



# The 4th Bioinformatics and Biodiversity Conference (BBC 2023)

Universitas Indonesia (Hybrid Conference)  
November 4-5 2023

# PROGRAM and ABSTRACT BOOK

*Exploring Biodiversity through Bioinformatics:  
Enhancing Our Understanding of the Biosphere  
and Environment*

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**\*Update per Day**



The 4th Bioinformatics and Biodiversity Conference (BBC)  
Hybrid (Lakarta, Indonesia) November 4-5, 2023

Main Room  
Hati Multidisciplinary Research Laboratory FMIPA UI - PT Pertamina

- BR 1-3 Online
- BR 4 R 202 (P) / 3rd floor of Multidisciplinary Research Laboratory FMIPA UI - PT Pertamina
- BR 5 R 502 (P) / 5th floor of Multidisciplinary Research Laboratory FMIPA UI - PT Pertamina
- BR 6 R 605 (P) / 6th floor of Multidisciplinary Research Laboratory FMIPA UI - PT Pertamina

Day/Date	Venue	Time (+7 GMT)		Duration	Program	PIC
		Start	End			
	Lobby - Hall	08.30	09.00	30'	Registration	Secretary
	Main Room - Hall	08.30	09.10	40'	Opening Ceremony by MC	MC: Dr. Yulia Mariana Teza Ayudita Putri
	Main Room - Hall	09.00	09.10	10'	Dring Honorary Rapsa together Praying	MC: Dr. Yulia Mariana Teza Ayudita Putri
	Main Room - Hall	09.10	09.15	5'	Mitigating Hospitality	K.S. UI
	Main Room - Hall	09.15	09.20	5'	Opening Remark by BBC 2023 Chair	Prof. Ahmad Budiman, S.S., M.Kom., Ph.D
	Main Room - Hall	09.20	09.25	5'	Opening Remark by MABBI Chair	Dr. Eng. Wini Ananta Kusuma, M.T
	Main Room - Hall	09.25	09.30	5'	Opening Ceremony - Virtual Greeting	Prof. Dedek Duhana, Ph.D
	Main Room - Hall	09.30	09.40	10'	Rantek Traditional Dance	MC: Dr. Yulia Mariana Teza Ayudita Putri
	Main Room - Hall	09.40	09.50	10'	Photo Session	MC: Dr. Yulia Mariana Teza Ayudita Putri
	Main Room - Hall	09.50	10.00	10'	Keynote Talk 1	Prof. Janna Supriatna, M.Sc., Ph.D
	Main Room - Hall	10.00	10.30	30'	Q&A	Session Chair: Dr. Eng. Wini Ananta Kusuma, M.T
	Main Room - Hall	10.30	10.40	10'	Coffee Break	
	Main Room - Hall	10.40	11.10	30'	Keynote Talk 2	Prof. Kevin Nadek, Ph.D
	Main Room - Hall	11.10	11.20	10'	Q&A	Session Chair: Husna Nugrahaprana, S.S., M.Sc., Ph.D
	Main Room - Hall	11.20	11.30	10'	Industry Talk - Seaweed	Seaweed
	Main Room - Hall	11.30	12.00	30'	Keynote Talk 3	Assoc. Prof. Babon Benyamin
	Main Room - Hall	12.00	12.10	10'	Q&A	Session Chair: Husna Nugrahaprana, S.S., M.Sc., Ph.D
	Main Room - Hall	12.10	12.15	5'	MC Break Announcement	MC: Dr. Yulia Mariana Teza Ayudita Putri
	Main Room - Hall	12.15	12.30	15'	VR Booth and Unmoderated Poster Session	Tech Admin (main hall and all BR)
	Main Room - Hall	12.30	13.00	30'	Lunch Break	
	Main Room - Hall	13.00	13.30	30'	Keynote Talk 4	Dr. Rosita Dindawati
	Main Room - Hall	13.30	13.40	10'	Q&A	Session Chair: Sahminia Sula Astuti, Ph.D
	Breakout Room	13.40	13.50	10'	Parallel session preparation	Online participant join to BR
	Breakout Room	13.50	17.10	150'	Oral parallel session (P) + Coffee break	Managerial control (oral speakers who are present are still here with their respective equipment in the committee room)

Day/Date	Venue	Time (+7 GMT)		Duration	Program	PIC
		Start	End			
Sat/Nov 4, 2023	Breakout Room	13.50	14.05	15'	Utilizing DNA Genetic Analysis on Hierarchical Clap from Jember, Bangkaya Belitung, Central Java, and East Java, Indonesia: A Phylogenetic Study	1
		14.05	14.20	15'	Enriched Analysis of Relationship Molecular Clones of Nite Trichia Varieties in Culturing, Using Next-Generation Sequencing	2
		14.20	14.35	15'	Genetic Diversity of Scomberomorus sp. from Java and Kalimantan based on Mitochondrial Cytochrome b Sub-Unit 1 Sequences	3
		14.35	14.50	15'	Identification of Gene Candidates in Mitochondrial Genomes of Curatoma 4 Using an mRNA Sequencing Approach	4
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		15.40	15.55	20'	The Potential of Staphylococcus pseudintermedius as a Bioindicator	8
		15.55	16.10	15'	Comparative Analysis of Tricollistin Using TMD, TROLLISTER and y-BTMAX on Three-Dimensional Hierarchical Data	9
		16.10	16.25	15'	Pharmacokinetic Insight into the Active Components of Stearopteryx adonidis (L.) R.M. King & H. Robinson Leaf using a Computational Approach	10
		16.25	16.40	15'	Regulation of Metastasis-Inducing Genes in Breast Cancer through Single-cell RNA Sequencing Analysis	11
		16.40	16.55	15'	Support Vector Machine (SVM) for Predicting Breast Cancer Severity	12

Day/Date	Venue	Time (+7 GMT)	Duration	Program	PIC
Sat/Nov 4, 2023	Virtual Poster Hall	08.30 - 18.00	450'	Open for all participant	Judge: Tech Admin
	Virtual Poster Hall	13.30 - 15.30	150'	Technical Admin Poster Hall and Judges	

Day/Date	Venue	Time (+7 GMT)		Duration	Program	PIC
		Start	End			
Sun/Nov 5, 2023	Main Room - Hall	08.30	09.00	30'	Registration	Secretary
		09.00	09.10	10'	Opening Ceremony by MC	MC: Abdulhadi Bramantyo, S.S.
		09.10	09.25	15'	Keynote Talk 5	Prof. Alhadi Bramantyo, S.S., M.Kom., Ph.D
		09.25	09.50	25'	Industry Talk - Kaboe Digital	Kaboe Digital
		09.50	10.00	10'	Q&A	Session Chair: Didi Huseino, Ph.D
		10.00	10.10	10'	Coffee Break	
		10.10	10.40	30'	Keynote Talk 6	Prof. Yoko Makita, Ph.D
		10.40	10.50	10'	Q&A	Session Chair: David Aguilawati, S.Kom., M.Sc., Ph.D
		10.50	11.20	30'	Keynote Talk 7	Ahmayyati, S.Kom., M.Sc., Ph.D
		11.20	11.30	10'	Q&A	Session Chair: David Aguilawati, S.Kom., M.Sc., Ph.D
		11.30	12.00	30'	Keynote Talk 8	Assoc. Prof. Dr. Triandri Sulandari
		12.00	12.10	10'	Q&A	Session Chair: Dr. Harshit Singh
12.10	12.15	5'	MC Break Announcement	MC: Abdulhadi Bramantyo, S.S.		
12.15	12.45	30'	Lunch - Break			
12.45	12.55	10'	Announcement (Self-Presenter oral (B topics and poster (1))	MC: Abdulhadi Bramantyo, S.S.		
12.55	13.00	5'	Closing Statements from BBC 2023 General Chair	Dr. Retno Listiyani, M.S.		
R 204 (2F)	13.00	16.00	180'	MABBI Gathering	Moderator: Didi Huseino, Ph.D	
R 204 (2F)	13.00	13.05	5'	Opening Ceremony	Moderator: Didi Huseino, Ph.D	
R 204 (2F)	13.05	13.25	20'	INPL Launching	Kholidi Azzah, Ph.D	
R 204 (2F)	13.25	13.35	10'	Briefing	MABBI Chair: Dr. Eng. Wini Ananta Kusuma, M.T	
R 204 (2F)	13.35	14.30	60'	Discussion: Each Task Force	MABBI Board	
R 204 (2F)	14.30	15.30	60'	Prenary Discussion	MABBI Board	
R 204 (2F)	15.30	15.45	15'	Conclusion	MABBI Board	
R 204 (2F)	15.45	15.60	15'	Closing Statements	MABBI Chair: Dr. Eng. Wini Ananta Kusuma, M.T	
R 204 (2F)	15.60	16.30	30'	After Event - Light Discussion	MABBI member	

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# Keynote Speakers



**Prof. Alhadi Bustamam,  
S.Si., M.Kom., Ph.D.**

Universitas Indonesia, Indonesia



**Prof. Jatna Supriatna, M.Sc.  
Ph.D.**

Universitas Indonesia, Indonesia



**Prof. Kenta Nakai, Ph.D.**

The University of Tokyo, Japan



**Prof. Yuko Makita, Ph.D.**

Maebashi Institute of Technology, Japan



**Assoc. Prof. Beben Benyamin**

University of South Australia, Australia



**Afiahayati, S.Kom., M.Cs,  
Ph.D.**

Universitas Gadjah Mada, Indonesia



**Assoc. Prof. Dr. Treenut  
Saithong**

KMUTT, Thailand



**Dr. Rosie Drinkwater**

Ludwig Maximilian University of Munich,  
Germany

# BR 5 | R 505 (5F)

## (Bioinformatics Solutions for Advancing Biomedical Science, Clinical Bioinformatics and Health Informatics)

13.50	14.05	Development of a Heart Rate Measurement System Based on Remote Photoplethysmography Techniques and Signal Processing Using Facial Video
		<b>Rinaldi Anwar</b>
14.05	14.20	Leprosy Early Detection Through Binary Segmentation Using ResU-Net
		<b>Andrew Jonathan Brahms S.</b>
14.20	14.35	SARS-CoV-2 variants genome analysis of Indonesian isolates and their responses to available vaccines
		<b>Kholis Abdurachim Audah</b>
14.35	14.50	Molecular Docking and Molecular Dynamic Simulation on the Interaction of Saffron's Active Compounds with p53 Protein
		<b>Sendi Handika Putra</b>
14.50	15.05	Study of Anti-Bacterial and Anti-Inflammatory Properties of Indonesia's Flora Biodiversity as Potential Drug Candidates for Diabetic Ulcer Through In Silico and In Vitro Approach
		<b>Cecilia Tiffanny</b>

# BR 5 | R 505 (5F)

## (Bioinformatics Solutions for Advancing Biomedical Science, Clinical Bioinformatics and Health Informatics)

15.25	15.40	Identification of Indonesian Ethnomedicinal Plants as Potential Drug Candidates for Acute Respiratory Infection Using Computer-Aided Drug Design and SIMRS Model
		<b>James Jackson</b>
15.40	15.55	ANALYSIS OF THE ABILITY OF THE NANO MIXTURE OF CORIANDER SEED ETHANOL EXTRACT ( <i>Coriandrum sativum</i> L.) AND ETHANOL EXTRACT OF PUTRI MALU LEAVES ( <i>Mimosa pudica</i> L) IN LOWERING BLOOD GLUCOSE LEVELS IN HYPERGL WISTAR RAT
		<b>Sri Wahjuni</b>
15.55	16.10	The Effect of Tilapia Skin and Pomegranate Flower Extract Hydrogel on the Expression of Angiogenic and Histopathological Factors in Second Degree Burn Injury
		<b>Jihan Nur Azizah</b>
16.10	16.25	Developing an Expert System for Web-Based Hepatitis Diagnosis Using the Certainty Factor Method
		<b>R. Rizal Isnanto</b>
16.25	16.40	The Effect of Sea Cucumber Extract as an Inhibitor of Adipogenesis Process in Mesenchymal Cells Derived from Umbilical Cord: Study on the Role of NF $\kappa$ B
		<b>Devana Alifia Afifah</b>
16.40	16.55	Metabolic Profile Analysis of Red Ginger ( <i>Zingiber officinale</i> var. <i>rubrum</i> Rosc.) by GC-MS and HPLC
		<b>Azminah</b>

## Metabolic Profile Analysis of Red Ginger (*Zingiber officinale* var. *rubrum* Rosc.) by GC-MS and HPLC

Azminah, Azminah<sup>1,\*</sup>; Kristina, Euodia Hana<sup>1</sup>; Salsabillah, Aisyah<sup>1</sup>; Yunita, Oeke<sup>1</sup>

<sup>1</sup>*Faculty of Pharmacy, University of Surabaya, Raya Kalirungkt Surabaya 60293, Indonesia*

Presenting Author: [azminah@staff.ubaya.ac.id](mailto:azminah@staff.ubaya.ac.id);

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This research aims to conduct a metabolic profile analysis of red ginger (*Zingiber officinale* var. *rubrum* Rosc.) using Gas Chromatography-Mass Spectrometry (GC-MS) and High-Performance Liquid Chromatography (HPLC). Red ginger is known to contain bioactive compounds that have the potential to provide significant health benefits. Red ginger samples were extracted using methanol as a solvent, and the compounds in the samples were analyzed using GC-MS and HPLC. The analysis results allowed for the identification of various metabolite compounds, including 6-gingerol, 8-gingerol 10-gingerol, 6-shogaol, and other related compounds. Additionally, this research also includes the relative quantification of these compounds in red ginger. The research results showed variations in the metabolic profile of red ginger, including significant differences in the types and amounts of detected compounds. This information can be valuable for a deeper understanding of the health potential and applications of red ginger in the pharmaceutical and nutraceutical fields. Furthermore, this research can serve as a basis for the development of cultivation and processing techniques for red ginger that can enhance the content of specific bioactive compounds. This research has important implications in supporting the development and utilization of red ginger as a source of natural compounds with the potential to benefit health and other industries.

**Keywords:** Red Ginger, GC-MS, HPLC, Metabolic Profile



## **Introduction**

In Indonesia, red ginger has been utilized for its antibacterial, antioxidant, and wound-healing properties [1]. The weekly demand for red ginger exports is about 4 tons [2]. Compared to other ginger types, red ginger (*Zingiber officinale* Roxb. var. *rubrum*, Zingiberaceae) has a distinctively stronger scent and hotter taste. Gingerol, a blend of several chemicals with antibacterial, anti-inflammatory, and antioxidant properties, is one of the primary phenolic components in ginger. The primary component of fresh ginger, 6-gingerol, has a variety of pharmacological actions in addition to 8-gingerol and 10-gingerol [3]. Rhizomes had a significantly larger gingerol content (104.39 µg/g) than stems (0.84 µg/g) and leaves (4.13 µg/g) [4]. According to Chumroenphat et al. (2018), peeled ginger rhizomes have 68.15 mg/100 g fresh weight of 6-gingerol, whereas fresh ginger rhizomes have the maximum concentration (75.25 mg/100 g fresh weight) [5].

The purpose of this study is to use High-Performance Liquid Chromatography (HPLC) and Gas Chromatography-Mass Spectrometry (GC-MS) to analyze the metabolic profile of red ginger (*Zingiber officinale* var. *rubrum* Rosc.).

## **Research Method**

### **GC-MS analysis**

Gas chromatography system used, GC-MS Brand:- SHIMADZU-AOC-20i, Auto Injector Model: QP-2010 SE, Column Length 60.0 m, Column Diameter 0.25 mm, Column Flow 0.4 mL/min, Column pressure 100 kPa. Oven temp 250°C, Interface temp 200°C, Low vacuum 8.9e+000 Pa, Ion Source voltage 70 V.

### **HPLC analysis**

Gradient elution was used for instrumental analysis in the following circumstances: The detection was performed at a wavelength of 280 nm while the system pressure was kept at 1000 Pa. Throughout the analysis, the flow rate was maintained at 1.2 mL/min. 0.1% trifluoroacetic acid (TFA) in water (A) and acetonitrile (B) made up the mobile phase. Twenty microliters was the injection volume. At 280 nm, UV detection was used to find the analytes. After gathering chromatographic data, peak analysis was carried out using peak area and retention time.

## Results and Discussion

The number of compounds contained in the test sample can be seen using the chromatogram in Figure 1, that is produced as a consequence of the GCMS analysis. More than 50 red ginger compounds were found to be prevalent in all test samples, according to the study of the chromatogram for each sample. The analysis's findings revealed that the terpenoid group contains a number of compounds, including monoterpenes, diterpenes, and sesquiterpenes, with sesquiterpene components being the most prevalent.

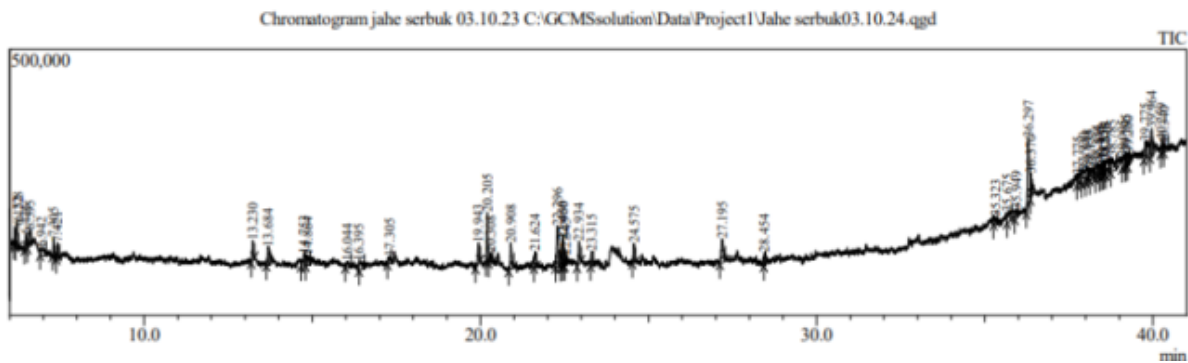


Figure 1. Chromatogram results of red ginger by GC-MS

The findings of this study are consistent with research [6] that yields a metabolite profile with the same primary component, namely the compounds  $\beta$ -Curcumene (15.88%),  $\beta$ -Sesquiphellandrene (15.57%),  $\beta$ -bisabolene (9.29%), and zingiberene (31.79%). The study's findings were reinforced by research findings [7] that indicate there is the molecules zingiberene,  $\beta$ -curcumene,  $\beta$ -sesquiphellandrene, and  $\beta$ -bisabolene are the six biggest components in their research.

Standard preparation was in accordance with previous research [8]. Conditions for gingerol analysis methods in culture of the red ginger used refers to the literature: 6-Gingerol  $\pm$  9.5 minutes, 8-Gingerol  $\pm$  20.0 minutes, 6-Shogaol  $\pm$  22.4 minutes, 10-Gingerol  $\pm$  33.8 minutes. The standard results for samples from ginger root 3 ppm, show Peak 2, 6-Gingerol  $\pm$  12.74 minutes, Peak 4, 8-Gingerol  $\pm$  20.0 minutes, Peak 5, 6-Shogaol  $\pm$  21.93 minutes, Peak 6, 10-Gingerol  $\pm$  36.4 minutes, as seen in Figure 2.

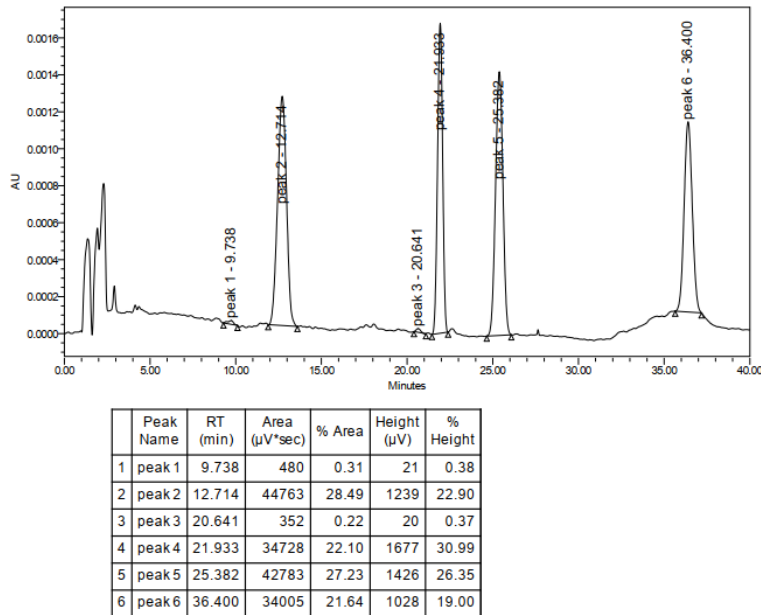


Figure 2. Results of analysis of red ginger root samples

## Conclusion

Furthermore, this research can serve as a basis for the development of cultivation and processing techniques for red ginger that can enhance the content of specific bioactive compounds. This research has important implications in supporting the development and utilization of red ginger as a source of natural compounds with the potential to benefit health and other industries.

## Reference

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