Table S2.

	COI	CR
Mutation model	SDR06	HKY+G+I
Molecular clock model	Strict	Relaxed uncorrelated logarithmic
Rate	$1.85\pm0.4\%.Myr^{-1}$	Estimated from COI
Tree prior		Bayesian skyline
Number of groups		10
Skyline model		Piecewise-linear
Length of chain		10'000'000
Burn-in		1'000'000