Oral presentation

Advances in the phylogenetic study of the tribe Cinchoneae (Rubiaceae) with emphasis on the genus *Cinchona* and *Ladenbergia*

Chilquillo E.^{1*}, Albán J.¹, Arakaki M.¹, Simões A.O.², Maldonado C.³, Romero P.⁴, Ramirez J.L.⁴, Cruz I.¹, Arieta L.¹ & Torres J.C.¹

¹Museo de Historia Natural, Universidad Nacional Mayor de San Marcos, Lima, Perú. ²Departamento de Biologia Vegetal, Instituto de Biologia, Universidade Estadual de Campinas, Campinas, SP, Brazil. ³Herbario Nacional de Bolivia, Instituto de Ecología, Universidad Mayor de San Andrés, La Paz, Bolivia. ⁴Facultad de Ciencias Biológicas, Universidad Nacional Mayor de San Marcos, Lima, Perú.

*Presenting author: Chilquillo E. E-mail: echilquillot@unmsm.edu.pe

In its current circumscription, the Cinchoneae tribe includes 9 genera of neotropical distribution. Its species occur mostly in the Andes in South America, with few reaching Central America. Preliminary phylogenetic studies have suggested the monophyly of Cinchoneae but intrageneric relationships are still debatable. In this study, we sampled 8 genera of Cinchoneae (with ca 50%) of species for *Cinchona* and *Ladenbergia*) and obtained multiple single-copy nuclear loci (ca 207 genes) by using the "Angiosperm353 universal probe set", which was complemented with a taxonomic review of Cinchoneae. Phylogenetic inferences were realized with multispecies pseudo-coalescent (ASTRAL III) and gene concatenation analysis (ML). Our results strongly support the monophyly of the tribe and most of the genera, except for *Ladenbergia*. Furthermore, Ciliosemina, Ladenbergia, and Remijia formed a clade, although the position of Ciliosemina (= *Remijia pedunculata*) and *Ladenbergia muzonensis* is still elusive. The position of *Ladenbergia* muzonensis is intriguing due to its intermediate floral morphology, which resembles both Remijia and *Ladenbergia* species. Additionally, our phylogeny also supports the recognition of a new species in Cinchona. Finally, our results show that sequencing data using the probe set designed for multiple gene capture is a useful tool for phylogenetic reconstructions in taxonomically complex groups.