

ID#	DGGE				Illumina
	Organism Detected	Accession #	Coverage	Identity	Abundance
21	<i>L. crispatus</i>	AY335496.1	100%	91%	29%
	* <i>L. iners</i>				40%
50	* <i>G. vaginalis</i>				50%
54	* <i>L. iners</i>				56%
63	* <i>L. iners</i>				69%
71	* <i>L. crispatus</i>				25%
	* <i>L. iners</i>				55%
86	<i>L. iners</i>	AY283272.1	100%	98%	78%
	* <i>G. vaginalis</i>				11%
6	<i>L. iners</i>	AY283272.1	100%	99%	60%
	* <i>G. vaginalis</i>				21%
18	* <i>L. iners</i>				54%
	<i>G. vaginalis</i>	CP001849.1	94%	100%	30%
49	<i>Streptococcus anginosus</i>	GU415393.1	99%	94%	28%
75	<i>L. iners</i>	AY283272.1	96%	97%	46%
	* <i>G. vaginalis</i>				33%
92	<i>Haemophilus haemolyticus</i>	GU561423.1	98%	100%	15%
	* <i>L. iners</i>				18%
	<i>G. vaginalis</i>	EF194095.1	99%	97%	27%
40	<i>Leptotrichia amnionii</i>	EF218612.1	99%	98%	15%
	* <i>L. iners</i>				18%
	<i>G. vaginalis</i>	EF194095.1	100%	95%	23%
48	* <i>L. crispatus</i>				13%
	* <i>L. iners</i>				22%
	<i>G. vaginalis</i>	CP001849.1	99%	97%	29%
57	<i>L. iners</i>	AY283272.1	99%	98%	55%
	<i>L. iners</i>	AY283272.1	99%	98%	55%
	<i>G. vaginalis</i>	EF194095.1	99%	97%	20%
60	<i>G. vaginalis</i>	CP001849.1	98%	95%	24%
67	<i>Streptococcus agalactiae</i>	AB297817.1	100%	97%	27%
	<i>G. vaginalis</i>	EF194095.1	99%	97%	54%
73	<i>Leptotrichia amnionii</i>	EF218612.1	100%	95%	2%
	<i>G. vaginalis</i>	EF194095.1	97%	98%	34%
83	<i>G. vaginalis</i>	CP001849.1	100%	96%	23%
89	<i>Prevotella amniotica</i>	AM422125.1	100%	91%	2%
	* <i>L. iners</i>				19%
	* <i>G. vaginalis</i>				42%
90	<i>G. vaginalis</i>	CP001849.1	100%	97%	60%

*Determined via band location on DGGE gel