

Table S4. Uncorrected p-distances (%) estimated from the 16Sar-16Sbr fragment between the lineages of *D. nanus* – *D. walfordi* species complex recognized in this work (Lineages A–E) and the *Dendropsophus* species analyzed by Guarnizo et al. (2012). Uncorrected p-distances identified within each species or lineage are also indicated (diagonal gray line).

	1	2	3	4	5	6	7	8
1 Lineage A	0.4							
2 Lineage B	1.0	0.2						
3 Lineage C	1.3	1.8	0.2					
4 Lineage D	1.9	1.9	1.8	0.6				
5 Lineage E	1.4	1.5	1.3	1.1	0.0			
6 <i>D. labialis</i>	12.3	12.3	12.0	11.9	11.7	#		
7 <i>D. ludekei</i>	12.7	12.6	12.3	12.2	11.9	2.5	#	
8 <i>D. meridensis</i>	12.1	12.1	11.7	11.6	11.4	1.3	1.8	#