Rashid Hussain

BS & MS, Bioinformatics

Ph.D. Computational Chemistry



Male, 35 years old, Pakistani national.



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https://rashid-bioinfo.github.io/

Research Experience

Jan 2022 – Apr 2022 Visiting Postgraduate Researcher – University of Manchester (UoM), UK

Project: VSpipe, an Integrated Resource for Virtual Screening and Hit Selection

Responsibilities: Successfully developed an open-source Python-based virtual screening

toolkit with a cross-platform user-interactive GUI interface

Jul 2016 – Dec 2019 Research Assistant – A.Z. Pharmaceuticals Company Limited, Pakistan

Focus: Research and management tasks

Responsibilities: Structure-based drug design of HCV NS3 protease genotype 3a

Ligand-based pharmacophore modeling

Coordination with universities for research facilitation

Monitoring of cGMP compliance and training of technical staff

Feb 2015 - Jan 2016 Research Associate - Biomedical Informatics Research Laboratory, Lahore University

of Management Sciences (LUMS), Pakistan

Projects: Higher Education Commission funded project on Hepatitis C Virus drug design

MATLAB-based toolbox for top-down proteomics data

Responsibilities: Ligand-based drug design

Designing GUI of top-down proteomics toolbox Lab management and compiling annual lab reports

Paper manuscript preparation

Jun 2014 – Jan 2015 Research Intern – National Center of Bioinformatics, Quaid-e-Azam University, Pak

Responsibilities: In-vivo testing on Zebrafish and hands-on experimental techniques

Use bioinformatics tools to find conserved regions in Zebrafish

Education

2017 - Present Ph.D. Chemistry – Forman Christian College (A Chartered University), Pakistan

> Thesis Title: Computer-aided drug design and synthesis of HCV NS3 protease inhibitors.

Supervisors: Dr. Hira Khalid and Dr. M. Qaiser Fatmi

2011 - 2013MS Bioinformatics - COMSATS University Islamabad, Pakistan

> Majors: Computer-Aided Drug Design (CADD)

Thesis Title: Rationalizing ligand-protein interactions and identifying cholinesterase

inhibitors using computational methods

Dr. M. Qaiser Fatmi Supervisor:

2006 - 2010BS Bioinformatics - COMSATS University Islamabad, Pakistan

> Majors: Phylogenetic analysis

Thesis Title: Phylogenetic analysis of major protein-coding genes of Geminiviridae:

A single-stranded DNA virus family

Supervisor: Dr. Muhammad Zeeshan Hyder

- 1. <u>Hussain, R.</u>, Khalid, H., Fatmi, M. Q. (2022). "HCV genotype-specific drug discovery through structure-based virtual screening." *Pure and Applied* Chemistry. DOI: 10.1515/pac-2021-1104. [Link]
- 2. <u>Hussain, R.</u>, Khalid, H., Fatmi, M. Q. (2021). "Molecular modelling approach of Serine Protease NS3-4A genotype 3a as a potential drug target of Hepatitis C Virus: Homology Modelling and Virtual Screening Study." *J. Comput. Biophys. Chem.*, Vol. 20, No. 06, pp. 631-639. [Link]
- 3. Khalid, H., <u>Hussain, R.</u>, & Hafeez, A. (2020). "Virtual screening of piperidine-based small molecules against COVID-19". *Lab-in-Silico*, Vol. 01, No. 02, pp. 50-55. DOI: 10.22034/lins20012050. [Link]
- Basharat, A. R., Iman, K., Bibi, Z., <u>Hussain, R.</u>, Kabir, H. G., Shahid, A., Humayun, M., Hayat, H. A., Mustafa, M., Shoaib, M. A., Ullah, Z., Zarina, S., Ahmed, S., Uddin, E., Hamera, S., Ahmad, F., & Chaudhary, S. U. (2019). "SPECTRUM A MATLAB toolbox for proteoform identification from top-down proteomics data." *Scientific Reports Nature*, Vol. 09, Issue 01, pp. 1 -14. DOI: 10.1038/s41598-019-47724-1. [Link]
- Ashraf, M. U., Iman, K., Khalid, M. F., Shafi, T., Salman, H. M., Rafi, M., Javaid, N., <u>Hussain</u>, <u>R</u>., Ahmad, F., Shahzad-ul-Hussan, S., Mirza, S., Shafiq, M., Afzal, S., Idrees, M., Hamera, S., Anwar, S., Qazi, R. Qureshi, S. A., Chaudhary, S. U. (2019). " Evolution of efficacious pangenotypic Hepatitis C Virus therapies." *Medicinal Research Reviews*, Vol. 39, No. 03, pp. 1091-1136. DOI: 10.1002/med.21554. [Link]
- Arfan, M., Siddiqui, S.Z., Abbasi, M.A., ur Rehman, A., Shah, S.A.A., Ashraf, M., Rehman, J., Saleem, R. S. Z., Khalid, H., <u>Hussain, R.</u>, Khan, U. (2018). "Synthesis, in vitro and silico studies of S-alkylated 5-(4-methoxyphenyl)-4-phenyl-4H-1, 2, 4-triazole-3-thiols as cholinesterase inhibitors." *Pak. J. Pharm. Sci*, Vol. 31, No. 6, pp. 2697-2708. [Link]
- 7. Khalid, H., Abbasi, M. A., <u>Hussain, R.</u>, Malik, A., Ashraf, M., & Fatmi, M. Q. (2017). "Synthesis, spectral analysis and biological evaluation of 5-Substituted 1,3,4-oxadiazole-2-yl-4-(piperidin-1-ylsulfonyl)benzyl sulfide." *Emerging Trends in Chemical Sciences*, Chapter-14 in Springer books, pp. 221-238. DOI: 10.1007/978-3-319-60408-4_14. [Link]
- 8. Mumtaz, S., <u>Hussain, R.</u>, Rauf, A., Fatmi, M. Q., Bokhari, H., Oelgemöller, M., & Qureshi, A. M. (2014). "Synthesis, molecular docking studies, and in vitro screening of barbiturates/thiobarbiturates as antibacterial and cholinesterase inhibitors." *Medicinal Chemistry Research*. Vol. 23, No. 06, pp. 2715-2726. DOI: 10.1007/s00044-013-0847-2. [Link]
- Khalid, H., Rehman, A. U., Abbasi, M. A., <u>Hussain, R.</u>, Khan, K. M., Ashraf, M., Ejaz, S.A., & Fatmi, M. Q. (2014). "Synthesis, biological evaluation, and molecular docking of N'-(Aryl/alkylsulfonyl)-1-(phenylsulfonyl) piperidine-4-carbohydrazide derivatives." *Turkish Journal of Chemistry*. Vol. 38, No. 02, pp. 189-201. DOI: 10.3906/kim-1303-89. [Link]

Manuscripts in Preparation

- 1. <u>Hussain, R.</u>, Hackett, A., Álvarez-Carretero, S., Khalid, H., Tabernero, L. (2022). "Vspipe 2.0, an integrated resource for virtual screening and hit selection with a graphical user interface."
- 2. <u>Hussain, R.</u>, Khalid, H., Fatmi, M. Q. (2022). "Computer-aided drug design and synthesis of HCV NS3 protease inhibitors".

Published Software

1. Basharat, A.R., Bibi, Z., Hussain, R., Kabir, H. G., Shahid, A., Humayun, M., Hayat, H. A., Mustafa, M., Shoaib, M. A., Ullah, Z., Zarina, S., Ahmed, S., Uddin, E., Hamera, S., Ahmad, F., & Chaudhary, S. U. SPECTRUM: A MATLAB Toolbox for Identifying Proteins from Top-down Proteomics Data, [Link]

Conferences Webinar/Talks/Posters

- 1. <u>Hussain, R.</u>, Khalid, H. (2020). "References made easy using Mendeley". Forman Christian College (A Charted University), Pakistan Webinar.
- 2. <u>Hussain, R.</u>, Khalid, H., Fatmi, M.Q. (2021). "Molecular Modelling Approach of Serine Protease NS3-4A Genotype 3a as a Potential Drug Target of Hepatitis C Virus". Virtual Conference on Chemistry and its Applications. The University of Mauritius. Mauritius Talk.
- 3. <u>Hussain, R.</u>, Khalid, H. (2018). "HCV genotype-specific drug discovery through Structure and Ligand-based Virtual Screening." 6th International Bau Drug Design Congress. Bahcesehir University School of Medicine, Istanbul, Turkey Talk.
- Khalid, H., ur Rehman, A., Ahmad, I., Arshad, S., Nadeem, M.H., <u>Hussain, R</u>. (2018). Synthesis of Potential Antibacterial Agents Derived from 5-[1-(Phenylsulfonyl) piperidin-4-yl]-1,3,4-oxadiazol-2-thiol. "International Conference on Chemical and Pharmaceutical Sciences." Forman Christian College (A Charted University), Lahore, Pakistan Talk.
- 5. Basharat, A.R., Bibi, Z., <u>Hussain, R.</u>, Kabir, H. G., Shahid, A., Humayun, M., Hayat, H. A., Mustafa, M., Shoaib, M. A., Ullah, Z., Zarina, S., Ahmed, S., Uddin, E., Hamera, S., Ahmad, F., & Chaudhary, S. U. (2017)." SPECTRUM: A MATLAB Toolbox for Identifying Proteins from Top-down Proteomics Data."16th Annual Human Proteome Organization World Congress (HUPO), Dublin, Ireland Talk.
- Abubakar, M., Bibi, A., <u>Hussain, R.</u>, Bibi, Z., Gul, A., Bashir, Z., Arshad, S. N., Uppal, M. A., & Chaudhary, S. U. (2016). "Towards Providing Full Spectrum Antenatal Health Care in Low and Middle-Income Countries." 9th International Joint Conference on Biomedical Engineering Systems and Technologies (BIOSTEC 2016), Vol 5 (HEALTHINF), pp. 478-483, 2016, Rome, Italy Talk.
- 7. <u>Hussain, R.</u>, Kabir, G. H., Chaudhary, S. U. (2015). "Towards an Accurate Measurement of Intact Protein Mass in High-Resolution Mass Spectrometry." 3rd National Computational Science Conference (NCSC), Islamabad, Pakistan. (2nd prize in project presentation competition) Poster.
- 8. <u>Hussain, R.</u>, Fatmi, M. Q. (2013). "Identification of Prospective Cholinesterase Inhibitors using Structure-Based Virtual Screening Approach." Poster presented at Chemistry Department, COMSATS Institute of Information Technology, Abbottabad, Pakistan Poster.

Projects

1. Machine Learning QSAR model for COVID19 Replicase Polyprotein to predict pIC50 of a given compound – Successfully developed and deployed. https://github.com/rashid-bioinfo/BioPredict

Honors and Awards

2021 – 22	International Research Support Initiative Program (IRSIP), \$9800 for six months by Higher Education Commission of Pakistan
2015	Best Poster Award, by Institute of Space Technology, Pakistan
2015	Marathon Race Winner, Lahore University of Management Science (LUMS), Pakistan

Professional Skills

Computer Programing: Python, Bash Scripting, GitHub (for version control) – Expert level

R, MATLAB, SQL – Proficient

C++, JAVA, LaTeX, Django, Bootstrap – Intermediate level

Operating System: Linux, Mac, Windows

Computational Chemistry: Docking – AD4, Autodock Vina, UCSF DOCK6

MD Simulations - GROMACS

Modeling – MODELLER, SWISS-MODEL

Visualization/Interactions – PyMOL, MOE, VMD, UCSF Chimera, BIOVIA DS

Other - Schrödinger Suite

Data Science &: Machine Learning Scikit-learn, Pandas, NumPy, KNIME, Streamlit, RDKit (Cheminformatics)

Teaching Experience

Aug 2021 – Present Voluntary Teaching Assistant – The Carpentries, California, USA. Remote

Course: High-Performance Computing (HPC) **Level**: Undergraduate

Responsibilities: Voluntary helper to mentor the learners and answer their questions

Mar 2019 – Feb 2020 Visiting Faculty – Minhaj University, Pakistan

Course: Bioinformatics **Level**: Undergraduate

Responsibilities: Designed and presented 30 lectures each to different batches

Conducted 16 labs of the course

Wrote and graded exams

Feb 2015 – Jan 2016 Teaching Assistant – Lahore University of Management Sciences (LUMS), Pakistan

Courses: Computational Biology and Protein Informatics Level: Undergraduate

Responsibilities: Conducted 16 labs, wrote and graded lab exams – Comp. Biology

Conducted 12 labs, wrote and graded lab exams – Protein Informatics

Graded class assignments for both courses

Supervisory and Mentoring Experience

Supervised four undergraduate summer intern students

Mentoring undergraduate/graduate research students in theses writeup/presentations

Voluntary Academic Service

As a team lead, trained teachers, students, and parents for taking online classes using Google Classroom, Google Meet, Zoom, etc. during COVID19

Recorded online tutorials on YouTube Channel, Computational Chemist on computer-aided drug design and for general public awareness, like traffic signs and rules

Memberships

American Chemical Society (ACS Member Number - 33206538)

Journal Reviewer

Springer - Molecular Diversity

Professional Courses and Certifications

2022	The Unix Workbench, Johns Hopkins University – Completed
2022	What is Data Science?, IBM – Completed
2022	Machine Learning, Datacamp – In progress
2022	Data Science, Online course — Completed
2021	Python, Online course, 1100 minutes – Completed
2020	Introduction to Molecular Modeling in Drug Discovery, Schrödinger online course, 1 month
2019	Computer-Aided Drug Design, NPTEL India online course, 900 minutes