

36th Malaysian Society of Pharmacology and Physiology Annual Scientific Meeting 2023

Santhra Segaran Balan*, Rusliza Basir, Yong Yoke Keong, Noor Aishah Mohammed Izham, Jasmine Siew Min Chia, Fezah Othman, Jonathan Chee Woei Lim, Siti Farah Md Tohid, Hasnah Bahari and Chau Ling Tham

*Correspondence: santhra@msu.edu.my

Received: 1 August 2023; Revised: 4 August 2023; Accepted: 6 August 2023; Published: 6 August 2023
DOI <https://doi.org/10.28916/lsm.7.1.2023.132>

ABSTRACT

The 36th Malaysian Society of Pharmacology and Physiology (MSPP) Annual Scientific Meeting was hosted in collaboration with the Faculty of Medicine and Health Sciences, Universiti Putra Malaysia from the 7th to 8th of August 2023 at Bangi Resort Hotel in Bangi, Malaysia. The event focused on the theme "*Pharmacology & Physiology Post-Millennial Era: Challenges & Opportunities*". This platform is one of the suitable avenues to facilitate the exchange of scientific endeavors and enhance networking among scientists, academicians, clinicians, and postgraduate students involved in the fields of pharmacology and physiology, as well as multidisciplinary areas such as drug discovery, omics approaches, stem cells, and regenerative medicine. The primary objective of the special issue is to revitalize recent advances and breakthroughs in pharmacology and physiology to enhance opportunities and vanquish challenges in the post-millennial era, especially in the wake of the COVID-19 pandemic. In light of this, the special issue compiles the conference proceeding, to provide opportunities for knowledge integration of research and innovations.

Keywords: *Pharmacology; physiology; drug discovery; omics approaches; stem cells and regenerative medicine*

ORGANIZED BY:

Malaysian Society of Pharmacology and Physiology (MSPP)
Department of Human Anatomy, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia

LEAD GUEST EDITORS

Dr. Santhra Segaran Balan

Department of Diagnostic and Allied Health Science, Faculty of Health and Life Sciences, Management and Science University, University Drive, Off Persiaran Olahraga, Section 13, 40100 Shah Alam, Selangor, Malaysia.

Email: santhra@msu.edu.my

Professor Dr. Rusliza Basir

Department of Human Anatomy, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: rusliza@upm.edu.my

Associate Professor Dr. Yoke Keong Yong

Department of Human Anatomy, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: yoke_keong@upm.edu.my

GUEST EDITORS

Dr. Noor Aishah Mohammed Izham

Department of Diagnostic and Allied Health Science, Faculty of Health and Life Sciences, Management and Science University, University Drive, Off Persiaran Olahraga, Section 13, 40100 Shah Alam, Selangor, Malaysia.

Email: noor_aishah@msu.edu.my

Dr. Jasmine Siew Min Chia

School of Pharmacy, Management and Science University, University Drive, Off Persiaran Olahraga, Section 13, 40100 Shah Alam, Selangor, Malaysia.

Email: jasmine_chia@msu.edu.my

Dr. Fezah Othman

Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: fezah@upm.edu.my

Dr. Jonathan Chee Woei Lim

Department of Medicine, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: cheewoei@upm.edu.my

Dr. Siti Farah Md Tohid

Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: sitifarah@upm.edu.my

Associate Professor Dr. Hasnah Bahari

Department of Human Anatomy, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: haba@upm.edu.my

Associate Professor Dr. Tham Chau Ling

Department of Biomedical Sciences, Faculty of Medicine and Health Sciences, Universiti Putra Malaysia, 43400 Serdang, Selangor, Malaysia.

Email: chauling@upm.edu.my

ABSTRACTS

All presented abstracts are listed from Page 4 to 60.

Optimization of a two-dimensional electrophoresis protocol for plasma proteomic profiling of obese schizophrenia patients

Siti Norain Mat Rasid^{1,*}, Nour El Huda Abd Rahim¹, Norbaiyah Mohamed Bakrim¹, Norlelawati A. Talib², Mohd Asyraf Abdull Jalil¹, Mohd Yusri Idorus³ and Ahmad Nabil Rosli⁴

¹Department of Basic Medical Sciences, Kulliyah of Medicine, IIUM, Kuantan, 25300, Pahang.

²Department of Pathology and Laboratory Medicine, Kulliyah of Medicine, IIUM, Kuantan, 25300, Pahang.

³Institute of Medical Molecular Biotechnology, Faculty of Medicine, UTM, Sungai Buloh, 47000, Selangor.

⁴Department of Psychiatry, Kulliyah of Medicine, IIUM, Kuantan, 25300, Pahang.

*Correspondence:

Department of Basic Medical Sciences, Kulliyah of Medicine, International Islamic University Malaysia, Kuantan, 25300, Pahang, Malaysia.

Email: siti.norain1109@gmail.com

DOI <https://doi.org/10.28916/lsm.7.1.2023.132>

Abstract

The proteomic approach is particularly effective for studying the association between obesity and schizophrenia. It allows for a comprehensive analysis of the complete proteome, leading to substantial breakthroughs in biomarker discovery and drug development. Isoelectric focusing (IEF) and SDS-PAGE procedures are combined in the proteomic approach known as two-dimensional electrophoresis (2-DE), which separates proteins according to their isoelectric point and mass. This study aimed to investigate optimized conditions for the 2-DE technique by focusing on the selection of an immobilized pH gradient (IPG) strip. Protein extraction was performed on pooled plasma samples from 10 obese schizophrenia patients. The extracted protein samples were loaded onto two different pH (7 cm) IPG strips. The pH ranges between (i) 3 – 10 and (ii) 4 – 7. IEF was conducted following the PROTEAN IEF Cell System protocol, followed by SDS-PAGE. The resulting gels were stained with BioSafe Coomassie stain and washed with milliQ water. The stained gels were scanned, and the images were analyzed using PD Quest software. High-abundance proteins with a molecular weight range of 60 – 80 kDa were detected on both IPG strips. The results showed that using a pH 3 – 10 IPG strip, 245 protein spots were detected and distributed throughout the gel, with a notable concentration in the middle. Whereas using a pH 4 – 7 IPG strip resulted in the detection of 321 protein spots, indicating a higher quantity of protein spots with increased intensity. This is attributed to the improved fractionation of proteins resulting from the narrower and more focused pH range. Thus, it can be inferred that utilizing this pH range will yield optimal outcomes in protein separation and analysis. This study suggests selecting a pH 4 – 7 IPG strip is the recommended choice to achieve enhanced resolution and precise detection of protein spots in plasma samples from obese schizophrenia patients when employing the 2-DE method.

Keywords: *Proteomics; 2-Dimensional electrophoresis; IPG strip; protein separation*

Citation:

Balan, S. S., Basir, R., Yong, Y. K., Mohammed Izham, N. A., Chia, J. S. M., Othman, F., Lim, J. C. W., Md Tohid, S. F., Bahari, H., & Tham, C. L. (2023). 36th Malaysian Society of Pharmacology and Physiology Annual Scientific Meeting 2023. *Life Sciences, Medicine and Biomedicine*, 7(1).
<https://doi.org/10.28916/l SMB.7.1.2023.132>



Life Sciences, Medicine and Biomedicine
ISSN: 2600-7207

Copyright © 2023 by the Author(s). Life Sciences, Medicine and Biomedicine (ISSN: 2600-7207) Published by Biome Journals - Biome Scientia Sdn Bhd. Attribution 4.0 International (CC BY 4.0). This open access article is distributed based on the terms and conditions of the Creative Commons Attribution license
<https://creativecommons.org/licenses/by/4.0/>