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## Phylogenetic signal of nematocysts in Staurozoa (Cnidaria)

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## **June 26<sup>th</sup> to 30<sup>th</sup>, 2015** Casa Grande Hotel Resort in Guarujá Brazil







Evolution 2015 is the joint annual meeting of the Society for the Study of Evolution (http://www.evolutionsociety.org/) (SSE), the Society of Systematic Biologists (http://systbiol.org/) (SSB), and the American Society of Naturalists (http://www.amnat.org/) (ASN). The meeting this year will be held on June 26-30 at the Casa Grande Resort

(http://www.casagrandehotel.com.br/pt/Eventos-e-Convencoes) in Guarujá, Brazil. Evolution 2015 in Brazil will be a great opportunity for the international scientific community of Evolutionary Biologists to interact with Brazilians to foster scientific cooperation programs and discuss their research.

This meeting is the premier annual opportunity for sharing scientific research related to evolution. Symposia presentations, concurrent contributed papers, and poster sessions will be presented by the 1,500 expected participants. Product and service providers will contribute to the meeting through their exhibits.

http://evolution2015.org/

## Sunday June 28, 19:00 - 21:00 PM

Suriday Julie 26, 19:00 - 21:00 Pivi	
DO CO3	Poster Session II
PO-603	Reduced cellular immune response in social insect lineages
DO 643	Margarita M López-Uribe
PO-612	Phylogenetic signal of nematocysts in Staurozoa (Cnidaria)
	Lucilia Souza Miranda
PO-615	Revisiting ecogeographic hypotheses behind latitudinal morphological variation: climatic correlates
	of body size in Torrent Ducks
	Natalia Gutierrez-Pinto
PO-637	BREEDING SYSTEM EVOLUTION INFLUENCED THE GEOGRAPHIC EXPANSION AND DIVERSIFICATION
	OF THE CORVIDES
	Petter Zahl Marki
PO-638	Breeding system transitions and the evolution of reproductive isolation in the large and diverse
	tropical genus Dendrobium (Orchidaceae)
	Fabio Pinheiro
PO-649	Diversification dynamics of Placentalia (Mammalia): integrating the fossil record with molecular
	phylogenies
	Mauro Toshiro Caiuby Sugawara
PO-651	Use of wing geometry to diagnose cryptic species of the Anopheles (Nyssorhynchus) strodei
	complex
	Flávia Virginio Fonseca
PO-666	Variability in the micro and macroevolutionary scales: a review of the patterns of morphological
	variability in the Cnidaria Medusozoa
	Amanda Ferreira e Cunha
PO-674	The turtle ants: understanding the diversification of a socially complex lineage and its associated
	microbes
	Shauna Price
PO-681	First molecular phylogeny of the sea-star Echinaster (Asteroidea: Echinodermata) reveals a non-
	monophyletic genus and insights for evolutionary history
	Elinia Medeiros Lopes
PO-683	Molecular evolution of Odorant Receptors proteins gene family in two closely related Anastrepha
	fruit flies .
	Diana Marcela Rojas Gallardo
PO-684	Scorched Mussels (Brachidontes spp.) from the Southwestern Atlantic: the role of the Isthmus of
	Panama and the Amazon River in their Speciation
	Berenice Trovant
PO-685	A BLASTP-free clustering method for classifying orthologs within a single gene family
	Stephen Bond
PO-686	Tempo and mode of climatic niche evolution in Primates
	Andressa Duran
PO-687	Molecular phylogeny of the Utricularia sect. Foliosa Kamiénski complex addressed using rps16,
	trnD-T, trnL-F and ITS sequences
	Paulo Cesar Baleeiro
PO-688	Diversification at large body sizes: Morphological novelty and ecological opportunity in Neotropical
	armoured catfishes (Loricariidae)
	Fabio F Roxo

Phylogenetic signal of nematocysts in Staurozoa (Cnidaria)

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Description of cnidome is considered important information for taxonomic studies of Cnidaria, although their value in identifying species has been recently questioned. Staurozoa has the least diverse cnidome among medusozoan cnidarians, with two types of nematocysts, euryteles and isorhizas. These nematocysts are concentrated in two main regions: the secondary tentacles and the white spots on the subumbrella. The aim of this study is to assess the taxonomic and evolutionary information contained by the cnidome from 17 species of Staurozoa. We employ our working hypothesis of evolutionary relationships within Staurozoa to comparatively analyze the phylogenetic signal of nematocysts from secondary tentacles and white spots of nematocysts, as measured with traditional and geometric morphometrics. Additionally, we generate a phylogenetic hypothesis based exclusively on continuous characters derived from nematocysts, as an independent test of the molecular hypothesis. Our results suggest that there is significant phylogenetic signal in the nematocysts from secondary tentacles, and the combined use of nematocysts from secondary tentacles and white spots can be useful to distinguish species. The possible relationship of phylogenetic signal and development of nematocysts in Staurozoa is discussed, as well as the possible evolutionary pressures involved in the presence and absence of phylogenetic signal.