

DISCIPLINE COUNCIL FOR CHEMICAL SCIENCES

mgr inż. Maria BZÓWKA

ANALYSIS OF MOLECULAR ASPECTS OF PROTEINS REGULATION CONSIDERING WATER MOLECULES AS A POTENTIAL MEDIATOR IN INTERMOLECULAR INTERACTIONS

ATTACHMENT TO THE DOCTORAL THESIS: CO-AUTHOR CONTRIBUTION STATEMENTS

Supervisor: dr hab. Artur GÓRA, prof. PŚ

Co-Author Contribution Statements for Paper 1

[1] **Paper 1:** Mitusińska K., Raczyńska A., **Bzówka M.**, Bagrowska W., Góra A. Applications of water molecules for analysis of macromolecule properties Computational and Structural Biotechnology Journal (2020) 18, 355-365 https://doi.org/10.1016/j.csbj.2020.02.001





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Maria Browks_

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Raczyńska A., Bzówka M., Bagrowska W., Góra A.

Applications of water molecules for analysis of macromolecule properties

Computational and Structural Biotechnology Journal (2020) 18, 355-365

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 15%

I was involved in the literature search and data organisation (particularly regarding the applicability and functionality of software reviewed in the second and third part of the article). I was involved in writing the manuscript, reviewing and editing the final version and providing answers to the reviewers' comments.







Dr Karolina Mitusińska PhD Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Raczyńska A., Bzówka M., Bagrowska W., Góra A.

Applications of water molecules for analysis of macromolecule properties

Computational and Structural Biotechnology Journal (2020) 18, 355-365

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 40%

I was involved in the literature search and data organisation. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Karolina Mimsinsko

Silesian University of Technology Biotechnology Centre









Agata Raczyńska MSc, BEng

20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Raczyńska A., Bzówka M., Bagrowska W., Góra A.

Applications of water molecules for analysis of macromolecule properties

Computational and Structural Biotechnology Journal (2020) 18, 355-365

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 20%

I was involved in the literature search and data organisation. I was involved in writing the manuscript and preparation of figures.









Weronika Bagrowska MSc, BEng

Gliwice, 17.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Raczyńska A., Bzówka M., Bagrowska W., Góra A.

Applications of water molecules for analysis of macromolecule properties

Computational and Structural Biotechnology Journal (2020) 18, 355-365

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 15%

I was involved in the literature search and data organisation. I was involved in writing the manuscript and preparation of figures.

Baprache Weroutez







Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Raczyńska A., Bzówka M., Bagrowska W., Góra A.

Applications of water molecules for analysis of macromolecule properties

Computational and Structural Biotechnology Journal (2020) 18, 355-365

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 10%

I conceived the project and contributed to its conceptualisation. I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.

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Co-Author Contribution Statements for Paper 2

[2] **Paper 2:** Magdziarz T., Mitusińska K., **Bzówka M.**, Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 16,5%

I was involved in software development by testing new features that were added to drivers *valve* and *pond*. I was working with the main developer of the software on the optimisation of the *pond* driver for local-distribution analysis. I was testing different modes of analysis (*time-window* and *sandwich*. I was working on the following case studies: human soluble epoxide hydrolase, potato soluble epoxide hydrolase, and haloalkane dehalogenase LinB. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewer's comments.

Maria Biówka



Tomasz Magdziarz, PhD

Email address: tomasz.magdziarz@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective
Bioinformatics (2020) 36 (8), 2599-2601
https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 20% I contributed to the conceptualisation of the project. I was the main developer of the computer code for AQUA-DUCT software. I was involved in writing the manuscript and responding to the Reviewers' comments.

Signature

Town Magalian





Dr **Karolina Mitusińska** PhD Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 18,5%

I was involved in the software development by running small-molecule tracking analysis for several different systems. I was involved in running benchmarking section. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Korolina Mimsili, te

Silesian University of Technology Biotechnology Centre









Agata Raczyńska MSc, BEng

20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 7,5%

I prepared various tutorials, e.g. for clustering, trimming paths, and smoothing paths. I was involved in writing the documentation and figures preparation.





Agnieszka Stańczak, mgr inż. agnieszka.stanczak@uochb.cas.cz Institute of Organic Chemistry and Biochemistry of the Czech Academy of Sciences Faculty of Science, Charles University, Czech Republic

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective Bioinformatics (2020) 36 (8), 2599-2601 https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 5 %

I performed the MD simulation of the haloalkane dehalogenase LinB with different substrates for consolidator mode usage (rare events analysis).

Signature

Apries la Feriul

Michał Banas, BEng

Email address: mich.banas0@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective
Bioinformatics (2020) 36 (8), 2599-2601
https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 7,5% I was the developer of the Graphical User Interface (GUI) version for AQUA-DUCT 1.0 and *Kraken* module for data visualisation.

Signature





Weronika Bagrowska MSc, BEng

Gliwice, 17.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A.

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 5%

I was testing different functionalities of the AQUA-DUCT 1.0 software.

Bugrowales Weronital







Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Magdziarz T., Mitusińska K., Bzówka M., Raczyńska A., Stańczak A., Banas M., Bagrowska W., Góra A

AQUA-DUCT 1.0: structural and functional analysis of macromolecules from an intramolecular voids perspective

Bioinformatics (2020) 36 (8), 2599-2601

https://doi.org/10.1093/bioinformatics/btz946

I declare the following contribution to this publication: 20%

I conceived the project and contributed to its conceptualisation. I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.

Silesian University of Technology Biotechnology Centre





Co-Author Contribution Statements for Paper 3

[3] **Paper 3: Bzówka M.*,** Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A. Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design International Journal of Molecular Sciences (2020) 21, 1-17 https://doi.org/10.3390/ijms21093099





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design

International Journal of Molecular Sciences (2020) 21, 1-17

https://doi.org/10.3390/ijms21093099

I declare the following contribution to this publication: 30%

I was responsible for running all mixed-solvent molecular dynamics simulations and their further analysis with AQUA-DUCT software, including the analysis of the intramolecular voids and hot-spots. Also, I was involved in conducting the same analysis with AQUA-DUCT software for classical MD simulations. I was involved in comparison of crystal structures of main proteases from SARS-CoV-2 and SARS-CoV and the analysis of the plasticity of their binding cavities. I was also involved in the analysis of the potential mutability of SARS-CoV-2. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewers' comments.

Maria Browke







Dr **Karolina Mitusińska PhD** Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design

International Journal of Molecular Sciences (2020) 21, 1-17

https://doi.org/10.1016/j.csbj.2020.02.001

I declare the following contribution to this publication: 30%

I was involved in conducting a set of classical molecular dynamics simulations and analysing them. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Koronina minstrike

Silesian University of Technology Biotechnology Centre









Agata Raczyńska MSc, BEng

20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design

International Journal of Molecular Sciences (2020) 21, 1-17

https://doi.org/10.3390/ijms21093099

I declare the following contribution to this publication: 10%

I performed the analysis of the energetic effect of amino acid substitutions.





Aleksandra Samol, MSc

Email address: 2aleksandrasamol@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design International Journal of Molecular Sciences (2020) 21, 1-17 https://doi.org/10.3390/ijms21093099

I declare the following contribution to this publication: 5% I was involved in conducting the evolutionary analysis.

Gome Signature

Professor Jack A. Tuszyński
Politecnico di Torino
University of Alberta
Email address: jacek.tuszynski@polito.it; jackt@ualberta.ca

Co-Author contribution statement

Hereby, I certify that I am the Co-Author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design International Journal of Molecular Sciences (2020) 21, 1-17 https://doi.org/10.3390/ijms21093099

I declare the following contribution to this publication: 5% I contributed to the project's conceptualization. I was also involved in the reviewing and editing of the manuscript.

Signature





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Samol A., Tuszyński J.A., Góra A.

Structural and Evolutionary Analysis Indicate That the SARS-CoV-2 Mpro Is a Challenging Target for Small-Molecule Inhibitor Design

International Journal of Molecular Sciences (2020) 21, 1-17

https://doi.org/10.3390/ijms21093099

I declare the following contribution to this publication: 20%

I conceived the project and contributed to its conceptualisation. I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.

Silesian University of Technology Biotechnology Centre



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Co-Author Contribution Statements for Paper 4

[4] **Paper 4:** Fischer A.*, Sellner M.*, Mitusińska K.*, **Bzówka M.***, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2 International Journal of Molecular Sciences (2021) 22 (4), 1-17 https://doi.org/10.3390/ijms22042065





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2

International Journal of Molecular Sciences (2021) 22 (4), 1-17

https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 20%

I was involved in the analysis of the similarity of the selected panel of proteases. I was involved in performing classical molecular dynamics simulations with the water molecules tracking and hydration sites identification of the selected panel of proteases. I was also involved in the analysis of the co-solvent sites. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewers' comments. I was responsible for coordinating the cooperation with international partners.

Maria Bzárka



Andre Fischer, PhD Scientist, dsm-firmenich Email address: andre fischer1994@hotmail.com

Co-Author contribution statement

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2

International Journal of Molecular Sciences (2021) 22 (4), 1-17

https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 20%

I contributed to the project's conceptualisation. I was involved in conducting cosolvent MD simulations, molecular docking, MD and MM/GBSA post-processing, and toxicity profiling. I was involved in writing the manuscript and responding to the reviewers' comments.

Signature

Manuel Sellner, Assistant / PhD candidate Computational Pharmacy Department of Pharmaceutical Sciences University of Basel, Switzerland Email address: manuel.sellner@unibas.ch

Co-Author contribution statement

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2 International Journal of Molecular Sciences (2021) 22 (4), 1-17 https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 20%

I was involved in molecular docking, MD and MM/GBSA post-processing, and toxicity profiling. I was involved in writing the manuscript.

Signature Signature





Dr **Karolina Mitusińska PhD** Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2

International Journal of Molecular Sciences (2021) 22 (4), 1-17

https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 20%

I was involved in conducting a set of classical molecular dynamics simulations and analysing them. Also, I was involved in the analysis of the similarity of the selected proteins. I was involved in writing the manuscript..

Yours sincerely,

Karohina Mimsimble

Silesian University of Technology Biotechnology Centre





Markus A. Lill, Prof. Dr.
Computational Pharmacy
Department of Pharmaceutical Sciences
University of Basel, Switzerland
Email address: markus.lill@unibas.ch

Co-Author contribution statement

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2 International Journal of Molecular Sciences (2021) 22 (4), 1-17 https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 6,67% I was involved in reviewing and editing the manuscript. I supervised the project.

Signature





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2

International Journal of Molecular Sciences (2021) 22 (4), 1-17

https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 6,67%

I was involved in editing and reviewing the manuscript. I supervised the project.

Silesian University of Technology Biotechnology Centre



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Martin Smieško, Prof. Dr. Computational Pharmacy Department of Pharmaceutical Sciences University of Basel, Switzerland Email address: martin.smiesko@unibas.ch

Co-Author contribution statement

Hereby, I certify that I am the co-author of the following publication:

Fischer A.*, Sellner M.*, Mitusińska K.*, Bzówka M.*, Lill M.A., Góra A., Smieško M.

Computational selectivity assessment of protease inhibitors against SARS-CoV-2 International Journal of Molecular Sciences (2021) 22 (4), 1-17 https://doi.org/10.3390/ijms22042065

I declare the following contribution to this publication: 6,67% I was involved in reviewing and editing the manuscript. I supervised the project.

Signature

Co-Author Contribution Statements for Paper 5

[5] Paper 5: Bzówka M., Mitusińska K., Hopko K., Góra A.

Computational insights into the known inhibitors of human soluble epoxide hydrolase

Drug Discovery Today (2021) 26 (8), 1914-1921

https://doi.org/10.1016/j.drudis.2021.05.017





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Mitusińska K., Hopko K., Góra A.

Computational insights into the known inhibitors of human soluble epoxide hydrolase

Drug Discovery Today (2021) 26 (8), 1914-1921

https://doi.org/10.1016/j.drudis.2021.05.017

I declare the following contribution to this publication: 60%

I contributed to the project's conceptualisation. I carried out all the analyses for known inhibitors co-crystallised with human soluble epoxide hydrolase. I performed the analysis of the interactions, including clustering of binding residues and inhibitors. I ran molecular dynamics simulations and further AQUA-DUCT analysis for the identification of tunnels and inner voids. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewers' comments.

Monia Brówka







Dr **Karolina Mitusińska PhD** Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Mitusińska K., Hopko K., Góra A.

Computational insights into the known inhibitors of human soluble epoxide hydrolase

Drug Discovery Today (2021) 26 (8), 1914-1921

https://doi.org/10.1016/j.drudis.2021.05.017

I declare the following contribution to this publication: 20%

I was involved in helping in data organisation and editing the manuscript.

Yours sincerely,

Karoline Mitaniako

Silesian University of Technology Biotechnology Centre









Biotechnology Centre

MSc, BEng **Katarzyna Żurawska** (**Hopko**)

Gliwice, 20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Mitusińska K., Hopko K., Góra A.

Computational insights into the known inhibitors of human soluble epoxide hydrolase

Drug Discovery Today (2021) 26 (8), 1914-1921 https://doi.org/10.1016/j.drudis.2021.05.017

I declare the following contribution to this publication: 5%

I did the general identification of chemical motives in inhibitors' structures.

Silesian University of Technology Biotechnology Centre

Krzywoustego 8, 44-100 Gliwice Katarzyna.Zurawska@polsl.pl

NIP PL631 020 07 36 ING Bank Śląski S.A. o/Gliwice 60 1050 1230 1000 0002 0211 3056



Matayus Lunevsha (Hopko)









Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Mitusińska K., Hopko K., Góra A.

Computational insights into the known inhibitors of human soluble epoxide hydrolase

Drug Discovery Today (2021) 26 (8), 1914-1921

https://doi.org/10.1016/j.drudis.2021.05.017

I declare the following contribution to this publication: 15%

I conceived the project and contributed to its conceptualisation. I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.

Silesian University of Technology Biotechnology Centre





Co-Author Contribution Statements for Paper 6

[6] **Paper 6:** Mitusińska K., Wojsa P., **Bzówka M.**, Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205 https://doi.org/10.1016/j.csbj.2021.10.042





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205

https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 17,5%

I was involved in analyses of the flow of water molecules for all the analysed structures, particularly the determination of the appropriate parameters for detecting tunnels and their subsequent characteristics. I also participated in a general comparison of the soluble epoxide hydrolase members by analysing their structural features. I was involved in data organisation, reviewing the manuscript and responding to the reviewers' comments.

Ylania Browka







Dr **Karolina Mitusińska PhD** Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205

https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 30%

I was involved in conducting a set of classical molecular dynamics simulations, analysing them and organising data. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Koroline Mimiliake





Piotr Wojsa, BEng

Email address: piotr.j.wojsa@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205 https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 17,5% I performed molecular dynamics simulation for *Vigna radiata* soluble epoxide hydrolase. I was involved in water path analysis for selected repetitions and figures preparation.

Signature

Note Man





Agata Raczyńska MSc, BEng

20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205

https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 5%

I conducted molecular dynamics simulation for human soluble epoxide hydrolase.









Weronika Bagrowska MSc, BEng

Gliwice, 17.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205

https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 5%

I conducted molecular dynamics simulation for *Bacillus Megaterium* soluble epoxide hydrolase. I was involved in preparation of figures.





Aleksandra Samol, MSc

Email address: 2aleksandrasamol@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205 https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 5% I performed molecular dynamics simulation for *Trichoderma reesei* soluble epoxide hydrolase.

Somol Helicenste

Patryk Kapica, MSc

Email address: Kapica. patnick@g mail. com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205 https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 5% I did a literature review on substrate specificity.

Signature





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K., Wojsa P., Bzówka M., Raczyńska A., Bagrowska W., Samol A., Kapica P., Góra A.

Structure-function relationship between soluble epoxide hydrolases structure and their tunnel network

Computational and Structural Biotechnology Journal (2022) 20, 193-205

https://doi.org/10.1016/j.csbj.2021.10.042

I declare the following contribution to this publication: 15%

I conceived the project and contributed to its conceptualisation. I was involved in writing and reviewing the manuscript. I participated in responding to the reviewers' comments. I was responsible for project administration and supervision.





Co-Author Contribution Statements for Paper 7

[7] Paper 7: Mitusińska K.*, Bzówka M.*, Magdziarz T., Góra A.

Geometry-Based versus Small-Molecule Tracking Method for Tunnel Identification: Benefits and Pitfalls

Journal of Chemical Information and Modeling (2022) 62 (24), 6803-6811 https://doi.org/10.1021/acs.jcim.2c00985





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K.*, Bzówka M.*, Magdziarz T., Góra A.

Geometry-Based versus Small-Molecule Tracking Method for Tunnel Identification: Benefits and Pitfalls

Journal of Chemical Information and Modeling (2022) 62 (24), 6803-6811

https://doi.org/10.1021/acs.jcim.2c00985

I declare the following contribution to this publication: 40%

I was involved in identifying tunnels in MD simulations using small-molecular tracking approach. I was partially involved in the identification of tunnels in crystal structures using a geometry-based approach. I was involved in refining the methods for comparing tunnels found in crystal structures with tunnels identified during MD simulations. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewers' comments.

Homa Blówka







Dr **Karolina Mitusińska PhD** Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K.*, Bzówka M.*, Magdziarz T., Góra A.

Geometry-Based versus Small-Molecule Tracking Method for Tunnel Identification: Benefits and Pitfalls

Journal of Chemical Information and Modeling (2022) 62 (24), 6803-6811

https://doi.org/10.1021/acs.jcim.2c00985

I declare the following contribution to this publication: 40%

I was involved in running geometry-based analysis for the set of the proteins. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Karolina Mitusinista





Tomasz Magdziarz, PhD

Email address: tomasz.magdziarz@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Mitusińska K.*, Bzówka M.*, Magdziarz T., Góra A.

Geometry-Based versus Small-Molecule Tracking Method for Tunnel Identification: Benefits and Pitfalls

Journal of Chemical Information and Modeling (2022) 62 (24), 6803-6811 https://doi.org/10.1021/acs.jcim.2c00985

I declare the following contribution to this publication: 5% I was involved in preparing scripts for the tunnel comparison.

Signature
Tomen Magshism





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Mitusińska K.*, Bzówka M.*, Magdziarz T., Góra A.

Geometry-Based versus Small-Molecule Tracking Method for Tunnel Identification: Benefits and Pitfalls

Journal of Chemical Information and Modeling (2022) 62 (24), 6803-6811

https://doi.org/10.1021/acs.jcim.2c00985

I declare the following contribution to this publication: 15%

I conceived the project and contributed to its conceptualisation I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.





Co-Author Contribution Statements for Paper 8

[8] **Paper 8: Bzówka M*.**, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M*., Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins-What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 25%

I performed the evolutionary analysis of the residues making up referential compartments and all tunnels in each of the selected member of soluble epoxide hydrolase subfamily. I participated in providing characteristic about the general evolutionary analysis of tunnels and a detailed analysis of the selected cases. I was involved in writing the manuscript, preparation of figures, reviewing and editing the final version and in providing answers to the reviewers' comments.

Monia Browke







Dr **Karolina Mitusińska** PhD Research assistant

Gliwice, 10.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M*., Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins-What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 25%

I was involved in conducting a set of classical molecular dynamics simulations. I was involved in writing the manuscript and responding to the Reviewers' comments.

Yours sincerely,

Kerollue Mi'msinika









Agata Raczyńska MSc, BEng

20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 15%

I participated in performing the variability analysis and characterisation of selected tunnels.







Tomasz Skalski PhD, DSc

Gliwice, 13.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 5%

I designed the statistical analysis workflow.

A.









Aleksandra Samol, MSc

Email address: 2aleksandrasamol@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 5%
I was involved in the preparation and adjusting of the Multiple Sequence Alignment.

Signature

Gomol Helewar





Weronika Bagrowska MSc, BEng

Gliwice, 17.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 5%

I conducted molecular dynamics simulation for *Bacillus Megaterium* soluble epoxide hydrolase.

Buprowska Weracika



Tomasz Magdziarz, PhD

Email address: tomasz.magdziarz@gmail.com

Co-author contribution statement

Hereby, I certify that I am the Co-author of the following publication:

Bzówka M.*, Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins - What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25 https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 5% I was involved in preparing scripts for the tunnel comparison.

Signature
Tomon Wagolian





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M*., Mitusińska K.*, Raczyńska A., Skalski T., Samol A., Bagrowska W., Magdziarz T., Góra A.

Evolution of tunnels in alpha/beta-hydrolase fold proteins-What can we learn from studying epoxide hydrolases?

PLoS Computational Biology (2022) 18 (5), 1-25

https://doi.org/10.1371/journal.pcbi.1010119

I declare the following contribution to this publication: 15%

I conceived the project and contributed to its conceptualisation I reviewed the manuscript and participated in responding to the reviewers' comments. I was responsible for project administration and supervision.





Co-Author Contribution Statements for Paper 9

[9] Paper 9: Bzówka M., Bagrowska W., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods Journal of Chemical Information and Modeling (2023) 63 (12), 3669–3687 https://doi.org/10.1021/acs.jcim.3c00419





Maria Bzówka MSc, BEng PhD student

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Bagrowska W., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods

Journal of Chemical Information and Modeling (2023) 63 (12), 3669–3687

https://doi.org/10.1021/acs.jcim.3c00419

I declare the following contribution to this publication: 60%

I conceived the project and contributed to its conceptualisation. I performed the literature revision and organised all the data. I was involved in writing the manuscript, reviewing and editing the final version and in providing answers to the reviewers' comments. I was responsible for project administration and supervision.

Mana Brówke







Weronika Bagrowska MSc, BEng

Gliwice, 17.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Bagrowska W., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods

Journal of Chemical Information and Modeling (2023), 63 (12) 3669-3687

https://doi.org/10.1021/acs.jcim.3c00419

I declare the following contribution to this publication: 30%

I was involved in the literature search. I was involved in writing the manuscript and preparation of figures.

Bogravola Weracika







Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 09.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Bagrowska W., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods

Journal of Chemical Information and Modeling (2023) 63 (12), 3669–3687

https://doi.org/10.1021/acs.jcim.3c00419

I declare the following contribution to this publication: 10%

I contributed to the project's conceptualisation. I reviewed the manuscript.

Silesian University of Technology Biotechnology Centre



CONTROL HOTCHROLOGY
Polyclast Styles

ELING GROUP

Co-Author Contribution Statements for Preprint 1

[10] **Preprint 1: Bzówka M.**, Szleper K., Stańczak A., Borowski T., Góra A. The proteolytic cleavage of TLR8 Z-loop by furin protease - molecular recognition, reaction mechanism and role of water molecules

Research Square Preprint (2023)

https://doi.org/10.21203/rs.3.rs-3590328/v1





Maria Bzówka MSc, BEng PhD student

Gliwice, 15.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following publication:

Bzówka M., Szleper K., Stańczak A., Borowski T., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods

Preprint deposited in the Research Square repository

https://doi.org/10.21203/rs.3.rs-3590328/v1

I declare the following contribution to this preprint: 55%

I conceived the project and contributed to its conceptualisation. I performed analyses related to the prediction of the complex. I ran all molecular dynamics simulations for each reactions species. I did the analysis of the interaction network and the analysis of the reorganisation of water molecules using AQUA-DUCT software. I was involved in writing the manuscript, preparation of figures, ad data organisation. I was responsible for project administration and supervision.

Maria Brówka







Doctoral School Biotechnology Centre

MSc, BEng Katarzyna Szleper

Gliwice, 20.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following preprint:

Bzówka M., Szleper K., Stańczak A., Borowski T., Góra A.,

The proteolytic cleavage of TLR8 Z-loop by furin protease - molecular recognition, reaction mechanism and role of water molecules

Preprint deposited in the Research Square repository htps://doi.org/10.21203/rs.3.rs-3590328/v1

I declare the following contribution to this preprint: 20%

I performed the QM/MM calculations and was involved in the parametrisation of non-standard residues for MD simulations. I was involved in data analysis, figures preparation and writing the manuscript.

Katariyna Szleper





Agnieszka Stańczak, mgr inż. agnieszka.stanczak@uochb.cas.cz Institute of Organic Chemistry and Biochemistry of the Czech Academy of Sciences Faculty of Science, Charles University, Czech Republic

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following preprint:

Bzówka M., Szleper K., Stańczak A., Borowski T., Góra A.

The proteolytic cleavage of TLR8 Z-loop by furin protease - molecular recognition, reaction mechanism and role of water molecules

Preprint deposited in the Research Square repository https://doi.org/10.21203/rs.3.rs-3590328/v1

I declare the following contribution to this preprint: 15 %

I performed the QM-cluster calculations for the TLR8-furin complex, analysed and interpreted data.

Signature

Apriesolu Steviny





Prof. dr hab. Tomasz Borowski

Jerzy Haber Institute of Catalysis and Surface Chemistry, Polish Academy of Sciences Email address: tomasz.borowski@ikifp.edu.pl

Co-Author contribution statement

Hereby, I certify that I am the co-author of the following preprint:

Bzówka M., Szleper K., Stańczak A., Borowski T., Góra A.,

The proteolytic cleavage of TLR8 Z-loop by furin protease - molecular recognition, reaction mechanism and role of water molecules

Preprint deposited in the Research Square repository

https://doi.org/10.21203/rs.3.rs-3590328/v1

I declare the following contribution to this preprint: 5%

I provided the methodological support to QM and QM/MM calculations. I participated in reviewing the manuscript.

Signature

Vomesz Boronylo





Artur Góra PhD, DSc Head of Tunneling Group

Gliwice, 15.11.2023

CO-AUTHOR CONTRIBUTION STATEMENT

Hereby, I certify that I am the co-author of the following preprint:

Bzówka M., Szleper K., Stańczak A., Borowski T., Góra A.

Recent advances in studying Toll-like receptors with the use of computational methods

Preprint deposited in the Research Square repository

https://doi.org/10.21203/rs.3.rs-3590328/v1

I declare the following contribution to this preprint: 5%

I contributed to the project's conceptualisation. I reviewed the manuscript.



