

Supplementary material. Table S2. Pairwise genetic distances between agathotanaids species for H3 and 18S sequences.

H3	<i>A. paleroi</i>			
	<i>A. frutosae</i>	0.148±0.020		
	<i>A. rushi</i>	0.084±0.017	0.174±0.023	
	<i>A. oharai</i>	0.059±0.014	0.161±0.022	0.054±0.012
18S	<i>A. jani</i>			
	<i>A. frutosae</i>	0.009±0.004		
	<i>A. paleroi</i>	0.04±0.009	0.031±0.008	