

DAFTAR PUSTAKA

- Andersson, A.A., Sands, A.F., Reid, K., Hains, T., Momigliano, P., Lee, J.G., Lee, G., Rheindt, F.E., Merilä, J. and Dingle, C., 2024. Museomics sheds light on evolutionary diversity in a critically endangered cockatoo species from Wallacea. *Molecular Ecology*, p.e17616. <https://doi.org/10.1111/mec.17616>
- Arias-Sosa, L.A., Del Valle-Useche, C.M., Brieva, C., Tusso, S. and Vargas-Ramírez, M., 2024. Utilising mitochondrial barcode sequencing to evaluate phylogeographical structure and guide the release of illegally traded Amazon parrots. *Bird Conservation International*, 34, p.e43. <https://doi.org/10.1017/S0959270924000339>
- Astuti, D. 2017a. Variasi gen mitokondria cytochrome b pada dua jenis burung kakatua putih (*Cacatua alba* dan *C. moluccensis*). *Jurnal Biologi Indonesia*, 7(2). 10.14203/jbi.v7i2.3113
- Astuti, D. 2017b. Struktur genetik populasi burung betet jawa (*Psittacula alexandri alexandri*) berdasarkan sekuen DNA mitokondria gen ND2. *Jurnal Biologi Indonesia*, 13(1). 10.14203/jbi.v13i1.3101
- Astuti, D., & Prijono, S. N. 2018. Diversity of the ornate lorikeet (*Trichoglossus ornatus*) birds based on mitochondrial DNA protein coding gene. *Biosaintifika: Journal of Biology & Biology Education*, 10(2), 464-470. DOI: <http://dx.doi.org/10.15294/biosaintifika.v10i2.13501>
- Astuti, D., & Sulandari, S. 2010. The DNA sequence performance of COI gene in white cockatoos (*Cacatua*, *Psittaciformes*). *Treubia*, 37, 1-14.
- Astuti, D., Irham, M. and Ashari, H., 2018. Variasi nukleotida gen COI pada Burung Sempidan (Genus *Lophura*). *Jurnal Biologi Indonesia*, 14(1). 10.14203/jbi.v14i1.3668
- Avise, J. C. 2012. *Molecular markers, natural history and evolution*. Springer Science & Business Media.
- Bandelt, H. J., Forster, P., & Röhl, A. 1999. Median-joining networks for inferring intraspecific phylogenies. *Molecular biology and evolution*, 16(1), 37-48. <https://doi.org/10.1093/oxfordjournals.molbev.a026036>
- Barreira, A. S., Lijtmaer, D. A., & Tubaro, P. L. 2016. The multiple applications of DNA barcodes in avian evolutionary studies. *Genome*, 59(11), 899-911. DOI: 10.1139/gen-2016-0086

- Bird, J.P., Martin, R., Akçakaya, H.R., Gilroy, J., Burfield, I.J., Garnett, S.G., Symes, A., Taylor, J., Şekercioğlu, Ç.H. and Butchart, S.H.M. 2020. Generation lengths of the world's birds and their implications for extinction risk. *Conservation Biology* 34(5): 1252-1261. <https://doi.org/10.1111/cobi.13486>
- BirdLife International. 2023. *Probosciger aterrimus*. *The IUCN Red List of Threatened Species* 2023:e.T22684723A221314898.<https://dx.doi.org/10.2305/IUCN.UK.20231.RLTS.T22684723A221314898.en>. Accessed on 08 June 2024.
- Brereton, S. R., & Brereton, J. E. 2020. Sixty years of collection planning: what species do zoos and aquariums keep?. *International Zoo Yearbook*, 54(1), 131-145. <https://doi.org/10.1111/izy.12264>
- Brown, G. G., Gadaleta, G., Pepe, G., Saccone, C., & SbisA, E. 1986. Structural conservation and variation in the D-loop-containing region of vertebrate mitochondrial DNA. *Journal of molecular biology*, 192(3), 503-511. [https://doi.org/10.1016/0022-2836\(86\)90272-X](https://doi.org/10.1016/0022-2836(86)90272-X)
- Bussière, H., Mulot, B., Ruivo, E. B., & Pasquet, E. 2014. Subspecies identification with mtDNA and morphometrics in captive palm cockatoos, *Probosciger aterrimus*. *Journal of Zoo and Aquarium Research*, 2(1), 1-7. <https://doi.org/10.19227/jzar.v2i1.60>
- Castillo-Chora, V. D. J., Zamudio-Beltrán, L. E., Pozo, C., & Hernández-Baños, B. E. 2021. Phylogeography of *Habia fuscicauda* (Cardinalidae) indicates population isolation, genetic divergence and demographic changes during the Quaternary climate shifts in the Mesoamerican rainforest. *Journal of Ornithology*, 162(4), 961-976. <https://doi.org/10.1007/s10336-021-01904-x>
- CITES Appendices I, II and III, valid from 14.02. 2021. Retrieved from <https://cites.org/sites/default/files/eng/app/2021/EAppendices-2021-02-14.pdf>
- Clayton, D. A. 1982. Replication of animal mitochondrial DNA. *Cell*, 28(4), 693-705. [https://doi.org/10.1016/0092-8674\(82\)90049-6](https://doi.org/10.1016/0092-8674(82)90049-6)
- Clayton, D. A. 1984. Transcription of the mammalian mitochondrial genome. *Annual review of biochemistry*, 53(1), 573-594.
- Clayton, D. A. 1991. Replication and transcription of vertebrate mitochondrial DNA. *Annual review of cell biology*, 7(1), 453-478. <https://doi.org/10.1146/annurev.cb.07.110191.002321>
- De Bruyn, A., Martin, D. P., & Lefeuvre, P. 2014. Phylogenetic reconstruction methods: an overview. *Molecular Plant Taxonomy: Methods and Protocols*, 257-277. DOI 10.1007/978-1-62703-767-9_13

- DeCicco, L. H., DeRaad, D. A., Ostrow, E. N., & Moyle, R. G. 2023. A complete species-level phylogeny of the *Erythrura* parrotfinches (Aves: Estrildidae). *Molecular Phylogenetics and Evolution*, 187, 107883. 10.1016/j.ympev.2023.107883
- Dharmayanti, N. I. 2011. Filogenetika molekular : Metode taksonomi organisme berdasarkan sejarah evolusi. *Filogenetika Molekular : Metode Taksonomi Organisme Berdasarkan Sejarah Evolusi*, 30, 1–10. <http://medpub.litbang.pertanian.go.id/index.php/wartazoa/article/view/948>
- Doda, J. N., Wright, C. T., & Clayton, D. A. 1981. Elongation of displacement-loop strands in human and mouse mitochondrial DNA is arrested near specific template sequences. *Proceedings of the National Academy of Sciences*, 78(10), 6116-6120. <https://doi.org/10.1073/pnas.78.10.6116>
- Doyle, J. M., Hacking, C. C., Willoughby, J. R., Sundaram, M., & DeWoody, J. A. 2015. Mammalian genetic diversity as a function of habitat, body size, trophic class, and conservation status. *Journal of Mammalogy*, 96(3), 564-572. <https://doi.org/10.1093/jmammal/gyv061>
- Dutoit, L., Burri, R., Nater, A., Mugal, C. F., & Ellegren, H. 2017. Genomic distribution and estimation of nucleotide diversity in natural populations: perspectives from the collared flycatcher (*Ficedula albicollis*) genome. *Molecular ecology resources*, 17(4), 586-597. 10.1111/1755-0998.12602
- Ellegren, H., & Galtier, N. 2016. Determinants of genetic diversity. *Nature Reviews Genetics*, 17(7), 422-433. 10.1038/nrg.2016.58
- Eyre-Walker, A. 1991. *Fundamentals of Molecular Evolution*. By Li Wen-Hsiung and Graur Dan. Sinauer Associates Inc. 1991. 284 pages. £16.95. ISBN 0 87893 452 9. *Genetical Research*, 58(3), 262–262. doi:10.1017/S0016672300030032
- Felsenstein, J. 1981. Evolutionary trees from DNA sequences: a maximum likelihood approach. *Journal of molecular evolution*, 17, 368-376.
- Felsenstein J. 1985. Confidence limits on phylogenies: an approach using the bootstrap. *Evolution* 39:783–791. <http://www.jstor.org/stable/2408678> .
- Fitch, W. M. 1971. Toward defining the course of evolution: minimum change for a specific tree topology. *Systematic Biology*, 20(4), 406-416. <https://doi.org/10.1093/sysbio/20.4.406>
- Forshaw, J. M. 1989. *Parrots of the World, 3rd ed.* Lansdowne, Willoughby, Australia.
- Forshaw, J. M. 2006. *Parrots of the World; an Identification Guide*. Illustrated by Frank Knight. Princeton University Press. ISBN 978-0-691-09251-5.

- Frankham, R. 1996. Relationship of genetic variation to population size in wildlife. *Conservation biology*, 10(6), 1500-1508.
- Frankham, R. 2010. Challenges and opportunities of genetic approaches to biological conservation. *Biological conservation*, 143(9), 1919-1927. <https://doi.org/10.1016/j.biocon.2010.05.011>
- Gaveau, D.L.A., Santos, L., Locatelli, B., Salim, M.A., Husnayaen, H., Meijaard, E., Heatubun, C. and Sheil, D. 2021. Forest loss in Indonesian New Guinea (2001–2019): Trends, drivers and outlook. *Biological Conservation* 261: 109225. <https://doi.org/10.1016/j.biocon.2021.109225>
- Gill, Frank; Donsker, David; Rasmussen, Pamela, eds. (January 2023). "Parrots, cockatoos". *IOC World Bird List Version 13.1*. International Ornithologists' Union. Retrieved 20 February 2023.
- Grant, P.R. (1968) Bill size, body size, and the ecological adaptations of bird species to competitive situations on islands. *Systematic Biology* 17 , 319–333.
- Hall, T. A. 1999. BioEdit: a user-friendly biological sequence alignment editor and analysis program for Windows 95/98/NT. In *Nucleic acids symposium series* (Vol. 41, No. 41, pp. 95-98).
- Hamilton, T.H. (1961) The adaptive significances of intraspecific trends of variation in wing length and body size among bird species. *Evolution* 15, 180–195.
- Haryoko, T., O'Hara, M., Mioduszewska, B., Sutrisno, H., Prasetyo, L.B. and Mardiasuti, A., 2021. Implementation of species protection act for the conservation of Tanimbar corella, *Cacatua goffiniana* (Roselaar & Michels, 2004). *Biodiversitas Journal of Biological Diversity*, 22(4).
- Havird, J. C., & Santos, S. R. 2014. Performance of single and concatenated sets of mitochondrial genes at inferring metazoan relationships relative to full mitogenome data. *PLoS One*, 9(1), e84080. <https://doi.org/10.1371/journal.pone.0084080>
- Heinsohn, R., Zdenek, C., Keighley, M.V., Murphy, S.A., Legge, S.M. & Garnett, S.T. 2021. Australian Palm Cockatoo *Probosciger aterrimus macgillivrayi*. In: S.T. Garnett and G.B. Baker (eds), *The Action Plan for Australian Birds 2020*, pp. 407-410. CSIRO Publishing, Melbourne.
- Huang, Z., Yang, C., & Ke, D. 2016. DNA barcoding and phylogenetic relationships in Anatidae. *Mitochondrial DNA Part A*, 27(2), 1042-1044. DOI: 10.3109/19401736.2014.926545

- Huang, Z., & Ke, D. 2016a. Structure and variation of the Anseriformes mitochondrial DNA control region. *Mitochondrial DNA Part A*, 27(3), 2036-2039. DOI: 10.3109/19401736.2014.974177
- Huang, Z., & Ke, D. 2016b. Structure and evolution of the Phasianidae mitochondrial DNA control region. *Mitochondrial DNA Part A*, 27(1), 350-354. DOI: 10.3109/19401736.2014.895987
- Igag, P., Mack, A.L., Legge, S. and Heinsohn, R. 2019. Breeding biology of three large, sympatric rainforest parrots in New Guinea: Palm Cockatoo, Pesquet's Parrot and Eclectus Parrot. *Emu - Austral Ornithology* 119(3): 196-204. <https://doi.org/10.1080/01584197.2019.1616211>
- Jarulis, J., Solihin, D. D., Mardiasuti, A., & Prasetyo, L. B. 2020. Characters of mitochondrial DNA d-loop hypervariable III fragments of Indonesian Rhinoceros Hornbill (*Buceros rhinoceros*) (Aves:Bucerotidae). *Treubia*, 47(2), 99-110. 10.14203/treubia.v47i2.3971
- Jarulis, J., Muslim, C., Kamilah, S. N., Utama, A. F., Permana, D., Sari, M. M., ... & Jannah, I. M. 2021. DNA barcode of Enggano hill myna, *Gracula religiosa enganensis* (Aves: Sturnidae) based on mitochondrial DNA cytochrome oxidase subunit I. *Biodiversitas Journal of Biological Diversity*, 22(4). DOI: 10.13057/biodiv/d220407
- Jobling, James A. 2010. *The Helm Dictionary of Scientific Bird Names*. London: Christopher Helm. pp. 317, 58. ISBN 978-1-4081-2501-4.
- Johnson, K.P. and Sorenson, M.D., 1998. Comparing molecular evolution in two mitochondrial protein coding genes (cytochromeband ND2) in the dabbling ducks (Tribe: Anatini). *Molecular phylogenetics and evolution*, 10(1), pp.82-94.
- Kangayan, M., De, A. K., Bhattacharya, D., Tamilvanan, S., Ponraj, P., Alyethodi, R. R., ... & Chakurkar, E. B. 2022. Genetic structure, population diversity and ancestry of Nicobari fowl based on mtDNA complete D-loop sequences. *Journal of Genetics*, 101(2), 31. <https://doi.org/10.1007/s12041-022-01372-z>
- Kartavtsev, Y.P., 2011. Divergence at Cyt-b and Co-1 mtDNA genes on different taxonomic levels and genetics of speciation in animals. *Mitochondrial DNA*, 22(3), pp.55-65.
- Kerr, K. C., Stoeckle, M. Y., Dove, C. J., Weigt, L. A., Francis, C. M., & Hebert, P. D. 2007. Comprehensive DNA barcode coverage of North American birds. *Molecular ecology notes*, 7(4), 535-543. doi: 10.1111/j.1471-8286.2006.01670.x

- Keighley, M. V., Heinsohn, R., Langmore, N. E., Murphy, S. A., & Peñalba, J. V. 2019. Genomic population structure aligns with vocal dialects in Palm Cockatoos (*Probosciger aterrimus*); evidence for refugial late-Quaternary distribution?. *Emu-Austral Ornithology*, 119(1), 24-37. <https://doi.org/10.1080/01584197.2018.1483731>
- Keighley, M.V., Haslett, S., Zdenek, C.N. and Heinsohn, R. 2021. Slow breeding rates and low population connectivity indicate Australian palm cockatoos are in severe decline. *Biological Conservation* 253: 108865.<https://doi.org/10.1016/j.biocon.2020.108865>
- Khan, H. A., Arif, I. A., Altwaijry, N. A., & Ahamed, A. 2023. DNA barcodes of Saudi Arabian birds: Implications for species identification and diversity analysis. *Journal of King Saud University-Science*, 35(8), 102887. <https://doi.org/10.1016/j.jksus.2023.102887>
- Kimura, M. 1980. A simple method for estimating evolutionary rate of base substitutions through comparative studies of nucleotide sequences. *Journal of Molecular Evolution*, 16, 111–120.
- Kim, J. I., Do, T. D., Choi, Y., Yeo, Y., & Kim, C. B. 2021. Characterization and comparative analysis of complete mitogenomes of three Cacatua Parrots (Psittaciformes: Cacatuidae). *Genes*, 12(2), 209.DOI: 10.3390/genes12020209
- Kress, W. J., García-Robledo, C., Uriarte, M., & Erickson, D. L. 2015. DNA barcodes for ecology, evolution, and conservation. *Trends in ecology & evolution*, 30(1), 25-35. DOI:<https://doi.org/10.1016/j.tree.2014.10.008>
- Kuhl, Heinrich. 1820. "Conspectus Psittacorum". *Verhandlungen der Kaiserlichen Leopoldinisch-Carolinische Akademie der Naturforscher* (in Latin). **10**: 1–104 [12].
- Kummer, E., & Ban, N. 2021. Mechanisms and regulation of protein synthesis in mitochondria. *Nature Reviews Molecular Cell Biology*, 22(5), 307-325. <https://doi.org/10.1038/s41580-021-00332-2>
- Lai, Y. T., Yeung, C. K., Omland, K. E., Pang, E. L., Hao, Y. U., Liao, B. Y., ... & Li, S. H. 2019. Standing genetic variation as the predominant source for adaptation of a songbird. *Proceedings of the National Academy of Sciences*, 116(6), 2152-2157. <https://doi.org/10.1073/pnas.1813597116>
- Librado, P., & Rozas, J. 2009. DnaSP v5: a software for comprehensive analysis of DNA polymorphism data. *Bioinformatics*, 25(11), 1451-1452. <https://doi.org/10.1093/bioinformatics/btp187>
- Luttrell, S. A., Drovetski, S., Dahlan, N. F., Eubanks, D., & Dove, C. J. 2020. ND2 as an additional genetic marker to improve identification of diving ducks

- involved in bird strikes. *Human-Wildlife Interactions*. <https://doi.org/10.26077/4DC6-AD95>
- Markert, J. A., Champlin, D. M., Gutjahr-Gobell, R., Grear, J. S., Kuhn, A., McGreevy, T. J., ... & Nacci, D. E. 2010. Population genetic diversity and fitness in multiple environments. *BMC evolutionary biology*, 10, 1-13.
- Marquardt, C. and K. Howard. 1998. Nutritional requirements of adult palm cockatoos In: Palm Cockatoo SSP Husbandry Manual. American Association of Zoos and Aquariums. Caloosahatchee Aviary and Botanical Gardens and the International Aviculturists Society
- Marshall, H. D., & Baker, A. J. 1997. Structural conservation and variation in the mitochondrial control region of fringilline finches (*Fringilla* spp.) and the greenfinch (*Carduelis chloris*). *Molecular Biology and evolution*, 14(2), 173-184. <https://doi.org/10.1093/oxfordjournals.molbev.a025750>
- McGowan, P. J., Traylor-Holzer, K., & Leus, K. 2017. IUCN guidelines for determining when and how ex situ management should be used in species conservation. *Conservation Letters*, 10(3), 361-366. <https://doi.org/10.1111/conl.12285>
- Mereu, P., Pirastru, M., Satta, V., Frongia, G. N., Kassinis, N., Papadopoulos, M., ... & Leoni, G. G. 2019. Mitochondrial D-loop Sequence Variability in Three Native Insular Griffon Vulture (*Gyps fulvus*) Populations from the Mediterranean Basin. *BioMed Research International*, 2019(1), 2073919. <https://doi.org/10.1155/2019/2073919>
- Murphy, S. A., Double, M. C., & Legge, S. M. 2007. The phylogeography of palm cockatoos, *Probosciger aterrimus*, in the dynamic Australo-Papuan region. *Journal of Biogeography*, 34(9), 1534-1545. <https://doi.org/10.1111/j.1365-2699.2007.01706.x>
- Nei, M. 1987. *Molecular evolutionary genetics*. Columbia university press.
- Nei, M., & Kumar, S. 2000. *Molecular evolution and phylogenetics*. Oxford University Press, USA.
- Nijman, V. and Shepherd, C., Oxford Brookes University–Research Archive and Digital Asset Repository (RADAR).
- Osawa, S., Z. Su, & Imura, Y. 2004. Molecular Phylogeny and Evolution of Carabid Graound Beetles. Springer-Verlag Tokyo: SNP Best-set Typesetter Ltd.
- Palacios, C., Campagna, L., Parra, J.L. and Cadena, C.D., 2023. Mito-nuclear discordance in the phenotypically variable Andean hummingbirds *Coeligena bonapartei* and *Coeligena helianthea* (Trochilidae). *Biological Journal of the Linnean Society*, 139(2), pp.145-157. <https://doi.org/10.1093/biolinnean/blad013>

- Peraturan Menteri Kehutanan. 2012. Peraturan Menteri Kehutanan Republik Indonesia P.31/MenhutII/2012 Tentang Lembaga Konservasi.
- Pratt, Thane K.; Bruce M. Beehler. 2015. *Birds of New Guinea: Second Edition*. Princeton University Press. p. 122. ISBN 9780691095639.
- Pulgarín-R, P. C., Olivera-Angel, M., Ortíz, L., Nanclares, D., Velásquez-Restrepo, S., & Díaz-Nieto, J. F. 2021. DNA barcodes of birds from northern Colombia. *Biodiversity Data Journal*, 9. doi: 10.3897/BDJ.9.e64842
- Rafiq, H., Nazir, F., & Khan, F. M. 2020. Molecular Taxonomy and Phylogenetic Analysis of Dove and Pigeon Species (Aves: Columbidae) of Pakistan, Based on COI Region of Mitochondrial DNA. *Journal of Bioresource Management*, 7(3), 3. DOI: <https://doi.org/10.35691/JBM.0202.0138>
- Randi, E., & Lucchini, V. 1998. Organization and evolution of the mitochondrial DNA control region in the avian genus *Alectoris*. *Journal of Molecular Evolution*, 47(4), 449-462. <https://doi.org/10.1007/PL00006402>
- Rowley, I. and G. M. Kirwan 2020. Palm Cockatoo (*Probosciger aterrimus*), version 1.0. In *Birds of the World* (J. del Hoyo, A. Elliott, J. Sargatal, D. A. Christie, and E. de Juana, Editors). Cornell Lab of Ornithology, Ithaca, NY, USA. <https://doi.org/10.2173/bow.palcoc1.01>
- Rozas, J., Sánchez-DelBarrio, J. C., Messeguer, X., & Rozas, R. 2003. DnaSP, DNA polymorphism analyses by the coalescent and other methods. *Bioinformatics*, 19(18), 2496-2497. <https://doi.org/10.1093/bioinformatics/btg359>
- Ruokonen, M., & Kvist, L. 2002. Structure and evolution of the avian mitochondrial control region. *Molecular Phylogenetics and Evolution*, 23(3), 422-432. [https://doi.org/10.1016/S1055-7903\(02\)00021-0](https://doi.org/10.1016/S1055-7903(02)00021-0)
- Saito, N. and Nei, M. 1987. The neighbor-joining method: A new method for reconstructing phylogenetic trees. *Molecular Biology and Evolution*, 4, 406-425.
- Sober, E. 1991. *Reconstructing the past: Parsimony, evolution, and inference*. MIT press.
- Sonet, G., Breman, F. C., Lenglet, G., Louette, M., Montañés, G., Nagy, Z., ... & Verheyen, E. 2011. Applicability of DNA barcoding to museum specimens of birds from the Democratic Republic of Congo. *Tropical Vertebrates in a Changing World*, 57, 117-208.
- Sorenson, M. D., Ast, J. C., Dimcheff, D. E., Yuri, T., & Mindell, D. P. 1999. Primers for a PCR-based approach to mitochondrial genome sequencing in birds and other vertebrates. *Molecular phylogenetics and evolution*, 12(2), 105-114. <https://doi.org/10.1006/mpev.1998.0602>

- Stiller, J., & Zhang, G. 2019. Comparative phylogenomics, a stepping stone for bird biodiversity studies. *Diversity*, 11(7), 115. doi:10.3390/d11070115
- Tamura, K., Stecher, G., & Kumar, S. 2021. MEGA11: molecular evolutionary genetics analysis version 11. *Molecular biology and evolution*, 38(7), 3022-3027. <https://doi.org/10.1093/molbev/msab120>
- Thompson, J. D., Gibson, T. J., Plewniak, F., Jeanmougin, F., & Higgins, D. G. 1997. The CLUSTAL_X windows interface: flexible strategies for multiple sequence alignment aided by quality analysis tools. *Nucleic acids research*, 25(24), 4876-4882. <https://doi.org/10.1093/nar/25.24.4876>
- Tvardikova, K. 2023. *Probosciger aterrimus*, Gmelin, JF, 1788. ew Guinea Birds online, <http://pngbirds.myspecies.info>, accessed on November 7, 2023.
- Urantówka, A.D., Krocak, A., Silva, T., Padrón, R.Z., Gallardo, N.F., Blanch, J., Blanch, B. and Mackiewicz, P., 2018. New insight into parrots' mitogenomes indicates that their ancestor contained a duplicated region. *Molecular biology and evolution*, 35(12), pp.2989-3009. <https://doi.org/10.1093/molbev/msy189>
- Van de Peer, Y., & Salemi, M. 2003. Phylogeny inference based on distance methods. *The Phylogenetic Handbook, a Practical Approach to DNA and Protein Phylogeny*, 101-136.
- Vázquez-López, M., Cortés-Rodríguez, N., Robles-Bello, S. M., Bueno-Hernández, A., Zamudio-Beltrán, L. E., Ruegg, K., & Hernández-Baños, B. E. 2021. Phylogeography and morphometric variation in the Cinnamon Hummingbird complex: *Amazilia rutila* (Aves: Trochilidae). *Avian Research*, 12, 1-17. 10.1186/s40657-021-00295-0
- Wenink, P. W., Baker, A. J., & Tilanus, M. G. 1994. Mitochondrial control-region sequences in two shorebird species, the turnstone and the dunlin, and their utility in population genetic studies. *Molecular Biology and evolution*, 11(1), 22-31. <https://doi.org/10.1093/oxfordjournals.molbev.a040089>
- White, N. E., Phillips, M. J., Gilbert, M. T. P., Alfaro-Núñez, A., Willerslev, E., Mawson, P. R., ... & Bunce, M. 2011. The evolutionary history of cockatoos (Aves: Psittaciformes: Cacatuidae). *Molecular phylogenetics and evolution*, 59(3), 615-622. <https://doi.org/10.1016/j.ympev.2011.03.011>
- Wright, T. F., Schirtzinger, E. E., Matsumoto, T., Eberhard, J. R., Graves, G. R., Sanchez, J. J., ... & Fleischer, R. C. 2008. A multilocus molecular phylogeny of the parrots (Psittaciformes): support for a Gondwanan origin during the Cretaceous. *Molecular biology and evolution*, 25(10), 2141-2156. <https://doi.org/10.1093/molbev/msn160>
- Yang, C., Lian, T., Wang, Q. X., Huang, Y., & Xiao, H. 2016. Structural characteristics of the Relict Gull (*Larus relictus*) mitochondrial DNA

control region and its comparison to other Laridae. *Mitochondrial DNA Part A*, 27(4), 2487-2491. 10.3109/19401736.2015.1033711

Zein, M. S. A., Haryoko, T., Fitriana, Y. S., Sulistyadi, E., & Prawiradilaga, D. M. 2017. Aplikasi kajian DNA molekuler dan fenotipik pada program pelepasliaran burung kakatua. *Jurnal Biologi Indonesia*, 13(1). 10.14203/jbi.v13i1.3105

