 1: **Profile of menstrual cup users** A) Odds ratio of covariates associated with the use of menstrual cups. Results show the odds ratio (OR) of the factors selected in the best generalised linear model against menstrual cup as the response variable and using an Akaike Information Criterion corrected for small sample size (AICc). All covariates included in the model selection process are listed in Table S3. 95% confidence intervals are shown in brackets. B) Multiple Correspondence Analysis of socio-economic variables clustering participant using tampons or menstrual cups. Ellipses indicate 95% confidence intervals. Blue indicates tampon users, whereas yellow indicates women using menstrual cup.

Table 2: Values of the main health metrics included in the study. We show the number of occurrences and associated percentages (in parentheses) for each metric in the two study populations of size n. 'p-value' indicates the outcome of a t-test.

	Tampon	Menstrual cup	p-value
$\overline{n}$	107	31	
Fungal genital infection (last 3 months) (%)	7(6.5)	7(22.6)	0.023
Genital infection (last 3 months) $(\%)$	5(4.7)	1(3.2)	1.000
Chlamydia infection (last 3 months)( $\%$ )	4(3.7)	1(3.2)	1.000
Vaginosis (last $3 \text{ months})(\%)$	1(0.9)	2(6.5)	0.248
Urinary tract infection (last $3 \text{ months})(\%)$	12(11.2)	6(19.4)	0.378
HPV infection (focal) (%)	49 (45.8)	10(32.3)	0.256

Table 3: Factors associated with the reporting of a fungal genital infection. Results show the odds ratios (OR) of the factors selected in the best generalised linear model with fungal genital infection as the response variable and using an Akaike Information Criterion corrected for small sample size (AICc). All covariates included in the model selection process are listed in Table 4. SE stands for standard 'error'. CI 2.5% and CI 97.5% indicate the lower and upper bounds of the 95% confidence interval.

Response = Fungal genital infection	OR	OR SE	CI $2.5\%$	CI 97.5 $\%$	p.value	
Menstrual cup Lifetime number of partners	$4.731 \\ 0.998$	$2.857 \\ 0.027$	$\begin{array}{c} 1.440\\ 0.938\end{array}$	$16.058 \\ 1.047$	$0.010 \\ 0.952$	*

\*: p < 0.05

Table 4: Frequency of occurrence of the covariates among the 80 best GLM models with fungal genital infection as a response variable (Model 2). The models were selected using AICc, assuming an AICc + 5 cutoff. The columns indicate the number and percentage of models that contain each covariate, as well as the percentage of models where the associated p-value is marginally significant (lower than 10%) or significant (lower than 5%).

Model [2]: Fungal genital infection	Presence in best models (#)	Presence in best models (%)	p.value $< 0.1$	$\mathrm{p.value} < 0.05$	
Menstrual cup	79	98.75	98.75	98.75	
Lifetime number of partners	79	98.75	0.00	0.00	
Age at first visit	39	48.75	0.00	0.00	
Intercourse with occasional partner (last two weeks)	16	20.00	0.00	0.00	
HPV positive (multiple infections)	16	20.00	0.00	0.00	
Vaccinated against HPV	16	20.00	0.00	0.00	
Intercourse with regular partner (last two weeks)	16	20.00	0.00	0.00	
Menses (last two weeks)	16	20.00	0.00	0.00	
Used Lubricant (last two weeks)	16	20.00	0.00	0.00	
Antibiotics (last two weeks)	16	20.00	0.00	0.00	
HPV positive (focal)	16	20.00	0.00	0.00	
Stress level (last week) (1, minimum)	2	2.50	0.00	0.00	
Stress level (last week) (2)	2	2.50	0.00	0.00	
Stress level (last week) (3, maximum)	2	2.50	0.00	0.00	
Smoking (Occasionally)	2	2.50	0.00	0.00	
Smoking (Often)	2	2.50	0.00	0.00	

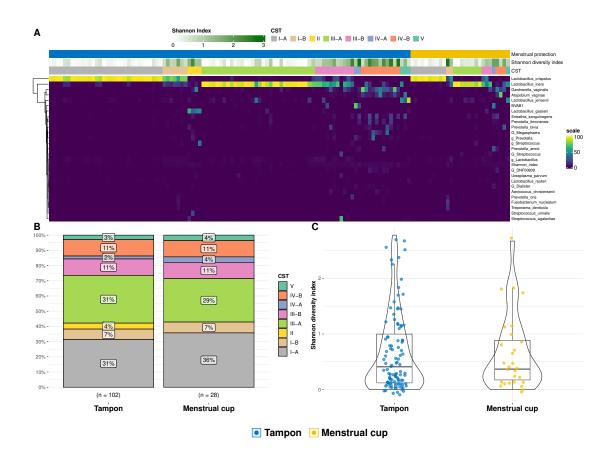


Figure 2: Metabarcoding differences between women using mainly menstrual cups or tampons. A) Abundance and diversity of the top 1% most abundant bacterial species found in participants, B) Community State Types (CST) distribution, C) Shannon diversity index. In A and B, colours show the type of menstrual product used (tampons in blue and cups in yellow).

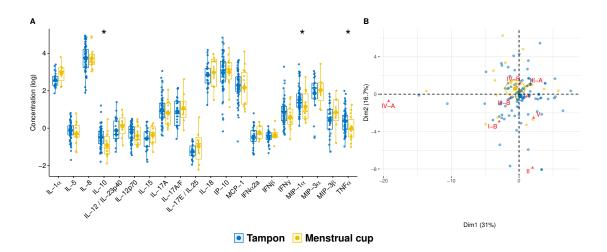


Figure 3: Immunological differences between women using mainly menstrual cups or tampons. A) Cytokines local concentrations (on a log scale), and B) Outcome of a multi-parametric clustering analysis using factor analysis of mixed data. In A, presence of a star indicates significance (p < 0.05). Ellipses indicate 95% confidence intervals. In A and B, colours show the type of menstrual product used (tampons in blue and cups in yellow). In B, the vaginal Community State Types (CSTs) are shown in red.