

**Table S3.** Nucleotide comparison of *cox1* sequences of *Lecithochirium* spp. based on 443 nt long alignment. P-distance (%) is given below diagonal and the number of variable nucleotides above diagonal.

		1	2	3	4	5	6	7	8
1	OP418194 <i>Lecithochirium synodi</i>		1	2	66	70	88	88	111
2	OP918024 <i>Lecithochirium synodi</i>	0.23		1	67	71	89	89	111
3	OP918026 <i>Lecithochirium synodi</i>	0.45	0.23		68	70	90	90	112
4	OP918021 <i>Lecithochirium microstomum</i>	14.90	15.12	15.35		5	102	102	111
5	OP918022 <i>Lecithochirium microstomum</i>	15.80	16.03	15.80	1.13		104	104	113
6	OP418195 <i>Lecithochirium floridense</i>	19.86	20.09	20.32	23.02	23.48		0	115
7	OP918025 <i>Lecithochirium floridense</i>	19.86	20.09	20.32	23.02	23.48	0		115
8	OP918023 <i>Lecithochirium muraenae</i>	25.06	25.06	25.28	25.06	25.51	25.96	25.96	