

Table S2. Nucleotide comparison of the partial ITS2 sequences of *Lecithochirium* spp. based on 416 nt long alignment. P-distance (%) is given below diagonal and the number of variable nucleotides above diagonal.

		1	2	3	4	5	6
1	OP918139 <i>Lecithochirium synodi</i>		0	8	56	58	100
2	OP458338 <i>Lecithochirium synodi</i>	0		8	56	58	100
3	OP918137 <i>Lecithochirium microstomum</i>	1.92	1.92		60	62	98
4	KU527429 <i>Lecithochirium floridense</i>	13.53	13.53	14.49		11	90
5	OP458339 <i>Lecithochirium floridense</i>	14.01	14.01	14.95	2.65		89
6	OP918139 <i>Lecithochirium</i> cf. <i>muraenae</i>	24.33	24.33	23.84	21.95	21.71	