webinar
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