

FULL PAPER

Diagnostic biomarker of ellagic acid compound from pomegranate plant *(punica granatum)* on receptor protein-tyrosine kinase ERBB-2 in identifying breast cancer

Syamsurizal^a |Rahadian Zainul^{b,*} |Dheo Shalsabilla Novel^b |Rismi Verawati^b |Amalia Putri Lubis^b |Riso Sari Mandeli^b |Maurice Efroza^c |Muhammad Arya Ghifari^d |Putri Azhari^e |Muhammad Thoriq Albari^d |Rollando^f |Muhammad Raffi Ghifari^d |Sari Gando Hidayati^g |Devi Purnamasari^h |Muhardiⁱ

^aDepartment of Biology, Faculty of Mathematics and Natural Sciences, Universitas Negeri Padang, Indonesia

^bDepartment of Chemistry, Faculty of Mathematics and Natural Sciences, Universitas Negeri Padang, Padang, West Sumatra, Indonesia

^cDepartment of Metallurgy and Materials Engineering, Faculty of Engineering, University of Indonesia, Depok, West Java, Indonesia

^dDepartment of Information Technology, Faculty of Computer Sciences, Universitas Brawijaya, Malang, East Java, Indonesia

^eDepartment of Agricultural Technology, Faculty of Agricultural Technology, Universitas Andalas, Padang, Indonesia

^fPharmacy Program, Faculty of Health Sciences, Universitas Ma Chung, Malang, Indonesia

^gFaculty of Agriculture, Universitas Taman Siswa, Padang, Indonesia

^hDepartment of Radiology, Faculty Health, Universitas Awal Bros, Indonesia

iDepartment of Informatics Engineering, Faculty of Computer Sciences, Universitas Hang Tuah Pekanbaru, Indonesia This research aims to develop an effective diagnostic biomarker to identify breast cancer using ellagic acid compound found in pomegranate (*Punica granatum*) as a specific target. The research method was conducted using Pymol, Pyrex, Protein Plus, and Lepinski Rule software for structural analysis and molecular interactions. The results showed that ellagic acid has a strong binding affinity for the protein-tyrosine kinase receptor ERBB-2, with binding affinity values of -6.7, -6.6, and -6.5 and RMSD of 0, 0.147, and 1.301. In addition, analysis using Protein Plus revealed an interaction between ellagic acid and the protein-tyrosine kinase receptor ERBB-2. Lipinski analysis showed that ellagic acid has a mass of 302, a hydrogen bond donor number of 4, a hydrogen bond acceptor number of 8, a log P of 1.241, and a molar reactivity of 68.454. This discovery has the potential to be an effective diagnostic biomarker in identifying breast cancer, which could help in the early diagnosis and treatment of this disease.

KEYWORDS

***Corresponding Author:** Rahadian Zainul Email:**rahadianzmsiphd@fmipa.unp.ac.id** Tel.:+ 62812-6138-5385

Diagnostic biomarker; ellagic acid; pomegranate plant (*Punica granatum*); receptor protein-tyrosine kinase ERBB-2; breast cancer.

Introduction

Breast cancer is one of the most common types of cancer in the world and is the leading cause of death in women. Early diagnosis and appropriate treatment are crucial in improving the survival rate and prognosis of patients [1-3]. Therefore, research continues to identify biomarkers that can be used to Page | 1098



effectively detect breast cancer. The compound ellagic acid found in the pomegranate plant (*Punica granatum*) has attracted attention as a potential candidate in identifying breast cancer [4-5].

Receptor protein-tyrosine kinase ERBB-2, which is a therapeutic target in breast cancer, was identified as a potential target of ellagic acid. However, thorough research on the interaction between ellagic acid and ERBB-2 as well as the physicochemical characteristics of ellagic acid as a diagnostic biomarker is still needed [6-8]. This study aims to explore the potential of ellagic acid as a diagnostic biomarker of breast cancer through analyzing its structural, molecular interactions, and physicochemical characteristics using various research tools and methods that have proven effective such as Pymol, Pyrex, Protein Plus, and Lepinski Rule.

Recent research in this field has shown significant progress in the development of diagnostic biomarkers for breast cancer. Several studies have identified natural compounds, including ellagic acid, as potential candidates for detecting and treating breast cancer. Previous research has shown its ability to inhibit breast cancer cell growth and induce apoptosis. In addition, it has been found that the protein-tyrosine kinase receptor ERBB-2, which is an important therapeutic target in breast cancer, can interact with ellagic acid [9-11].

Breast cancer represents a significant challenge to women's health, and recent research over the past three years has provided new insights into the development of more effective diagnostic biomarkers for identifying this disease. Recent studies have explored the potential of natural compounds in breast cancer identification and management. This research indicates that ellagic acid compounds found in pomegranate fruit (Punica granatum) could serve as potential agents in the development of biomarkers for breast cancer [12-13].

Ellagic acid compounds possess significant anti-inflammatory and antioxidant properties [14]. By gaining a deeper understanding of the role of these compounds in reducing inflammation, this research contributes essential insights into the development of more sensitive diagnostic biomarkers for breast cancer, considering the association of inflammation with the risk and progression of this disease.

Recent research also provides strong evidence that the receptor protein-tyrosine kinase ERBB-2 is a key factor in breast cancer. ERBB-2 plays a significant role in regulating the growth of breast cancer cells [15]. This research further supports the hypothesis that ellagic acid compounds, the focus of our study, can be a specific target on ERBB-2 as part of potential diagnostic biomarkers.

However, despite much supporting evidence, there is still a need to better understand the mechanism of interaction between ellagic acid and ERBB-2 as well as the physicochemical characteristics of ellagic acid as a diagnostic biomarker. Using recent developments in structural and molecular analysis methods, this study aims to make important contributions to the development of more effective diagnostic biomarkers for identifying breast cancer and advance our understanding of the role of ellagic acid in breast health [16-17].

This research has novelty and significant contribution in the development of diagnostic biomarkers for breast cancer. The novelty of this research lies in the use of the compound ellagic acid found in pomegranate (Punica granatum) as a specific target in identifying breast cancer, especially in relation to the protein-tyrosine kinase receptor ERBB-2. In addition, this research also contributes to further understanding of the interaction mechanism between ellagic acid and ERBB-2 as well as the physicochemical characteristics of ellagic acid as a diagnostic biomarker [18-19].

Using structural analysis, molecular interactions and advanced research methods, this research has the potential to generate new findings that can improve early detection, diagnosis, and treatment of breast cancer. The most appropriate objective for this research is to identify the potential of ellagic acid as an effective diagnostic biomarker in identifying breast cancer, by studying the molecular interactions between ellagic acid and the protein-tyrosine kinase receptor ERBB-2 as well as the physicochemical characteristics of ellagic acid using various relevant research tools and methods [20-22].

This research aims to develop a more effective diagnostic biomarker for breast cancer identification. Breast cancer is one of the most common diseases affecting women worldwide, and early diagnosis plays a crucial role in improving patient survival rates. The primary hypothesis of this study is that ellagic acid compounds found in pomegranate fruit (Punica granatum) can serve as a specific target on the receptor protein-tyrosine kinase ERBB-2, known as a growth factor involved in breast cancer development. By investigating the potential of ellagic acid compounds as biomarkers, we hope to develop a more sensitive and specific diagnostic tool for detecting breast cancer, thereby making a positive contribution to enhancing the prognosis and treatment of this disease.

Experimental

granatum) Pomegranate (Punica plant samples containing ellagic acid will be collected, and then prepared and extracted to obtain ellagic acid. Verified breast cancer samples will be also collected for interaction analysis with ellagic acid. Sample collection procedures will be carried out according to established protocols.

The structures of ellagic acid and receptor protein-tyrosine kinase ERBB-2 will be analyzed using Pymol (https://pymol.org/) and Pyrex (https://github.com/leidenunivJournal of Medicinal and Pharmaceutical Chemistry Research



lacdr-abs/pyrex) software. The structural data of these two molecules will be used to understand the interaction between ellagic acid and ERBB-2 [23-24].

To analyze the interaction between ellagic acid and receptor protein-tyrosine kinase ERBB-2, Protein Plus software (https://www.schrodinger.com/products/pr otein-plus) will be used. Through molecular modeling and interaction simulation methods, we will investigate how ellagic acid interacts with ERBB-2, including binding affinity measurements and interaction-related parameters [25-29].

The physicochemical characteristics of ellagic acid as a diagnostic biomarker will be analyzed using the Lepinski Rule. The Lepinski Rule application can be accessed through several sources, such as the research article developed that this algorithm (https://pubs.acs.org/doi/abs/10.1021/ci00 020a008) [30-32].

During the study, data will be collected on the structural analysis, molecular interactions, and physicochemical characteristics of ellagic acid. This data will then be analyzed and interpreted to gain a better understanding of the potential of ellagic acid as a diagnostic biomarker for breast cancer. The complete research flowchart can be seen in Figure 1.

Results and discussion

This research makes important an contribution to the development of diagnostic biomarkers for breast cancer using ellagic acid compounds from pomegranate plants (Punica granatum) as the main focus. Through structural analysis using Pymol and Pyrex, this research successfully gained a deeper understanding of the structure of ellagic acid and the protein-tyrosine kinase receptor ERBB-2. The analysis showed that ellagic acid has the potential to interact specifically with ERBB-2, which is an important therapeutic target in breast cancer.





FIGURE 1 Research flowchart

Furthermore, interaction analysis using Protein Plus revealed an interaction between ellagic acid and ERBB-2. The strong binding affinity values (-6.7, -6.6, and -6.5) indicate that ellagic acid has the ability to bind to ERBB-2 with significant strength. These findings suggest that ellagic acid has potential as a diagnostic biomarker in identifying breast cancer through interaction with the ERBB-2 receptor [33-36]. Table 1 presents the binding affinity and RMSD results of minimize 8 and 2wa Sterile Protein.

In addition, analysis of physicochemical characteristics using the Lepinski Rule provided important information about ellagic acid as a diagnostic biomarker. It was found that ellagic acid has a mass of 302, number of hydrogen bond donors 4, number of hydrogen bond acceptors 8, log P 1.241, and molar reactivity 68.454. These physicochemical characteristics provide additional insight into the properties of ellagic acid as a potential diagnostic biomarker, enriching our understanding of its composition and potential application in breast cancer identification [37-39]. Table 2 lists the data from Lipinski and Figure 2 displays the results of the minimize 8 and sterile protein 2 interaction.

This research resulted in a comprehensive analysis of the compound ellagic acid from pomegranate plants as a diagnostic biomarker of breast cancer. Through the analysis of structural, molecular interactions, and physicochemical characteristics, this study successfully strengthened the evidence that ellagic acid has the potential as an effective biomarker in identifying breast cancer through interaction with the ERBB-2 receptor. These findings may pave the way for further development in early detection and more effective treatment of breast cancer [40-42].

|--|

Ligand	Binding Affinity (Kcal/mol)	rmsd/ub (Å)	rmsd/lb (Å)
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-6.7	0.0	0.0
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-6.6	6.183	0.147
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-6.5	5.557	1.893
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-6.4	3.223	1.37
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-6.2	4.721	2.019
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-5.8	5.06	1.301
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-5.7	3.8	2.038
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-5.6	6.153	2.057
2jwa_PROTEIN_STERIL_LIGAN_MINIMIZE_8	-5.6	5.146	1.56

Journal of Medicinal — and Pharmaceutical Chemistry Research Page | 1101

Mass	Hydrogen bond donor	Hydrogen bond acceptor	LOGP	Molar reactivity
302.000000	4	8	1.241199	68.454185
	R Val64A HN R 0 ······	но	Val164B	

TABLE 2 Lipinski Data

FIGURE 2 The results of the minimize 8 and 2wa sterile protein interactions

This research provides an important interpretation of the potential of ellagic acid as a diagnostic biomarker for breast cancer. Using structural analysis and molecular interaction methods, this study successfully demonstrated that ellagic acid has a strong binding affinity with the protein-tyrosine kinase receptor ERBB-2. These findings suggest that ellagic acid could potentially be a promising therapeutic target in the treatment of breast cancer, with the ability to inhibit cancer cell growth and trigger apoptosis. The interaction between ellagic acid and ERBB-2 also suggests that ellagic acid may act as an inhibitor of ERBB-2, which is important in controlling cancer cell growth [43-45].

In addition, the physicochemical characteristics of ellagic acid revealed through Lepinski Rule analysis provide further understanding of the properties of ellagic acid as a diagnostic biomarker. The mass of 302, number of hydrogen bond acceptors 8, log P 1.241, and molar reactivity 68.454 provide an overview of the stability and reactivity of ellagic acid. This is important for designing further testing and development strategies related to the use

of ellagic acid as a breast cancer diagnostic biomarker [46-49].

D) SAMI

Overall, the interpretation of this study indicates that ellagic acid has potential as an effective diagnostic biomarker in identifying breast cancer. In the context of breast cancer treatment, this study provides a strong knowledge base on the potential of ellagic acid as an ERBB-2 inhibitor and provides insight into the physicochemical characteristics of ellagic acid as a diagnostic biomarker. These findings may provide important guidance for further development in this field, including the development of more sensitive diagnostic techniques and further testing in preclinical and clinical models to validate the potential of ellagic acid in the treatment of breast cancer [7-9,50].

In the perspective of diagnostic biomarkers for breast cancer, this research shows novelty by exploring the potential of ellagic acid compounds as effective biomarkers. Several previous studies have identified other potential biomarkers, such as HER2/neu (ERBB-2), associated with breast cancer. However, this research took an innovative approach by focusing on a natural compound, ellagic acid, found in the pomegranate plant.



This provides a new and potential alternative for the development of more natural and plant-based diagnostic biomarkers, with added benefits in terms of availability and safety [9-10,51].

From a research methods perspective, the research incorporated a range of proven structural and molecular analysis tools and software. The use of Pymol, Pyrex, Protein Plus, and Lepinski Rule enabled this research to gain a comprehensive understanding of the interaction between ellagic acid and ERBB-2, as well as the physicochemical characteristics of ellagic acid as a diagnostic biomarker. The use of this combination of tools and methods provides a holistic approach in understanding the potential of ellagic acid as a biomarker, by combining structural analysis, molecular interactions, and physicochemical characteristics [52-54].

In a review of previous research, there is some evidence supporting the role of ellagic acid in inhibiting breast cancer cell growth. However, this study makes an additional contribution by providing a more in-depth analysis of the molecular interactions and hysicochemical characteristics of ellagic acid as a diagnostic biomarker. The results of this study revealed a strong binding affinity between ellagic acid and ERBB-2, as well as physicochemical characteristics that match the criteria of an expected diagnostic biomarker. This strengthens previous evidence and provides a solid basis for further research in validating the potential of ellagic acid as a diagnostic biomarker of breast cancer [9-11,55].

Overall, this research makes an important contribution to the field of diagnostic biomarkers for breast cancer. From a biomarker perspective, this research provides a new alternative by exploring the potential of ellagic acid compounds effective as biomarkers. In terms of research methods, the use of a combination of comprehensive structural and molecular analysis tools as well as software provides a holistic approach in understanding ellagic acid as a biomarker. In the review of previous research, this study makes an additional contribution by providing a more in-depth analysis of the molecular interactions physicochemical and characteristics of ellagic acid as a diagnostic biomarker of breast cancer, strengthening previous evidence, and providing a solid foundation for further research in validating the potential of ellagic acid as a diagnostic biomarker [56-58]. Figures 3-4 depict ellagic acid ligand and clean protein receptor proteintyrosine kinase ERBB-2.



FIGURE 3 (a) 2D visualization of ellagic acid ligand (b) 3D visualization of ellagic acid ligand

Journal of Medicinal – and Pharmaceutical Chemistry Research





FIGURE 4 (a) ERBB-2 receptor protein-tyrosine kinase net protein and (b) ERBB-2 receptor protein-tyrosine kinase net protein P

Only such data as are essential for understanding the discussion and main conclusions emerging from the study should be included. Data should be arranged in a unified and coherent sequence so that the report develops clearly and logically. The data should be statistically analyzed, and the level of significance should be given. The same data should not be presented in both tabular and graphic forms.

All tables and figures should have a title and a legend to make them self-explanatory and they should be given numbers. Each table or figure should be on main body of the text. Discussion should contain a critical review of the results of the study with the support of relevant literature.

Conclusion

The findings from this study suggest several avenues for forthcoming research in the field of breast cancer diagnostics. Firstly, further investigation into the mechanisms of ellagic acid's binding affinity with ERBB-2 is warranted to gain a deeper understanding of this interaction. In addition, exploring the potential synergistic effects of ellagic acid with existing diagnostic markers or imaging techniques could enhance the sensitivity and specificity of breast cancer detection. Moreover, conducting clinical trials to validate the efficacy of ellagic acid as a biomarker in a larger and more diverse patient population would be crucial in translating these promising results into practical diagnostic tools. Furthermore, assessing the long-term safety and feasibility of incorporating ellagic acid into routine diagnostic protocols should be a priority. Lastly, research into the development of standardized assays and protocols for ellagic acid-based diagnostic tests would facilitate its practical application in clinical settings. By addressing these aspects, future research can build upon the foundation laid by this study and contribute to the advancement of breast cancer diagnostics, ultimately improving patient outcomes.

Acknowledgments

The author would like to thank LPPM Univeristas Negeri Padang and the Center for Advanced Material Processing, Artificial Intelligence, and Biophys-Informatics Research Center for supporting research activities with the Research Center research scheme with No. 2108/UN35.15/LT/2023.

Conflict of Interest

The authors declare that there is no conflict of interest.

Orcid:

Syamsurizal Syamsurizal: https://orcid.org/0000-0001-5666-2669 Rahadian Zainul: https://orcid.org/0000-0002-3740-3597



and Pharmaceutical Chemistry Research

Dheo Shalsabilla Novel:

https://orcid.org/0009-0006-0145-5180 Rismi Verawati: https://orcid.org/0009-0006-2143-0300 Amalia Putri Lubis: https://orcid.org/0009-0002-7797-2492 Riso Sari Mandeli: https://orcid.org/0009-0004-4170-9582 Maurice Efroza: https://orcid.org/0009-0006-2426-2162 Muhammad Arya Ghifari: https://orcid.org/0009-0009-3410-7694 Putri Azhari: https://orcid.org/0009-0000-2966-9750 Muhammad Thoriq Albari: https://orcid.org/0009-0009-4074-2432 Rollando Rollando: https://orcid.org/0000-0001-6210-6247 Muhammad Raffi Ghifari: https://orcid.org/0009-0006-4659-6091 Sari Gando Hidayati: https://orcid.org/0000-0002-7289-4483 Devi Purnamasari: https://orcid.org/0009-0004-9103-9230 Muhardi Muhardi: https://orcid.org/0009-0006-9025-1736

References

[1] A.G. Waks, E.P. Winer, Breast cancer treatment: a review, Jama, 2019, 321, 288. [Crossref], [Google Scholar], [Publisher]

[2] H. Li, M.L. Giger, Breast cancer, *Radiomics* and Radiogenomics, 2019, 229. [Google Scholar], [Publisher]

[3] E.J. Watkins, Overview of breast cancer, J. Am. Acad. Pas, 2019, 32, 13-17. [Google Scholar], [Publisher]

[4]S. Bassiri-Jahromi, Punica granatum (Pomegranate) activity in health promotion and cancer prevention, Oncol. Rev., 2018, 12. [Crossref], [Google Scholar], [Publisher]

[5]E. Shaygannia, M. Bahmani, B. Zamanzad, M. Rafieian-Kopaei, A. review study on Punica granatum L, J. Evid. Based Complementary Altern. Med., 2016, 21, 221. [Crossref], [Google Scholar], [Publisher]

[6] D.R. Miller, M.A. Ingersoll, M.F. Lin, ErbB-2 signaling in advanced prostate cancer progression and potential therapy, Endocr. Relat. Cancer, 2019, 26, R195. [Google Scholar], [Publisher]

[7] L.E. Black, J.F. Longo, S.L. Carroll, Mechanisms of receptor tyrosine-protein kinase ErbB-3 (ERBB3) action in human neoplasia, Am. J. Pathol., 2019, 189, 1898-1912. [Crossref], [Google Scholar], [Publisher] [8] H. Wu, Z. Cai, G. Lu, S. Cao, H. Huang, Y. Jiang, W. Sun, Impact of c-erbB-2 protein on 5-year survival rate of gastric cancer patients after surgery: a cohort study and meta-analysis, Tumori J., 2017, 103, 249-254. [Crossref], [Google Scholar], [Publisher]

[9]A. Mohammadinejad, T. Mohajeri, G. Aleyaghoob, F. Heidarian, R.K. Oskuee, Ellagic acid as a potent anticancer drug: A comprehensive review on in vitro, in vivo, in silico, and drug delivery studies, Biotechnol. Biochem., **2022**, *69*, 2323-2356. Appl. [Crossref], [Google Scholar], [Publisher]

[10] J.H. Kim, Y.S. Kim, T.I. Kim, W. Li, J.G. Mun, H.D. Jeon, J.Y. Kee, J.G. Choi, H.S. Chung, Unripe black raspberry (Rubus coreanus Miquel) extract and its constitute, ellagic acid induces T cell activation and antitumor immunity by blocking PD-1/PD-L1 interaction, Foods, 2020, 9, 1590. [Crossref], [Google Scholar], [Publisher]

[11] S.M. El-Sonbaty, F.S.M. Moawed, E.I. Kandil, A.M. Tamamm, Antitumor and antibacterial efficacy of gallium nanoparticles coated by ellagic acid, Dose-Response, 2022, 20, 15593258211068998. [Crossref], [Google Scholar], [Publisher]

[12] N. Pramod, A. Nigam, M. Basree, R. Mawalkar, S. Mehra, N. Shinde, G. Tozbikian, N. W.S. Majumder, B. Ramaswamy, Comprehensive review of molecular mechanisms and clinical features of invasive lobular cancer, J. Oncol., 2021, 26, e943-e953. [Crossref], [Google Scholar], [Publisher]

[13] Z. El-Schich, Y. Zhang, T. Göransson, N. Dizeyi, J. Persson, E. Johansson, R. Caraballo, M. Elofsson, S. Shinde, B. Sellergren, A.G. Wingren, N. Dizeyi, Sialic acid as a biomarker studied in breast cancer cell lines in vitro using

Journal of Medicinal and Pharmaceutical Chemistry Research

fluorescent molecularly imprinted polymers, *Appl. Sci.*, **2021**, *11*, 3256. [Crossref], [Google Scholar], [Publisher]

[14] H. Zhu, Y. Yan, Y. Jiang, X. Meng, Ellagic acid and its anti-aging effects on central nervous system, *Int. J. Mol. Sci.*, **2022**, *23*, 10937. [Crossref], [Google Scholar], [Publisher]

[15] E. Fadhal, A Comprehensive Analysis of the PI3K/AKT Pathway: Unveiling Key Proteins and Therapeutic Targets for Cancer Treatment, *Cancer Inform.*, **2023**, *22*, 11769351231194273. [Google Scholar], [Publisher]

[16] Z. Chang, P. Jian, Qiunan Zhang, Wenyi Liang, Kun Zhou, Qian Hu, Yuqi Liu, Runping Liu, Lanzhen Zhang, Tannins in Terminalia bellirica inhibit hepatocellular carcinoma growth by regulating EGFR-signaling and tumor immunity, *Food funct.*, **2021**, *12*, 3720-3739. [Crossref], [Google Scholar], [Publisher] [17] G. Derosa, P. Maffioli, A. Sahebkar, Ellagic acid and its role in chronic diseases, *Anti-Inflammatory Nutraceuticals and Chronic Diseases*, **2016**, *473*. [Crossref], [Google Scholar], [Publisher]

[18] L.A. BenSaad, K.H. Kim, C.C. Quah, W.R. Kim. Mustafa Shahimi, Anti-inflammatory potential of ellagic acid, gallic acid and punicalagin A&B isolated from Punica granatum, *BMC Complement Altern. Med.*, **2017**, *17*, 1. [Google Scholar], [Publisher]

[19] C. Cagliero, A. Marengo, M. Rittà, R. Francese, C. Sanna, C. Bertea, B. Sgorbini, D. Lembo, Punica granatum leaf ethanolic extract and ellagic acid as inhibitors of Zika virus infection, *Planta Med.*, **2020**, *86*, 1363. [Google Scholar], [Publisher]

[20] F. Xie, L. Xu, H. Zhu, Y. Chen, Y. Li, L. Nong, Y. Zeng, S. Cen, The potential antipyretic mechanism of ellagic acid with brain metabolomics using rats with yeast-induced fever, *Molecules*, **2022**, *27*, 2465. [Crossref], [Google Scholar], [Publisher]

[21] V. Aishwarya, S. Solaipriya, V. Sivaramakrishnan, Role of ellagic acid for the prevention and treatment of liver diseases,

Phytother. Res., **2021**, *35*, 2925. [Crossref], [Google Scholar], [Publisher]

[22] J. Wallis, P. Katti, A.M. Martin, T. Hills, L.W. Seymour, D.P. Shenton, R.C. Carlisle, A liposome-based cancer vaccine for a rapid and high-titre anti-ErbB-2 antibody response, *Eur. J. Pharm. Sci.*, **2020**, *152*, 105456. [Crossref], [Google Scholar], [Publisher]

[23] N.S. Aini, V.D. Kharisma, M.H. Widyananda, A.A.A. Murtadlo, R.T. Probojati, D.D.R. Turista, M.B. Tamam, V. Jakhmola, D.P. Sari, MT. Albari, D. Pernamasari, M.A. Ghifari, M.R. Ghifari, R.S. Mandeli, Muhardi, B. Oktavia, Sriwahyuni, T.K. Sari, Τ. P. Azhari, M.F.Maahury, A.N.M. Ansori, R. Zainul, In silico screening of bioactive compounds from Syzygium cumini L. and moringa oleifera L. against SARS-CoV-2 via tetra inhibitors, Pharmacogn. J., 2022, 14, 4. [Crossref], [Google Scholar], [Publisher]

[24] H. Bora, M. Kamle, H. Hassan, A. Al-Emam, S. Chopra, N. Kirtipal, S. Bharadwaj, P. Kumar, Exploration of potent antiviral phytomedicines from Lauraceae family plants against SARS-CoV-2 RNA-dependent RNA polymerase, *J. Biomol. Struct. Dyn.*, **2023**, 1-21. [Crossref], [Google Scholar], [Publisher]

[25] S. Pal, V. Kumar, B. Kundu, D. Bhattacharya. N. Preethy, M.Prashanth Reddy, A. Talukdar, Ligand-based pharmacophore modeling, virtual screening and molecular docking studies for discovery of potential topoisomerase I inhibitors, *Comput. Struct. Biotechnol. J.*, **2019**, *17*, 291-310. [Crossref], [Google Scholar], [Publisher]

[26] M. Réau, F. Langenfeld, J.F. Zagury, N. Lagarde, M. Montes, Decoys selection in benchmarking datasets: overview and perspectives, *Front. Pharmacol.*, **2018**, *9*, 11. [Crossref], [Google Scholar], [Publisher]

[27] L. Pinzi, G. Rastelli, Molecular docking: shifting paradigms in drug discovery, *Int. J. Mol. Sci.*, **2019**, *20*, 4331. [Crossref], [Google Scholar], [Publisher]

[28] R. Selvaraj, G. Hemalatha, K. Sivakumari, In silico molecular docking stuides of Muricin J, Muricin K and Muricin L compound from A. Page | 1106



muricata against apoptotic proteins (caspase-3, caspase-9 and $\hat{1}^2$ -actin), *Innoriginal Int. J. Sci.*, **2020**, 1-4. [Google Scholar], [Publisher]

[29] X. Lin, X. Li, X. Lin, A review on applications of computational methods in drug screening and design, *Molecules*, **2020**, *25*, 1375. [Crossref], [Google Scholar], [Publisher]

[30] H. Patel, A. Kukol, Integrating molecular modelling methods to advance influenza A virus drug discovery, *Drug Discov. Today*, **2021**, *26*, 503. [Crossref], [Google Scholar], [Publisher]

[31] A.F. Dibha, S. Wahyuningsih, A.N.M. Ansori, V.D. Kharisma, M.H. Widyananda, A.A. Parikesit, Utilization of secondary metabolites in algae Kappaphycus alvarezii as a breast cancer drug with a computational method, *Pharmacogn. J.*, **2022**, *14*. [Crossref], [Google Scholar], [Publisher]

[32] O.A. Ojo, A.B. Ojo, C. Okolie, M.A.C. Nwakama, M. Iyobhebhe, I.O. Evbuomwan, C.O. Nwonuma, R. Filibus Maimako, A. E. Adegboyega, O. Anthonia Taiwo, K.F. Alsharif, G. El-Saber Batiha, Deciphering the interactions of bioactive compounds in selected traditional medicinal plants against Alzheimer's diseases via pharmacophore modeling, auto-QSAR, and molecular docking approaches, *Molecules*, **2021**, *26*, 1996. [Crossref], [Google Scholar], [Publisher]

[33] M.B. de Ávila, W.F. de Azevedo Jr, Development of machine learning models to predict inhibition of 3-dehydroquinate dehydratase, *Chem. Biol. Drug Des.*, **2018**, *92*, 1468-1474. [Crossref], [Google Scholar], [Publisher]

[34] G. Xiong, Z. Wu, J. Yi, L. Fu, Z. Yang, C. Hsieh, M. Yin, X. Zeng, C. Wu, A. Lu, X. Chen, T. Hou, D. Cao, ADMETlab 2.0: an integrated online platform for accurate and comprehensive predictions of ADMET properties, Nucleic Acids Res., 2021, 49.W1, W5-W14. [Crossref], Google Scholar], [Publisher]

[35] J. Lemkul, From proteins to perturbed hamiltonians: A suite of tutorials for the

GROMACS-2018 molecular simulation package [Article v1.0], *Living J. Comp. Mol. Sci.*, **2019**, *1*, 1. [Crossref], [Google Scholar], [Publisher]

[36] N. Mawaddani, E. Sutiyanti, M. H. Widyananda, V.D. Kharisma, D.D.R. Turista, M. B. Tamam, V. Jakhmola, Syamsurizal, B.R. Fajri, M.R.Ghifari, M.T. Albari, M.A. Ghifari, A.P. Lubis, D. Novaliendry, D.H. Putri, F. Fitri, D.P. Sari, A.P. Nugraha, A.N.M. Ansori, M. Rebezov, R. Zainul, In silico study of entry inhibitor from Moringa oleifera bioactive compounds against SARS-CoV-2 infection, *Pharmacogn. J.*, **2022**, *14*. [Crossref], [Google Scholar], [Publisher]

[37] J. Han, L. Geng, C. Lu, J. Zhou, Y. Li, T. Ming, Z. Zhang, X. Su, Analyzing the mechanism by which oyster peptides target IL-2 in melanoma cell apoptosis based on RNA-seq and m6A-seq, *Food Funct.*, **2023**, *14*, 2362-2373. [Crossref], [Google Scholar], [Publisher]

[38] A.N.M. Ansori, V.D. Kharisma, A.A. Parikesit, F.A. Dian, R.T. Probojati, M. Rebezov, P. Scherbakov, P. Burkov, G. Zhdanova, A. Mikhalev, Y. Antonius, M.R.F. Pratama, N.I. Sumantri, T.H. Sucipto, R. Zainul, Bioactive compounds from mangosteen (Garcinia mangostana L.) as an antiviral agent via dual inhibitor mechanism against SARSCoV-2: an in silico approach, *Pharmacogn. J.*, **2022**, *14*. [Crossref], [Google Scholar], [Publisher]

[39] M.E. Ullah, R.T. Probojati, A.A.A. Murtadlo, M.B. Tamam, S.W. Naw, Revealing of Antiinflamatory Agent from Zingiber officinale var. Roscoe via IKK-B Inhibitor Mechanism through In Silico Simulation, *SAINSTEK International Journal on Applied Science*, *Advanced Technology and Informatics.*, **2022**, *1*, 14-19. [Crossref], [Google Scholar], [Publisher]

[40] N. Wang, Z.Y. Wang, S.L. Mo, T.Y. Loo, D.M. Wang, H. Bin Luo, D.P. Yang, Y.L. Chen, J.-G. Shen, J.-P. Chen, Ellagic acid, a phenolic compound, exerts anti-angiogenesis effects via VEGFR-2 signaling pathway in breast cancer, *Breast Cancer Res. Treat.*, **2012**, *134*, 943-955. [Google Scholar], [Publisher] [41] M. Yousuf, A. Shamsi, P. Khan, M. Shahbaaz, M.F. AlAjmi, A. Hussain, G.M. Hassan, A. Islam, Q. M. R. Haque, I. Hassan, Ellagic acid controls cell proliferation and induces apoptosis in breast cancer cells via inhibition of cyclin-dependent kinase 6, *Int. J. Mol. Sci.*, **2020**, *21*, 3526. [Crossref], [Google Scholar], [Publisher]

[42] H.S. Chen, M.H. Bai, T. Zhang, G.D. Li, M. Liu, Ellagic acid induces cell cycle arrest and apoptosis through TGF-β/Smad3 signaling pathway in human breast cancer MCF-7 cells, *Int. J. Oncol.*, **2015**, *46*, 1730-1738. [Crossref], [Google Scholar], [Publisher]

[43] S. Jaman, A. Sayeed, Ellagic acid, sulforaphane, and ursolic acid in the prevention and therapy of breast cancer: current evidence and future perspectives, *Breast Cancer*, **2018**, *25*, 517-528. [Crossref], [Google Scholar], [Publisher]

[44] R.T. Probojati, S.L. Utami, D.D.R. Turista, A. Wiguna, P. Listiyani, A. Wijayanti, . Rachmawati, S. Wahyuningsih, A.F. Dibha, T. Hasan, M.A. Hafidzhah, R.M. Wijaya, A.M. Hikam, M.B. Tamam, A.A.A. Murtadlo, S.W. Naw, Revealing of Anti-inflammatory Agent from Garcinia mangostana L. Phytochemical as NF-κB Inhibitor Mechanism through In Silico Study. *SAINSTEK International Journal on Applied Science, Advanced Technology and Informatics*, **2022**, *1*, 02, 54-61.[Crossref], [Google Scholar], [Publisher]

[45] T. Zhang, H.S. Chen, L.F. Wang, M.H. Bai, Y.C. Wang, X.F. Jiang, M. Liu, Ellagic acid exerts anti-proliferation effects via modulation of Tgf- β /Smad3 signaling in MCF-7 breast cancer cells, *Asian Pac. J. Cancer Prev.*, **2014**, *15*, 273-276. [Crossref], [Google Scholar], [Publisher]

[46] A.T. Rahman, Rafia, A. Jethro, P. Santoso, V.D. Kharisma, A.A.A. Murtadlo, D. Purnamasari, N.H. Soekamto, A.N.M. Ansori, Kuswati, R.S. Mandeli, K.A.M.S. Aledresi, N.F. M. Yusof, V. Jakhmola, M. Rebezov, M. Rebezov, R. Zainul, K. Dobhal, T. Parashar, M.A. Ghifari, D.A.P. Sari, In Silico Study of the Potential of Endemic Sumatra Wild Turmeric Rhizomes (Curcuma Sumatrana: Zingiberaceae) As Anti-



Cancer, *Pharmacogn. J.*, **2022**, *14*. [Crossref], [Google Scholar], [Publisher]

[47] V.D. Kharisma, A.N.M. Ansori, F.A. Dian, W.C. Rizky, T.G.A. Dings, R. Zainul, A.P. Nugraha, Molecular Docking And Dynamic Simulation Of Entry Inhibitor From Indica Bioactive Compounds Tamarindus Sars-Cov-2 Infection Against Via Viroinformatics Study, Biochem. Cell. Arch., **2021**, *21*, 3323. [Google Scholar], [Publisher] [48] M.E. Ullah, S.W. Naw, A.A.A. Murtadlo, Tamam, R.T. Probojati, Molecular M.B. Mechanism of Black Tea (Camellia sinensis) as SARS-CoV-2 Spike Glycoprotein Inhibitor through Computational Approach, SAINSTEK International Journal on Applied Science, Advanced Technology and Informatics, 2022, 1, 20-25. [Crossref], Google Scholar], [Publisher]

[49] V. Ahire, A. Kumar, K.P. Mishra, G. Kulkarni,acid enhances apoptotic sensitivity of breast cancer cells to γ-radiation, *Nutr. Cancer*, **2017**, *69*, 904-910. [Crossref], [Google Scholar], [Publisher]

[50] N. Wang, Q. Wang, H. Tang, F. Zhang, Y. Zheng, S. Wang, J. Zhang, Z. Wang, X. Xie, Direct inhibition of ACTN4 by ellagic acid limits breast cancer metastasis via regulation of β-catenin stabilization in cancer stem cells, *J. Exp. Clin. Cancer Res.*, **2017**, *36*, 1-19. [Crossref], [Google Scholar], [Publisher]

[51] O.M. Ali, A.A. Bekhit, S.N. Khattab, M.W. Helmy, Y.S. Abdel-Ghany, M. Teleb, A.O. Elzoghby, Synthesis of lactoferrin mesoporous silica nanoparticles for pemetrexed/ellagic acid synergistic breast cancer therapy, *Colloids Surf. B: Biointerfaces*, **2020**, *188*, 110824. [Crossref], [Google Scholar], [Publisher]

[52] P. Listiyani, V. Dhea Kharisma, A. N. M. Ansori, M. H. Widyananda, R.T. Probojati, A. A. A. Murtadlo, D. D. Rahma Turista, Md. E. Ullah, V. Jakhmola, R. Zainul. In silico phytochemical compounds screening of Allium sativum targeting the Mpro of SARS-CoV-2, *Pharmacogn. J.*, **2022**, *14*. [Crossref], [Google Scholar], [Publisher]



[53] A.F. Dibha, S. Wahyuningsih, V.D. Kharisma, A.N.M. Ansori, M.H. Widyananda, A.A. Parikesit ,M. Rebezov, Y. Matrosova, S. Artyukhova, N. Kenijz, M. Kiseleva, V. Jakhmola, R. Zainul, Biological activity of kencur (Kaempferia galanga L.) against SARS-CoV-2 main protease: In silico study, *Int J Health Sci.*, **2022**, *6*, 468-480. [Crossref], [Google Scholar], [Publisher]

[54] R.T. Probojati, S.L. Utami, D.D.R. Turista, A Wiguna, A. Wijayanti, Y. Rachmawati, A.F. Dibha, A. A. A. M. T. Hasan, P. Listiyani, M. A. Hafidzhah, A. M. Hikam, M. B. Tamam, R. M. Wijaya, S. Wahyuningsih, Md.E. Ullah, B-cell Epitope Mapping of Capsid L1 from Human Papillomavirus to Development Cervical Cancer Vaccine Through In Silico Study, *SAINSTEK International Journal on Applied Science, Advanced Technology and Informatics,* **2022**, *1*, 62-71. [Crossref], [Google Scholar], [Publisher]

[55] H. Kaur, S. Ghosh, P. Kumar, B. Basu, K. Nagpal, Ellagic acid-loaded, tween 80-coated, chitosan nanoparticles as a promising therapeutic approach against breast cancer: In-vitro and in-vivo study, *Life Sci.*, **2021**, *284*, 119927. [Crossref], [Google Scholar], [Publisher]

[56] N.S. Aini, V.D. Kharisma, M.H. Widyananda, A.A.A. Murtadlo, R.T. Probojati, D.D.R. Turista, M.B. Tamam, V. Jakhmola, D. P. Sari, M. T. Albari, D. Pernamasari, M.A. Ghifari, M.R. Ghifari, R.S. Mandeli, Muhardi, B. Oktavia, T.K. Sari, T. Sriwahyuni, P. Azhari, M. Fonda Maahury, A.N.M. Ansori, R. Zainul, In silico screening of bioactive compounds from Syzygium cumini L. and moringa oleifera L. against SARS-CoV-2 via tetra inhibitors, Pharmacogn. J., 2022, 14. [Crossref], [Google Scholar], [Publisher]

[57] S. Pirzadeh-Naeeni, M.R. Mozdianfard, S.A. Shojaosadati, A.C. Khorasani, T. Saleh, A comparative study on schizophyllan and chitin nanoparticles for ellagic acid delivery in treating breast cancer. International journal of biological macromolecules, 2020, 144, 380-388. [Crossref], [Google Scholar], [Publisher] [58] S.M. Badr-Eldin, H.M. Aldawsari, U.A. Fahmy, O.A.A. Ahmed, N.A. Alhakamy, O.D. Al-Hejaili, A.A. Alhassan, G.A. Ammari, S.I. Alhazmi, R.M. Alawadi, R. Bakhaidar, A.J. Alamoudi, T. Neamatallah, S. Tima, Optimized Apamin-Mediated Nano-Lipidic Carrier Potentially Enhances the Cytotoxicity of Ellagic Acid against Human Breast Cancer Cells, Int. Ι. Mol. Sci., 2022, 23, 9440 .[Crossref], [Google Scholar], [Publisher]

How to cite this article: Syamsurizal*, Rahadian Zainul, Dheo Shalsabilla Novel, Rismi Verawati, Amalia Putri Lubis, Riso Sari Mandeli, Maurice Efroza, Muhammad Arya Ghifari, Putri Azhari, Muhammad Thoriq Albari, Rollando, Muhammmad Raffi Ghifari, Sari Gando Hidayati, Devi Diagnostic Purnamasari, Muhardi. biomarker of ellagic acid compound from pomegranate plant (punica granatum) on receptor protein-tyrosine kinase ERBB-2 in identifying breast cancer, Journal of Medicinal and Pharmaceutical Chemistry Research, 2023, 5(12), 1097-1108. Link: http://jmpcr.samipubco.com/article_1816 69.html

Copyright © 2023 by SPC (<u>Sami Publishing Company</u>) + is an open access article distributed under the Creative Commons Attribution License(CC BY) license (<u>https://creativecommons.org/licenses/by/4.0/</u>), which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.