

Vorläufiges Tagungsprogramm der 16. Jahrestagung der Gesellschaft für Biologische Systematik (GfBS)

(*Stand vom 17.03.2015. Änderungen vorbehalten)

	Zeiten	Veranstaltung (Redezeit)	Redner	Thema	Chair	Raum
Mittwoch						
	14.00- 18.00	Treffen GfBS Vorstand				Seminarraum
Mi., 18.03.	18.00- 22.00	Registrierung und Icebreaker		-	-	Empfangshalle/ Savanne
Donnerstag						
Do., 19.03.	8.00- 9.10	Registrierung			-	Empfangshalle
	9.10- 9.30	Eröffnung	J. W. Wägele, U. Fritz	Begrüßung, Organisatorisches, Historisches	-	Hörsaal
	9.30- 10.15	Invited Speaker (40+5min)	Wolfgang Weisser	Ecology and Systematics – how to form a new partnership	J. W. Wägele	Hörsaal
	10.15- 11.00	Invited Speaker (40+5min)	Florian Leese	Improvement of freshwater ecosystem assessment with DNA barcoding	J. W. Wägele	Hörsaal
	11.00- 11.30-	Pause (30 min)				
	11.30- 12.00	Invited Speaker(25+5min)	HaJo Mader	Monitoring in wilderness areas	J.W. Wägele	Hörsaal
	12.00- 13.30	Mittagspause (90 min) oder Besichtigung des Biohistoricum/Sammlungsführung				
	13.30- 13.45	Kurzvortrag (12+3)	Alexander M. Weigand, Daniel Grabner, Martina Weiss, Daniel Hering, Bernd Sures, Ralph	DNA barcodes of freshwater amphipods as a springboard for investigating biological processes in natural and novel freshwater ecosystems		Hörsaal

Do., 19.03.		Tollrian & Florian Leese		F. Leese	
	13.45- 14.00	Kurzvortrag (12+3)	<u>Matthias Geiger</u>		
	14.00- 14.15	Kurzvortrag (12+3)	<u>Elise Laetz, Leif Moritz, André Haubrich & Heike Wägele</u>		
	14.15- 14.30	Kurzvortrag (12+3)	<u>Jahn R., Abarca N., Enke N., Kusber W.-H., Skibbe O. & Zimmermann J.</u>		
	14.30- 14.45	kurze Pause (20 min)			
	14.50- 15.05	4 Kurzvorträge (12+3min)	<u>Caroline Ring & Michael Ohl</u>	T. Töpfer	Hörsaal
	15.05- 15.20		<u>Constantin Zohner</u>		
	15.20- 15.35		<u>Alexander Rockinger & Susanne S. Renner</u>		
	15.35- 16.00	kurze Pause (25 min)			
	16.00- 16.20	Verleihung der Ehrennadel			Hörsaal
	16.20- 16.35	3 Kurzvorträge (12+3min)	<u>Rachel C. M. Warnock, Jonas Eberle & Dirk Ahrens</u>	T. Wesener	Hörsaal
	16.35- 16.50		<u>Dirk Ahrens, Tomochika Fujisawa, Hans-Joachim Krammer, Jonas Eberle, Silvia Fabrizi & Alfried P. Vogler</u>		
	16.50- 17.05		<u>Jan Phillip Oyen</u>		
	17.05-	Pause			

Do., 19.03.	17.30	(25 min)			
	17.30- 19.30	Treffen der AG Junge Systematiker (JuSys) (offen für Interessenten)		AG Treffen	CCF Schinkel Seminarraum
	17.30- 19.30	Treffen der AG Kuratoren (offen für Interessenten) Thema: Preserving Tissues, Molecules, and other odd specimens – a practical look at special collections	Peter Giere & Peter Michalik	New and uncommon collection types – an introduction	P. Giere Festsaal
			Dirk Neumann	DNA and Tissue storage – from individual trials and chaos towards modern cross-linked Biorepositories	
			Brian Tindall		
			Jonas Astrin	Biobank networks (mit anschließender Labor Führung)	
ab 20.00 Konferenz-Dinner (!!verschoben von Fr. auf Do.!!) Abendessen in gemütlicher Atmosphäre Individuelle Bestellung à la carte					Restaurant ENTE in der Bonner City

Freitag

Fr., 20.03.	9.00- 09.30	Invited Speaker(25+5min)	Maximilian Weigend	Critical plant distribution data as a basis for mesoscale conservation planning - examples from the South American Andes	J.W. Wägele	Hörsaal	
	09.30- 10.00	Invited Speaker(25+5min)	Peter Zulka	On subspecies, cryptic species and synonyms – taxonomic impediment in the national Red List assessment process.			
	10.00- 10.15	Kurzvortrag (12+3min)	Omid Paknia, Hossein Rajaei Sh. & André Koch	Lack of well-maintained natural history collections and taxonomists in megadiverse developing countries hampers global biodiversity exploration			
	10.15- 10.30	Kurzvortrag (12+3min)	Thomas Wesener	The GBOL - Myriapoda project: 1000 sequences reveal insights and shortcomings of the Barcoding of myriapods			
	10.30- 11.00	Pause (30 min)					
	11.00- 12.00	20 Posterkurzvorträge (2-3min)				Hörsaal	
	12.00- 13.30	Mittagspause (90 min) oder Tour durch die Sammlungen des Museums					

Fr., 20.03.	13.30- 13.45	4 Kurzvorträge (12+3min)[parallel]	<u>Thorleif H. Dörfel</u> & Michael Ohl	The Australian digger wasps of the genus <i>Sphex</i> – a taxonomic revision	R. Peters	Hörsaal
	13.45- 14.00		<u>Ximo Mengual</u> , Gunilla Ståhls & Santos Rojo	Systematics of pipizines (Diptera: Syrphidae) with implications for the evolution of aphidophagy		
	14.00- 14.15		<u>Ralph S. Peters</u> , Oliver Niehuis & Lars Krogmann	Phylogenomic data and a strikingly rich fossil record illuminate the evolution of jewel wasps (Hymenoptera: Chalcidoidea)		
	14.15- 14.30		<u>Lena Lutz</u> , Hannes Baur & Ralph S. Peters	Taxonomy of some common Western European jewel wasps (Hymenoptera: Chalcidoidea: Pteromalidae) - from integrative to desperate and back		
	13.30- 13.45	4 Kurzvorträge (12+3min)[parallel]	Jonas Astrin	Trends in DNA barcoding and biodiversity research - a bibliometric perspective	J. Astrin	Festsaal
	13.45- 14.00		Gerhard Haszprunar	Use, misuse, and criteria for „a DNA-barcoded species“		
	14.00- 14.15		<u>Harald Letsch</u> , Brigitte Gottsberger & Jessica Ware	Differential diversification in anisopterous dragonflies (Odonata): a result of different habitat use and climate?		
	14.15- 14.30		Werner Kunz	400 new bird species - the "Tobias-criteria" for species delimitation		
	14.30- 15.00	Pause (30 min)				
	15.00- 15.15	3 Kurzvorträge (12+3min)[parallel]	Till Töpfer	Acquisition, loss and re-activation of pathways of plumage colour synthesis in bullfinches (Aves: Fringillidae: <i>Pyrrhula</i>)	C. Koch	Hörsaal
	15.15- 15.30		Markus Lambertz	Pulmonary complexity as the starting point for amniote evolution		
	15.30- 15.45		<u>Timo Hartmann</u> , Flora Ihlow, Morris Flecks, Andreas Schmitz & Dennis Rödder	Underestimated diversity in <i>Calotes mystaceus</i> (Squamata: Agamidae) - An integrative approach to unravel a putative species complex		
	15.00- 15.15	3 Kurzvorträge (12+3min)[parallel]	<u>Anja C. Schunke</u> & Anna Anschütz	Lost variation - The hyoid bone in rodents	T. Struck	Festsaal
	15.15- 15.30		<u>Michael Weidhase</u> & Christoph Bleidorn	Regeneration in annelids and its relationship to phylogeny		
	15.30- 15.45		<u>Ekin Tilic</u> & Thomas Bartolomaeus	Chaetae and chaetogenesis as a character complex to analyze annelid evolution		

	15.45- 16.05	kurze Pause (20 min)				
Fr., 20.03.	16.05- 16.20	3 Kurzvorträge (12+3min)[parallel]	Morris Flecks, Lasse Lemm, Timo Hartmann, Jonas Astrin, Nikolay A. Poyarkov, Sebastian Steinfartz, Wolfgang Böhme & Dennis Rödder	How to identify species? An integrative approach to delimiting species in the threatened Middle Eastern mountain newts	C. Koch 	Hörsaal
	16.20- 16.35		Dennis Rödder, Eva Meyers, Flora Ihlow, Morris Flecks, Timo Hartmann, Tanja Ziesmann, Igor Oliveira, Faraham Ahmadzadeh & Ralph S. Peters	Identifying the gaps: status of molecular exploration of global amphibian diversity		
	16.35- 16.50		Felix Pokrant & Wolfgang Böhme	Integrative Taxonomy of the Grass Snake (<i>Natrix</i> <i>natrix</i>)		
	16.05- 16.20	3 Kurzvorträge (12+3min)[parallel]	Juliane Kretschmann, Marc Gottschling & Malte Elbrächter	The epitype - an important tool for reliable species identification in protists such as dinophytes	D. Ahrens 	Festsaal
	16.20- 16.35		Alexander Riedel, Thomas van de Kamp, Rene Tänzler & Michael Balke	A species description pipeline for the hyperdiverse weevil genus <i>Trigonopterus</i>		
	16.35- 16.50		Natalie Cusimano & Susanne S. Renner	Does tree shape matters in ancestral state reconstructions?		
	16.50- 17.00	kurze Pause (10 min)				
	ab 17.00	Mitgliederversammlung		- Bericht des Vorstands - Rensch-Preis Verleihung - Laudatio - Studenten Poster Preise	-	Hörsaal
	ab 20.00	Abend zur freien Verfügung				
Samstag						
Sa., 21.03.	9.00- ca. 18.00	Workshops 1	Oliver Niehuis, Malte Petersen, Christoph Mayer, Vera Batista & Manuela Sann	Target DNA Enrichment – Methodology, Software, Practical Experience Report	Kursraum Neubau	

	9.00- ca. 16.00	Workshops 2	Matthias Bernt & Alexander Donath	“MITOS” Mitogenom-Annotation	Seminarraum (Hauptgebäude)
	9.00- 18.00	Workshop 3	A. Blanke	3D-Rekonstruktion – High Quality 3D Reconstruction for Everyone: open-Source Solutions and their Workflow to analyse and publish MicroCT Data	Museumsschule (neben Seminarraum)