

Laboratories of Analytical Biology Publications from 2015

Publications listed in alphabetical order.

Arcila, D, Alexander Pyron, R., Tyler, J.C., **G Ortí & R Betancur-R**. 2015. An evaluation of fossil tip-dating versus node-age calibrations in tetraodontiform fishes (Teleostei: Percomorphaceae). *Molecular Phylogenetics & Evolution* 82: 131-145.

Angulo A, **CC Baldwin & DR Robertson**. In Press. A new species of *Leptoderma* Vaillant, 1886 (Osmeriformes: Alepocephalidae) from the eastern Pacific Ocean. *Zootaxa*.

Attaran-Farimani G, S Estekani, VG Springer, O Crimmen, GD Johnson & **CC Baldwin**. In press. Validation of the synonymy of the teleost blenniid fish species *Salarias phantasticus* Boulenger 1897 and *Salarias anomalous* Regan 1905 with *Salarias pulcher* Murray 1887 (= *Ecsenius pulcher*) based on DNA barcoding and morphology. *Zootaxa*.

Baldwin, CC & DR Robertson. 2015. A new, mesophotic *Coryphopterus* goby (Teleostei: Gobiidae) from the southern Caribbean, with comments on relationships and depth distributions within the genus. *Zookeys* 513:123–142.

Bennett, KL, **Y-M Linton**, F Shija, M Kaddumukasa, R Djouaka, G Misinzo, J Lutwama, Y-M Huang, **LB Mitchell**, M Richards & C Walton. 2015. Molecular differentiation of the African yellow fever vector *Aedes bromeliae* (Diptera: Culicidae) from its sympatric non-vector sister species, *Aedes lili*. *PLoS Neglected Tropical Diseases* 9(12): e0004250. [doi:10.1371/journal.pntd.0004250](https://doi.org/10.1371/journal.pntd.0004250)

Betancur-R. R, G Ortí, & AR Pyron. 2015. In press. Fossil-based comparative analyses reveal ancient marine ancestry erased by extinction in ray-finned fishes. *Ecology Letters*.

Blaimer, B, SG Brady, TR Schultz & BL Fisher. 2015. Functional and phylogenetic approaches reveal the evolution of diversity in a hyper diverse biota. *Ecography* 38: 901-912. [doi:10.1111/ecog.01370](https://doi.org/10.1111/ecog.01370)

Blaimer, BB, SG Brady, TR Schultz, MW Lloyd, BL Fisher & PS Ward. 2015. Phylogenomic methods outperform traditional multi-locus approaches in resolving deep evolutionary history: a case study of formicine ants. *BMC Evolutionary Biology* 15:271. [doi:10.1186/s12862-015-0552-5](https://doi.org/10.1186/s12862-015-0552-5)

Blakeslee, AMH. 2015. Parasites and Genetics in Marine Invertebrate Introductions: Signatures of Diversity Declines across Systems. In: *Biological Invasions in Changing Ecosystems: Vectors, Ecological Impacts, Management and Predictions* (Ed. J. Canning Clode). De Gruyter Open (Warsaw, Poland). pp 138-182.

Bohu T, C Santelli, DM Akob, T Neu, V Ciobota, J Popp, S Nietzsche & K Küsel. 2015. *Mesorhizobium australicum* T-G1 initiated bixbyite formation at acidic pH. *Frontiers in Microbiology* 17 July 2015 [doi:10.3389/fmicb.2015.00734](https://doi.org/10.3389/fmicb.2015.00734)

Borges, R, I Khan, **WE Johnson**, MTP Gilbert, G Zhang, ED Jarvis, SJ O'Brien, & A Antunes. 2015. Gene loss, adaptive evolution, and the co-evolution of plumage coloration genes with opsins in birds. *BMC Genomics* 16:751.

Bowles, DE, SC Britch, KJ Linthicum, RN Johnson, Y-M Linton & GB White. 2015. *Sand Flies: Significance, Surveillance, and Control in Contingency Operations*. Armed Forces Pest Management Board Technical Guide 49: 1-72.

Budd, C, EA Zimmer & JR Freeland. 2015. Conservation genetics of *Magnolia acuminata*, an endangered species in Canada: Can genetic diversity be maintained in fragmented, peripheral populations? *Conservation Genetics*. [doi:10.1007/s10592-015-0746-9](https://doi.org/10.1007/s10592-015-0746-9)

Cairns, SD & HH Wirshing. 2015. Phylogenetic reconstruction of scleraxonian octocorals supports the resurrection of the family Spongiodermidae (Cnidaria, Alcyonacea). *Invertebrate Systematics* 29(4): 345-368. [doi:10.1071/IS14063](https://doi.org/10.1071/IS14063)

Chen, J, J Zhao, DL Erickson, N Xia & WJ Kress. 2015. Testing DNA barcodes in closely related species of *Curcuma* (Zingiberaceae) from Myanmar and China. *Molecular Ecology Resources*, 15(2): 337-348. [doi:10.1111/1755-0998.12319](https://doi.org/10.1111/1755-0998.12319)

De Smet WH, G Melone, D Fontaneto & F Leasi. 2015. Marine Rotifera – Fauna d’Italia Vol. L. Ed. Calderini, Bologna. Pp: 272. ISBN 978-88-506-5490-1.

Díaz MC, RW Thacker, NE Redmond, T Perez & AG Collins. 2015. *Vansoestia caribensis* gen. nov., sp. nov.: first report of the family Ianthellidae (Verongida, Demospongiae) in the Caribbean. *Zootaxa* 3956: 403-412. [doi:10.11646/zootaxa.3956.3.5](https://doi.org/10.11646/zootaxa.3956.3.5)

Dobrynin, P, S Liu, G Tamazian, Z Xiong, A Yurchenko, K Krasheninnikova, S Kliver, A Schmidt-Kuntzel, K-P Koepfli, WE Johnson, L Kuderna, R García-Pérez, M deManuel, R Godinez, A Komissarov, A Makunin, V Brukhin, W Qiu, L Zhou, F Li, J Yi, C Driscoll, A Antunes, TK Oleksyk, E Eizirik, P Perelman, M Roelke, D Wildt, M Diekhans, T Marques-Bonet, L Marker, J Bhak, J Wang, G Zhang & SJ O’Brien. 2015. Genomic Legacy of the African Cheetah, *Acinonyx jubatus*. *Genome Biology* 16:277.

Dominguez J, M Aira, JW Breinholt, M Stojanovic, SW James & M Pérez-Losada. 2015. Underground evolution: new roots for the old tree of lumbricid earthworms. *Molecular Phylogenetics & Evolution* 83: 7–19.

Duan, L, J Wen, X Yang, P-L Liu, E Arslan, K Ertugrul & Z-Y Chang. 2015. Phylogeny of *Hedysarum* and tribe Hedysareae (Leguminosae: Papilionoideae) inferred from sequence data of ITS, matK, trnL-F and psbA-trnH. *Taxon* 64(1): 49-64. [doi:10.12705/641.26](https://doi.org/10.12705/641.26)

Dziki, A, G Binford, JA Coddington, I Agnarsson. 2015. *Spintharus flavidus* in the Caribbean—a 30 million year biogeographical history and radiation of a ‘widespread species’ *PeerJ*. e1422.

Engene, N, A Tronholm, L Salvador-Reyes, H Luesch & VJ Paul. 2015. *Caldora penicillata* gen. nov., comb. nov. (Cyanobacteria), a pantropical marine species with biomedical relevance. *Journal of Phycology*, 51(4): 670-681. [doi:10.1111/jpy.12309](https://doi.org/10.1111/jpy.12309)

Faircloth, BC, MG Branstetter, ND White & SG Brady. 2015. Target enrichment of ultraconserved elements from arthropods provides a genomic perspective on relationships among Hymenoptera. *Molecular Ecology Resources* 15: 489–501. [doi:10.1111/1755-0998.12328](https://doi.org/10.1111/1755-0998.12328)

Fowler AE, **AMH Blakeslee**, J Canning-Clode, MF Repetto, AM Phillip, JT Carlton, FC Moser, **GM Ruiz & AW Miller**. 2015. Opening Pandora's bait box: a potent vector for biological invasions of live marine species. *Diversity and Distributions*. [doi:10.1111/ddi.12376](https://doi.org/10.1111/ddi.12376)

García-Robledo, C, CL Staines & WJ Kress. 2015. A new species of bromeliad-feeding Cephaloleia Chevrolat (Coleoptera, Chrysomelidae, Cassidinae) from Costa Rica: evidence from DNA barcodes, larval and adult morphology and insect diets. *ZooKeys*, 477: 143-155. [doi:10.3897/zookeys.477.8220](https://doi.org/10.3897/zookeys.477.8220)

García-Robledo, C, EK Kuprewicz, CL Staines, TL Erwin & WJ Kress. 2015. Limited tolerance by insects to high temperatures across tropical elevational gradients and the implications of global warming for extinction. *Proceedings of the National Academy of Science*: early view.

Goodheart, JA, AL Bazinet, **AG Collins** & MP Cummings. 2015. Relationships within Cladobranchia (Gastropoda: Nudibranchia) based on RNA-Seq data: An initial investigation. *Royal Society Open Science* 150196. [doi:10.1098/rsos.150196](https://doi.org/10.1098/rsos.150196)

Gunay, F, B Alten, F Simsek, A Aldemir & **Y-M Linton**. 2015. Barcoding Turkish *Culex* mosquitoes to facilitate arbovirus vector incrimination studies reveals hidden diversity and new potential vectors. *Acta Tropica* 143: 112-120. [doi:10.1016/j.actatropica.2014.10.013](https://doi.org/10.1016/j.actatropica.2014.10.013)

Harasewych, MG. Systematics and Phylogeography of *Cerion sensu stricto* (Mollusca: Gastropoda: Pulmonata: Cerionidae) from Aruba, Curaçao and Bonaire. *Journal of Molluscan Studies*. 81: 66-84. electronic 2014. Hard copy 2015. [doi:10.1093/mollus/eyu062](https://doi.org/10.1093/mollus/eyu062)

Harasewych, MG & I Témkin. 2015. The Miocene Genus *Mantellina*(Bivalvia: Limidae) discovered living in the deep reefs off Curaçao, with the description of a new species. *Journal of Molluscan Studies*, 81: 442-454. [doi:10.1093/mollus/eyv021](https://doi.org/10.1093/mollus/eyv021)

Harasewych, MG, AM Windsor, E Lopez-Vera & FG Thompson. On the phylogenetic relationships of the genus *Mexitrophia* and of the family Cerionidae (Gastropoda: Eupulmonata). *The Nautilus* 129:156-162.

Harris, AJ, **S Lutz, P Acevedo-Rodríguez & J Wen**. 2015. The utility of the morphological variation of pollen for resolving the evolutionary history of *Billia* (subfam. Hippocastanoideae, Sapindaceae). *Journal of Systematics and Evolution* 53(3): 228-238. [doi:10.1111/jse.12130](https://doi.org/10.1111/jse.12130)

Hazelton, ELG, **MK McCormick**, M Sievers, KM Kettenring & DF Whigham. 2015. Stand age is associated with clonal diversity, but not vigor, community structure, or insect herbivory in Chesapeake Bay *Phragmites australis*. *Wetlands* 35: 877-888.

Hostetler, JA & RB Chandler. 2015. Improved state-space models for inference about spatial and temporal variation in abundance from count data. *Ecology*: in press. [doi:10.1890/14-1487.1](https://doi.org/10.1890/14-1487.1)

Ickert-Bond, S, JM Gerrath, U Poslusny & **J Wen**. 2015. Inflorescence development in the *Vitis-Ampelocissus* clade of Vitaceae: the unusual lamellate inflorescence of *Pterisanthes*. *Botanical Journal of the Linnean Society* 179(4): 725-741. [doi:10.1111/boj.12348](https://doi.org/10.1111/boj.12348)

Jezkova, T, JR Jaeger, V Olah-Hemmings, KB Jones, RA Lara-Resendiz, **DG Mulcahy** & BR Riddle. 2015. Range and niche shifts in response to past climate change in the desert horned lizard *Phrynosoma platyrhinos*. *Ecography* Early View: [doi:10.1111/ecog.01464](https://doi.org/10.1111/ecog.01464) *Smithsonian Science News*

Kartzinel, TR, PA Chen, TC Coverdale, **DL Erickson, WJ Kress, ML Kuzmina**, DI Rubenstein, W Wang & RM Pringle. 2015. DNA metabarcoding illuminates dietary niche partitioning by African large herbivores. *Proceedings of the National Academy of Sciences*, 112(26): 8019-8024. [doi:10.1073/pnas.1503283112](https://doi.org/10.1073/pnas.1503283112)

Kayal E, B Bentlage, P Cartwright, A Yanagihara, DJ Lindsay, RR Hopcroft & **AG Collins**. 2015. Phylogenetic analysis of higher-level relationships within Hydroidolina (Cnidaria: Hydrozoa) using mitochondrial genome data and insight into their mitochondrial transcription. *PeerJ*. 3: e1403. [doi:10.7717/peerj.1403](https://doi.org/10.7717/peerj.1403)

Kearns, AM, L Joseph, LC White, JJ Austin, C Baker, **AC Driskell**, JF Malloy & KE Omland. 2015. *In press*. Norfolk Island Robins are a distinct and endangered species: ancient DNA unlocks surprising relationships and phenotypic discordance within the Australo-Pacific Robin. *Conservation Genetics*.

Koepfli, K-P, B Paten, A Antunes, K Belov, C Bustamante, TA Castoe, H Clawson, AJ Crawford, M Diekhans, D Distel, R Durbin, D Earl, MK Fujita, T Gamble, A Georges, N Gemmell, MTP Gilbert, J Marshall Graves, RE Green, G Hickey, E Jarvis, **WE Johnson**, A Komissarov, I Korf, R Kuhn, DM Larkin, H Lewin, JV Lopez, J Ma, T Marques-Bonet, W Miller, R Murphy, P Pevsner, B Shapiro, C Steiner, G Tamazian, B Venkatesh, J Wang, R Wayne, E Wiley, H Yang, G Zhang, D Haussler, O Ryder, & SJ O'Brien. 2015. The Genome 10K Project - A way forward. *Annual Reviews of Animal Biosciences*. 3:22.1-22.55.

Koepfli, K-P, J Pollinger, J Robinson, A Lea, S Hendricks, R Schweizer, O Thalmann, P Silva, Z Fan, A Yurchenko, P Dobrynin, A Makunin, JA Cahill, B Shapiro, E Geffen, F Zachos, R Godinho, JA Leonard, **WE Johnson**, SJ O'Brien, B Van Valkenburgh, & RK Wayne. 2015. Parallel evolution of a Golden Jackal-like morphology in North and East Africa. *Current Biology*.

Kress, WJ, C García-Robledo, M Uriarte, & DL Erickson. 2015. DNA barcodes for ecology, evolution, and conservation. *Trends in Ecology & Evolution*, 30(1): 25-35. [doi:10.1016/j.tree.2014.10.008](https://doi.org/10.1016/j.tree.2014.10.008)

Kuntner, M, M Năpăruș, D Li & JA Coddington. 2014. Phylogeny predicts future habitat shifts due to climate change. *PloS One* 9 (6), e98907.

Laporta, GZ, **Y-M Linton, RC Wilkerson**, ES Bergo, SS Nagaki, C Sant'Ana, & MAM Sallum. 2015. Malaria vectors in South America: current and future scenarios. *Parasites & Vectors* 8: 426. [doi:10.1186/s13071-015-1038-4](https://doi.org/10.1186/s13071-015-1038-4)

Leasi, F, C Gaynus, A Mahardini, NT Moore, **JL Norenburg** & HP Barber. 2015. Spatial and ecological distribution of neglected microinvertebrate communities across endangered ecosystems: Meiofauna in Bali (Indonesia). *Marine Ecology* in press.

Leasi, F & JL Norenburg. 2015. At least some meiofauna species are not everywhere. Indication of ecological and geological barriers affecting the dispersion of species of *Ototyphlonemertes*(Nemertea, Hoplonemertea). *Molecular Ecology* in press.

Lee, JL, MK Thura, **DG Mulcahy & GR Zug**. 2015. Three Colubrid snakes new to Myanmar. *Herpetology Notes* 8:217–220.

Leray M & N Knowlton. 2015. DNA barcoding and metabarcoding of standardized samples reveal patterns of marine benthic diversity. *Proceedings of the National Academy of Sciences* 112:2076-2081.

Leray M, CP Meyer & SC Mills. 2015. Metabarcoding dietary analysis of coral dwelling predatory fish demonstrates the minor contribution of coral mutualists to their highly partitioned, generalist diet. *PeerJ*. 3:e1047

Lim, HC, F Zou & FH Sheldon. 2015. Genetic differentiation in two widespread, open-forest bird species of Southeast Asia (*Copsychus saularis* and *Megalaima haemacephala*): Insights from ecological niche modeling. *Current Zoology* 61:922-934.

Lindsay DJ, MM Grossmann, J Nishikawa, B Bentlage & AG Collins. 2015. DNA barcoding of pelagic cnidarians: current status and future prospects. *Bulletin of the Plankton Society of Japan*: in press.

Liu, X-Q, S Ickert-Bond, Z Nie, , Z Zhou, L-Q Chen & J Wen. 2015. Phylogeny of the *Ampelocissus*–*Vitis* clade in Vitaceae supports the New World origin of the grape genus. *Molecular Phylogenetics & Evolution*, [doi:10.1016/j.ympev.2015.10.013](https://doi.org/10.1016/j.ympev.2015.10.013)

Lohan KMP, KM Hill, RC Fleischer, ME Torchin, EE Strong & GM Ruiz. 2015. Molecular phylogenetics reveals first record and invasion of *Saccostrea* species in the Caribbean. *Marine Biology* 162: 957-968.

Lu, J-M, N Zhang, X-Y Du, J Wen & D-Z Li. 2015. Chloroplast phylogenomics resolves key relationships in ferns. *Journal of Systematics and Evolution* 53(5): 448-457. [doi:10.1111/jse.12180](https://doi.org/10.1111/jse.12180)

Mandel, JR, RB Dikow & VA Funk. 2015. Using phylogenomics to resolve mega-families: an example from Compositae. *Journal of Systematics and Evolution*. [doi:10.1111/jse.12167](https://doi.org/10.1111/jse.12167)

Menezes, RST, SG Brady, AF Carvalho, MA Del Lama, & MA Costa. 2105. Recent diversification of the Neotropical swarm-founding social wasp genus *Synoeca* de Saussure, 1852 (Hymenoptera: Vespidae: Polistinae): biogeographic and systematic implications. *PLoS ONE* 10(3): e0119151. [doi:10.1371/journal.pone.0119151](https://doi.org/10.1371/journal.pone.0119151)

Mengual, X, G Stahls & S Rojo. 2015. Phylogenetic relationships and taxonomic ranking of pipizine flower flies (Diptera: Syrphidae) with implications for the evolution of aphidophagy. *Cladistics* 2015:1-18. [doi:10.1111/cla.12105](https://doi.org/10.1111/cla.12105)

Muscarella, R, M Uriarte, DL Erickson, NG Swenson, JK Zimmerman & WJ Kress. 2015. Climate and biodiversity effects on standing biomass in Puerto Rican forests. *Caribbean Naturalist*: in press.

Nie, Z-L, VA Funk, Y Meng, T Deng, H Sun & J Wen. 2015. Recent assembly of the global herbaceous flora: evidence from the paper daisies (Asteraceae: Gnaphalieae). *New Phytologist*. [doi:10.1111/nph.13740](https://doi.org/10.1111/nph.13740)

Pei, N, DL Erickson, B Chen, X Ge, X Mi, NG Swenson, J-L Zhang, FA Jones, C-L Huang, W Ye, Z Hao, C-F Hsieh, S Lum, NA Bourg, JD Parker, JK Zimmerman, WJ McShea, IC Lopez, IF Sun, SJ Davies, K Ma & WJ Kress. 2015. Closely-related taxa influence woody species discrimination via DNA barcoding: evidence from global forest dynamics plots. *Scientific Reports*, 5. [doi:10.1038/srep15127](https://doi.org/10.1038/srep15127)

Pérez-Losada, M, JW Breinholt, M Aira & J Domínguez. 2015. An updated multilocus phylogeny of the Lumbricidae (Annelida: Clitellata: Oligochaeta) earthworms. *Phylogenetics & Evolutionary Biology* 3:1.

Peterson, AT, RG Moyle, F Lei, LC Campillo, PA Hosner, LB Klicka, **HC Lim**, AS Nyari, Y QU, S Reddy, FH Sheldon & F Zou. 2015. Avian evolution and speciation in the Southeast Asian tropics. *Current Zoology* 61:898-900.

Podar, M, **CC Gilmour**, CC Brandt, A Soren, SD Brown, BR Crable, AV Palumbo, AC Somenahally & DA Elias. 2015. Global prevalence and distribution of genes and microorganisms involved in mercury methylation. *Science Advances* 1:e500675. [doi:10.1126/sciadv.1500675](https://doi.org/10.1126/sciadv.1500675)

Reeder, TW, TM Townsend, **DG Mulcahy**, BP Noonan, PL Wood, Jr., JW Sites, Jr & JJ Wiens. 2015. Integrated Analyses Resolve Conflicts over Squamate Reptile Phylogeny and Reveal Unexpected Placements for Fossil Taxa. *PLOS ONE* 10(3): e0118199. [doi:10.1371/journal.pone.0118199](https://doi.org/10.1371/journal.pone.0118199)

Salvador-Reyes, L, N Engene, **VJ Paul** & H Luesch. 2015. Targeted natural products discovery from marine Cyanobacteria using combined phylogenetic and mass spectrometric evaluation. *Journal of Natural Products*, 78(3): 486-492. [doi:10.1021/np500931q](https://doi.org/10.1021/np500931q)

Santos, S, ML Bartholomei-Santos, G Bond-Buckup, L Buckup, KA Crandall, CG Jara & M Pérez-Losada. 2015. Three new species of Aeglidae (AeglaLeach, 1820) from Paraná State, Brazil. *Journal of Crustacean Biology*. [doi:10.1163/1937240X-00002376](https://doi.org/10.1163/1937240X-00002376)

Schachat, S, **DG Mulcahy** & JR Mendelson III. 2015. Conservation threats and the utility of IUCN Red List rankings in *Incilius* toads. *Conservation Biology* Early View: [doi:10.1111/cobi.12567](https://doi.org/10.1111/cobi.12567)

Schultz, T, J Sosa-Calvo, SG Brady, CT Lopes, UG Mueller, H Vasconcelos, M Bacci, & A Rodrigues. 2015. The most relictual fungus-farming ant species cultivates the most recently evolved and most highly domesticated fungal symbiont. *American Naturalist* 185: 693–703. [doi:10.1086/680501](https://doi.org/10.1086/680501)

Shapcott, Alison, Forster, Paul I., Guymer, Gordon P., McDonald, William J. F., Faith, Daniel P., **Erickson, David L.** and **Kress, W. John**. 2015. Mapping Biodiversity and Setting Conservation Priorities for SE Queensland's Rainforests Using DNA Barcoding. *PloS One*, 10(3) [doi:10.1371/journal.pone.0122164](https://doi.org/10.1371/journal.pone.0122164)

Sheldon, FH, **HC Lim** & RG Moyle. 2015. Return to the Malay Archipelago: the biogeography of Sundaic rainforest birds. *Journal of Ornithology* 156:91-113.

Shrestha, Y, HH Wirshing & MG Harasewych. 2015. The Genus *Cerion* (Gastropoda: Cerionidae) in the Florida Keys. *PLoS ONE*, 10(9): 1-26. [doi:10.1371/journal.pone.0137325](https://doi.org/10.1371/journal.pone.0137325)

Smith, CR, SH Cahan, C Kemeny, **SG Brady**, W Yang, E Bornberg-Bauer, T Eriksson, J Gadau, M Helmpkampf, D Gotzek, MO Miyakawa, A Suarez & A Mikheyev. 2015. How do genomes create novel phenotypes? Insights from the loss of the worker caste in ant social parasites. *Molecular Biology and Evolution* 32: 2919–2931. [doi:10.1093/molbev/msv165](https://doi.org/10.1093/molbev/msv165)

Tepolt, CK & SR Palumbi. 2015. Transcriptome sequencing reveals both neutral and adaptive divergence in a marine invader. *Molecular Ecology* 24: 4145–4158. [doi:10.1111/mec.13294](https://doi.org/10.1111/mec.13294)

Tepolt, CK. 2015. Adaptation in marine invasions: a genetic approach. *Biological Invasions* 17(3): 887-903. [doi:10.1007/s10530-014-0825-8](https://doi.org/10.1007/s10530-014-0825-8)

Todaro MA, M Dal Zotto & **F Leasi**. 2015. An integrated morphological and molecular approach to the description and systematisation of a novel genus and species of *Macrodasyida* (Gastrotricha). *PLoS ONE* 10: e0130278.

Tornabene L, JL Van Tassell, RG Gilmore, DR Robertson, F Young & **CC Baldwin**. In Press. Molecular phylogeny, analysis of discrete character evolution, and submersible collections facilitate a new classification for a diverse group of gobies (Teleostei: Gobiidae: Gobiosomatini: Nessubgroup), with descriptions of nine new species. *Zoological Journal of the Linnean Society*.

Tu, T-H, C-F Dai & M-S Jeng. 2015. Phylogeny and systematics of deep-sea precious corals (Anthozoa: Octocorallia: Coralliidae). *Molecular Phylogenetics & Evolution* 84: 173-184. [doi:10.1016/j.ympev.2014.09.031](https://doi.org/10.1016/j.ympev.2014.09.031)

Villarreal-Treviño, C, GM Vasquez, V Lopez-Sifuentes, K Escobedo-Vargas, A Huayanay-Repetto, **Y-M Linton**, C Flores-Mendoza, AG Lescano & FM Stell. 2015. First free-mating, long-standing and highly productive laboratory colony of *Anopheles darlingi* from the Peruvian Amazon. *Malaria Journal* 14: 227. [doi:10.1186/s12936-015-0733-0](https://doi.org/10.1186/s12936-015-0733-0)

Ward, PS, **SG Brady**, BL Fisher & **TR Schultz**. 2015. The evolution of myrmicine ants: Phylogeny and biogeography of a hyperdiverse ant clade. *Systematic Entomology* 40: 61–81. [doi:10.1111/syen.12090](https://doi.org/10.1111/syen.12090)

Wen, J, J Liu, S Ge, Q-Y Xiang & **EA Zimmer**. 2015. Phylogenomic approaches to deciphering the tree of life. *Journal of Systematics and Evolution* 53: 369-370.

Wen, J, A Egan, R Dikow & EA Zimmer. 2015. Utility of transcriptome sequencing for phylogenetic inference and character evolution. Hörandl, E. & Appelhans, M.S. (eds.), Next-Generation Sequencing in Plant Systematics. Chapter 2. International Association for Plant Taxonomy (IAPT), 2015.

Whatton, JF, CJ Dove & R Greenwood. 2013. Birdstrike in Georgia documents mixed-species flock of nocturnal migrating vertebrates. *The Oriole* 78: 1-6.

Wilkerson, RC & Y-M Linton. 2015. Elevation of *Pseudoskusea*, *Rusticoidus* and *Protomacleaya* to valid subgenera in the mosquito genus *Aedes* based on taxon naming criteria recently applied to other members of the Tribe Aedini (Diptera: Culicidae). *Parasites & Vectors* 8:668. [doi:10.1186/s13071-015-1247-x](https://doi.org/10.1186/s13071-015-1247-x)

Wilkerson, R, Y-M Linton, D Fonseca, T Schultz, D Price & D Strickman. 2015. Making mosquito taxonomy useful: A stable classification of Tribe Aedini that balances utility with current knowledge of evolutionary relationships. *PLoS One* 10(7): e0133602. [doi:10.1371/journal.pone.0133602](https://doi.org/10.1371/journal.pone.0133602)

Williams, JT & K.E. Carpenter. 2015. A new fish species of the subfamily Serraninae (Perciformes: Serranidae) from the Philippines. *Zootaxa* 3911 (2): 287-293. [doi:10.11646/zootaxa.3911.2.10](https://doi.org/10.11646/zootaxa.3911.2.10)

Xiang, J-Y, **J Wen** & H Peng. 2015. Evolution of the eastern Asian–North American biogeographic disjunctions in ferns and lycophytes. *Journal of Systematics and Evolution*, 53(1): 2-32. [doi:10.1111/jse.12141](https://doi.org/10.1111/jse.12141)

Xu, X-W, J-W Wu, M-X Qi, Q-X Lu, PF Lee, **S Lutz**, S Ge & **J Wen**. 2015. Comparative phylogeography of the wild-rice genus *Zizania* (Poaceae) in eastern Asia and North America. *American Journal of Botany* 102(2): 239-247. [doi:10.3732/ajb.1400323](https://doi.org/10.3732/ajb.1400323)

Yi, T-S, G-H Jin & J Wen. 2015. Chloroplast capture and intra- and inter-continental biogeographic diversification in the Asian – New World disjunct plant genus *Osmorrhiza* (Apiaceae). *Molecular Phylogenetics & Evolution* 85: 10-21. [doi:10.1016/j.ympev.2014.09.028](https://doi.org/10.1016/j.ympev.2014.09.028)

Zapata F, FE Goetz, SA Smith, M Howison, S Siebert , SH Church, SM Sanders, CL Ames, CS McFadden, SC France, M Daly, AG Collins, SHD Haddock, CW Dunn & P Cartwright. 2015. Phylogenomic analyses support traditional relationships within Cnidaria. *PLoS ONE*. 10(10): e0139068. [doi:10.1371/journal.pone.0139068](https://doi.org/10.1371/journal.pone.0139068)

Zapico, SC and DH Ubelaker. 2015. Relationship between mitochondrial DNA mutations and aging: estimation of age-at-death. *J Gerontology A: Biological Science and Medical Sciences* 2015. [doi:10.1093/gerona/glv115](https://doi.org/10.1093/gerona/glv115)

Zhang, G, C Li, Q Li, B Li, DM Larkin, C Lee, JF Storz, A Antunes, MJ Greenwold, RW Meredith, A Ödeen, J Cui, Q Zhou, L Xu, H Pan, Z Wang, L Jin, P Zhang, H Hu, W Yang, J Hu, J Xiao, Z Yang, Y Liu, Q Xie, H Yu, J Lian, P Wen, F Zhang, H Li, Y Zeng, Z Xiong, S Liu, L Zhou, Z Huang, N An, J Wang, Q Zheng, Y Xiong, G Wang, B Wang, J Wang, Y Fan, RR da Fonseca, A Alfaro-Núñez, M Schubert, L Orlando, T Mourier, JT Howard, G Ganapathy, A Pfenning, O Whitney, MV Rivas, E Hara, J Smith, M Farré, J Narayan, G Slavov, MN Romanov, R Borges, J Paulo Machado, I Khan, MS Springer, J Gatesy, FG Hoffmann, JC Opazo, O Håstad, RH Sawyer, H Kim, K-W Kim, HJ Kim, S Cho, N Li, Y Huang, MW Bruford, X Zhan, A Dixon, MF Bertelsen, E Derryberry, W Warren, RK Wilson, S Li, DA Ray, RE Green, SJ O'Brien, D Griffin, WE Johnson, D Haussler, OA Ryder, E Willerslev, GR Graves, P Alström, J Fjeldså, DP Mindell, SV Edwards, EL Braun, C Rahbek, DW Burt, P Houde, Y Zhang, H Yang, J Wang, Avian Genome Consortium, ED Jarvis, M Thomas P Gilbert & J Wang. 2014. Comparative genomics across modern bird species reveal insights into avian genome evolution and adaptation. *Science*. 12 December 2014: 1311-1320.

Zhang, N, J Wen, and EA Zimmer. 2015. Expression patterns of *AP1*, *FUL*, *FT* and *LEAFY* orthologs in Vitaceae support the homology of tendrils and inflorescences throughout the grape family. *Journal of Systematics and Evolution*. [doi:10.1111/jse.12138](https://doi.org/10.1111/jse.12138)

Zhang, N, J Wen & EA Zimmer. 2015. Congruent deep relationships in the grape family (Vitaceae) based on sequences of chloroplast genomes and mitochondrial genes via genome skimming. *PLoS ONE* [doi:10.1371/journal.pone.0144701](https://doi.org/10.1371/journal.pone.0144701)

Zhang, J-Q, S-Y Meng, J Wen & G-Y Rao. 2015. DNA barcoding of *Rhodiola* (Crassulaceae): a case study on a group of recently diversified medicinal plants from the Qinghai-Tibetan plateau. *PloS One* 10(3): 1-15. [doi:10.1371/journal.pone.0119921](https://doi.org/10.1371/journal.pone.0119921)

Zhao, J-L, Y-M Xia, WJ Kress, CH Cannon & Q-J Li. 2015. Multi-stage uplift of Himalayan-Tibetan Plateau and rapid extrusion of Indochinese: Clues from alpine gingers' divergence. *Gondwana Research*: in press.

Zimmer, EA and J Wen. 2015. Using nuclear gene data for plant phylogenetics: Progress and prospects II. Next-gen approaches." *Journal of Systematics and Evolution* 53: 371-379.

Zuo, Y-J, J Wen, J-S Ma, & S-L Zhou. 2015. Evolutionary radiation of the *Panax bipinnatifidus* species complex (Araliaceae) in the Sino-Himalayan region of eastern Asia as inferred from AFLP analysis. *Journal of Systematics and Evolution* 15(3): 210-220. [doi:10.1111/jse.12119](https://doi.org/10.1111/jse.12119)

Andrade, SCS, H Montenegro, M Strand, M Schwartz, H Kajihara, JL Norenburg, JM Turbeville, P Sundberg, & G Giribet. 2014. Phylogenomics of ribbon worms (phylum Nemertea). *Molecular Biology & Evolution*. [doi:10.1093/molbev/msu253](https://doi.org/10.1093/molbev/msu253)

Appelhans, MS, J Wen & WL Wagner. 2014. A molecular phylogeny of *Acronychia*, *Euodia*, *Melicope* and relatives (Rutaceae) reveals polyphyletic genera and key innovations for species richness. *Molecular Phylogenetics & Evolution*: in press.

Appelhans, MS, J Wen , KR Wood, GJ Allan, EA Zimmer & WL Wagner. 2014. Molecular phylogenetic analysis of Hawaiian Rutaceae (*Melicope*, *Platydesma* & *Zanthoxylum*) and their different colonization patterns. *Botanical Journal of the Linnean Society* 174(3): 425-448. [doi:10.1111/boj.12123](https://doi.org/10.1111/boj.12123)

Baeza, JA, RT Bauer, J Okuno & M Thiel. 2014. Molecular phylogeny of hinge-beak shrimps (Decapoda: Caridea: *Rhynchocinetes* and *Cinetorhynchus*) and allies: a formal test of familiar and generic monophly using a multilocus phylogeny. *Zoological Journal of the Linnean Society*: in press.

Baldwin, CC & GD Johnson. 2014. Connectivity across the Caribbean Sea: DNA barcoding and morphology unite an enigmatic fish larva from the Florida Straits with a new species of sea bass from deep reefs off Curaçao. *PLoS ONE* 9(5): e97661.

Baldwin, CC & DR Robertson. 2014. A new *Liopropoma* sea bass (Serranidae: Epinephelinae: Liopropomini) from deep reefs off Curaçao, southern Caribbean, with comments on depth distributions of western Atlantic liopropomins. *Zookeys* 409: 71-92.

Betancur-R R & G Ortí. 2014. Molecular evidence for the monophly of flatfishes (Carangimorpharia: Pleuronectiformes). *Molecular Phylogenetics & Evolution* 73: 18-22. [doi:10.1016/j.ympev.2014.01.006](https://doi.org/10.1016/j.ympev.2014.01.006)

Betancur-R R, G Naylor, & G Ortí. 2014. Conserved genes, sampling error, and phylogenomic inference. *Systematic Biology*. [doi: 10.1093/sysbio/syt073](https://doi.org/10.1093/sysbio/syt073)

Bowen, BW, K Shanker, N Yasuda, MCMD Malay, S von der Heyden, G Paulay, LA Rocha, KA Selkoe, PH Barber, ST Williams, HA Lessios, ED Crandall, G Bernardi, CP Meyer , KE Carpenter & RJ Toonen. 2014. Phylogeography unplugged: comparative surveys in the genomic era. *Bulletin of Marine Science* 90(1): 13-46.

Bracken-Grissom, HD, ST Ahyong, RD Wilkinson, RM Feldmann, CE Schweitzer, JW Breinholt, M Bendall, F Palero, T-Y Chan, DL Felder, R Robles, K-H Chu, L-M Tsang, D Kim, JW Martin, & KA Crandall. 2014. The emergence of lobsters: Phylogenetic relationships, morphological evolution and divergence time comparisons of an ancient group (Decapoda: Achelata, Astacidea, Glypheidea, Polychelida). *Systematic Biology* 63(4):457-479. [doi:10.1093/sysbio/syu008](https://doi.org/10.1093/sysbio/syu008)

Brady, SG, BL Fisher, TR Schultz, & PS Ward. 2014. The rise of the army ants: Diversification of the specialized predatory doryline ants. *BMC Evolutionary Biology* 14:e93. [doi: 10.1186/1471-2148-14-93](https://doi.org/10.1186/1471-2148-14-93)

Brix, S, F Leese, T Riehl & TC Kihara. 2014. A new genus and new species of Desmosomatidae Sars, 1897 (Isopoda) from the east South-Atlantic abyss described by means of integrative taxonomy. *Marine Biodiversity*: online 3 May 2014. [DOI: 10.1007/s12526-014-0218-3.](https://doi.org/10.1007/s12526-014-0218-3)

Buffington, ML, E Perkovsky, & SG Brady . 2014. The description of *Rovnoeucoila typanomorpha*Buffington & Perkovsky, a new genus and species of fossil eucoiline, with observations on

the asynchronous evolution of Diglyphosematini (Hymenoptera: Figitidae: Eucoilinae). *Proceedings of the Entomological Society of Washington* 116: 243–254. [doi: 10.4289/0013-8797.116.3.243](https://doi.org/10.4289/0013-8797.116.3.243)

Callicrate, T, RB Dikow, JW Thomas, JC Mullikin, ED Jarvis, **RC Fleischer** & NISC Comparative Sequencing Program. 2014. Genomic resources for the endangered Hawaiian honeycreepers. *BMC Genomics* 15: 1098. [doi: 10.1186/1471-2164-15-1098](https://doi.org/10.1186/1471-2164-15-1098).

Chen, H, S Sun, **JL Norenburg**, & P Sundberg. 2014. Mutation and selection cause codon usage and bias in mitochondrial genomes of ribbon worms (Nemertea). *PLoS ONE* 9: e85631. [doi:10.1371/journal.pone.0085631](https://doi.org/10.1371/journal.pone.0085631)

Chen, J, J Zhao, **DL Erickson**, N Xia & **WJ Kress**. 2014. Testing DNA barcodes in closely related species of *Curcuma* (Zingiberaceae) from Myanmar and China. *Molecular Ecology Resources*: early view. [doi: 10.1111/1755-0998.12319](https://doi.org/10.1111/1755-0998.12319).

Chin, S-W, J Shaw, R Haberle, **J Wen** & D Potter. 2014. Diversification of almonds, peaches, plums and cherries - molecular systematics and biogeographic history of *Prunus* (Rosaceae). *Molecular Phylogenetics & Evolution* 76: 34-48. [doi:10.1016/j.ympev.2014.02.024](https://doi.org/10.1016/j.ympev.2014.02.024)

Cialdella, AM, SM Sede, **K Romaschenko**, **PM Peterson**, **RJ Soreng**, FO Zuloaga & O Morrone. 2014. Phylogeny of *Nassella* (Stipeae, Pooideae, Poaceae) based on analyses of chloroplast and nuclear ribosomal DNA and morphology. *Systematic Botany* 39: 814–828. dx.doi.org/10.1600/036364414X681419

Cochran, AT, J Prado & **E Schuettelpelz**. 2014. *Tryonia*, a new taenitidoid fern genus segregated from *Jamesonia* and *Eriosorus* (Pteridaceae). *PhytoKeys* 35: 23–43.

Connolly, SR, MA MacNeil, MJ Caley, E Cripps, M Hisano, LM Thibaut, BD Bhattacharya, L Benedetti-Cecchi, RE Brainard, A Brandt, F Bulleri, KE Ellingsen, S Kaiser, **N Knowlton**, I Kröncke, K Linse, E Maggi, TD O'Hara, **L Plaisance**, GCB Poore, SK Sarkar, KK Satpathy, U Schückel, A Williams & RS Wilson. 2014. Commonness and rarity in the marine biosphere. *Proceedings of the National Academy of Sciences*, USA: in press.

Conway, KW, **CC Baldwin** & MD White. 2014. Cryptic diversity and venom glands in western Atlantic clingfishes of the genus *Acyrtus* (Teleostei: Gobiesocidae). *PLoS ONE* 9(5): e97664.

Davies, N, D Field, L Amaral-Zettler, MS Clark, J Deck, A Drummond, DP Faith, J Geller, J Gilbert, FO Glockner, PR Hirsch, J-A Leong, **CP Meyer**, et al & GOs-COS. 2014. The founding charter of the Genomic Observatories Network. *GigaScience* 3:2.

Deeds, JR, SM Handy, F Fry Jr., H Granade, **JT Williams**, M Powers, R Shipp & **LA Weigt**. 2014. Protocol for building a reference standard sequence library for DNA-based seafood identification. *Journal of AOAC International* 97(6): 1-14. [doi: 10.5740/jaoacint.14-111](https://doi.org/10.5740/jaoacint.14-111)

Delrieu-Trottin, E, **JT Williams** & S Planes. 2014. *Macropharyngodon pakoko*, a new species of wrasse (Teleostei: Labridae) endemic to the Marquesas Islands, French Polynesia. *Zootaxa* 3857 (3): 433–443. <https://dx.doi.org/10.11646/zootaxa.3857.3.6>

Erickson, DL, FA Jones, NG Swenson, N Pei, N Bourg, W Chen, **SJ Davies**, X Ge, Z Hao, RW Howe, C-L Huang, A Larson, S Lum, JA Lutz, K Ma, M Meegaskumbura, X Mi, JD Parker, I Fang-Sun, **J Wright**, AT

Wolf, W Ye, D Xing, JK Zimmerman & **WJ Kress**. 2014. Comparative evolutionary diversity and phylogenetic structure across multiple forest dynamics plots: a mega-phylogeny approach. *Frontiers in Genetics*: [doi: 10.3389/fgene.2014.00358](https://doi.org/10.3389/fgene.2014.00358).

Escobar, JE, R González, ML Quiñones, **RC Wilkerson, F Ruiz**, BA Harrison. 2014. Morphology of the larvae, male genitalia and DNA sequences of *Anopheles (Kerteszia) pholidotus* (Diptera: Culicidae) from Colombia. *Mem Inst Oswaldo Cruz, Rio de Janeiro*: 1-7, 2014.

Faircloth, BC, **MG Branstetter, ND White & SG Brady**. 2014. Target enrichment of ultraconserved elements from arthropods provides a genomic perspective on relationships among Hymenoptera. *Molecular Ecology Resources* [DOI: 10.1111/1755-0998.12328](https://doi.org/10.1111/1755-0998.12328).

Faulde, MK, **LM Rueda** & BA Kaireh. 2014. First record of the Asian malaria vector *Anopheles stephensi* and its possible role in resurgence of malaria in Djibouti, Horn of Africa (Diptera, Culicidae). *Acta Tropica* 139: 39-43.

Foley, DH, M-Y Linton, JF Ruiz-Lopez, JE Conn, MAM Sallum, MM Póvoa, E Bergo, I Sucupira & RC Wilkerson. 2014. Using occurrence data to explore the evolution and disease importance of species within the Neotropical *Anopheles albitarsis* group (Diptera, Culicidae). *Journal of Vector Ecology* 39(1): 168-181. [doi:10.1111/j.1948-7134.2014.12084.x](https://doi.org/10.1111/j.1948-7134.2014.12084.x)

Funk, VA, G Sancho, N Roque, CL Kelloff, I Ventosa-Rodríguez, M Diazgranado, JM Bonifacino & R Chan. 2014. A Phylogeny of the Gochnatiaeae: Understanding a critically placed tribe in the Compositae. *Taxon*. 63(4): 859–882.

Geller, J, **CP Meyer, M Parker** & H Hawk. 2014. Redesign of PCR primers for mitochondrial cytochrome c oxidase subunit I for marine invertebrates and application in all-taxon biotic surveys. *Molecular Ecology Resources*: In press.

Grossmann MM, **AG Collins** & DJ Lindsay. 2014. Description of the eudoxid stages of *Lensia havock* and *Lensia leloupi* (Cnidaria: Siphonophora: Calycophorae), with a review of all known *Lensia* eudoxid bracts. *Biodiversity & Systematics*: in press.

Gunay, F, B Alten, F Simsek, A Aldemir & **Y-M Linton**. 2014. Barcoding Turkish *Culex* mosquitoes to facilitate arbovirus vector incrimination studies reveals hidden diversity and new potential vectors. *Acta Tropica*; (available online). [DOI: 10.1016/j.actatropica.2014.10.013](https://doi.org/10.1016/j.actatropica.2014.10.013)

Harasewych, MG. 2014. Systematics and phylogeography of *Cerion sensu stricto* (Mollusca: Gastropoda: Pulmonata: Cerionidae) from Aruba, Curaçao and Bonaire. *Journal of Molluscan Studies*: in press.

Harasewych, MG & RE Petit. 2014. Morphology and systematic relationships of *Brocchinia clenchi*Petit, 1986 (Mollusca, Caenogastropoda, Cancellariidae) from the Azores. In: Martins, AMF (ed.) The Marine Fauna and Flora of the Azores (Proceedings of the 4rd International Workshop of Malacology and Marine Biology, São Miguel, 2011). *Açoreana, Supplement*: in press.

Horn, JW, Z Xi, R Riina, JA Peirson, Y Yang, BL Dorsey, PE Berry, CC Davis & KJ Wurdack. 2014. Evolutionary bursts in *Euphorbia* (Euphorbiaceae) are linked with photosynthetic pathway. *Evolution* 68(12): 3485-3504. [doi:10.1111/evo.12534](https://doi.org/10.1111/evo.12534)

Hubert, N, B Espiau, **C Meyer** & S Planes. 2014. Identifying ichthyoplankton of a coral reef using DNA barcoding. *Molecular Ecology Resource*. [doi: 10.1111/1755-0998.12293](https://doi.org/10.1111/1755-0998.12293)

Jarvis, ED, et al., **A Driskell**, et al., **M Braun**, et al., **W Johnson**, et al., **GR Graves**, et al. 2014. Whole-genome analyses resolve early branches in the tree of life of modern birds. *Science* 346(6215): 1320-1331.

Keyse, J, ED Crandall, RJ Toonen, **CP Meyer**, EA Treml & C Riginos. 2014. The scope of published population genetic data for Indo-Pacific marine fauna and future research opportunities in the region. *Bulletin of Marine Science* 90(1): 47-78.

Kingston, SE, AG Navarro-Sigüenza, EA García-Trejo, H Vázquez-Miranda, WF Fagan, & **MJ Braun**. 2014. Genetic differentiation and habitat connectivity across towhee hybrid zones in Mexico. *Evolutionary Ecology* 28(2): 277-297.

Kress, WJ, C García-Robledo, M Uriarte & **DL Erickson**. 2014. DNA Barcodes for Ecology, Evolution, and Conservation. *TREE* 30: 25-35. <https://dx.doi.org/10.1016/j.tree.2014.10.008>.

Lasky, JR, M Uriarte, V Boukili, **DL Erickson**, **WJ Kress**, RL Chazdon.. 2014. The relationship between tree biodiversity and biomass dynamics changes with tropical forest succession. *Ecology Letters* 17: 1158-1167.

Leasi F, & JL Norenburg. 2014. The necessity of DNA taxonomy to reveal cryptic diversity and spatial distribution of meiofauna, with a focus on Nemertea. *PLoS ONE* 9: e104385. [DOI: 10.1371/journal.pone.0104385](https://doi.org/10.1371/journal.pone.0104385)

Lemer, S, GY Kawauchi, SC Andrade, SC, **VL González**, MJ Boyle & G Giribet. 2014. Re-evaluating the phylogeny of Sipuncula through transcriptomics. *Molecular Phylogenetics & Evolution* 83C:174-183. [doi: 10.1016/j.ympev.2014.10.019](https://doi.org/10.1016/j.ympev.2014.10.019)

Liu, Q, B Jiang, **J Wen** & **PM Peterson**. 2014. Low-copy nuclear gene and McGISH resolves polyploid history of *Eleusine coracana* and morphological character evolution in *Eleusine*. *Turkish Journal of Botany* 38: 1-12. [doi:10.3906/bot-1305-12](https://doi.org/10.3906/bot-1305-12)

Liu, Q, H Liu, **J Wen** & **PM Peterson**. 2014. Infrageneric phylogeny and temporal divergence of *Sorghum* (Andropogoneae, Poaceae) based on low-copy nuclear and plastid sequences. *PLoS ONE* 9(8): 1-14. doi.org/10.1371/journal.pone.0104933

Mandel, JR, **RB Dikow**, **VA Funk**, R. Masalia, SE Staton, A Kozik, RW Michelmore, LH Rieseberg, & JM Burke. 2014. Target enrichment for next-gen sequencing and phylogenetic analyses: Lessons from the Compositae. *Applications in Plant Sciences (APPS)*: 2(2): 1300085. [doi:10.3732/apps.1300085](https://doi.org/10.3732/apps.1300085)

Martin, AR, **DL Erickson**, **WJ Kress** & SC Thomas. 2014. Wood nitrogen concentrations in tropical trees: phylogenetic patterns and ecological correlates. *New Phytologist*: [doi: 10.1111/nph.12943](https://doi.org/10.1111/nph.12943)

Martin, MD, EA Zimmer, MT Olsen, AD Foote, MTP Gilbert & GS Brush. 2014. Herbarium specimens reveal a historical shift in phylogeographic structure of common ragweed during native range disturbance. *Molecular Ecology* 23: 1701–1716. [doi: 10.1111/mec.12675](https://doi.org/10.1111/mec.12675)

Masiulionis, VE, C Rabeling, HH De Fine Licht, **T Schultz**, M Bacci Jr, CM Santos Bezerra & FC Pagnocca. 2014. The asexual fungus-growing ant *Mycocepurus smithii* (Formicidae, Myrmicinae, Attini) cultivates morphologically and genetically diverse fungal symbionts with gongylidia-like structures. *PLoS ONE* 9(8): e103800. [doi: 10.1371/journal.pone.0103800](https://doi.org/10.1371/journal.pone.0103800)

Morrone, O, **K Romaschenko**, M Giusanni & **PM Peterson**. 2014. Phylogeny of *Nassella* (Poaceae: Stipeae) based on four plastid and nuclear DNA ITS regions. *Systematic Botany* 39(2): xxx-xxx. (in press)

Mulcahy, DG, JE Martínez-Gómez, G Aguirre-León, JA Cervantes Pasqualli & **GR Zug**. 2014. Rediscovery of an endemic vertebrate from the remote Islas Revillagigedo in the eastern Pacific Ocean: The Clarión Nightsnake, with conservation and systematic implications. *PLoS ONE*: 9(5): e97682. [doi:10.1371/journal.pone.0097682](https://doi.org/10.1371/journal.pone.0097682)

Muscarella, R, M Uriarte, **DL Erickson**, NG Swenson, JK Zimmerman & **WJ Kress**. 2014. A well-resolved phylogeny of the trees of Puerto Rico on DNA barcode sequence data. *PLoS ONE* 9(11): e112843. [doi: 10.1371/journal.pone.0112843](https://doi.org/10.1371/journal.pone.0112843)

Pérez-Losada M, JT Høeg, N Simon-Blecher, Y Achituv, D Jones, **KA Crandall**. 2014. Molecular phylogeny, systematics and morphological evolution of the acorn barnacles (Thoracica: Sessilia: Balanomorpha). *Molecular Phylogenetics & Evolution* 81:147–158.

Peterson, PM, **K Romaschenko** & Y Herrera Arrieta. 2014. A molecular phylogeny and classification of the Cteniinae, Farraginiinae, Gouiniinae, Gymnopogoninae, Perotidinae, and Trichoneurinae (Poaceae: Chloridoideae: Cynodontae). *Taxon* 63: 275–286. doi.org/10.12705/632.35

Peterson, PM, **K Romaschenko**, Y Herrera Arrieta & JM Saarela. 2014. A molecular phylogeny and new subgeneric classification of *Sporobolus* (Poaceae: Chloridoideae: Sporobolinae). *Taxon* 63(6): 1212–1243. doi.org/10.12705/636.19

Peterson, PM, **K Romaschenko**, Y Herrera Arrieta & JM Saarela. 2014. (2332) Proposal to conserve the name *Sporobolus* against *Spartina*, *Crypsis*, *Ponceletia*, and *Heleochochloa* (Poaceae, Chloridoideae, Sporobolinae). *Taxon* 63(6): 1373–1374. doi.org/10.12705/636.23

Peterson, P M, **K Romaschenko** & **RJ Soreng**. 2014. A laboratory guide for generating DNA barcodes in grasses: A case study of *Leptochloa* s.l. (Poaceae: Chloridoideae). *Webbia* 69(1): 1–12. doi.org/10.1080/00837792.2014.927555

Pons, J-M, S. Sonstagen, **C. Dove**, and P-A Crochet. 2014. Extensive mitochondrial introgression in North American Great Black-backed Gulls (*Larus marinus*) from the American Herring Gull (*Larus smithsonianus*) with little nuclear DNA impact. *Heredity* 2013: 1- 14. [Doi:10.1038/hdy.2013.98](https://doi.org/10.1038/hdy.2013.98)

Puebla O, **E Birmingham**, **WO McMillan**. 2014. Genomic atolls of differentiation in coral reef fishes (*Hypoplectrus* spp, Serranidae). *Molecular Ecology* 23: 5291–5303. [doi:10.1111/mec.12926](https://doi.org/10.1111/mec.12926)

Puillandre, N, TF Duda, **C Meyer**, BM Olivera & P Bouchet. 2014. One, four or 100 genera? Classification of the cones. *Journal of Molluscan Studies*. eyu055

Puillandre, N, P Bouchet, TF Duda, S Kauferstein, AJ Kohn, BM Olivera, M Watkins & **CG Meyer**. 2014. Molecular phylogeny and evolution of the cone snails (Gastropoda, Conoidea). *Molecular Phylogenetics & Evolution* 78: 290-303. [doi:10.1016/j.ympev.2014.05.023](https://doi.org/10.1016/j.ympev.2014.05.023)

Rabeling, CR, **TR Schultz**, NE Pierce & M Bacci, Jr. 2014. A social parasite evolved reproductive isolation from its fungus-growing ant host in sympatry. *Current Biology* 24: 1-6. [doi:10.1016/j.cub.2014.07.048](https://doi.org/10.1016/j.cub.2014.07.048).

Riehl, T, N Brenke, **S Brix**, **A Driskell**, S Kaiser & A Brandt. 2014. Field and laboratory methods for DNA studies on deep-sea isopod crustaceans. *Polish Polar Research* 35(2): 203-224.

Robinson, H & VA Funk. 2014. *Dysaster cajamarcensis*, a new shrubby genus and species of Astereae from Peru (Asteraceae). *PhytoKeys* 36: 35-40. [doi: 10.3897/phytokeys.36.7393](https://doi.org/10.3897/phytokeys.36.7393)

Robinson, H & VA Funk. 2014. *Gymnanthemum koekemoeri* (Compositae: Vernonieae), a new species from South Africa. *PhytoKeys* 36: 59-65.

Robison, HW, CT McAllister, JW Breinholt & **K.A. Crandall**. 2014. Status, distribution and genetics of Blair's fencing crayfish, *Faxonella blairi* (Decapoda: Cambaridae). *The Southwestern Naturalist* 59(2):185-191.

Romaschenko, K, N Garcia-Jacas, **PM Peterson**, **RJ Soreng**, R Vilatersana & A Susanna. 2014. Miocene–Pliocene speciation, introgression, and migration of *Patis* and *Ptilagrostis* (Poaceae: Stipeae). *Molecular Phylogenetics & Evolution* 70: 244–259. doi.org/10.1016/j.ympev.2013.09.018

Rosenthal, MA, SR Rosenthal, **G Johnson**, WC Taylor & **EA Zimmer**. 2014. *Isoetes viridimontana*: a previously unrecognized quillwort from Vermont, USA. *Amer. Fern Jnl.*: in press.

Rothfels, CJ & **E Schuettpelz**. 2014. Accelerated rate of molecular evolution for vittarioid ferns is strong and not due to selection. *Systematic Biology* 63: 31–54.

Scarpa, F, P Cossu, D Sanna, T Lai, **JL Norenburg**, M Curini-Galletti, & M Casu. 2014. An 18S and 28S-based clock calibration for marine Proseriata (Platyhelminthes). *Journal of Experimental Marine Biology and Ecology* 463: 22-31. [doi:10.1016/j.jembe.2014.10.020](https://doi.org/10.1016/j.jembe.2014.10.020)

Schuettpelz, E, A Davila, J Prado, RY Hirai & G Yatskievych. 2014. Molecular phylogenetic and morphological affinities of *Adiantum senae* (Pteridaceae). *Taxon* 63: 258–264.

Ssymank, A & **X Mengual**. 2014. *Paragus caligneus* sp. nov., a new Afrotropical species of flower fly (Diptera: Syrphidae). *Acta Entomologica Musei Nationalis Pragae*: in press.

Thompson, AW, **R Betancur-R**, H Lopez-Fernandez, & **G Ortí**. 2014. A time-calibrated, multi-locus phylogeny of piranhas and pacus (Characiformes: Serrasalmidae) and a comparison of species tree methods. *Molecular Phylogenetics & Evolution* 81: 242-257.

Todaro MA, **Leasi F**, Hochberg R. 2014. A new species, genus and family of marine Gastrotricha from Jamaica, with a phylogenetic analysis of Macrodasysida based on the 18S rDNA gene. *Systematics and Biodiversity* 12: 473-488. [DOI: 10.1080/14772000.2014.942718](https://doi.org/10.1080/14772000.2014.942718)

Toledo-Hernández, C, X Vélez-Zuazo, CP Ruiz-Díaz, AR Patricio, P Mège, M Navarro, AB Sabat, **R Betancur-R** & R Papa. 2014. Population ecology and genetics of the invasive lionfish in Puerto Rico. *Aquatic Invasions* 9: 227-237. <https://dx.doi.org/10.3391/ai.2014.9.2.12>

Valcárcel, V, O Fiz-Palacios & **J Wen**. 2014. The origin of the early differentiation of Ivis (*Hedera*L.) and the radiation of the Asian Palmate group (Araliaceae). *Molecular Phylogenetics & Evolution* 70: 492-503. [doi:10.1016/j.ympev.2013.10.016](https://doi.org/10.1016/j.ympev.2013.10.016)

Ward, PS, **SG Brady**, BL Fisher, & **TR Schultz**. 2014. The evolution of myrmicine ants: Phylogeny and biogeography of a hyperdiverse ant clade. *Systematic Entomology* 40: 61–81. [doi: 10.1111/syen.12090](https://doi.org/10.1111/syen.12090)

Wen, J, J-Q Zhang, Z-L Nie, Y Zhong & H Sun. 2014. Evolutionary diversifications of plants on the Qinghai-Tibetan Plateau. *Frontiers in Genetics* 5. [doi:10.3389/fgene.2014.00004](https://doi.org/10.3389/fgene.2014.00004)

Windsor, AM & DL Felder. 2014. Molecular phylogenetics and taxonomic reanalysis of the family Mithracidae MacLeay (Decapoda: Brachyura: Majoidea). *Invertebrate Systematics* 28(2): in press.

Zarowiecki, M, **YM Linton**, RJ Post, MJ Bangs, PT Htun, T Hlaing, CM Seng, V Baimai, DH Ding, T Sochantha & C Walton. 2014. Repeated landmass reformation limits diversification in the widespread littoral zone mosquito *Anopheles sundaicus* sensu lato in the Indo-Oriental Region. *Molecular Ecology* 23: 2573–2589. [doi: 10.1111/mec.12761](https://doi.org/10.1111/mec.12761)