DNA Barcoding of ants from the Galápagos: searching native and introduced species

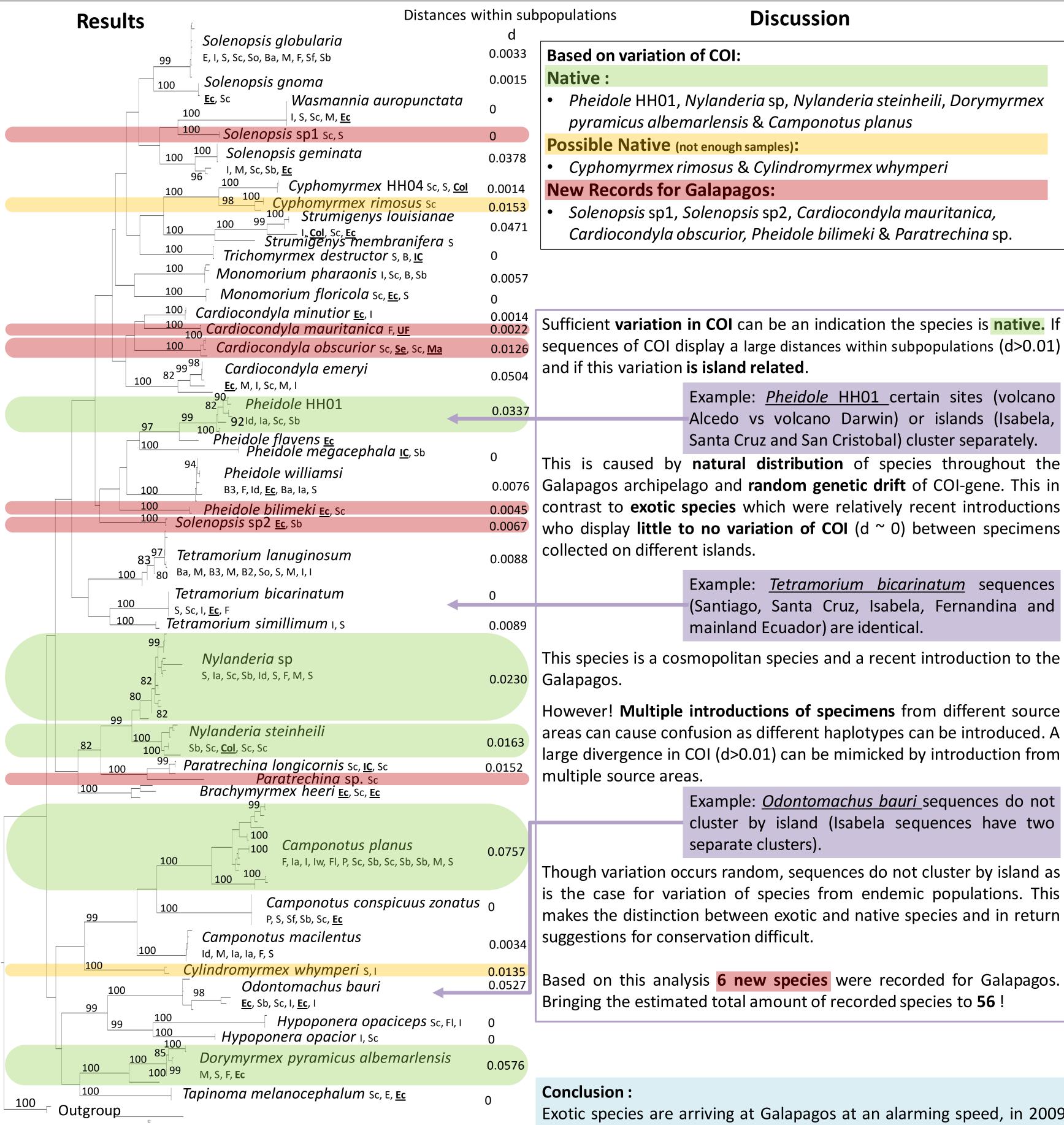
Vanderheyden A.^{1,2}, Sonet G.², Herrera H.W. ^{2,3,8}, Delsinne T.⁴, Donoso D.⁵, Lattke J.⁶, De Busschere C. ², Wittocx H. ^{1,2}, Hendrickx F. ^{2,7}, Leponce M. ², Pauly A ², Backeljau T. ^{1,2} and Dekoninck W. ^{2*}

Abstract

Until now 50 ant species have been recorded from the Galápagos Archipelago. Yet, for 26 of them it is still unclear if they are native to the Galápagos. This uncertainty is due to the fact that the ant fauna of mainland South America is too poorly known to unequivocally infer the status of species in the Galápagos. Here we explore the possibility to use amounts of COI sequence variation (a DNA barcoding fragment) as an indicator to distinguish between introduced and native species.

For the presumed endemic species C. planus, Dorymyrmex pyramicus albemarlensis and N. steinheili, large intraspecific nucleotide diversity was observed that was structured according to island geography. We observed a similar pattern of variation for two species with an unknown status i.e. Pheiodole HH1 and Nylanderia sp., which suggests that these are probably native species. Camponotus conspicuous zonatus, Pheidole megacephala, Hypoponera opacior, Hypoponera opacipes, Monomorium floricola, Cardiocondyla emery and Strumigenys louisianae, showed no COI variation, which might indicate that they are recently introduced species.

Significance: Our results stress that future studies should include a sufficient number of distinct populations from the archipelago and from areas where the species are native to increase confidence in the status of a species. If these conditions are met, our initial results showed that COI may serve as an indicative tool to distinguish native from introduced species, even if mainland relatives are unknown. Nevertheless lack of variation within COI might also be caused by other factors than recent introduction and this will be discussed and illustrated.



ML Tree of ants collected from Galápagos and Ecuador, and some other countries Galápagos Islands: Ba = Bartholomew, B2 = Beagle 2, B3 = Beagle 3, S = Santiago, E = Espanola, F = Fernandina, FI = Floreana, I = Isabela (a:Alcedo-, d:Darwin-, w:Wolf- volcano), M = Marchena, P = Pinzon, Sb = San Cristobal, Sc = Santa Cruz, Sf = Santa Fe. **BLACK/BOLD and underlined=** Extracted from specimen collected in mainland Ecuador (Ec = Ecuador) or collected elsewhere or downloaded from BOLD (Col = Columbia, IC = Ivory Coast, Se = Seychelles, UF = USA Florida and **Ma** = Madagascar,)

Affiliations

¹ Evolutionary Ecology Group, University of Antwerp, Universiteitsplein 1, B-2610 Antwerp, Belgium. ² Royal Belgian Institute of Natural Sciences, Vautierstraat 29, B-1000 Brussels, Belgium. ³ Charles Darwin Research Station, Puerto Ayora, Santa Cruz – Galápagos, Casilla 17-01-3891. Société d'Histoire Naturelle Alcide-d'Orbigny, 57 rue de Gergovie, 63170 Aubière, France. ⁵ Instituto de Ciencias Biológicas, Escuela Politécnica Nacional, Av. Ladrón de Guevara E11-253, Apdo 17-01-2759, Quito, Ecuador ⁶ Departamento de Zoologia, Universidade Federal do Paraná, Caixa Postal 19020, CEP 81531-980, Curitiba, Brazil. ⁷ Terrestrial Ecology Unit (TEREC), Biology Department, Ghent University, K.L. Ledeganckstraat 35, B-9000 Ghent, Belgium. 8 Facultad de Recursos Naturales, Departamento de Entomología, Escuela Superior Politécnica de Chimborazo, Panamericana Sur km 1 ½, Riobamba, Ecuador.

* Corresponding author. E-mail: wdekonink@naturalsciences.be







GALÁPAGOS





Conclusion:

Exotic species are arriving at Galapagos at an alarming speed, in 2009 alone 9 new records were made. Hence there are 15 new species introductions in less than 10 years! Status of species cannot always be correctly identified by use of COI-variation. These results stress the need for continued monitoring and correct species identification in Galapagos. This way potentially harmful species can be discovered and possibly eradicated.

Discussion

Solenopsis sp1, Solenopsis sp2, Cardiocondyla mauritanica,

Cardiocondyla obscurior, Pheidole bilimeki & Paratrechina sp.

Example: <u>Pheidole HH01</u> certain sites (volcano

Alcedo vs volcano Darwin) or islands (Isabela,

Example: <u>Tetramorium bicarinatum</u> sequences

(Santiago, Santa Cruz, Isabela, Fernandina and

Example: <u>Odontomachus bauri</u> sequences do not

cluster by island (Isabela sequences have two

mainland Ecuador) are identical.

separate clusters).

Santa Cruz and San Cristobal) cluster separately.

pyramicus albemarlensis & Camponotus planus

Funding Sources/Acknowledgements:

This research was done in the context of the FWO research community GO9.9.11N and was funded by FWO grant G0D2915N. We would like to thank the Charles Darwin Research Station (CDRS) and the Galápagos National Park for providing permits and field support, the Belgian Worktable for financial and logistic support at the CDRS, the Léopold III Founding and the Entomology Department of RBINS for their financial supports. This project is part of the project 'Control of invasive invertebrates' of the Charles Darwin foundation. This research also benefited from the financial support from the Belgian Directorate-General for Development Cooperation (DGD),

partim Global Taxonomy Initiative, within the framework of the CEBioS programme.