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Welcome to the 34th MSPP Scientific Meeting 2021

The Malaysian Society of Pharmacology and Physiology (MSPP) has been the primary Malaysia learned society of pharmacologists and physiologists for more than 33 years. The Society has been organizing its own Scientific Meetings as a platform for researchers to share their research findings and build networking and collaborations. To date, MSPP is proud to have organized 34 scientific meetings. Due to the pandemic, for the first time the 34th Scientific Meeting of MSPP was held virtually on July 15-17, 2021. The meeting which was organized by Monash University Malaysia together with MSPP in collaboration with several local and private universities in Malaysia had attracted 700 delegates from locally and abroad and received more than 160 abstracts. The scientific programme this year focused on translational research under the theme 'Trailblazing the Translational Research'. The three-day conference provided opportunities for sharing and exchanging evidence-based practices and scientific endeavors among physiologists, pharmacologists, physicians, general practitioners, scientists and allied health professionals.

LIST OF ORGANIZERS

Jeffrey Cheah School of Medicine and Health Science, Monash University Malaysia Malaysian Society of Pharmacology and Physiology (MSPP)



34th MSPP 2021 Programme "Trailblazing the Translational Research"

Day 1: 15th July 2021

Programme

8.30 am – 9.15 am	Opening Ceremony
	Speakers Prof. Dr. Nafeeza Hj Mohd Ismail (President, MSPP; International Medical University)
	Prof. Dr. Shajahan Yasin (Head, JCSMHS, Monash University Malaysia)
	Assoc. Prof. Dr. Mohd. Farooq Shaikh (Chair, 34th MSPP 2021; Monash University Malaysia)
9.15 am – 10.00 am	Keynote Address Prof. Dr. Gordon Harold Williams Professor of Medicine, Harvard Medical School, USA
	Title: Personalized Medicine: The Link Between People, Animals and Cells
10.00 am – 10.30 am	
10.00 am – 10.30 am 10.30 am – 11.00 am	Between People, Animals and Cells
	Between People, Animals and Cells Break Plenary Lecture 1 Prof. Dr. Wayne Hodgson

12.00 pm – 2.00 pm	Lunch Break
2.00 pm – 3.00 pm	MSPP Young Investigator Award 2021
3.00 pm – 3.30 pm	Break
3.30 pm – 5.30 pm	MSPP Annual General Meeting
Day 2: 16 th July 2021	
9.00 am– 9.30 am	Plenary Lecture 2 Assoc. Prof. Dr. Elizabeth Davis Monash University, Australia
	Title: Developing and Incorporating Learning Resources to Facilitate Active Learning in Pharmacology
9.30 am – 9.35 am	Break
9.35 am – 10.00 am	Symposium 1: Drug Metabolism Research
	Invited speaker: Prof. Dr. Ong Chin Eng International Medical University
	Title: Structure-Function Analysis of Cytochrome P450 Enzymes: Kinetic and Computational Modelling for Drug Clearance Prediction
	Symposium 2: Natural Products and Integrative Medicine
	Invited speaker: Assoc. Prof. Dr. Wan Amir Nizam Wan Ahmad Universiti Sains Malaysia
	Title: Flavonoids and Metainflammation in Diabetes
10.00 am – 10.15 am	Break



10.15 am – 11.15 am	Finalist 1: Oral presentations
	Finalist 2: Oral presentations
11.15 am – 11.30 am	Break
11.30 am – 12. 30 pm	Finalist 1: Shortlisted Poster presentations
	Finalist 2: Shortlisted Poster presentations
12.30 pm – 2.00 pm	Lunch Break
2.30 pm – 2.50 pm	Symposium 3: Cardiovascular Research
	Invited speaker: Assoc. Prof. Dr. Satirah Zainalabidin Universiti Kebangsaan Malaysia
	Title: The Action of SAC on Cardiac Remodelling in Myocardial Injury
	Symposium 4: Cancer Research
	Invited speaker: Dr. Alan Yiu Wah Lee Monash University, Malaysia
	Title: Exploration of the Tumor Suppressor Function of Semaphorin 5A in Brain Cancer Treatment
2.50 pm – 3.00 pm	Break
3.00 pm – 4.00 pm	Finalist 3: Oral presentations
	Finalist 4: Oral presentations
4.00 pm – 4.15 pm	Break
4.15 pm – 5.15 pm	Finalist 3: Shortlisted Poster presentations
	Finalist 4: Shortlisted Poster presentations

Day	3:	17^{th}	July	2021
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9.00 am – 9.30 am	Plenary Talk 3: Prof. Dr. Michael Fenech University of South Australia, Australia
	Title: Nutrigenomics from Laboratory to Clinical Practice
9.30 am – 9.50 am	Symposium 5: Neurological Diseases
	Invited speaker: Prof. Dr. Renu Agarwal International Medical University
	Title: Glaucoma and Neuroinflammation: Role of Magnesium-Based Compounds as Therapeutic Instruments
	Symposium 6: Metabolic Diseases
	Invited speaker: Assoc. Prof. Dr. Naguib Salleh University of Malaya
	Title: Ion channels in Endometrium: Their Roles in Uterine Fluid Regulation and Embryo Implantation
9.50 am – 10:30 am	Finalist 5: Oral presentations
	Finalist 6: Oral presentations
10.30 am – 11.30 pm	Break
11.30 pm – 12.00 pm	Awards Presentation
	Closing ceremony and remarks

PNPIM-20

Microarray data analysis to identify differentially expressed genes and biological pathways associated with tualang honey supplementation in NASH animal model

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Tualang Honey (TH) is a multifloral jungle honey that has been shown to have anti-inflammatory and antioxidant properties. Thus, it has been proposed as a potential supplement in the management of non-alcoholic steatohepatitis (NASH). This study aimed to identify potential key genes and biological process associated with the therapeutic effects of TH in NASH animal model. Twenty Sprague Dawley rats were used in this experiment. Five rats were given standard commercial pellets throughout the experiment and acted as controls. Fifteen rats were subjected to the 12% high-cholesterol diet (HCD) for sixteen weeks to induce NASH. They were then divided into 3 groups, with each group receiving 3 different dosages of TH supplementation for four weeks. RNA was extracted from serum using RNeasy extraction kit (Qiagen) and microarray analysis was performed. A total of 3062 differentially expressed genes (DEGs) were identified including 2788 upregulated genes and 274 downregulated genes in all treated groups. There were 15 genes significantly upregulated including aldehyde dehydrogenase family, cytochrome P450 and nuclear transcription factor. Whilst, 13 genes were significantly downregulated such as apelin, interferon regulatory factor 2 and stearoyl-Coenzyme A. Following pathway enrichment analysis, 34 biological processes were significantly modified (p < 0.05). Among the pathways implicated in the effects of TH were those involved in the inflammatory response, lipid biosynthetic process and cholesterol metabolic process. The present



study demonstrated that the DEGs involved in the therapeutic effects of TH were enriched in several pathways and further validation is required to confirm their specific roles.

Keywords: Tualang honey, NASH, DEG, Microarray, Pathway analysis

