	1	100	200	300	400	500	600	700	800	900	1,000	1,100	1,200	1,300	1,400	1,500	1,600	1,685
FWD AB085191 FWD AY693840							- '											
FWD U09833			-1					_ i										
FWD AY789075		- 1					- 11 -							I-	- 1			1
FWD AB071177		- 1												1-	- 1		- 1	-
FWD AB219802 FWD AB032434			•											- '!'	_			
FWD AB050732		0075							- 1					- '.				
FWD KC478600		- 11					п н							- 1				
FWD AF231349																		
FWD AB241633		- 11					1 F							1	-			
FWD AB119446								11						-				
FWD AB242176 FWD EF413181	11.1		-1					11						- '				
FWD Bab2																		
REV Bab3																		
FWD F109		-																
FWD DLP133 REV R176																		
FWD hah8fwd																		
FWD bab8fwd FWD bab8mgb REV bab8rev																		
REV bab8rev																		
FWD KebabF FWD KebabP																		
REV KebabR								8111										
FWD Bm18Sf								110										
FWD Bm18Sp																		
REV Bm18Sr																		
FWD smbaJF REV smbaKR															_			
FWD 18S-F																		
REV 18S-R																		
FWD 18S-probe																	-	

Supplemental Figure 1. Sequence alignment of partial 18S rRNA sequences (labeled with their GenBank accession numbers) and oligonucleotides from the seven real-time PCR assays targeting the 18S rRNA gene that were included in this study. Vertical lines indicate point mutations and horizontal lines indicate gaps. References: Bab2/Bab3—Bloch and others; F109/DLP133/R176—Tonnetti and others; bab8fwd/bab8mgb/bab8rev—Teal and others; KebabF/KebabP/KebabR—Bown and others; Bm18Sf/Bm18Sp/Bm18Sr—Rollend and others; smbaJK/smbaKR—Hersh and others; 18S-F/18S-probe/18S-R—Hojgaard and others.