

Table S3. Taxa that are differentially abundant in urethral samples in day-0 vs day-8 specimens and day-0 vs endpoint specimens (ALDEX2 analysis)

Taxon ^a	Mean relative abundance at Day 0 (%)	Mean relative abundance at Day-8 (%)	Mean relative abundance at Endpoint (%)	Day 0 vs Day-8			Day-0 vs Endpoint		
				p-value ^b	FDR-p ^c	Effect size ^d	p-value ^e	FDR-p ^f	Effect size ^g
<i>Anaerococcus</i> *	1.47	0.00	2.88	1.49E-04	2.82E-03	-1.05	3.15E-01	7.00E-01	0.21
<i>Finegoldia</i> *	1.86	0.04	4.55	1.60E-04	3.19E-03	-1.00	2.09E-01	6.32E-01	0.31
<i>Peptoniphilus</i> *	2.42	0.04	3.62	2.41E-04	4.36E-03	-1.00	6.40E-01	8.72E-01	0.11
<i>Prevotella</i> spp. unclassified*	3.89	0.04	1.33	9.12E-04	1.10E-02	-0.79	1.16E-01	4.96E-01	-0.25
<i>Dialister</i> *	1.31	0.03	0.67	1.17E-03	1.21E-02	-0.79	1.64E-01	5.42E-01	-0.18
<i>Escherichia/Shigella</i> *	4.10	14.87	3.79	3.89E-03	2.90E-02	0.50	6.03E-01	8.59E-01	0.08
<i>Staphylococcus</i> *	4.77	15.28	4.60	1.64E-03	1.81E-02	0.65	1.76E-03	7.92E-02	0.37
<i>Porphyromonas</i>	1.22	0.00	0.27	1.76E-02	6.41E-02	-0.55	4.09E-01	7.34E-01	-0.12
<i>Campylobacter</i>	0.41	0.00	0.28	1.86E-02	6.64E-02	-0.55	5.44E-01	8.11E-01	-0.07
<i>Rothia</i>	0.22	0.31	0.01	2.55E-02	7.49E-02	0.44	6.21E-01	8.49E-01	0.08
<i>Atopobium vaginae</i>	0.65	0.17	0.00	2.28E-02	7.67E-02	-0.54	2.68E-03	6.98E-02	-0.69
<i>Streptococcus</i>	15.86	3.94	17.70	3.35E-02	1.11E-01	-0.32	5.34E-01	8.34E-01	0.10
<i>Veillonella</i>	3.23	0.39	0.51	3.22E-02	9.91E-02	-0.46	1.22E-01	5.14E-01	-0.33
<i>Gardnerella</i>	7.51	4.68	6.02	1.86E-02	7.30E-02	-0.46	4.81E-01	7.99E-01	-0.16
<i>Prevotella disiens</i>	1.75	0.00	0.50	2.15E-02	7.22E-02	-0.51	3.48E-01	6.86E-01	-0.13
<i>Prevotella timonensis</i>	0.59	0.01	0.76	1.35E-02	5.30E-02	-0.61	3.05E-01	6.65E-01	-0.19
<i>Sneathia amnii</i>	4.90	0.03	0.83	3.48E-02	1.02E-01	-0.41	1.42E-01	5.15E-01	-0.34
<i>Sneathia sanguinegens</i>	4.54	0.00	0.31	2.35E-02	7.81E-02	-0.45	7.00E-02	3.90E-01	-0.35
<i>BVAB-2</i>	0.32	0.00	0.00	1.40E-01	2.42E-01	-0.28	3.99E-02	2.70E-01	-0.42
<i>Lactobacillus iners</i>	8.96	10.78	18.02	1.93E-01	3.31E-01	0.13	3.21E-02	3.06E-01	0.25
<i>Corynebacterium</i>	3.69	21.37	7.59	2.80E-01	4.31E-01	0.31	3.25E-02	3.26E-01	0.26
<i>Parvimonas</i>	0.95	0.06	0.06	5.91E-02	1.41E-01	-0.43	3.56E-02	2.82E-01	-0.46

^aOnly those taxa with a p-value <0.05 for either the Day-0 vs Day-8 or Day-0 vs Endpoint comparison are shown in this table. * FDR-p<0.05 for Day-0 vs Day-8 comparison

^bExpected P value of Wilcoxon rank test

^cExpected Benjamini-Hochberg corrected P value of Wilcoxon test. Bold indicates statistically significant difference at FDR-p<0.05

^dMedian difference in CLR transformed abundance between Day-0 vs Day-8 groups /median of the largest difference in CLR transformed abundance within Day-0 and Day-8 groups

^eExpected P value of Wilcoxon rank test

^fExpected Benjamini-Hochberg corrected P value of Wilcoxon test. Bold indicates statistically significant difference at FDR-p<0.05

^gMedian difference in CLR transformed abundance between Day-0 vs Endpoint groups /median of the largest difference in CLR transformed abundance within Day-0 and Endpoint groups