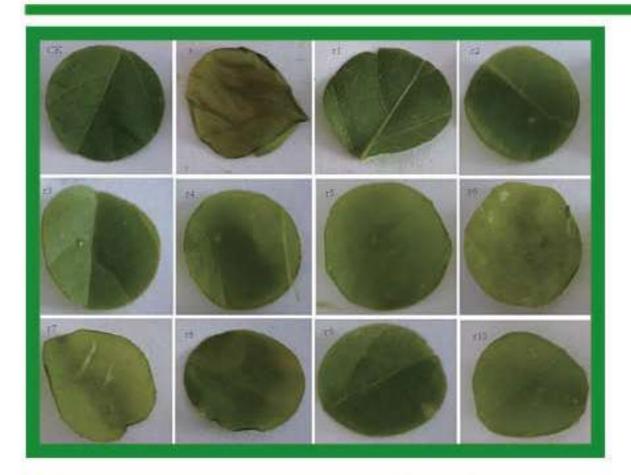




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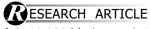
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Dietary flavonoids against various breast cancer subtypes: a molecular docking study

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ABSTRACT: Breast cancer is female most frequent diagnosed cancer and the leading cause of cancer death. The consumption of dietary flavonoids is reported to cause significant breast cancer risk reduction. In vitro studies often used aglycone flavonoids rather than its conjugated form that actually present in human body. Thus its mechanism against breast cancer has not been elucidated completely. The present study aimed to investigate the possible mechanism of dietary flavonoids against breast cancer by in silico study. Conjugated flavonoids were docked to ER (estrogen receptor), HER2 (human epidermal growth factor receptor 2) and EGFR (epidermal growth factor receptor) kinase domains. The molecular docking of 22 flavonoid conjugates towards EGFR and HER2 kinase domain, and ER was successfully performed. Potential binders to proteins: epicatechin conjugates to ER (–8.7 kcal/mol), isoflavone conjugates to HER2 kinase domain (–10.7 kcal/mol), and epigallocatechin and epicatechin conjugates to EGFR kinase domain (–9.2 kcal/mol), were suggested. Supported by other studies, conjugated flavonoids may exert similar inhibitory and agonistic properties to their parent flavonoids. Taken together, the present study showed possible effects of dietary flavonoids against various breast cancer subtypes.

KEYWORDS: molecular docking simulation, diet, flavonoids, antineoplastic agents, breast neoplasms

INTRODUCTION

According to GLOBOCAN 2018 database, breast cancer is female most diagnosed cancer and the leading cause of cancer death ¹. Breast cancer is heterogenous disease, comprised of various subtypes observable by the presence of the predictive molecular markers. Breast cancer can be categorized into: Luminal A (ER+, PR+/-, HER-), Luminal B (ER+, PR^{+/-}, HER2⁺), HER2 (ER⁻, PR⁻, HER2⁺), basal like and claudin low (triple negatives)². Each has a different prognosis and responds differently to cancer treatment. Luminal A has the best prognosis, while HER2⁺ and triple negative breast cancer (TNBC) have the poorest³. Today breast cancer endorsed to be treated with endocrine theraphy, targeted theraphy, and cytotoxic chemotheraphy⁴. Breast cancer with high expression of estrogen receptor

(ER) and progesterone receptor (PR) is sensitive against endocrine theraphy. ER inhibition treatment in Luminal A and B breast cancer was proven to be effective and safe⁵. While HER2⁺ breast cancer was effectively treated using targeted theraphy with trastuzumab or lapatinib⁶. But different from previous subtypes, triple negative breast cancer is not responded very well to hormone treatment and HER2 antibody, and often treated with systemic chemotherapy. Previous study found that epidermal growth factor receptor (EGFR) kinase inhibitor, gefitinib, was able to halt the TNBC cell outgrowth in vitro⁷. Thus ER, HER2, EGFR served as important targets in breast cancer treatment. It is also possible other chemical compounds found in food may also interacts with these particular proteins. Flavonoid is the most common phytochemical compound found

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ubiquitously in human diet^{8,9} and has huge impact in human health. In vitro studies showed that flavonoids have wide range of biological activity antioxidant, anti-inflammatory, anti-microbial, antifungal, antiviral, and anti-cancer 9, 10. Consumption of flavonoids is related with less risk of cardiovascular diseases and stroke 11,12. Other studies found that the intake of flavonoids improved the outcome of the gastric and lung cancer ^{13, 14}. Human study on consumption of food rich in flavonoids, green tea, against breast cancer showed mixed results. Case studies has shown green tea intake was correlated with significant breast cancer risk reduction 15-17. while recent prospective cohort studies showed no correlation ¹⁸. Soy products were rich in isoflavone and high soy intake was modestly associated with reduced breast cancer risk 19.

It is important to note that flavonoid is quickly metabolized in the body. After ingested, glycoside flavonoid found in plant materials is subjected to deglycosylation, releasing aglycone compound that readily absorbed by the intestine lining 20. Once entered circulatory system, flavonoid is immediately transported to liver and undergoes extensive metabolism. Phase metabolism II transformed free aglycone onto flavonoid conjugates by adding glucuronides and sulphate moiety²¹. Because of this, aglycone flavonoids are rarely found in plasma. Previous in vitro studies often used aglycone flavonoids rather than its conjugated forms that present in human body. Thus its mechanism against breast cancer has not been elucidated completely, since conjugation may affect how the molecules behave. To address this, conjugated flavonoids found in plasma after ingesting food rich in flavonoid was subjected to molecular docking against ER, HER2, EGFR. The present study describes possible mechanism how the dietary flavonoids may contribute against breast cancer.

MATERIALS AND METHODS

Molecular docking towards EGFR, HER2, and ER

Conjugated flavonoids found in plasma after ingesting food rich in flavonoids as previously reported from other studies were used ^{22–24}. Structural data of conjugated flavonoids were retrieved from PubChem database (Fig. 1) (pubchem.ncbi. nlm.nih.gov). Co-crystallized structures of ER-4-hydroxytamoxifen (PDB ID: 3ERT) ²⁵, HER2-SYR127063 (PDB ID: 3PP0) ²⁶, and EGFR-gefitinib (PDB ID: 4WKQ) were obtained from RCSB database (rcsb.org). Crystal structure data were

Table 1 Predicted binding affinity (kcal/mol) of flavonoid conjugates and known inhibitors towards ER, HER2, and EGFR.

Flavonoid conjugate	Binding affinity			
	ER	HER2	EGFR	
Gefitinib	_	_	-8.8	
SYR127063	_	-11.0	_	
4-hydroxytamoxifen	-9.7	_	_	
(-)-Epicatechin-3-gallate	-8.7	-8.5	-8.6	
(-)-Epigallocatechin-3-gallate	-7.0	-8.6	-9.2	
(-)-Epigallocatechin-7-gallate	-7.4	-8.9	-8.0	
4'-Methylepicatechin-5-sulfate	-7.2	-8.8	-7.7	
4'-Methylepicatechin-7-sulfate	-7.9	-9.6	-7.2	
4'-Methyl-epigallocatechin- 3'-glucuronide	-7.9	-9.9	-9.1	
4'-Methyl-epigallocatechin- 7-glucuronide	-6.5	-9.0	-8.2	
Daidzein-4'-sulfate	-7.1	-9.4	-8.5	
Daidzein-7-sulfate	-6.6	-9.2	-8.4	
Epicatechin-3'-glucuronide	-8.5	-9.5	-9.1	
Epicatechin-3'-sulfate	-7.7	-9.1	-8.2	
Epicatechin-5-sulfate	-7.1	-8.6	-8.4	
Epicatechin-7-lucuronide	-7.5	-9.3	-8.3	
Epigallocatechin-3'-glucuronide	-7.7	-9.7	-9.1	
Epigallocatechin-7-glucuronide	-7.2	-8.3	-8.3	
Genistein-4'-O-glucuronide	-8.3	-10.7	-8.0	
Genistein-4'-sulfate	-7.3	-9.7	-8.3	
Genistein-7-O-glucuronide	-6.8	-9.4	-7.7	
Genistein-7-sulfate	-6.4	-9.1	-8.6	
Isorhamnetin-3-O-glucuronide	-7.6	-8.3	-8.8	
Quercetin-3'-glucuronide	-8.7	-8.8	-8.7	
Quercetin-3'-sulfate	-8.1	-9.5	-8.4	

prepared by removing solvent and extracting bound ligand. AutoDock vina was used in molecular docking under default settings. The docking methodology was validated by redocking the extracted bound ligand. Chimera was used on visualization in this study. Intramolecular analysis was performed using Pose View, available at Protein Plus (proteins. plus) ²⁷.

RESULTS

Molecular docking analysis

Redocking was performed to evaluate software and docking parameters used. The root mean square deviation between docked and crystal compounds was less than 2Å except for EGFR bound ligand (gefitinib). This is due to the 6-propylmorpholino moiety of gefitinib sticking out to solvent and able to move freely²⁸. Thus, AutoDock Vina has favourable accuracy and proceeds the docking of flavonoid-

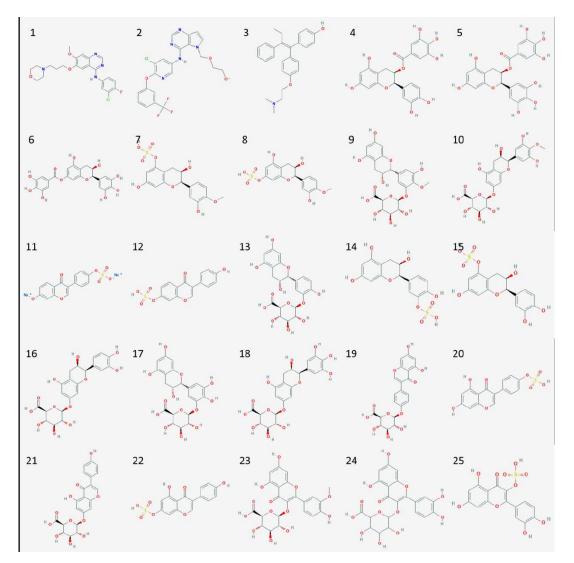


Fig. 1 The 2 dimensional structure of all studied ligands: (1) gefitinib, (2) SYR127063, (3) 4-hydroxytamoxifen, (4) (-)-epicatechin-3-gallate, (5) (-)-epigallocatechin-3-gallate, (6) (-)-epigallocatechin-7-gallate, (7) 4'-methylepicatechin-5-sulfate, (8) 4'-methylepicatechin-7-sulfate, (9) 4'-methyl-epigallocatechin-3'-glucuronide, (10) 4'-methyl-epigallocatechin-7-glucuronide, (11) daidzein-4'-sulfate, (12) daidzein-7-sulfate, (13) epicatechin-3'-glucuronide, (14) epicatechin-3'-sulfate, (15) epicatechin-5-sulfate, (16) epicatechin-7-glucuronide, (17) epigallocatechin-3'-glucuronide, (18) epigallocatechin-7-glucuronide, (19) genistein-4'-O-glucuronide, (20) genistein-4'-sulfate, (21) genistein-7-O-glucuronide, (22) genistein-7-sulfate, (23) isorhamnetin-3-O-glucuronide, (24) quercetin-3'-glucuronide, (25) quercetin-3'-sulfate.

conjugates. The molecular docking was performed to assess possible binding conformation of flavonoid conjugates towards receptors and possible biological actions of these compounds. The molecular docking of 22 flavonoid conjugates towards EGFR and HER2 kinase domain, and ER was successfully performed. The predicted binding affinity value of flavonoid conjugates was compared to each other and receptor bound ligand (Table 1).

In the present study, based on molecular docking, epichatehin and quercetin conjugates ((-)-epicatechin-3-gallate, quercetin-3'-glucuronide, and epicatechin-3'-glucuronide) were predicted as a potential binder towards estrogen receptor. HER2 kinase domain was predicted to interact strongly towards genistein and epigallocatechin conjugates (genistein-4'-O-glucuronide, genistein-4'-sulfate, epigallocatechin-3'-glucuronide,

Table 2 Hydrogen bond formed by flavonoids conjugates and known inhibitors towards EGFR, HER2, and ER.

Flavonoid conjugate	Receptor	Hydrogen bond
4-hydroxytamoxifen	ED	Asp394 Glu353 Leu346 Thr347 Asp351
(-)-Epicatechin-3-gallate Quercetin-3'-glucuronide	ER	Leu387 Glue419 Gly420 Asp351 Glu353
SYR127063 Genistein-4'-O-glucuronide	HER2	Met801 Asp863 Ser783 Thr798
Gefitinib (-)-Epigallocatechin-3- gallate	EGFR	Met793 Glu762 Leu788 Met793 Arg841

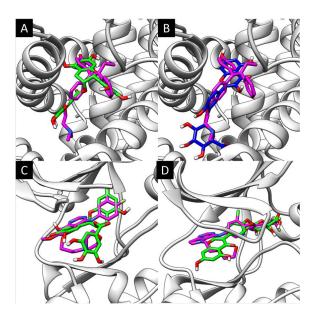


Fig. 2 Superimposed binding mode between flavonoid conjugate and known inhibitor towards ER, HER2, EGFR. (A) and (B) the binding mode of (-)-epicatechin-3-gallate (green), quercetin-3'-glucuronide (blue), and 4-hydroxytamoxifen towards (magenta) ER (light gray). (C) the binding mode of genistein-4'-O-glucuronide (green) and SYR127063 (magenta) towards EGFR kinase domain (light gray). (D) the binding mode of (-)-epigallocatechin-3-gallate (green) and gefitinib (magenta) towards HER2 kinase domain (light gray). The protein represented as ribbon, the compound as stick, and heteroatom represented in a different color from the carbon atom.

and 4'-methyl-epigallocatechin-3'-glucuronide) compared to other compounds. While several catechin derivatives ((-)-epigallocatechin-3-gallate, 4'-methyl-epigallocatechin-3'-glucuronide, epicatechin-3'-glucuronide, and epigallocatechin-3'-glucuronide) were predicted with a high affinity toward EGFR kinase domain as these compounds

surpassed known inhibitor binding score.

Binding mode of the compound with the highest predicted binding affinity was visualized and superimposed with the receptor known inhibitor (Fig. 2). All compounds were found to occupy the active site of the protein. (-)-Epigallocatechin-3gallate resembled similar binding mode with gefitinib, whereas its core structure aligned with the quinazoline and aniline moiety. While genistein-4'-O-glucuronide, (-)-epicatechin-3-gallate, quercetin-3'-glucuronide only shared small geometrical similarity when compared with the bound ligand. Furthermore, hydrogen bond inferred by Pose View was compared and presented in (Table 2). Similar to gefitinib, (-)-epigallocatechin-3-gallate formed hydrogen bond with Met793. SYR127063 and genistein-4'-O-glucuronide shared no similar intermolecular interaction. Quercetin-3'-glucuronide had similar interaction with 4-hydroxytamoxifen at Glu353. (-)-Epicatechin-3-gallate had the most hydrogen bond towards ESR1.

DISCUSSION

In present study, predicted binding affinity and binding mode of conjugated flavonoids present in plasma after ingestion of dietary flavonoids against ER, EGFR kinase domain, and HER2 kinase domain were characterized *in silico*. The result shows that most compounds with predicted high binding affinity were glucuronide flavonoid conjugates. It is interesting to point out that the predominant flavonoid metabolite found in plasma after an hour ingestion of radiolabelled epicatechin was its glucuronide conjugates²⁹. Thus, the potential compounds found in this study were likely exist in large concentration in plasma after consumption of dietary flavonoids.

More than 70% diagnosed breast cancer was the overexpressed ER³. ER plays important role in development and progression of breast cancer, since ER drives proliferation of mammary cells upon binding with estrogenic hormone³⁰. ER⁺ breast cancer is sensitive against endocrine therapy and is effectively treated using selective estrogen receptor modulators such as tamoxifen³¹. This study found that (-)-epicatechin-3-gallate was a potential inhibitor of ER, because of its high predicted binding affinity and similar binding mode when compared to active metabolite of tamoxifen (4hydroxytamoxifen). Tamoxifen interacts with several amino acid residues inside the binding pocket, including Leu346, Thr347 and Leu387, forming a van der walls interaction that stabilize the com-

plex ²⁵. Epicatechin conjugates was predicted to be interacted with similar manner. This finding confirmed by other studies whereas epicatechin gallate was able to hamper ER activity through direct inhibition ^{32,33}. Quercetin conjugates also have high predicted binding affinity towards ER. But quercetin conjugates may act as an agonist rather than antagonist, since previous study found that aglycone quercetin induced cell proliferation of ER-positive breast cancer cell line through ER stimulation ³⁴.

HER2 is a receptor tyrosine kinase which is overexpressed in 30% human breast cancer³. HER2⁺ breast cancer characterized by its aggressive phenotype: high tumorigenicity and invaseness³⁵. The treatment involved is either by targeting the extracellular domain using trastuzumab or its kinase domain using lapatinib 4,36. Previous study showed that flavonoid compounds were able to inhibit human kinases³⁷. Molecular docking study reported that genistein-4'-O-glucuronide had predicted binding affinity close to SYR127063. SYR127063 itself is a potent HER2 kinase domain inhibitor at IC50 of 11 nM²⁶. From the experimental study, genistein was able to attenuate HER2 phosphorylation in BT474 cell line through tyrosine kinase inhibition, thus supported present finding ³⁸.

TNBC occurs approximately 10% in breast cancer cases3. TNBC is biologically aggressive and has the poorest prognosis when compared to other subtypes³. Previous study reported that EGFR is a potential target for TNBC⁷. In this work, epigallocatechin and epicatechin metabolites had notable predicted binding affinity towards EGFR kinase domain. This finding supported by another study where epigallocatechin-3-gallate was able to inhibit EGFR activity³⁹. Inhibition of EGFR by epigallocatechin conjugate may also affect HER2 activation, since both proteins are able to form heterodimeric complex and activate each other.

CONCLUSION

Binding affinity and binding mode between conjugated flavonoids found in plasma against ER, HER2, EGFR had been characterized in silico. Supported by other studies, conjugated flavonoids may exert similar inhibitory and agonistic properties to their parent flavonoids. Our study thus confirm and offer possible explanations how dietary flavonoids act against various breast cancer subtypes.

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Dear Author,

This is to acknowledge the receipt of your manuscript entitled: **Dietary Flavonoids Against Various Breast Cancer Subtypes: A Molecular Docking Study** by **Wahyu Widowati,Diana Krisanti Jasaputra,Yusuf Heriady,Ahmad Faried,Rizal Rizal,Wahyu Setia Widodo,Satrio Haryo Benowo Wibowo,Hanna Sari Widya Kusuma,Ermi Girsang,I Nyoman Ehrich Lister**, for consideration for publication in *ScienceAsia*.

The manuscript is under editorial review to consider its suitability for publication in *ScienceAsia* with respect to research areas, novelty, conformity to journal guidelines, etc. You will soon receive an email informing you of the Editor's opinion.

Regards, ScienceAsia Editorial

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October 23, 2019

MS Code No: 2019-0552 Password: wyy4db

Dr. Wahyu Widowati Jl. Surya Sumantri No. 65 Bandung 40164, West Java, Indonesia myoungtanurat@gmail.com

Re: Revision 1 of MS 2019-0552 Dear Dr. Wahyu Widowati,

Your manuscript (code 2019-0552) entitled Dietary Flavonoids Against Various Breast Cancer Subtypes: A Molecular Docking Study by Wahyu Widowati, Diana Krisanti Jasaputra, Yusuf Heriady, Ahmad Faried, Rizal Rizal, Wahyu Setia Widodo, Satrio Haryo Benowo Wibowo, Hanna Sari Widya Kusuma, Ermi Girsang, I Nyoman Ehrich Lister, submitted on July 14, 2019 for consideration for publication in *ScienceAsia*, has been sent for review and comments from the reviewers received are given below for your consideration. You are required to clarify these queries and comments in your revised version by copying and pasting the reviewers queries into your reply, and answer each of the points one by one. To expedite the publication process, please make sure that the references are in the correct format by checking the following points:

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Re-submit the revised manuscript, with a document indicating responses or changes in the manuscript against the referees amendments to ScienceAsia within 1 week. Kindly visit www.scienceasia.org, click the Author login menu and use the MS Code No. and password provided at the top-right corner of this email to submit your revised manuscript.

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Yours sincerely, Prof. Supa Hannongbua Editor, ScienceAsia

REVIEWERS COMMENT

Scientific validity: not sound, in part
Scientific interest: above average
Presentation: not clear

Comment : Comments for Dietary Flavonoids Against Various Breast Cancer Subtypes: A Molecular

Docking Study

This manuscript presents the computational study about the interactions between

flavonoid conjugate compounds and breast cancer related proteins, ER, HER2 and EGFR by

using molecular docking.

COMMENTS TO THE AUTHORS:

Comment 1: The abstract should be concise and clear.

Comment 2: More details of docking protocol are required. PDB id of the structure EGFR-gefitinib (lines 87-88) should be provided.

Comment 3: The sentence The 3-dimensional structure data for every compound was obtained from PubChem database (https://pubchem.ncbi.nlm.nih.gov) (lines 86-87) is a bit awkward and should be rewritten.

Comment 4: The paragraph in lines 107-116 should be rewritten.

Comment 5: (-)-Epicatechin-3-gallate showed binding energy with ER similar to 4-hydroxytamoxifen. More discussion on which amino acid residues interact with compound (lines 142-144) in comparison to 4-hydroxytamoxifen is required.

Comment 6: For superimposed binding mode between flavonoid conjugate and known inhibitor in Figure 1, heteroatom should be presented in a different color from the carbon atom and figure caption should be rewritten for better understanding. In addition, the 2D structure of all studied ligands should be added.

Comment 7: The reference format must be rechecked. Grammar and typo errors throughout the manuscript must be corrected.

EDITORS COMMENT

According to the comments from the reviewer, this manuscript can be accepted after minor revsion. Please inform the authors and ask them to response with clear indication about the revision. The generation of the line number can be done in order to making the clear response. Also, the authors shall follow the instruction for authors and follow the criteria of the journal. Figure and Table should be less than 8.

EDITORIAL OFFICE: c/o Science Society of Thailand, Faculty of Science, Chulalongkorn University PhyaThai Road, Patumwan, Bangkok 10330, Thailand

Tel:+662 218 5245 Fax: +662 252 7987 E-mail: editor@scienceasia.org

Dear Author.

We are still waiting to hear from you after our email of 5 November 2019, please send back as soon.

Regards,

Mongkon Youngtanurat

Dr. Wahyu Widowati

Jl. Surya Sumantri No. 65 Bandung 40164, West Java, Indonesia

wahyu_w60@yahoo.com, wahyu@amubbrc.co.id

Re: Acceptance for MS 2019-0552

Dear Dr. Wahyu Widowati,

I am pleased to inform you that your manuscript (code 2019-0552) entitled **Dietary Flavonoids Against Various Breast Cancer Subtypes:** A Molecular Docking Study by Wahyu Widowati, Diana Krisanti Jasaputra, Yusuf Heriady, Ahmad Faried, Rizal Rizal, Wahyu Setia Widodo, Satrio Haryo Benowo Wibowo, Hanna Sari Widya Kusuma, Ermi Girsang, I Nyoman Ehrich Lister, submitted on July 14, 2019 has been considered to be accepted for publication in Science Asia. However, this letter is not an ACCEPTANCE LETTER. Prior to issuing a formal ACCEPTANCE LETTER, we would like to remind you that papers recommended to be published in Science Asia are required to pay for publication fee of 500 USD. The charge excludes printing of graphics or figures in full colour in the hard copy version which will cost an additional of 1,000 USD per paper. Printing of graphics in full colour is optional and is at the author's discretion. In all cases, the payment must be settled soon after the authors have been notified that the paper is recommended to be published. Delay in receiving the publication fee from you may mean that the journal will have to postpone the publication of your article.

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Emeritus Prof. Piamsook Pongsawasdi

Editor-in-chief, ScienceAsia

Dr. Wahyu Widowati

Jl. Surya Sumantri No. 65 Bandung 40164, West Java, Indonesia wahyu w60@yahoo.com, wahyu@amubbrc.co.id

Subject: Re: 2019-0552 proof of your article

Entitled: Dietary Flavonoids Against Various Breast Cancer Subtypes: A Molecular Docking Study Author(s): Wahyu Widowati,Diana Krisanti Jasaputra,Yusuf Heriady,Ahmad Faried,Rizal Rizal,Wahyu Setia Widodo,Satrio Haryo Benowo Wibowo,Hanna Sari Widya Kusuma,Ermi Girsang,I Nyoman Ehrich Lister

Dear Dr. Wahyu Widowati,

The proof version of your manuscript is available at (http://scienceasia.org/manuscript/msfiles/2019-0552/2019-0552P1.pdf) Please read the proof very carefully. It is journal policy to edit manuscripts to ensure that they conform to the journal style and also in some cases to improve conciseness, clarity, and accuracy. These changes are based on the editors' interpretation of what was originally written -- it is possible that the editors have misunderstood some parts. Any sections highlighted in yellow indicate words that we have added, changed, or are unsure about that we would like you to check in particular. Sections highlighted in red definitely need a response from you -- e.g., the definition of an abbreviation is not given, or we couldn't understand at all what is meant. ??? signifies a missing value or word. If there is an error in a citation number in the text (not the references), give us the corrected citation number in terms of the existing numbers in the list of references. If necessary we will then reorder the list of references and make the corresponding adjustments to the citation numbers throughout.

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Yours sincerely, Asst. Prof. Dr. Khamron Mekchay Managing Editor, *ScienceAsia*

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KARYA ILMIAH: JURNAL ILMIAH

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REVIEWER 1

(Prof. Dr. Chrismis Novalinda Ginting, M.Kes) NK: 0115127801 UNIVERSITAS PRIMA INDONESIA

LEMBAR HASIL PENILAIAN SEJAWAT SEBIDANG atau *PEER REVIEW*

KARYA ILMIAH: JURNAL ILMIAH

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REVIEWER 2

(Prof. Dr. Ermi Girsang, M. Kes)

NIK: 0117057501

UNIVERSITAS PRIMA INDONESIA

LEMBAR HASIL PENILAIAN SEJAWAT SEBIDANG atau *PEER REVIEW*

KARYA ILMIAH: JURNAL ILMIAH

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Medan, Reviewer 2

(Prof. Dr. Ermi Girsang, M.Kes)

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Medan, Reviewer 1

(Prof. Dr. Chrismis Novalinda Ginting, M.Kes) NIK: 011517801

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