

Articles & Theses Citing StrAuto (2011–2016)

Last updated: April 16, 2017

1. Larsen R.J. *et al* 2017. Patterns of morphological and molecular evolution in the Antillean tree bat, *Ardops nichollsi* (Chiroptera: Phyllostomidae). Mammology Papers: University of Nebraska State Museum. doi: <http://digitalcommons.unl.edu/museummammalogy/269/>
2. Harrison T. *et al* 2017. Geographically structured genetic variation in the *Medicago lupulina-Ensifer* mutualism. BioRxiv doi: <https://doi.org/10.1101/117192>
3. Hoyos-Villegas V. *et al* 2016. Genome-wide Association Analysis for Drought Tolerance and Associated Traits in Common Bean. The Plant Genome, DOI: <http://dx.doi.org/10.3835/plantgenome2015.12.0122>
4. Tay Y.C. *et al* 2016. Beyond the Coral Triangle: High genetic diversity and near panmixia in Singapore's populations of the broadcast spawning sea star *Protoreaster nodosus*. *Royal Society Open Science* 3(8):160253. DOI: <http://dx.doi.org/10.1098/rsos.160253>
5. Plumer L. *et al* 2016. Wolves Recolonizing Islands: Genetic Consequences and Implications for Conservation and Management. *PLoS ONE* 11(7): e0158911. DOI: <http://dx.doi.org/10.1371/journal.pone.0158911>
6. Hubbard A. *et al* 2015. Field pathogenomics reveals the emergence of a diverse wheat yellow rust population. *Genome Biology*, 16:23. DOI: <http://dx.doi.org/10.1186/s13059-015-0590-8>
7. Patel S. *et al* 2015 Characterizing a hybrid zone between a cryptic species pair of freshwater snails. *Molecular Ecology*, 24:643–655. DOI: <http://dx.doi.org/10.1111/mec.13049>
8. Putman A.I. & Carbone, I. 2015 Challenges in analysis and interpretation of microsatellite data for population genetic studies. *Ecology & Evolution*, 4(22):4399–4428. DOI: <http://dx.doi.org/10.1002/ece3.1305> **Review Article**
9. Besnier F. & Glover K.A. 2013. ParallelStructure: A R package to distribute parallel runs of the population genetics program STRUCTURE on multi-core computers. *PLoS ONE*, 8(7): e70651. DOI: <http://dx.doi.org/10.1371/journal.pone.0070651>
10. Porras-Hurtado *et al* 2013 An overview of STRUCTURE: applications, parameter settings, and supporting software. *Frontiers in Genetics*, 4(98)1:13. DOI: <http://dx.doi.org/10.3389/fgene.2013.00098> **Review Article**
11. Lah L. *et al* 2014 Investigating harbor porpoise (*Phocoena phocoena*) population differentiation using RAD-tag genotyping by sequencing. FAO AGRIS Database Working Paper. <http://goo.gl/NVah2D>.
12. Huang M. 2012 Genetic Analysis of Quality Traits for Food-Grade Soybean (*Glycine max* L. Merr.) in a Breeding Population. A Master's Thesis in Horticulture and Crop Science at the Ohio State University.