

	<i>L. boulardi</i>	<i>L. clavipes</i>	<i>L. decemflagella</i> *	<i>L. heterotoma</i>	<i>L. leipsi</i> *	<i>L. maia</i> *
<i>L. boulardi</i> (n=18)	0.11 (0-0.64)					
<i>L. clavipes</i> (n=3)	16.41 (14.88-17.49)	1.51 (0.45-2.26)				
<i>L. decemflagella</i> * (n=4)	16.68 (16.12-17.95)	18.63 (16.58-20.25)	0.54 (0-0.84)			
<i>L. heterotoma</i> (n=7)	14.62 (13.77-15.42)	15.17 (14.46-15.6)	18.77 (17.95-19.15)	0.2 (0-0.38)		
<i>L. leipsi</i> * (n=4)	15.07 (14.15-16.2)	8.21 (7.43-8.64)	16.7 (16.01-17.67)	15.14 (14.63-15.58)	0.37 (0.16-0.6)	
<i>L. maia</i> * (n=3)	15.82 (14.74-16.74)	9.18 (8.35-10.22)	18.06 (17.5-19.04)	14.56 (14.1-14.87)	9.42 (8.83-10.29)	0
<i>L. victoria</i> (n=1)	13.47 (13.21-13.93)	14.73 (14.35-15.03)	18.73 (18.53-19.19)	11.04 (10.93-11.16)	17.01 (16.32-17.68)	16.29 (16.09-16.63)

Table 1. Percent sequence difference among species at the barcode region of the cytochrome oxidase subunit I gene. Values in bold are the percent sequence difference within each species. Sample sizes are in parentheses in the left-most column. The asterisk indicates the newly described species of *Leptopilina*.