



Correction

Correction: Shoombuatong, W., et al. iQSP: A Sequence-Based Tool for the Prediction and Analysis of Quorum Sensing Peptides via Chou's 5-Steps Rule and Informative Physicochemical Properties. *Int. J. Mol. Sci.* 2020, 21, 75

Phasit Charoenkwan ¹, Nalini Schaduagratt ², Chanin Nantasenamatt ² , Theeraphon Piacham ³ and Watshara Shoombuatong ^{2,*}

¹ College of Arts, Media and Technology, Chiang Mai University, Chiang Mai 50200, Thailand; phasit.c@cmu.ac.th

² Center of Data Mining and Biomedical Informatics, Faculty of Medical Technology, Mahidol University, Bangkok 10700, Thailand; nalini.sch@mahidol.edu (N.S.); chanin.nan@mahidol.edu (C.N.)

³ Department of Clinical Microbiology and Applied Technology, Faculty of Medical Technology, Mahidol University, Bangkok 10700, Thailand; theeraphon.pia@mahidol.ac.th

* Correspondence: watshara.sho@mahidol.ac.th; Tel.: +66-2-441-4371 (ext. 2715)

Received: 7 April 2020; Accepted: 8 April 2020; Published: 10 April 2020



The authors wish to make the following corrections to this paper [1]:

Please note that all references orders have been re-numbered in the corrected paper.

1. Change in Title

We have found the following error in the title of this article which was recently published in *Int. J. Mol. Sci.* [1]:

The correct title should be: iQSP: A Sequence-Based Tool for the Prediction and Analysis of Quorum Sensing Peptides Using Informative Physicochemical Properties

2. Change in Main Body Paragraphs

We have found four inadvertent errors in our paper published in *Int. J. Mol. Sci.* [1].

In “a series of recent publications [67–72] . . .”, we deleted the references [68,69] and their citations, and renumbered the references orders.

In “As described in [35,85] and investigated by Equation (50) of [86], among those testing methods, the jackknife test is considered as one of the most rigorous that can provide a unique result for a given benchmark dataset.”, we deleted the sentence “As described in [35,85] and investigated by Equation (50) of [86],”. We deleted the references [85,86] and their citations, moved the citation of reference [35] to the end of the sentence, and renumbered all references orders.

In “into an unprecedented revolution [34,94–107].”, we deleted the the references [94–107] and their citations.

In “S-palmitoylation sites in proteins [68], lysine crotonylation sites [108], and phosphotyrosine sites [109]. Furthermore, our method could be integrated with other beneficial peptide features such as pseudo amino acid composition [110–112] or amphiphilic pseudo amino acid composition as proposed [113] by Chou [35,114] for further improving the QSP prediction.”, we deleted the references [68], [108–114] and their citations. We deleted the sentence “as proposed [113] by Chou [35,114]”.

These changes have no material impact on the conclusions of our paper.

The authors would like to apologize for any inconvenience caused to the readers by these changes.

Reference

1. Charoenkwan, P.; Schaduangrat, N.; Nantasenamat, C.; Piacham, T.; Shoombuatong, W. iQSP: A Sequence-Based Tool for the Prediction and Analysis of Quorum Sensing Peptides Using Informative Physicochemical Properties. *Int. J. Mol. Sci.* **2020**, *21*, 75. [[CrossRef](#)] [[PubMed](#)]



© 2020 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (<http://creativecommons.org/licenses/by/4.0/>).