

Adalgisa Caccone's Publications

August 2017

Submitted

- Richardson, J., K.-Y. Lee, P. Mireji, J. Enyaru, M. Sistrom, S. Aksoy, H. Zhao, A. CACCOME. Genomic analyses of African *Trypanosoma* strains to assess evolutionary relationships and identify markers for strain identification. Plos NTD (in review)
- Gloria-Soria, A, W.A. Dunn, WA, X Yu, A Vigneron A, K-Y Lee, BL Weiss, H. Zhao , S. Aksoy, S., and A. CACCOME. Harnessing genomic data from closely related taxa to uncover gene-phenotype associations in non-model organisms: genomic regions associated with *Trypanosoma* infections in wild populations of the tsetse fly *Glossina fuscipes*. G3 (in review).
- Jensen, E.L, D.L. Edwards, R.C. Garrick, J. M. Miller, J. P. Gibbs, L.J. Cayot, W. Tapi, A. CACCOME and M.A. Russello. Population genomics through time corroborates bottleneck theory in Pinzón Galápagos giant tortoises. Nature (in review).
- Havill, N.P, S. Gaimari, and A. CACCOME. Cryptic east-west divergence and molecular diagnostics for two species of silver flies (Diptera: Chamaemyiidae: *Leucopis*) from North American being evaluated for biological control of hemlock woolly adelgid. Biological Invasions (submitted),
- Burak, M., N. Saarman, R. Opiro, C. Hyseni, R. Echodu, E. Opiyo, K. Dion, A. Dunn, S. Aksoy and A. CACCOME. A spatial genetics approach to inform vector control of tsetse flies (*Glossina fuscipes fuscipes*) in Northern Uganda. Scientific Reports (in review).
- Chiari, Y, A. van der Meijden, A. CACCOME, J. Claude, B. Gilles. Different performance in rolling – over may relate to shell shape evolution in Galápagos giant tortoises". Scientific reports (in review).
- Marshall, J.C., E. Bastiaans, A. CACCOME, A. Camargo, M. Moranda, M. Niemiller, M. Pabijan, M. Russello, B. Sinervo, J. W. Sites, Jr., M. Vences, J. J. Wiens, F. P. Werweck, K. C. Wollenberg Valero, S. Steinfartz. Inferring processes and mechanisms of speciation in reptiles and amphibians: a synopsis. Amphibia-Reptilia (submitted).
- Orr, A., J.E. Quagrainne, P. Suwondo, F.P. Dornas, D. Humphries, L.M. Harrison, B. Evans, A. CACCOME, M. D. Wilson, M. Cappello. Genetic markers of benzimidazole resistance among human hookworms (*Necator americanus*) in Kintampo North Municipality, Ghana. J. of Infectious Diseases (submitted).

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215. Gaughran, S.J., M.C. Quinzin, J.M. Miller, R.C. Garrick, D.L. Edwards, M.A. Russello, N. Poulakakis, C. Ciofi⁷, L.B. Beheregaray, A. CACCOME. Theory, practice, and conservation in the age of genomics: the Galápagos giant tortoise as a case study. Evolutionary Application (in press).
214. Okeyo, W.O., N. P. Saarman, M. Mengual, K. Dion, R. Bateta, P. O. Mireji, S. Okoth, J. O. Ouma, C. Ouma, J. Ochieng, G. Murilla, S. Aksoy and A. CACCOME. Genetic diversity and temporal instability in *Glossina pallidipes* tsetse fly populations in Kenya. Parasites and vectors (in press).
213. Kotsakiozi, P., J. Richardson, G. Favia, A. J. Martins, S. Urbanelli, P. Armbruster, A. CACCOME. Population genomics of the Asian tiger mosquito, *Aedes albopictus*: insights into the recent worldwide invasion. Ecology and Evolution (in press)
212. Walter, K.S., G. Carpi, A. CACCOME*, M.A. Diuk-Wasser*. 2017. Genomic insights into the ancient spread of Lyme disease across North America. Nature genetics (in press). *authors contributed equally.
211. Miller, J.M., M.C. Quinzin, N. Poulakakis, L. B. Beheregaray, J.P. Gibbs, C. Garrick, M. A. Russello, C. Ciofi, D. L. Edwards, E. A. Hunter, W. Tapia, D. Rueda, J. Carrión, A.A. Valdivieso, A. Caccone. 2017. Reviving a Lost Species: The Case of the Floreana Galápagos Giant Tortoise *Chelonoidis elephantopus*. Scientific Reports (in press).
210. Kamidi, C.M, N.P. Saarman, K. Dion, P.O. Mireji, C. Ouma, G. Murilla, S. Aksoy, A. Schnaufer, A. CACCOME. 2017. Multiple evolutionary origins of *Trypanosoma evansi* in Kenya. Plos Neglected diseases (in press).
209. Kotsakiozi, P., A. Gloria-Soria, A. CACCOME, B. Evans, R. Schama, A. J. Martins and J. R. Powell. 2017. Tracking the return of *Aedes aegypti* to Brazil, the major vector of the dengue, chikungunya and Zika viruses. Plos NTD (in Press).
208. Opiro, R, N. P. Saarman, R. Echodu, E. A. Opiyo, K. Dion, A. Halyard, A. W. Dunn, S. Aksoy and A. CACCOME. 2017. Genetic Diversity and Population Structure of the Tsetse fly *Glossina fuscipes fuscipes* (Diptera: Glossinidae) in Northern Uganda: Implications for Vector Control. Plos Neglected

Diseases Apr 28;11(4):e0005485. doi: 10.1371/journal.pntd.0005485. [Epub ahead of print]. PMID: 28453513.

207. Manangwa O, N. Gamba, J.O. Ouma, F. Mramba, I. Malele, K. Dion, M. Sistrom, F. Khan, S. Aksoy S and A. CACCOME. 2017. Genetic diversity of *Glossina fuscipes fuscipes* along the shores of Lake Victoria in Tanzania and Kenya: implication for management. Parasites & Vectors 2017, 10:268. DOI: 10.1186/s13071-017-2201-X. PMID :28558831.
206. Andersen, J.C., N. P. Havill A. CACCOME, and J. S. Elkinton. 2017. Post-glacial recolonization shaped the genetic diversity of the winter moth (*Operophtera brumata*) in Europe. Ecology and Evolution (2017). DOI: 10.1002/ece3.2860. PMID:28515868.
205. Puckett, E.E., J. Park, M. Combs, M. J. Blum, J. E. Bryant, A. CACCOME, F. Costa, E. Deinum, A. Esther, C. G. Himsworth, P. Keightley, A. Ko, Å. Lundkvist, L. M. McElhinney, S. Morand, J. Robins, J. Russell, T. M. Strand, O. Suarez, L. Yon, and J. Munshi-South. Global population divergence and admixture of the brown rat (*Rattus norvegicus*). 2017. Proceedings of the Royal Society B: Biological Sciences. Evol Appl. 2017 Feb 23;10(4):323-337. doi: 10.1111/eva.12449. PMID:28352293.

2016

204. Richardson, J.L., M.K. Burak, C. Hernandez, J. M. Shirvell, C. Mariani, T. S. A. Carvalho-Pereira, A. C. Pertile, J. A. Panti-May, G. G. Pedra, S. Serrano, J. Taylor, M. Carvalho, G. Rodrigues, F. Costa, J. E. Childs, A. I. Ko, and A. CACCOME. Using fine scale spatial genetics of Norway rats to improve control efforts and reduce leptospirosis risk in urban slum environments. Evol Appl. 2017 Feb 23;10(4):323-337. doi: 10.1111/eva.12449. eCollection 2017 Apr. PMID:28352293.
203. Havill, N.P, J. Elkinton, J.C. Andersen, S.B. Hagen, H. J. Broadley, G.J. Boettner, A. CACCOME. Asymmetric hybridization between non-native winter moth, *Operophtera brumata* (Lepidoptera: Geometridae), and native Bruce spanworm, *O. bruceata*, in the northeastern United States, assessed with novel microsatellites and SNPs. Bulletin of Entomological Research, pp. 1–10. doi: 10.1017/S0007485316000857.
202. Carpi, G., K. S. Walter, C. Ben Mamoun, P. J. Krause, A. Kitchen; T. Leopore; Ankit Dwivedi; E. Cornillot; A. CACCOME, and M. Diuk-Wasser. *Babesia microti* from humans and ticks hold a genomic signature of strong population structure in the United States. BMC Genomics 17:888. PMID: 27821055.
201. Jensen E.L., A.Ø. Mooers, A. CACCOME, and M.A. Russello MA. (2016) I-HEDGE: determining the optimum complementary sets of taxa for conservation using evolutionary isolation. PeerJ 4:e2350 <https://doi.org/10.7717/peerj.2350>.
200. Walter, K. S., G. Carpi, B.R. Evans, A. CACCOME, M.A. Diuk-Wasser. Vectors as epidemiological sentinels: Patterns of within-tick *Borrelia burgdorferi* diversity. PLoS Pathog 12(7): e1005759. doi:10.1371/journal.ppat.1005759.
199. Guevara, E. E., C.C. Veilleux, K. Saltonstall, A. CACCOME, N.I. Mundy, N. I. and B. Bradley, B. J. (2016), Potential arms race in the coevolution of primates and angiosperms: brazzein sweet proteins and gorilla taste receptors. Am. J. Phys. Anthropol., 161: 181–185. doi:10.1002/ajpa.23046.
198. Opiro, R., N.P. Saarman, R. Echodu, E. A. Opiyo, K. Dion, A. Haylard, S. Aksoy and A. Caccone. Evidence of temporal stability in allelic and mitochondrial haplotype diversity in populations of *Glossina fuscipes fuscipes* (Diptera: Glossinidae) in northern Uganda. Parasites and Vectors. 9:258.
197. Gloria-Soria, A, WA Dunn WA, EL Telleria, BR Evans, L. Okedi L, R. Echodu, WC Warren, S. Aksoy S, A. CACCOME. Patterns of genome-wide variation in *Glossina fuscipes fuscipes* tsetse flies from Uganda. G3: Genes|Genomes|Genetics Early Online, published on April 13, 2016 as doi:10.1534/g3.116.027235.
196. Graf, F.E., P. Ludin, J. C. Munday, J. Krezdorn, N. Schaub, C. Kunz-Renggli, O. Balmer, A. CACCOME, H. P. de Koning and P. Mäser Comparative genomics of drug resistance of the sleeping sickness parasite *Trypanosoma brucei rhodesiense*. Cell. Mol. Life Sci. DOI 10.1007/s00018-016-2173-6.
195. Chiari, Y., S. Glaberman, P. Tarroso, A. CACCOME, J. Claude. Ecological and phylogeographic influences on body size and shape in the Galápagos marine iguana (*Amblyrhynchus cristatus*). Oecologia. DOI 10.1007/s00442-016-3618-1.
194. Havill, N.P., S. Shiyake, A.L. Galloway, R.G. Foottit, G. Yu, A. Paradis, J. Elkinton, M.M. Montgomery, M. Sano, and A. CACCOME. Ancient and modern colonization of North America by hemlock woolly adelgid (*Adelges tsugae* Annand) (Hemiptera: Aphidoidea), an invasive insect from Asia. Molecular Ecology, first published online: 28 MAR 2016 | DOI: 10.1111/mec.13589.
193. Sistrom, M.J., B. Evans, J. Benoit, O. Balmer, S. Aksoy and A. CACCOME. *De novo* genome assembly shows genome wide similarity between *Trypanosoma brucei brucei* and *Trypanosoma brucei rhodesiense*. PlosOne, Published: February 24, 2016. <http://dx.doi.org/10.1371/journal.pone.0147660>.
192. Glass, J.R., M. Davis, T.J. Walsh, E. J. Sargis, and A. CACCOME. Was frozen mammoth or giant ground sloth served for dinner at The Explorers Club? PlosOne Published February 3, 2016. DOI: 10.1371/journal.pone.0146825.

2015

191. Richardson, JB, BR Evans, Pati Pyana P, N. Van Reet, M. Sistrom, P. Buscher, S. Aksoy, and A CACCOME. Whole genome sequencing shows sleeping sickness relapse is due to parasite regrowth and not reinfection. *Evolutionary Application*: 9: 381–393.
190. Costa, F., J. Richardson, K. Dion, C. Mariani, A. C. Pertile, J. E. Childs, A. I. Ko, and A. CACCOME. Multiple paternity in the Norway rat, *Rattus norvegicus*, from urban slums in Salvador, Brazil. *J. of Heredity* 107 (2): 181–186.
189. Poulakakis, N., D.L Edwards, Y.Chiari, R.C Garrick, E. Benavides, M. A. Russello, G. J. Watkins-Colwell, S. Glaberman, W. Tapia, J. P Gibbs, L. J. Cayot, A. CACCOME. Description of a new Galapagos Giant Tortoise Species (*Chelonoidis*; Testudines: Testudinidae) from Cerro Fatal on Santa Cruz Island. *PlosOne*. Published: October 21, 2015. DOI: 10.1371/journal.pone.0138779.
188. Carpi, G. K. S. Walter, S. J. Bent, A. G. Hoen, M. Diuk-Wasser and A. CACCOME. Whole genome capture of human pathogens from field-collected disease vectors. *BMC Genomics* *BMC Genomics* 2015, 16:434 (6 June 2015).
187. MacLeod, A., A. Rodríguez, M. Vences, P. Orozco-terWengel, C. García, F. Trillmich, G. Gentile, A. CACCOME, G. Quezada, S. Steinfartz. Hybridization masks speciation in the evolutionary history of the Galápagos marine iguanas. Published 3 June 2015.DOI: 10.1098/rspb.2015.0425.
186. Jensen, E.L., W. Tapia, A. CACCOME, M. Russello. Genetics of a head-start program to guide conservation of an endangered Galápagos tortoise (*Chelonoidis ephippium*). *Conservation genetics* 4 (2): 31–46.
185. Echodu, R., M.Sistrom, R. Bateta, G. Murilla, L. Okedi, S. Aksoy, C. Enyioha, J. Enyaru, E. Opiyo, W. Gibson, and A. CACCOME. Genetic diversity and population structure of *Trypanosoma brucei* in Uganda: implications for the epidemiology of sleeping sickness and Nagana. *PLOS Neglected Tropical Diseases* 10.1371/journal.pntd.0003353. PMID:25695634.
184. Garrick, R.C., B. Kajdacsy, M.I A. Russello, J.P. Gibbs, W. Tapia and A. CACCOME. Naturally rare versus newly rare: demographic inferences on two timescales inform conservation of Galapagos giant tortoises. *Ecology and Evolution* 2015; 5(3): 676–694.

2014

183. Sistrom, M. B. Evans, R. Bjornson, W. Gibson, O. Balmer, P. Mäser, S. Aksoy & A. CACCOME. Comparative genomics reveals multiple genetic backgrounds of human pathogenicity in the *Trypanosoma brucei* complex. *Genome Biol. Evol.* 6(10):2811–2819. PMID:25287146.
182. Garrick, R.C., E. Benavides, M. A. Russello, C. Hyseni, D. L. Edwards, J. P. Gibbs, W. Tapia, C. Ciofi and A. CACCOME. Lineage fusion in Galápagos giant tortoises. *Molecular Ecology*, 23: 5276–5290. PMID: 25223395.
181. Bayha, K.M., M. H. Chang, C. L. Mariani, J. L. Richardson, D. L. Edwards, T. S. DeBoer, C. Moseley, E. Aksoy,, M. B. Decker, P. M. Gaffney, G. R. Harbison, J. H. McDonald, A. CACCOME Worldwide phylogeography of the invasive ctenophore *Mnemiopsis leidyi* (Ctenophora) based on nuclear and mitochondrial DNA data. *Biological Invasions*. DOI 10.1007/s10530-014-0770-6. Publ. on line Aug. 31, 2014.
180. Edwards, D.L., Ryan C. Garrick, W.H. Tapia, and A. CACCOME. Cryptic structure between ecologically distinct genetic clusters in threatened giant Galápagos tortoises from southern Isabela Island. *Conservation Genetics*, 15: 1357–1369.
179. Aksoy, E., E. Telleria, R. Echodu, Y. Wu, L. Okedi, B. Weiss, S. Aksoy and A. CACCOME. Analysis of multiple tsetse fly populations in Uganda reveals restricted and species-specific gut microbiomes. *Applied and Environmental Microbiology*, 80:18 5844–5853. PMID:24814785.
178. Federman, S., C. Hyseni, W. Clement, Mi. Oatham, A. CACCOME.Habitat fragmentation and the genetic structure of wild populations of the Amazonian palm *Mauritia flexuosa* L.f. (Arecaceae) on the island of Trinidad". *Conservation Genetics* 15:355–362.

2013

177. Brown, J. E., B. R. Evans, W. Zheng, V. Obas, L. Barrera-Martinez. A. Egizi, H. Zhao, A. CACCOME, and J.R. Powell. Human impacts have shaped historical and recent evolution in *Aedes aegypti*, the dengue and yellow fever mosquito. *Evolution* 68: 514–525. PMCID 24111703 [PubMed - in process].
176. Kajdacsy B., F. Costa, C. Hyseni, F. Porter, J. Brown, G. Rodrigues, H. Farias, M. G. Reis, J. E. Childs, A. I. Ko, and A. CACCOME. Urban population genetics of slum-dwelling rats (*Rattus norvegicus*) in Salvador, Brazil. *Molecular Ecology*, 22: 5056–5070. doi: 10.1111/mec.12455. PMID: 24118116.
175. Hodges, T.K, G. Athrey, K. C. Deitz, H. J. Overgaard, A. Matias, A. CACCOME and M. A. Slotman. Large fluctuations in the effective population size of the malaria mosquito *Anopheles gambiae* s.s. during vector control cycle. *Evolutionary Applications*. doi: 10.1111/eva.12094. PMID:24478799.
174. Pinto, J., A.E. Yawson, J.L. Vicente, B. Gomes, F. Santolamazza, M. Moreno, J.D. Charlwood, F. Simard, N. Elissa, D. Weetman, M.J. Donnelly, A. CACCOME and A. della Torre. Geographic population structure of the African malaria vector *Anopheles gambiae* suggests a role for the forest-savannah biome transition as a barrier to gene flow. *Evolutionary Applications*, 6: 910–924. PMID:24062800.
173. Echodu, R., M. Sistrom, C. Hyseni, J. Enjaru, L. M. Okedi, S. Aksoy, and A. CACCOME Genetically distinct *Glossina fuscipes fuscipes* populations in the Lake Kyoga region of Uganda and its relevance

- for Human Africa trypanosomiasis. *Biomed Res Int.* 2013;2013:614721. doi: 10.1155/2013/614721. PMID:24199195.
172. Aksoy, S. A. CACCOME, A.P. Galvani and L. M. Okedi *Glossina fuscipes* populations provide insights for Human African Trypanosomiasis transmission in Uganda. *Trends in Parasitology* August 2013, 29 (8): 394-406. PMID: 23845311.
 171. Rehman A.M., A. G Mann, C. Schwabe, M. R. Reddy, I. R. Gomes, M. A Slotman, L. Yellott, A. Matias, A. CACCOME, G. Nseng Nchama, I. Kleinschmidt. Five years of malaria control in the continental region, Equatorial Guinea. *Malar J.* 2013 May 7;12:154. doi: 10.1186/1475-2875-12-154. PMID:23651490.
 170. Ketmaier, V. and A. CACCOME. Twenty Years of Molecular Biogeography in the West Mediterranean Islands of Corsica and Sardinia: Lessons Learnt and Future Prospects, *Current Progress in Biological Research*, Dr. Marina Silva-Oppa (Ed.), ISBN: 978-953-51-1097-2, InTech, DOI: 10.5772/55458. pp.71-93.
 169. Reddy MR, Godoy A, Dion K, Atue M, Matias A, Abaga S, Callender, K Kiszevski AE, A. CACCOME, M.A. Slotman. 2013. Insecticide Resistance Allele Frequencies in *Anopheles gambiae* before and after Anti-Vector Interventions in Continental Equatorial Guinea. *Am. J. Trop. Med. Hyg.*, 88(5): 897–907. PMID:23438768.
 168. Symula, R.E, U. Alam, C. Brelsfoard, Y. Wu, S. Aksoy, A. CACCOME. *Wolbachia* association with the tsetse fly, *Glossina fuscipes fuscipes*, reveals high levels of genetic diversity and complex evolutionary dynamics. *BMC Evolutionary Biology* 2013, 13:31 doi:10.1186/1471-2148-13-31. PMID: 23384159.
 167. Edwards, D.L., E. Benavides, R. C. Garrick, J. P. Gibbs, M.A. Russello, K. B. Dion, C. Hyseni, J. P. Flanagan, W. Tapia, A. CACCOME. The genetic legacy of Lonesome George survives: giant tortoises with Pinta Island ancestry identified in Galápagos. *Biological Conservation* 157: 225-228.

2012

166. Aksoy, S., V. M. F. Almeida-Val, V. C. R. Azevedo, R. Baucom, P. Bazaga, L. B. Beheregaray, Je. L. Bennetzen et al. Permanent Genetic Resources added to Molecular Ecology Resources Database 1 October 2012.30 November 2012." *Molecular ecology resources* (2013).
165. Alter, S.E., H.C. Rosenbaum, L.D. Postma, P. Whitridge, C. Gaines, D. Weber, M.G. Egan, M. Lindsay, G. Amato, L. Dueck, R.L. Brownell Jr., M. Heide-Jorgensen, K.L. Laidre, A. CACCOME, B. Hancock. "Gene flow on ice: The role of sea ice and whaling in shaping Holarctic genetic diversity and population differentiation in bowhead whales (*Balaena mysticetus*)."*Ecology and Evolution*. 2: 2895–2911.
164. Milinkovitch. M.C., R. Kanitz, R. Tiedemann, W. Tapia, F. Llerena, A. CACCOME, J. P. Gibbs and J.R. Powell. Recovery of a Nearly Extinct Galápagos Tortoise Despite Minimal Genetic Variation. *Evolutionary Applications*, 6: 377–383.
163. Athrey, G., T. K. Hodges, M. Reddy, H. Overgaard, A. Matias, F. Ridl, I. Kleinschmidt, A. CACCOME, M. A. Slotman. The Effective Population Size of Malaria Mosquitoes: Large Impact of Vector Control. *Plos Genetics*, 8 (12): e1003097. PMID: 23271973
162. Havill, N., G. Davis, D. L. Mausel, J. Klein; R. McDonald, C. Jones, M. Fischer, S. Salom;and A. CACCOME. Hybridization between a native and introduced predator of Adelgidae: An unintended result of classical biological control. *Biological Control*, 63 (3): 359-369.
161. Hyseni, C., Kato, A., L.M. Okedi, C. Masembe, J.O. Ouma, S. Aksoy, A. CACCOME. Genetic diversity and structure of *Glossina fuscipes fuscipes* Uganda populations from the lake Victoria basin, insights for vector control and monitoring. *Parasites and Vectors* 4;5(1): 222. [Epub ahead of print].PMID:23036153.
160. Deitz, K.C, Gi. Athrey, M. R. Reddy, H. J Overgaard, A. M Arnez, J.Musa, A. della Torre, J. Pinto, A. Kiszevski, C. Costantini, A. CACCOME and M. A. Slotman. Genetic Isolation within the malaria mosquito *Anopheles melas*. *Molecular Ecology* 21(18): 4498-513.PMID:22882458
159. Symula, R.E., J. S. Beadell, M. Sistrom1, K. Agbebakun1 , O.Balmer, W. Gibson, S. Aksoyand A. CACCOME. *Trypanosoma brucei gambiense* group 1 is distinguished by a unique amino acid substitution in the HpHb receptor implicated in human serum resistance. *PLoS Negl Trop Dis.*, 6(7):e1728. Epub 2012 Jul 10.160. PMID:22802982.
158. Alam, U., C. Hyseni, R/ E. Symula, C. Brelsfoard, Y. Wu,O. Kruglov, R. Echodu, V. Alioni, L. M. Okedi, A. CACCOME and S. Aksoy. Implications of microfauna-host interactions for trypanosome transmission dynamics in *Glossina fuscipes fuscipes* in Uganda. *Appl Environ Microbiol.* 78 (13): 4627-37. PMID: 22544247.
157. Deitz, K.C., V. P. Reddy, M. Reddy, N. Satyanarayanan, M. Lindsey, H. J. Overgaard, J. Musa, A. CACCOME and M. A. Slotman. Limited Usefulness of Microsatellite Markers between the Closely Related Malaria Vectors *Anopheles gambiae* and *Anopheles melas*. *Journal of Heredity*, 103(4): 585-93. PMID: 22593601
156. Garrick, R.C., E. Benavides, M.A. Russello, J.P. Gibbs, N. Poulikakis, K.B. Dion, C. Hyseni, B. Kajdacsy, L. Márquez, S. Bahan, C. Ciofi, W. Tapia, A. CACCOME. Genetic rediscovery of an 'extinct' Galápagos giant tortoise species". *Current Biology* 22: R1-R10. PMID:22240469

155. Poulakakis, N., M. A. Russello, D. Geist, and A. CACCOME. Unraveling the Peculiarities of Island Life: Vicariance, Dispersal and the Diversification of the Extinct and Extant Giant Galápagos Tortoises. *Molecular Ecology* 21, 160–173.

2011

154. Ouma JO, Beadell JS, Hyseni C, Okedi LM, Krafsur ES, Aksoy S, A. CACCOME. Genetic diversity and population structure of *Glossina pallidipes* in Uganda and western Kenya. *Parasit Vectors*. 2011 Jun 28;4:122. PMID: 21711519/
153. Glaberman, S., C. Hyseni, W. Clement, and A. CACCOME. Isolation of 13 novel highly polymorphic microsatellite loci for the Amazonian Palm *Mauritia flexuosa* L.f. (Arecaceae). *Conservation Genetics Resources*: DOI 10.1007/s12686-011-9547-8).
152. Symula RE, I. Marpuri, RD Bjornson, L. Okedi, J. Beadell, U. Alam, S. Aksoy, A. CACCOME. 2011. Influence of host phylogeographic patterns and incomplete lineage sorting on within-species genetic variability in *Wigglesworthia* species, obligate symbionts of tsetse flies. *Appl Environ Microbiol*. Dec;77(23):8400-8. doi: 10.1128/AEM.05688-11. Epub 2011 Sep 23. PMID: 21948847
151. Reddy, M.R, H. J. Overgaard, S. Abaga, V. P. Reddy, A. CACCOME, A. E. Kiszewski, M. Slotman. Outdoor host-seeking behavior of *Anopheles gambiae* s.l. mosquitoes following initiation of malaria vector control on Bioko Island, Equatorial Guinea. *Malaria Journal* 2011, **10**:184 (7 July 2011).
150. Bonomi A, Bassetti F, Gabrieli P, Beadell J, Falchetto M, et al. 2011 Polyandry Is a Common Event in Wild Populations of the Tsetse Fly *Glossina fuscipes fuscipes* and May Impact Population Reduction Measures. *PLoS Negl Trop Dis* 5(6): e1190. doi:10.1371/journal.pntd.0001190.
149. Benavides E., M. Russello, D. Boyer, B. Wiese, B. Kajdacsy, L. Marquez, R. Garrick, A. Caccone. Lineage Identification and Genealogical Relationships Among Captive Galápagos Tortoise. *Zoo Biology*, 31: 107–120.
148. Davis, G., N. P. Havill, Z. N. Adelman, A. CACCOME, L.T. Kok, S. M Salom. DNA barcodes and molecular diagnostics to distinguish between an introduced and native *Laricobius* (Coleoptera: Derodontidae) species in eastern North America. *Biological Control*, Volume 58, Issue 1, July 2011, Pages 53-59149-2011.
147. Hyseni, C., J. Beadell, Z. Ocampo Gomez, J. Ouma, L. Okedi, M. Gaunt, A.CACCOME. The G.m. morsitans (Diptera: Glossinidae) genome as a source of microsatellite markers for other tsetse fly (*Glossina*) species. *Molecular Ecology Resources* (2011) 11, 586–589 doi: 10.1111/j.1755-0998.2011.03004.x
146. Echodu, R., J. S. Beadell, L. M. Okedi, C. Hyseni, S. Aksoy, A. CACCOME. Temporal stability of *Glossina fuscipes fuscipes* populations in Uganda. *Parasites & Vectors* 2011, 4:19.
145. Balmer , O., Beadell, J.S. Gibson W., A. CACCOME. Phylogeography and Taxonomy of *Trypanosoma brucei*. *PLoS Negl Trop Dis* 5(2): e961. doi:10.1371/journal.pntd.0000961
144. Pappa, V., Reddy M., Overgaard H.J., Abaga S., A. CACCOME. Estimation of the Human Blood Index in malaria mosquito vectors in Equatorial Guinea following indoor anti-vector interventions. *Am. J. Trop. Med. Hyg.*, 84(2): 298-301. doi:10.4269/ajtmh.2011.10-0463.

2010

143. Klein, J. Caccone, A. N. Havill, Polymorphic microsatellite loci for *Laricobius nigrinus* and *L. rubidus* (Coleoptera: Derodontidae), predators of the hemlock woolly adelgid. *Molecular Ecology Resources*, 10:751–754.
142. Balmer, O., C. Ciofi, D. A. Galbraith, I. R. Swingland, G. R. Zug and A. CACCOME. Population genetic structure in Aldabra giant tortoises. *Journal of Heredity* 2010; doi: 10.1093/jhered/esq096.
141. Lanterbecq D., S. Glaberman1, M. N. Vitousek, S. Steinfartz, M. Wikelski, and A. CACCOME. Genetic Differentiation between Marine Iguana Breeding Sites on the Island of Santa Fé (Galápagos Archipelago). *Journal of Heredity* 2010;101(6):663–675.
140. Garrick R.C., CACCOME, A., Sunnucks, P. Inference of population divergence history by coupling exploratory and model-driven phylogeographic analyses: Assessing concordance and integrating outcomes. *International Journal of Molecular Sciences* (Cladistic Analysis & Molecular Evolution special issue). *International Journal of Molecular Sciences* 11:1190–1227.
139. Beadell, J.S, C. Hyseni, P. Abila, J. C. K. Enyaru, J. Ouma, Y. O. Mohammed, L. M. Okedi, S. Aksoy, A. CACCOME Gene flow among populations of *Glossina fuscipes* at multiple scales: implications for control of tsetse in Uganda. *PLoS Neglected Tropical Diseases* 2010;4(3):e636.
138. Parmakelis, A., M. Moustaka, N. Poulakakis, C. Louis, M. A. Slotman, J. C. Marshall, P.H. Awono-Ambene, C. Antonio-Nkondjio, F. Simard, A. CACCOME, J. R. Powell. *Anopheles* Immune Genes and Amino Acid Sites Evolving Under the Effect of Positive Selection *Anopheles* genes and selection. *Plos ONE* 5:e8885.
137. Steinfartz, S., S. Glaberman, D. Lanterbecq, M. Russello, S. Rosa, T. C. Hanley, C. Marquez , H.L. Snell, H. M. Snell, G. Gentile, G. dell'Olmo, A. M. Powell and A. CACCOME. Progressive colonization and

restricted gene flow shape island-dependent population structure in Galápagos marine iguanas (*Amblyrhynchus cristatus*). BMC Evolutionary Biology 9: 297.

136. Russello, M.A., N. Poulakakis, J. P. Gibbs, W. Tapia, J. R. Powell, and A. CACCONE. Ex situ conservation in crisis: an “extinct” species of Galápagos tortoise identified in captivity. PLoS ONE 5(1): e8683.

2009

135. Henry, P., D. Miquelle, T. Sugimoto; D. McCullough, Dale; A. CACCONE, Russello, Michael. In situ population structure and ex situ representation of the endangered Amur tiger. Mol. Ecol. doi: 10.1111/j.1365-294X.2009.04266.x.
134. Chiari, Y., C. Hyseni, T. H. Fritts, S. Glaberman, C. Marquez, J. P. Gibbs, J. Claude, A. CACCONE. Morphometrics parallel genetics in a newly discovered and endangered taxon of Galápagos tortoise. PLoS ONE 4(7): e6272. doi:10.1371/journal.pone.0006272.
133. Glaberman S, Moreno MA, CACCONE A. Characterization and Evolution of MHC Class II B Genes in Galapagos Marine Iguanas. Developmental and Comparative Immunology, 33: 939-947.

2008

132. Glaberman S, Du Pasquier L, A. CACCONE. 2008 Characterization of a Nonclassical Class I MHC Gene in a Reptile, the Galápagos Marine Iguana (*Amblyrhynchus cristatus*). PLoS ONE 3(8): e2859.
131. Poulakakis, N., S.Glaberman, M.Russello*, L. B. Beheregaray, C. Ciofi, J. R. Powell, and A. CACCONE. Rediscovery of an extinct species of giant Galápagos tortoise. PNAS,105: 15464-15469.
130. Marshall, J.C., J. Pinto, J. D. Charlwood, G. Gentile, F. Santolamazza, F. Simard, A. della Torre, M. J. Donnelly, and A. CACCONE. Exploring the origin and degree of genetic isolation of *An. gambiae* from the islands of São Tomé and Príncipe: implications for testing transgenic-based vector control. Evolutionary Applications:1 (4): 631-644.
129. Balmer O. and A. CACCONE. Frequency of multiple-strain infections in *Trypanosoma brucei*. Acta Tropica,107(3): 275-279
128. Brown, J.E. .K.J. Komatsu1, P.P. Abila, A.S. Robinson, L.M. Okedi, Naomi Dyer, M.J. Donnelly, M.A. Slotman and A. CACCONE. Polymorphic microsatellite markers for the tsetse fly *Glossina fuscipes fuscipes* (Diptera: glossinidae), a vector of human African trypanosomiasis. Molecular Ecology Resources 8 (6): 1506-1508.
127. Oliveira, E., Salgueiro P., Pálsson K., Vicente J.L., Arez A.P., Jaenson T., do Rosário V., CACCONE A., Pinto J. High Levels of hybridisation between molecular forms of *Anopheles gambiae* from Guinea Bissau. J.of Med. Entomol, 45 (6): 1057-1063.
126. Chiari, Y., B. Wang, H. Rushmeier, and A. CACCONE. Using digital images to reconstruct 3D biological forms: a new tool for morphometric studies. Biol. J. of the Linnean Soc., 95 (2): 425-436.
125. Gentile, G., C. Vernesi, S. Vicario, E. Pecchioli, G. Bertorelle, A. CACCONE, and V. Sbordoni. MtDNA variation in roe deer from Italy (*Capreolus capreolus*): evidence of admixture in one of the last *C.c.italicus* populations from central-southern Italy. J. of Italian Zoology iFirst Article. doi:10.1080/11250000802018725.
124. Parent, C, E. A. CACCONE and K. Petren. Colonization and diversification of Galápagos terrestrial fauna: a phylogenetic and biogeographical synthesis. Philos Trans R Soc Lond B Biol Sci. 2008 Oct 27;363(1508):3347-61. Review.
123. Ciofi, C., A. CACCONE, L. B. Beheregaray, M. C. Milinkovitch, M. Russello and J. R. Powell. “Genetics and conservation on islands: the Galápagos giant tortoise as a case study” in “Population Genetics for Animal Conservation” (Bertorelle, Bruford, Chemini, Hauffe, and Vernesi, eds). Cambridge University Press, pages: 269-293.
122. Glaberman, S. and A. CACCONE. Species-specific evolution of class I MHC genes of iguanas (Order: Squamata; Subfamily: Iguaninae). Immunogenetics 60(7): 371-82.
121. Powell J.R. and A. CACCONE. CSI tortoise: unraveling the mystery of mysteries. Galapagos News. 26:8-9.
120. Abila, P.A., M. A. Slotman, A. Parmakelis, K. B. Dion, A. S. Robinson, V. B. Muwanika, J. CK Enyaru, L.M. Okedi, S. Aksoy and A. CACCONE. High levels of genetic differentiation between Ugandan *Glossina fuscipes fuscipes* populations separated by Lake Kyoga. PLOS Neglected Tropical Diseases 2(5): e242
119. Parmakelis, A., M.A. Slotman, J.C. Marshall, P.H. Awono-Ambene, C. Antonio- Nkondjio, F. Simard, A. CACCONE, and J.R. Powell The molecular evolution of four anti-malarial immune genes in the *Anopheles gambiae* species complex. BMC Evolutionary Biology 2008, 8:79 (06 Mar 2008).
118. Palkovacs,E.P., K. B. Dion, D. M. Post, and A. CACCONE. Independent evolution of landlocked alewife populations and rapid, parallel evolution of phenotypic traits. Molecular Ecology, Volume 17: 582-597.

117. Santolamazza F., M. Calzetta, J. Etang, E. Barrese, I. Dia, A. CACCONE, M.J. Donnelly, V. Petrarca, F. Simard, J. Pinto, A. della Torre. Distribution of *knock-down resistance* mutations in *Anopheles gambiae* (Diptera: Culicidae) molecular forms in west and central-west Africa. *Malaria Journal* 2008, 7:74.
116. Parmakelis, A., M. A. Russello, A. CACCONE, C. Brisola Marcondes, J. Costa, O. P. Forattini, M. Sallum, R. C. Wilkerson, and J. R. Powell. Historical analysis of a near disaster: *anopheles gambiae* in Brazil. *Am J. of Trop. Med and Hyg.*, 78 (1): 176-178.

2007

115. Pinto, J., A. Lynd, J.L. Vicente, F. Santolamazza, N.P. Randle, G. Gentile, M. Moreno, F. Simard, J.D. Charlwood, A. CACCONE, A. della Torre and M. J. Donnelly. Multiple origin of knockdown resistance in the Afro-tropical mosquito vector *Anopheles gambiae*. *Plos ONE* 11:e1243.
114. Moreno M, P. Salgueiro, J.L. Vicente, J. Cano, P.J. Berzosa, A. De Lucio A. F. Simard, CACCONE A, V.E. Do Rosario, J. Pinto J and A. Benito. Genetic population structure of *Anopheles gambiae* in Equatorial Guinea. *Malaria Journal* 2007, 6:137 (15 October 2007).
113. Beheregaray,L. and A. CACCONE. Cryptic Biodiversity in a Changing World. Invited minireview. *J. of Biology*, 6:9. Highly Assessed label of BioMed central.
112. Steinfartz, S., S. Glaberman, D. Lanterbecq, C. Marquez, K. Rassmann, and A. CACCONE. Genetic impact of a severe El Niño event on Galápagos marine iguanas. *PloS ONE* 2: e1285.
111. Slotman, M. A. Parmakelis, J. Marshall, P. Awono-Ambene, C. Antonio-Nkondjio, F. Simard, A. CACCONE, and J.R. Powell. Adaptive Evolution in the Anti-Malarial Immune Gene *LRIM1* in *Anopheles arabiensis*. *PloS One Genetics* (August 2007, issue 8, e793).
110. Schielke, E., C. Costantini, G. Carchini, N'fale Sagnon, J.R.Powell, and A. CACCONE. Development of a molecular assay to detect predation on *Anopheles gambiae* complex larval stages. *Am J. of Trop. Med and Hyg* Sep 2007; 77: 464 - 466.
109. Russello,M.A., C. Hyseni, J. P. Gibbs, S. Cruz, C. Marquez, W. Tapia, P. Velensky, J. R. Powell and A. CACCONE. Lineage identification of Galápagos tortoises in captivity worldwide. *Animal Conservation* 10: 304-311.
108. Russello, M.A., L. B. Beheregaray, J. P. Gibbs, T. Fritts, N. Havill, J. R. Powell, and A. CACCONE. Lonesome George is not alone among Galápagos Tortoises. *Current Biology* 17:317-318.
107. Milinkovitch, M.C., D. Monteyne, M. Russello, J. P. Gibbs, H. L. Snell, W. Tapia, C. Marquez, A. CACCONE and J. R. Powell. Giant Galápagos Tortoises: Molecular Genetic Analysis Reveals Contamination in a Repatriation Program of an Endangered Taxon. *BMC Ecology* 2007, 7:2 Published on line 15 February 2007.

2006

106. Russello, M.A., V. Saranathan, S. Buhrman-Deever, J. Eberhard and A. CACCONE. Characterization of polymorphic microsatellite loci for the invasive Monk Parakeet (*Myiopsitta monachus*). *Mol. Ecol. Notes* Volume 6, Issue 2: 390-392.
105. Steinfartz, S. and A. CACCONE. A set of highly discriminating microsatellite for the Galápagos marine iguana *Amblyrhynchus cristatus*. *Molecular Ecology Notes*, 6: 927-929.
104. Poulakakis N., A. Parmakelis, P. Lymberakis, M. Mylonas, E. Zouros, D. S. Reese, S. Glaberman, A. CACCONE. Reply. It remains a mammoth DNA fragment. *Biology Letters*. 3, 60-63.
103. Russello M.A., P. Brazaitis, J. Gratten, G.J. Watkins-Colwell, and A. CACCONE. Molecular assessment of the genetic integrity, distinctiveness and phylogeographic context of the saltwater crocodile (*Crocodylus porosus*) on Palau. *Conservation Genetics*, 8: 777-787.
102. Slotman, M.A., N. B. Kelly, L. C. Harrington, S. Kitthawee, J. W. Jones, T. W. Scott, A. Caccione, and J.R. Powell. Polymorphic microsatellites markers for studies of *Aedes aegypti* (Diptera Culicidae), the vector of dengue and yellow fever. *Molecular Ecology Notes*, 7: 168-171.
101. Steinfartz, S, S. Vicario, J. W. Arntzen and A. CACCONE. A Bayesian approach on molecules, morphology and behavior: reconsidering evolutionary patterns in *Triturus* newts (Amphibia: Salamandridae)". *J. of Exp. Zoology (Mol.Dev.Evol)*, 306B: 1-24.
100. Garren, M., S. M.Walsh, A. CACCONE, and N. Knowlton. Patterns of association between *Symbiodinium* and members of the *Montastraea annularis* species complex on spatial scales ranging from within colonies to between geographic regions. *Coral Reefs*, 25: 503-512.
99. Halverson, M.A, D. K. Skelly, and A. CACCONE. Inbreeding linked to amphibian survival the wild but not in the laboratory. *J. of Heredity* 97: 499-507.
98. Beheregaray, L.B., M. Piggott,N.L. Chao, and A. CACCONE. Development and characterization of microsatellite markers for the Amazonian blackwing hatchetfish *Carnegiella marthae* (Teleostei, Gasteropelecidae). *Mol. Ecol. Notes*. 6: 787-788.
97. Suatoni, E. S. Vicario, S. Rice, T. Snell, and A. CACCONE. An analysis of species boundary and biogeographic patterns in the salt water rotifer, *Brachionus plicatilis*. *Mol. Phylogen. and Evol.* 41: 86-98

96. Jounhyoung Lee, Jonathan Marshall, Oswald J. Schmitz, and A. CACCONE. Genetic divergence of Connecticut *Melanoplus femur-rubrum* populations. *J. of Heredity* 97:290-293.
95. Poulakakis N., A. Parmakelis, P. Lymberakis, M. Mylonas, E. Zouros, D. S. Reese, S. Glaberman, A. CACCONE. Ancient DNA forces reconsideration of evolutionary history of Mediterranean pygmy elephantids. *Biology Letters*, 2 451-454.
94. Olson, M.-A., R. Zajac, A. CACCONE, and M. A. Russello. Characterization of polymorphic microsatellite loci for the polychaete tubeworm *Hobsonia florida*. *Mol. Ecol. Notes* 6: 390-392.
93. Havill, N.P., M. E. Montgomery, G. Yu, S. Shiyake, A. CACCONE. Mitochondrial DNA from hemlock woolly adelgids, *Adelges tsugae* (Hemiptera, Adelgidae), suggests cryptic speciation and pinpoints the source of introduction to eastern North America. *Ann. of the Entomol. Soc. of America*, 99(2): 195-203.
92. Balmer, O., C. Palma, A. MacLeod, A. CACCONE. Characterization of di-, tri-, and tetranucleotide microsatellite markers with perfect repeats for *Trypanosoma brucei* and related species. *Molecular Ecology Notes* 6:508-510.
91. Wolf, J.B., D. Tautz, A. CACCONE, and S. Steinfartz. Development of new microsatellite loci and evaluation of loci from other pinniped species for the Galapagos sea lion (*Zalophus californianus wollebaeki*). *Conservation Genetics*, 7: 461-465
90. Ciofi, c., K. A. Wilson, L. B. Beheregaray, C. Marquez, J. P. Gibbs, W. Tapia, H. L. Snell, A. CACCONE and J. R. Powell Phylogeographic history and gene flow among giant Galápagos tortoises on southern Isabela island. *Genetics* 172: 1727-1744.
89. Slotman, M. A., M. Mendez, A. dellaTorre, Y. Toure', G. Dolo, and A. CACCONE. AFLP analysis indicates genetic differentiation between the Bamako and Savanna chromosomal forms of *Anopheles gambiae*. *Am. J. of Trop. Med and Hyg.*, 74:641-648.
88. Ketmaier, V., F. Giusti, A. CACCONE. Molecular phylogeny and historical biogeography of the land snail genus *Solatopupa* (Pulmonata) in the peri-Tyrrhenian area. *Mol. Phylog. And Evol.*, 39:439-451.
87. Powell, J.R. and A. CACCONE. A quick guide to Galapagos tortoises. *Current Biology*, 16: R144-145.
86. Halverson, M.A, D. K. Skelly, and A. CACCONE. Kin Distribution among wood frog (*Rana sylvatica*) larvae in the wild. *Molecular Ecology* 15: 1139-1146.
85. Pinto, J., A. Lynd, N. Elissa, M.J. Donnelly, C. Costa, G. Gentile, A. CACCONE, and V.E. do Rosário.: Co-occurrence of East and West African *kdr* mutations suggests high levels of resistance to pyrethroid insecticides in *Anopheles gambiae* from Libreville, Gabon. *Med. and Vet. Entomol*, 20:27-32.

2005

84. Russello, M.A., K. Lin, G. Amato, and A. CACCONE. (2005) Additional microsatellite loci for the endangered St.Vincent parrot, *Amazonia guilingii*. *Conservation Genetics*, 6: 643-645.
83. Marshall, J. C., J.R. Powell, and A. CACCONE. Phylogenetic relationships of five major malaria (*Plasmodium falciparum*) vectors within the mosquito subgenus Cellia. *Am. J. of Trop. Med and Hyg.* 73: 749-752.
82. Hanley, T., and A. CACCONE. Development of primers for the characterization of the mitochondrial control region of Galápagos land and marine iguanas (*Conolophus* and *Amblyrhynchus*). *Molecular Ecology Notes* 5:599-601.
81. Beheregaray LB, Chae J, Chao NL, A. CACCONE. Characterization of microsatellite loci for the rummy nose tetra *Hemigrammus bleheri* (Characidae), an Amazonian flooded forest fish. *Molecular Ecology Notes* 5 (3): 536-537.
80. Russello, M, S, Glaberman, , C. Marquez, J.R. Powell, and A. CACCONE. A novel taxon of Giant tortoises in conservation peril. *Biology Letters* 1(3): 287-290.
79. Leonard, J.A., N. Rohland, S. Glaberman, R. C. Fleischer, S. Pääbo, A. CACCONE and M. Hofreiter. How a Zebra Lost its Stripes: 20 Years of Ancient DNA and the Origin of the Quagga. *Biology Letters* 1(3): 291-295.
78. Karanth, P.K., S. Glaberman, E. Palkovacs, A. CACCONE, and A. Yoder. Native Seychelles tortoises or Aldabra imports? The importance of radiocarbon dating for ancient DNA studies this issue. *Amphibia and Reptilia*, 26: 116-121.

2004

77. Russello, MA, E. Gladyshev, D. Miquelle, and A. CACCONE. Potential genetic consequences of a recent bottleneck in the Amur tiger of the Russian Far East. *Conservation Genetics* 5:707-713.
76. Beheregaray LB, Schwartz T, Möller LM, D. Call, Chao NL, CACCONE AA set of microsatellite markers for the one-lined pencil-fish *Nannostomus unifasciatus*, an Amazonian flooded forest fish. *Molecular Ecology Notes*, 4: 333-335.

75. Beheregaray LB, Schwartz T, Möller LM, Chao NL, CACCOME A. Microsatellite markers for the cardinal tetra *Paracheirodon axelrodi*, an economically important fish from central Amazonia. *Molecular Ecology Notes*, 4: 330-332.
74. Gentile G, Santolamazza F, Fanello C, Petrarca V, CACCOME A, Della Torre A variation in an intron sequence of the voltage gated sodium channel gene correlates with genetic differentiation between *Anopheles gambiae* s.s. molecular forms. *Insect Mol Biol*: 13: 371-377.
73. Beheregaray, LB, JP Gibbs, N Havill, TH Fritts, JR Powell, and A CACCOME. Giant tortoises are not so slow: Rapid diversification by recent volcanism in Galápagos. *PNAS* 101: 6514-6519.
72. Palkovacs, EP, AJ. Oppenheimer, E. Gladyshev, JE Toepfer, G. Amato, T. Chase, and A. CACCOME 2004. Phylogeography of the Greater Prairie Chicken (*Tympanuchus cupido*) Including the Extinct Heath Hen: Potential Implications for Conservation and Management. *Molecular Ecology* 13: 1759-1769.
71. Pinou, D., S. Vicario, M. Mendez, and A. CACCOME. Relict Snakes of North America and their relationship to the Caenophidia using Likelihood-Based Bayesian Methods on Mitochondrial Sequences. *Mol. Phylog. and Evol.*, 32:563-574.
70. Milinkovitch, MC,, A. CACCOME, and G.Amato. Molecular Phylogenetic Analyses Indicate Extensive Morphological Convergence between the "Yeti" and Primates. *Mol. Phylog. and Evol.* 31:1-3.
69. Santolamazza F A. della Torre and A. CACCOME. A new PCR-RFLP method to identify *An. arabiensis* from *An. gambiae* and its two molecular forms from DNA templates of poor quality. *Am. J. of Trop. Med. and Hyg.*, 70: 604-606.
68. CACCOME, A., G. Gentile, C. Burns, E. Sezzi, W. Bergman, and J.R. Powell. Extreme difference in rate of mitochondrial and nuclear DNA evolution in a large ectotherm, Galápagos tortoises. *Molecular Phylogenetics and Evolution* 31: 794-798.

2003

67. Milinkovitch, M., D. Monteyne, J.P. Gibbs, T.H. Fritts, W. Tapia, H. L. Snell, R. Tiedemann, A. CACCOME and J.R. Powell. Genetic analysis of a successful repatriation program: Giant Galápagos tortoises. *Proc. Royal. Soc. London B*, 271:341-345.
66. Powell, J.R., E. Sezzi, E. Moriyama, J. Gleason, and A. CACCOME. Analysis of a shift in codon usage in *Drosophila*. *J. Mol. Evol.* 57: 214-225.
65. Burns, C.T., Ciofi, C., L.B. Beheregaray, T.H. Fritts, J.P. Gibbs., C. Marquez, M. C. Milinkovitch, J.R. Powell and A. CACCOME The origin of captive Galápagos tortoises based on DNA analysis: Implications for the management of natural populations. *Animal Conservation*, 6(4): 329-37.
64. Roth R., C. Dick, M. R. Lemes, C. Navarro, A. CACCOME, and E. Bermingham. Genetic structure of Mesoamerican populations of Big-leaf mahogany (*Swietenia macrophylla*) inferred from microsatellite analysis. *Molecular Ecology*, 12: 2885-2894.
63. Beheregaray, LB, Ciofi, C, Geist, D., Gibbs, J, Powell, J.R, and A. CACCOME. Genes record a prehistorical volcano eruption in the Galápagos. *Science*, 302: 75.
62. Palkovacs, E.P., M. Mendez, C. Ciofi, J. Gerlach, and A. CACCOME. Are the native Giant tortoises from the Seychelles really extinct? A genetic perspective based on mtDNA and microsatellite data. *Molecular Ecology* 12: 1403-1413.
61. Vicario, S., A. CACCOME, and J. Gauthier. Xantusiid "Night" lizards: a puzzling phylogenetic problem revisited using likelihood-based Bayesian methods on mtDNA sequences. *Molecular Phylogenetics and Evolution*, 26: 243-261.
60. Ketmaier, V., R. Argano, and A. CACCOME. Phylogeography and molecular rates of subterranean aquatic Stenasellid Isopods with a peri-Tyrrhenian distribution. *Molecular Ecology* 12: 547-555.
59. Beherengary, L. B., C. Ciofi, A. CACCOME, J. P. Gibbs, and J. R. Powell. Genetic divergence, phylogeography and conservation units of Giant tortoises from Santa Cruz and Pinzon, Galápagos islands. *Conservation Genetics*, 4:31-46.

2002

58. Ciofi, C., Milinkovitch M, Gibbs J.P., CACCOME A, Powell J. R. Nuclear DNA microsatellite analysis of genetic divergence among and within island populations of giant Galápagos tortoises. *Molecular Ecology* 11:2265-2283.
57. CACCOME, A., G. Gentile, J. P. Gibbs, T. H. Fritts, H. L. Snell, and J. R. Powell. Phylogeography and history of Giant Galapagos Tortoises. *Evolution* 56, 2052-2066.
56. della Torre, A., C. Costantini, N.J. Besansky, A. CACCOME, V. Petrarca, J.R. Powell, and M. Coluzzi (2002). Speciation Within *Anopheles gambiae*: The Glass is Half Full.*Science* 248: 115-117.
55. Palkovacs, E., J. Gerlach, and A. CACCOME. The evolutionary origin of Indian Ocean tortoises (*Dypsochelys*). *Molecular Phylogenetics and Evolution*, 24: 216-227.

54. Gentile, G., A. Della Torre, B. Maegga, J.R. Powell, and A. CACCOME. Genetic differentiation in the African malaria vector, *Anopheles gambiae* s.s., and the problem of taxonomic status. *Genetics*, 161:1561-1578.

2001

53. Gentile G., M. Slotman, V. Ketmaier, J.R. Powell, and A. CACCOME. Attempts to molecularly distinguish cryptic taxa in *Anopheles gambiae* s.s. *Insect Molecular Biology* 10 (1): 25-32.
52. CACCOME, A. and Sbordoni, V. Molecular biogeography, evolutionary rates, and morphological adaptation to cave life: a case study using Bathysciine beetles and sequence data from the mitochondrial COI gene. *Evolution* 55(10: 122-130).

1999

51. Powell, J. R., A. CACCOME, V. Petrarca, A. della Torre, and M. Coluzzi. Population structure, speciation, and introgression in the *Anopheles gambiae* complex. *Parassitologia* 41:101-113.
50. CACCOME A, J.P. Gibbs, V. Ketmaier, El. Suatoni, J. R. Powell. Origin and evolutionary relationships of giant Gálapagos tortoises. *Proceedings National Academy of Sciences (PNAS) USA*, 96: 13223-13228.
49. Allegrucci G., A. CACCOME, V. Sbordoni. Cytochrome *b* sequence divergence in the European Sea Bass (*Dicentrarchus labrax*) and phylogenetic relationships among some Perciformes species. *J. of Zool. Syst. and Evol. Res.* , 37:149-156.
48. CACCOME, A., G. Amato, O. C. Gratry, J. Behler, J.R. Powell. A molecular phylogeny of four endangered Madagascar tortoises based on mtDNA sequences. *Molecular Phylogenetics and Evolution*, 12: 1-9.

1998

47. CACCOME A., B.A. García, K. D. Mathiopoulos, Gi-Sik Min, E. N. Moriyama, J.R. Powell. Characterization of the soluble guanylyl cyclase -subunit gene in *Anopheles gambiae* and evidence of *Pegasus-27* transposable element insertion. *Insect Molecular Biology*, 8 (1): 23-30.
46. CACCOME A., Gi-Sik Min, J.R. Powell. Multiple origins of cytologically identical chromosome inversions in the *Anopheles gambiae* complex. *Genetics*, 150: 807-814.

1997

45. CACCOME A., G. Allegrucci, C. Fortunato, V. Sbordoni. Genetic differentiation within the European sea bass (*Dicentrarchus labrax*) as revealed by RAPD-PCR assays. *Journal of Heredity*, 88 (4): 316-324.
44. Gleason JH., A. CACCOME, E.N. Moriyama, K.P. White, J.R. Powell. Mitochondrial DNA phylogenies in the *Drosophila obscura* group. *Evolution* , 51(2):433-440.
43. CACCOME, A., M. Milinkovitch, V. Sbordoni, and J.R. Powell. Phylogeny, biogeography, and molecular rates in European newts (genera *Euproctus* and *Triturus*), inferred from mitochondrial DNA sequences. *Systematic Biology*, 46 (10):126-144.

1996

42. CACCOME A., E. N. Moriyama, J. M. Gleason, L. Nigro, and J.R. Powell. A molecular Phylogeny or the *Drosophila melanogaster* subgroup. *Molecular Biology and Evolution* 13:1224-1232.
41. García, B. A., A. CACCOME, K. Mathiopoulos, and J. R. Powell. Inversion monophyly in African malaria vectors. *Genetics* 143:1313-1320.
40. CACCOME, A., B. A. García, and J. R. Powell. Evolution of the mitochondrial DNA control region in the *Anopheles gambiae* complex. *Journal of Insect Molecular Biology*, 5:51-59.

1995

39. Allegrucci, G., A. CACCOME, S. Cataudella, J. R. Powell, and V. Sbordoni. Acclimation of the European sea bass to freshwater: Monitoring genetic changes by RAPD polymerase chain reaction to detect DNA polymorphisms. *Marine Biology* 121:591-599.

1994

38. Besansky, N. J., J. R. Powell, A. CACCOME, D. M. Hamm, J. M. Scott, and F. H. Collins. Molecular phylogeny of the *Anopheles gambiae* complex suggests genetic introgression between principal malaria vectors. *Proceedings National Academy of Sciences (PNAS) USA*, 91:6885-6888.
37. CACCOME, A., M. C. Milinkovitch, V. Sbordoni, and J. R. Powell. Molecular biogeography: Calibrating mitochondrial rDNA evolutionary rates in mountain newts (*Euproctus*) using the Corsica-Sardinia microplate disjunction. *Journal of Evolutionary Biology* 7:227-245.

1993

36. Powell, J. R., A. CACCOME, J. M. Gleason, and L. Nigro. Rates of DNA evolution in *Drosophila* depend on function and developmental stage of expression. *Genetics*, 133:291-298.

1992

35. Sbordoni V., F. Baldari, and A. CACCOME. Tempi relativi e assoluti dell'evoluzione. Contributi Centro Linceo Interdisciplinare B. Segre, Accademia Nazionale dei Lincei, 85: 197-244.
34. CACCOME, A., J. M. Gleason, and J. R. Powell. Complementary DNA-DNA hybridization in *Drosophila*. *J. of Mol. Evol.*, 34:130-140.

1991

33. Allegrucci, G., A. CACCOME, D. Cesaroni, and V. Sbordoni. Evolutionary divergence in European cave crickets: a comparison of scDNA data with allozymic and morphometric distances. *Journal of Evolutionary Biology*, 5: 121-148.
32. CACCOME, A. Man and molecular phylogeny. *Bollettino di Zoologia*, 58: 379-384.
31. Powell, J. R., and A. CACCOME. DNA-DNA hybridization: principles and results. In: G. M. Hewitt (ed) Molecular techniques in taxonomy, NATO ASI Series. Springer, Berlin Heidelberg New York. pp. 117-130.

1990

30. CACCOME, A., and J.R. Powell. A protocol for the TEACL method of DNA-DNA hybridization. In: GM Hewitt (ed) Molecular techniques in taxonomy, NATO ASI Series. Springer, Berlin Heidelberg New York. pp. 385-407.
29. Powell, J.R., and A. CACCOME. The TEACL method of DNA-DNA hybridization: Technical Considerations. *J. of Mol. Evol.*, 30: 267-272.
28. CACCOME, A., and J.R. Powell. Extreme rates and heterogeneity in Insects DNA evolution. *J. of Mol. Evol.*, 30:273-280.
27. Goddard K.A., A. CACCOME, and J.R. Powell. Evolutionary implications of DNA divergence in the *Drosophila obscura* group. *Evolution*, 44: 1656-1670.
26. Sbordoni, V., A. CACCOME, G. Allegrucci, and D. Cesaroni. Molecular Island Biogeography, p.55-83. In: A. Azzaroli (ed.), " Biogeographical aspects of insularity". Atti dei Convegni Lincei, vol. 85, Accademia Nazionale dei Lincei, Roma, Italy, 841pp.

1989

25. Powell, J.R. and A. CACCOME. Intra- and interspecific genetic variation in *Drosophila*. *Genome*, 31:233-238.
24. CACCOME A. and J.R. Powell. DNA divergence among hominoids. *Evolution*, 43 (5):925-942.

1988

23. CACCOME A., G.D. Amato and J.R. Powell. Rates and patterns of scnDNA and mtDNA divergence within the *Drosophila melanogaster* subgroup. *Genetics*, 118:671-683.
22. CACCOME A., R. DeSalle and J.R. Powell. Calibration of the change in thermal stability of DNA duplexes and degree of base-pair mismatch. *J. of Mol. Evol.*, 27: 212-216.

1987

21. Sbordoni V., G. Allegrucci, A. CACCOME, G. Carchini and D. Cesaroni. Microevolutionary studies. In Dolichopodinae cave crickets, pp. 514-540. In "Evolutionary Biology of Orthopteroid Insects", B. Baccetti ed., Horwood Ltd. Publ., Chichester, UK.
20. CACCOME, A. and J.R. Powell. Molecular evolutionary divergence among North American cave crickets. II. DNA-DNA hybridization. *Evolution*, 41: 1215-1238.
19. CACCOME, A. and V. Sbordoni. Molecular evolutionary divergence among North American cave crickets. I. Allozyme variation. *Evolution*, 41: 1198-1214.
18. CACCOME, A., G. D. Amato and J. R. Powell. Intraspecific DNA divergence in *Drosophila*: a study on parthenogenetic *D. mercatorum*. *Molecular Biology and Evolution*, 4: 343-350.

1986

17. Powell, J.R., A. CACCOME, G. D. Amato and C. Yoon. Rates of nucleotide substitutions in *Drosophila* mitochondrial DNA and nuclear DNA are similar. *Proceedings National Academy of Sciences USA (PNAS)*, 83: 9090-9093.

16. CACCONE, A. Molecular studies of evolutionary divergence within and among North American cave crickets. *Phylosophical Doctorate Thesis*. Yale University (USA). Abstract.
15. CACCONE, A., G. Allegrucci, D. Cesaroni, M. Cobolli, E. De Matthaeis, G. La Rosa, V. Sbordoni. Genetic variability and divergence between cave dwelling populations of *Typhlocirolana* from Majorca and Sicily. *Biochemical Systematics and Ecology*, 14: 215-221.

1985

14. CACCONE, A. Gene flow in cave organisms: a qualitative and quantitative approach. *Evolution*, 39: 1223-1235.

1983

13. Cesaroni, D., V. Sbordoni, G. Allegrucci, A. CACCONE, M. Cobolli, E. De Matthaeis. Analisi multivariata dei dati sul differenziamento genetico tra popolazioni e specie di *Dolichopoda* (Orthoptera, Raphidophoridae) dell'Italia centro-meridionale. *Atti XII Congr. Naz. Entomol. Roma*, vol. II: 231-234.
12. De Matthaeis, E., M. Cobolli, M. Mattoccia, G. Allegrucci, A. CACCONE, D. Cesaroni, M. Rampini e V. Sbordoni. Struttura genetica di popolazioni cavernicole di *Bathysciola derosasi* (Coleoptera, Catopidae): bottleneck e divergenza genetica. *Atti XII Congr. Naz. Ital. Entomol. Roma*, vol. II: 253-254.
11. Allegrucci, G., A. CACCONE, D. Cesaroni, M. Cobolli, E. De Matthaeis e V. Sbordoni. Ibridazione naturale e sperimentale tra specie di *Dolichopoda* (Orthoptera). *Atti XII Congr. Naz. Ital. Entomol.*, Roma, vol. I: 225-230.
10. De Matthaeis, E., G. Allegrucci, A. CACCONE, D. Cesaroni, M. Cobolli. Genetic differentiation between *Penaeus kerathurus* and *P. japonicus* (Crustacea, Decapoda). *Marine Ecology Progress Series*, 12: 191-197.

1982

9. CACCONE, A., M. Cobolli, E. De Matthaeis, V. Sbordoni. Una datazione su base genetico molecolare della divergenza tra specie cavernicole e marine di Sferomidi (gen. *Monolistra* e *Sphaeroma*, Crustacea, Isopoda). *Lavori Società Italiana di Biogeografia* (n.s.) 7: 853-867.
8. Allegrucci, G., A. CACCONE, D. Cesaroni, M. Cobolli, E. De Matthaeis, V. Sbordoni. Natural and experimental interspecific hybridization between populations of *Dolichopoda* cave crickets. *Experientia*, 38: 96-98.

1981

7. De Matthaeis, E., M. Cobolli, A. CACCONE, G. Allegrucci, D. Cesaroni, V. Sbordoni. Diversità genetica e flusso genico in ragni cavernicoli del genere *Nesticus*. In: "Ecologia", SITE, Atti 1:275-279.
6. Cobolli, M., G. Allegrucci, A. CACCONE, D. Cesaroni, E. De Matthaeis, V. Sbordoni. Variabilità genetica in popolazioni cavernicole del genere *Troglophilus* (Orth. Rhaph.): influenza di fattori stocastici e deterministici. In: "Ecologia", SITE, Atti 1:269-272.
5. Cesaroni, D., G. Allegrucci, A. CACCONE, M. Cobolli, E. De Matthaeis, M. Di Rao. Genetic variability and divergence between population and species of *Nesticus* cave spiders. *Genetica*, 56: 81-92.
4. Sbordoni, V., G. Allegrucci, A. CACCONE, D. Cesaroni, M. Cobolli, E. De Matthaeis. Genetic variability and divergence in cave populations of *Troglophilus cavicola* and *T. andreinii* (Orthoptera, Raphidophoridae). *Evolution*, 35 (1): 226-233.

1980

3. Sbordoni, V., G. Allegrucci, A. CACCONE, D. Cesaroni, M. Cobolli, E. De Matthaeis. A preliminary report of the genetic variability levels in troglobitic Bathysciinae: *Leptodirus hohenwarti* and two *Orostygia* species (Coleoptera, Catopidae). *Fragmenta Entomologica*, 15: 327-336.
2. Sbordoni, V., A. CACCONE, E. De Matthaeis, M. Cobolli. Biochemical divergence between cavernicolous and marine Sphaeromidae and the Mediterranean salinity crisis. *Experientia*, 36: 48-49.

1978

1. Sbordoni, V., A. CACCONE, E. De Matthaeis, M. Cobolli. Una datazione su base genetico-molecolare della divergenza tra specie cavernicole di *Monolistra* (Isopoda, Sphaeromidae). In "Biogeografia delle caverne italiane". XXII Congresso Nazionale Società Italiana di Biogeografia, Verona ITALY: 91-94.