

Table S13 Description of all scenarios used in the Approximate Bayesian Computation analysis in DIYABC v2.0.4 to test the hybrid origin.

	Scenario	Posterior probability	Credibility interval
	1	0.978	(0.9594, 0.9966)
SSR	2	0.004	(0.0000, 0.0010)
	3	0.0216	(0.0030, 0.0420)
Nuclear genes	1	0.382	(0.0000, 0.8079)
	2	0.28	(0.0000, 0.6736)
	3	0.338	(0.0000, 0.7526)