



Genetics and Genomics in Wildlife and Human Sciences

Organised by Anja Widdig from UL-MPI-EVA and Julie Duboscq from CNRS-MNHN-Université de Paris
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Interested in how to use genetic and genomic tools to answer questions about animal social behaviour, genetic diversity of endangered population or human – nature relations?

When: 3rd of December 2020

What time: 9:00 – 12:00 Paris time / 15:00 – 18:00 Jakarta time

Where: on [Zoom](#) (Meeting ID: 614 674 6081; Passcode: 1FNCA4)

The webinar will feature the following speakers:

Rika Raffuidin (IPB) *Morphology, nest structure, and molecular characteristics of stingless bees from South and West Sulawesi*

Nicolas Césard (MNHN), Rika Raffuidin (IPB), Hélène Quach (CNRS) *Is traditional honey harvesting of the Giant Honeybee sustainable? A collaborative project between IPB and MNHN to understand the role of rafters on bee migration using genetic tools*

Hélène Quach (CNRS) *Using whole genome enrichment of endogenous DNA from non-invasive samples as a tool to study the genetic diversity of wild animal populations*

Dyah Perwitasari-Farajallah (IPB) *Madurese people cattle-raising: local wisdom and molecular approach*

Eleonora Neugebauer (UL) *Analyzing the genetic diversity of different populations of the critically endangered crested macaques (*Macaca nigra*) of Sulawesi, Indonesia*

Anja Widdig (UL), Julie Duboscq (CNRS) *Reproductive skew, genetic relatedness and social behaviour in crested macaques*

