

Urogenital microbiome composition test
Androflor®

Date 1 June 2017, 9:57:15
Number of tube
Patient name
Sex
Age
Organization
Clinician name
Comments

Information about laboratory

Sample ID: Sample_1

№	Test title	Result			% of TMD
		Quantitative	Relative Lg (X/TMD)		
	Human DNA	10 ^{5.0}			
1	Total Bacterial Mass	10 ^{4.3}			
Transit microorganisms					
2	Lactobacillus spp.	not detected			
Commensals					
3	Staphylococcus spp.	not detected			
4	Streptococcus spp.	not detected			
5	Corynebacterium spp.	not detected			
	Commensals, sum	not detected			
BV-associated opportunistic microorganisms					
6	Gardnerella vaginalis	not detected			
7	Megasphaera spp. / Veillonella spp. / Dialister spp.	10 ^{3.2}	-1.6 (2.0-2.7 %)		
8	Sneathia spp. / Leptotrichia spp. / Fusobacterium spp.	not detected			
9	Ureaplasma urealyticum*	not detected			
10	Ureaplasma parvum*	not detected			
11	Mycoplasma hominis*	not detected			
12	Atopobium cluster	10 ^{3.3}	-0.9 (10-14 %)		
	BV-associated opportunistic microorganisms, sum	10 ^{4.0}	-0.8 (12-17 %)		
Opportunistic anaerobes					
13	Bacteroides spp. / Porphyromonas spp. / Prevotella spp.	10 ^{4.3}	-0.5 (27-36 %)		
14	Anaerococcus spp.	10 ^{3.4}	-1.4 (3.4 %)		
15	Peptostreptococcus spp. / Parvimonas spp.	10 ^{3.2}	-1.6 (1.9-2.6 %)		
16	Eubacterium spp.	10 ^{3.2}	-1.6 (2.0-2.7 %)		
	Opportunistic anaerobes, sum	10 ^{4.4}	-0.2 (57-77 %)		
Opportunistic microorganisms Haemophilus spp.					
17	Haemophilus spp.	not detected			
Opportunistic microorganisms Pseudomonas aeruginosa / Ralstonia spp. / Burkholderia spp.					
18	Pseudomonas aeruginosa / Ralstonia spp. / Burkholderia spp.	not detected			
Opportunistic microorganisms Enterobacteriaceae / Enterococcus spp.					
19	Enterobacteriaceae / Enterococcus spp.	not detected			
Yeast-like fungi					
20	Candida spp.*	not detected			
Sexually transmitted infections					
21	Mycoplasma genitalium**	detected			
22	Trichomonas vaginalis**	detected			
23	Neisseria gonorrhoeae**	not detected			
24	Chlamydia trachomatis**	not detected			

* Quantitative Analysis Lg(X)
** Qualitative Analysis

logarithmic scale

Conclusion:
DETECTED: Mycoplasma genitalium, Trichomonas vaginalis.
Dysbiosis mixed etiology. The severity of dysbiosis is indicated at a level of total bacterial mass greater than 10⁵.

Version e024-1



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Femoflor® developers



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The winner
of Prizvanie national
medical prize, 2014



The winner of Prix
Galien Russia 2016
in the category of
"Best Russian Product"

DETECTED: *Mycoplasma genitalium*. The quantitative proportions of commensals are not violated