HYBRID CONFERENCE

South African Genetics Society & South African Society for Bioinformatics JOINT CONFERENCE

BIO2022 Bioscience, Big Data & The 4th Industrial Revolution

PROGRAMME

SUNDAY 24 APRIL

16:00 - 18:00	Welcome Reception (STIAS)	
	Registration, canapés, red & white wine, soft drinks	

DAY 1| monday 25 April

07:30 - 08:45	Registration (STIAS foyer)			
08:45	Welcome & Presidential Address SAGS/SASBi (Clint Rhode, Gerard Tromp, Sanushka Naidoo)			
09:15 - 10:30	JOINT PLENARY SESSION (Auditorium 1 & 2) Chair: Dr Clint Rhode			
09:15 - 10:00	Invited Plenary Speaker: Prof Chandra Verma Bioinformatics Institute, The Agency for Science, Technology and Research, Singapore Translating biomolecular modelling to industry and the clinic			
10:00 -10:30	Invited Keynote Speaker (Biomedical Sciences): Prof Soraya Bardien Division of Molecular Biology & Human Genetics, Stellenbosch University Parkinson's disease: a genetic conundrum			
10:30 - 11:00	TEA			
11:00 - 12:45	JOINT SESSION: Biomedical Sciences (Auditorium 1 & 2) Chair: Prof Helena Kuivaniemi			
11:00 - 11:15	Moela Pontsho Retinoblastoma Binding Protein 6 (RBBP6) expression elicits the sensitization of cervical cancer cells to cisplatin treatment			
11:15 - 11:30	Bolu Oladunjoye Structure-based virtual screening of selected malaria box compounds against falstatin, a multi- staged protein in <i>Plasmodium falciparum</i>			
11:30 - 11:45	Adebowale Emmanuel Aladejana In vitro evaluation of the anti-diabetic potential of Helichrysum petiolare Hilliard & B.L. Burtt using HepG2 and L6 cell lines			
11:45 - 12:00	Kathryn Step Parkinson's disease, plink, and putative disease susceptibility variants			
12:00 - 12:15	Fourie Joubert Germline sequence variants contributing to cancer susceptibility in South African breast cancer patients of African ancestry			
12:15 - 12:30	Graeme Ford Pharmacogenetics of CYP2a6, CYP2b6 and ugt2b7: relevance to HIV treatment in African populations			
12:30 - 12:45	Natasha Kitchin The gut microbiota in foetal alcohol spectrum disorders			
12:45 - 14:00	LUNCH			
14:00 - 15:45	JOINT SESSION: Bio-Economy, Industries & Technologies (Auditorium 1 & 2) Chair: Willem Botes			
14:00- 14:15	Gold Sponsor presentation - DIPLOMICS			
14:15 - 14:30	Oleg Reva Genomic and epigenetic comparison of Bacillus strains suitable for plant protection			
14:30 - 14:45	Dave Berger RNAi "vaccines" for plants: grey leaf spot disease control in maize			
14:45 - 15:00	Demissew Teshome Transcriptional re-programming during recovery from short term drought stress in Eucalyptus grandis			
15:00 - 15:15	Anneri Lotter Phased genome assembly and haplogenome comparison in an F1 hybrid of <i>Eucalyptus urophylla</i> and <i>E. grandis</i>			
15:15 - 15:30	Nanette Christie Data driven molecular breeding of tropical pines			
15:30 - 15:45	Masethabela Maphatsoe Exploring the role of microbial cars as potential biocatalysts, and its potential contribution to South Africa's bioeconomy			
15:45 - 16:15	TEA			
16:15 - 17:00	JOINT SESSION: Poster Flash Presentations (Auditorium 1 & 2)			
17:00 - 17:45	SASBi AGM			





DAY 2| tuesday 26 april

07:30 - 08:45	Registration (STIAS foyer)			
08:45	Welcome & housekeeping (Clint Rhode, Gerard Tromp)			
09:00 - 10:20	JOINT KEYNOTE SESSION: Ecology & Evolution (Auditorium 1 & 2) Chair: Dr Ruben Cloete			
09:00 - 09:35	Invited Keynote Speaker (Ecology and Evolution): Peter van Heusden South African National Bioinformatics Institute Realising the potential for pathogen genomics in advancing public health in Africa			
09:35 - 09:50	Olga Bochkareva Detection of genome rearrangements re	esponsible for bacterial phenotype switching		
09:50 - 10:05	Michal Slupski Phylogenetic analysis of circadian clock genes in Chrysoporthe species			
10:05 - 10:20	Cassandra Bianca Schoeman The Encephalartos natalensis-cyanobacterial coralloid root partnership for nitrogen acquisition			
10:20 - 11:00	TEA			
11:00 - 12:45	PARALLEL SESSION: Ecology & Evolution (Auditorium 1 & 2) Chair: Prof Irene Barnes	PARALLEL SESSION: Biomedical Sciences (Manor House) Chair: Dr Rencia Van Der Sluis		
11:00 - 11:15	Sumari Venter Multi-omics approaches to studying host-pathogen interactions between the forest pest, <i>Sirex</i> <i>noctilio</i> and its biocontrol agent, <i>Deladenus siricidicola</i> <i>noctilio</i>	Bonginkosi Shabangu Genomic characterisation of <i>Acinetobacter baumannii</i> associated with neonatal sepsis and stillbirths in a South African population		
11:15 - 11:30	Tanya Welgemoed Pangenome of African populations of the maize fungal pathogen <i>Cercospora zeina</i>	Souiai Oussema Longitudinal and comparative analysis of Tunisian newborns gut microbiota according to delivery mode		
11:30 - 11:45	Khajamohiddin Syed Impact of lifestyle on cytochrome p450 monooxygenase evolution is clearly evident in bacteria	Sikozile Ncembu Scalable production of h22(scfv)-eta' targeting cd64 in acute myeloid leukemia (AML)		
11:45 - 12:00	Leandri Klynsmith Tissue- and sex-specific odour coding for ecological niche adaptation in the forest pest, <i>Sirex noctilio</i>	Nikita Simone Pillay Novel Gene Discovery in a Xhosa Family with Parkinson's Disease		
12:00 - 12:15	Shannon Bennet Fusarium soil survey from the Kalahari	Chrystal Steyl Host genetic factors contributing to susceptibility to Covid-19		
12:15 - 12:30	Shailesh Joshi Evaluation of gene flow from commercial sugarcane hybrids to compatible wild relatives	Caitlin Uren The Northern Cape tuberculosis case- control consortium (NCTBC3)		
12:30 - 12:45	Lindokuhle Emmanuel Interaction of <i>Paraburkholderia</i> species with different agricultural and acacia legumes in South Africa	Kili James Identification of Genetic Variants in Human- Mouse Orthologous Genes in Patients Diagnosed with Non-syndromic Hearing Impairment		
12:45 - 13:45	LUNCH			
13:45 - 15:05	JOINT SESSION: Functional Biology (Auditorium 1 & 2) Chair: Dr Marlon le Roux			
13:45 - 14:20	Invited Keynote Speaker: Prof Sanjib Kumar Panda Department of Biochemistry, Central University of Rajasthan Functional genomics for abiotic stress tolerance in crops			
14:20 - 14:35	Julia Candotti Towards haplotype and structural variant based genetic dissection of complex traits in <i>Eucalyptus</i> hybrids			
14:35 - 14:50	Medha Sood The ultrastructure and role of plastids in carbon partitioning during <i>xylogenesis</i> in VND7-inducible Arabidopsis			
14:50 - 15:05	Robyn Jacob Comparing the resistant and susceptible defense response of sugarcane challenged with <i>Eldana</i> saccharina			
15:05 - 15:30	TEA			
15:30 - 16:30	JOINT SESSION: Functional Biology (Auditorium 1 & 2) Chair: Dr Cedric Werely			
15:30 - 15:45	Kelvin Hull Functional genomic processes of early domestication and artificial selection in black soldier fly colonies			
15:45 - 16:00	Deborah Narh Mensah NRPS-dependent siderophore synthetase gene clusters and characteristics of NRPS siderophore synthetase genes in Armillaria and other species in the <i>Physalacriaceae</i>			
16:00 - 16:15	Tassin Jackson Draft genome assembly and annotation of <i>Argyrosomus japonicus</i> in South Africa, provides insight into the evolutionary characteristics of this species			
16:15 - 16:30	Darisia Moonsamy Expression profiling of interferon-stimulated genes in peripheral blood mononuclear cells from healthy and SARS-COV-2 infected individuals			
16:30 - 17:00	JOINT SESSION: Poster Flash Presentations (Auditorium 1 & 2)			
17:00 - 17:45	SAGS AGM			
19:00	Wine Pairing Social Event at Lanzerac Wine Estate			

$DAY\ 3$ | wednesday 27 April

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07:30 - 08:45	Registration (STIAS foyer)			
08:45	Welcome & housekeeping (Clint Rhode, Gera	ard Tromp)		
09:00 - 10:30	JOINT KEYNOTE SESSION: Statistical & Computational Methods (Auditorium 1 & 2) Chair: Prof Gerard Tromp			
09:00 - 09:45	Invited Keynote Speaker: Prof Fyodor Kondrashov Evolutionary Genomics, Institute of Science and Technology Austria (IST Austria)			
09:45 - 10:00	Ozlem Tastan Bishop Role of variations in modern drug research and development – Way forward via dynamic residue networks			
10:00 - 10:15	Shae Swanepoel The in planta expression of Austropuccinia psidii in resistant and susceptible Eucalyptus grandis			
10:15 - 10:30	Hannes Strydom Modelling the population dynamics of CRISP-CAS9 gene drive systems in Sirex noctilio			
10:30 - 11:00	TEA			
11:00 - 12:30	PARALLEL SESSION: Statistical & Computational Methods (Auditorium 1 & 2) Chair: Dr Alisa Postma	11:00 - 12:30	PARALLEL SESSION: Conservation & Environmental Sustainability (Manor House) Chair: Prof Aletta van der Merwe	
11:00 - 11:15	Tiego Mohlaba Development of comprehensive single-cell RNA sequencing pipeline: pre-processing, quality control and identification of outliers	11:00 - 11:30	Keynote address: Dr Simo Maduna (NIBIO, Norway) Genetic monitoring for biodiversity conservation and environmental sustainability – Examples from the Norwegian Barents Region	
11:15 - 11:30	Ruben Cloete Molecular docking of zinc database natural compounds to SARS- COV-2 coronavirus proteins to identify novel inhibitors with antiviral activity	11:30 - 11:45	Emma Hubble Population dynamics of two endemic catshark species inferred from molecular and tag-recapture data	
11:30 - 11:45	Matthew Adeleke Designing t- cell epitope-based vaccine against Eimeria from microneme protein 2 (MIC2) antigen using immunoinformatics approach	11:45 - 12:00	Mondi Xaba Long read de novo assembly of the <i>Acacia mearnsii</i> genome	
11:45 - 12:00	Keaghan Brown Automated computational workflow to prioritize potential resistance variants identified in HIV Integrase Subtype C and CRF02_AG°	12:00 - 12:15	Mia Groeneveld Towards resolving genetic differences and population distributions of rhino rays from the Southwest Indian Ocean region	
12:00 - 12:15	Carla Louw Wastewater-based epidemiology as a viable minimum data strategy for antimicrobial resistance surveillance in a peri-urban setting	12:15 - 12:30	Anandi Bierman Phylogeography of the invasive Ambrosia beetle, <i>Euwallacea fornicatus</i> - a prequel to fine-scale landscape genomics for understanding invasiveness	
12:15 - 12:30	Shailesh Joshi Evaluation of gene flow from commercial sugarcane hybrids to compatible wild relatives			
12:30 - 13:45	LUNCH			
13:45 - 15:15	JOINT SESSION: Functional Biology (Auditor	ium 1 & 2) Cha	ir: Prof Sanushka Naidoo	
13:45 - 14:00	Bulelani Sizani Structural and functional characterization of NLR proteins differentially expressed in Cassava plant Infected with SACMV			
14:00 - 14:15	Lazarus Takawira Inferring R2R3-MYB family transcription factor gene targets in <i>Eucalyptus grandis</i> using DNA Affinity Purification sequencing and machine learning			
14:15 - 14:30	Carla Buitendag The maize fungal pathogen Cercospora zeina ecp2 virulence effector causes necrosis in tobacco			
14:30 - 14:45	Tendo Stanley Tshilate Genome comparison	in abalone spec	ies to provides insights into growth related genes	
14:45 - 15:00	Rencia van der Sluis Critical analyses of the literature: In isovaleryl acidemia, is isovalerylglycine a product fromed by glycine-n acyltransferase?			
15:00 - 15:15	Kelda Perumal Differences and commonalities induced by pma and 1,25(OH)2d3 in THP-1 cells during monocyte-to- macrophage differentiation			
15:15 - 16:00	TEA			
16:00 - 16:30	JOINT SESSION: E-posters Presentations (Auditorium 1 & 2)			
19:00	Gala Dinner at STIAS			

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Arranged by