Table 4. Representation of 41 bacterial genera detected in four human distal esophageal biopsies

	0	Number (%) of clones										
	Genus detected (n = 41)		A (n = 186)		B (n = 205)		C (n = 264)		D (n = 245)		Total (n = 900)	
	Streptococcus	77	(41.4)	66	(32.2)	84	(31.8)	147	(60.0)	374	(38.6)	
All four specimens $(n = 13)$	Prevotella	30	(16.1)	55	(26.8)	44	(16.7)	28	(11.4)	157	(17.4)	
	Veillonella	34	(18.3)	37	(18.0)	39	(14.7)	18	(7.3)	128	(14.2)	
	Rothia	1	(0.5)	2	(1.0)	14	(5.3)	5	(2.0)	22	(2.4)	
	Megasphaera	3	(1.6)	8	(3.9)	7	(2.7)	1	(0.4)	19	(2.1)	
	Granulicatella	7	(3.8)	3	(1.5)	2	(8.0)	5	(2.0)	17	(1.9)	
nens (Gemella	6	(3.2)	1	(0.5)	1	(0.4)	6	(2.4)	14	(1.6)	
pecin	TM7	4	(1.1)	2	(1.0)	5	(1.9)	2	(8.0)	13	(1.4)	
onr s	Actinomyces	2	(1.1)	1	(0.5)	7	(2.7)	2	(8.0)	12	(1.3)	
Ā	Bacteroides	1	(0.5)	6	(2.9)	1	(0.4)	3	(1.2)	11	(1.2)	
	Clostridium	1	(0.5)	2	(1.0)	7	(2.7)	1	(0.4)	11	(1.2)	
	Haemophilus	2	(1.1)	1	(0.5)	2	(8.0)	2	(8.0)	7	(8.0)	
	Bulleidia	1	(0.5)	1	(0.5)	2	(8.0)	1	(0.4)	5	(0.6)	
	Subtotal	169	(90.9)	185	(90.2)	215	(81.4)	221	(90.2)	790	(87.8)	
	Lactobacillus	0		1	(0.5)	25	(9.5)	0		26	(2.9)	
	Fusobacterium	6	(3.2)	0		2	(8.0)	2	(8.0)	10	(1.1)	
	Leptotrichia	0		7	(3.4)	0		3	(1.2)	10	(1.1)	
	Porphyromonas	3	(1.6)	0		0		6	(2.4)	9	(1.0)	
	Clostridiales AF287770	1	(0.5)	3	(1.5)	4	(1.5)	0		8	(0.9)	
	Atopobium	0		0		4	(1.5)	1	(0.4)	5	(0.6)	
	Bacteroidetes AJ289174	0		0		4	(1.5)	0		4	(0.4)	
	Campylobacter	1	(0.5)	0		3	(1.1)	0		4	(0.4)	
	Clostridiales AF201981	0		0		2	(8.0)	2	(8.0)	4	(0.4)	
	Selenomonas	0		2	(1.0)	0		2	(8.0)	4	(0.4)	
	Neisseria	2	(1.1)	0		0		1	(0.4)	3	(0.3)	

Subtotal	17	(9.1)	20	(9.8)	49	(18.6)	24	(9.8)	110	(12.2
Yersinia krist	1	(0.5)	0		0		0		1	(0.1
Tennerella	1	(0.5)	0		0		0		1	(0.1
Shewanella baltica	0		0		0		1	(0.4)	1	(0.1
Selenomonas 9-1- 7(92%)	0		1	(0.5)	0		0		1	(0.1
Paenibacillus	0		0		1	(0.4)	0		1	(0.1
Mogibacterium	0		0		1	(0.4)	0		1	(0.1
Filifactor alocis	0		0		0		1	(0.4)	1	(0.1
Comamonas	0		1	(0.5)	0		0		1	(0.
Clostridiales AF385508	0		1	(0.5)	0		0		1	(0.
Clostridiales AF287772	0		0		0		1	(0.4)	1	(0.1
Centipeda	0		1	(0.5)	0		0		1	(0.
Achromobacter	0		1	(0.5)	0		0		1	(0.
Peptostreptococcus	0		0		0		2	(8.0)	2	(0.2
Eubacterium	0		1	(0.5)	0		1	(0.4)	2	(0.2
Delftia	0		1	(0.5)	0		1	(0.4)	2	(0.2
Clostridiales AF385510	2	(1.1)	0		0		0		2	(0.
(95%)	0		0		3	(1.1)	0		3	(0.3