Table S1. Framework adopted in this study for taxonomic assessment and decision making, based on Schlick-Steiner *et al.* (2010).

Sources of information	Criteria	Single source	Integrated sources
Pelage pigmentation data Pelage pigmentation pattern	Phenotypic distinctness	H0 – H1	
Phylogenomic data Genome-wide sampling of nuclear DNA	Monophyly	H0 – H1	H0 – H1
Distribution data Localities of occurrence	Allopatry/ Parapatry	H0 – H1	