

# Mohamed Marzouk Sobaih, Ph.D.

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🔗 <https://github.com/mohamedmarzouk22>



## Career Progression

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| April 2024 – present    | 📌 <b>Specially Appointed Researcher</b> , WPI Premium Research Institute for Human Metaverse Medicine (WPI-PRIME), Osaka University, Osaka, Japan.          |
| April 2023 – March 2024 | 📌 <b>Lecturer</b><br>Physics Department, Faculty of Science, Ain Shams University, Cairo, Egypt.  |
| April 2022 – March 2023 | 📌 <b>Postdoctoral researcher, Molecular Modeling and Simulation (MMS) Team</b> , National Institute for Quantum Science and Technology (QST), Chiba, Japan. |
| Dec. 2016 – April 2019  | 📌 <b>Senior Teaching/Research Assistant</b><br>Physics Department, Faculty of Science, Ain Shams University, Cairo, Egypt.                                  |
| March 2010 – Dec. 2016  | 📌 <b>Junior Teaching/Research Assistant</b><br>Physics Department, Faculty of Science, Ain Shams University, Cairo, Egypt.                                  |

## Education

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| April 2019 – March 2022    | 📌 <b>Ph.D., School of Life Science and Technology, Tokyo Institute of Technology</b> in Computational Biology.<br>Thesis title: <i>Investigating Dissociation Process and Binding Free Energy of P53-DBD/DNA Complex by PaCS-MD and MSM.</i> |
| June 2012 – October 2016   | 📌 <b>M.Sc., Faculty of Science, Ain Shams University</b> in Biophysics.<br>Thesis title: <i>Genetic and Physiological Effects of Ultraviolet Radiation on Tomato Plant (solanum lycopersicum).</i>   |
| September 2005 – June 2009 | 📌 <b>B.Sc., Faculty of Science, Ain Shams University</b> in Biophysics.  |

## Research Publications

### Journal Articles


- 1 Sobeh, M. M., & Kitao, A. (2022). Dissociation pathways of the p53 DNA binding domain from DNA and critical roles of key residues elucidated by dPaCS-MD/MSM. *Journal of Chemical Information and Modeling*, 62(5), 1294–1307. 🔗 doi:10.1021/acs.jcim.1c01508
- 2 Hata, H., Tran, D. P., Sobeh, M. M., & Kitao, A. (2021). Binding free energy of protein/ligand complexes calculated using dissociation parallel cascade selection molecular dynamics and markov state model. *Biophysics and Physicobiology*, 18(0), 305–316. 🔗 doi:10.2142/biophysico.bppb-v18.037

### Conference Proceedings


- 1 Sobeh, M. M., Kono, H., & Sakuraba, S. (2023.2.18). Predicting the Salt Bridges Contribution to Protein Stability using the Free-Energy Perturbation and Enhanced Sampling Approach. In *Proceedings of the 67th Annual Meeting of the Biophysical Society*, Poster, San Diego, California, US.

- 2 Sobeh, M. M. (2021.9.16). Investigating the dissociation process and binding free energy of the p53-DBD/DNA complex by PaCS-MD and MSM. In *Proceedings of The Third International Online Conference on Molecular Modeling and Spectroscopy*, Short presentation, Online Conference, Giza, Egypt. Retrieved from <https://www.youtube.com/watch?v=wroOofDsxos>
- 3 Sobeh, M. M., & Kitao, A. (2021.4.28). Investigating the dissociation process and binding energy of the p53-DBD/DNA complex by PaCS-MD and MSM. In *Proceedings of WE-Heraeus-Seminar, Advanced Physical and Computational Techniques to Investigate Protein Dynamics*, Poster, Online Conference via MeetAnyway, Freie Universität Berlin, Germany. Retrieved from <https://shortest.link/2mnF>
- 4 Sobeh, M. M., & Kitao, A. (2021.6.16). Investigating the dissociation process and binding free energy of the p53-DBD/DNA complex by PaCS-MD and MSM. In *Proceedings of the 21st Annual Meeting of the Protein Science Society of Japan*, Poster, Online Conference, Tokyo, Japan.





## Invited Talks / Presentations

Feb., 2025  **Accelerating Biomolecular Simulations with Enhanced Sampling Techniques.** 2nd Serial Networking of Foreign Researchers: Sysmex - Osaka University, Osaka, Japan. (10-minute Presentation)







## Grants

March, 2023  **Early-Career Scientists, KAKENHI, Japan, (Principal Investigator).** Molecular Dynamics Simulation for Association/Dissociation of Protein-Ligand and Protein-DNA Complexes by Advanced Enhanced Sampling Technique, **3 years (5 000 000 Yen.)**

## Research Skills

MD Simulation	 Molecular dynamics (MD) simulations for protein-DNA, Protein-Protein complexes.
Enhanced sampling MD	 Enhanced sampling MD simulations using techniques such as using PaCS-MD, REMD, gREST, and REUS.
Binding Free energy	 Binding Free energy predictions using MSM, FEP, and MMBPSA.
Protein predictions	 Protein and peptide structure predictions using Modeller, SWISS-MODEL, and AlphaFold.




## Other Skills

Languages	 English: strong reading, writing, and speaking. Japanese: basic reading, writing, and speaking (JLPT N5).
Coding	 Python, Bash scripting, Tcl, C++, LaTeX, Git.
MD Packages	 AMBER, GROMACS, GENESIS, Autodock vina, Rosetta, PyRosetta, Modeller.
MD Analysis	 VMD, PyMOL, ChimeraX, PLIP, PyEMMA, Mdtraj, MDAAnalysis.
Machine Learning	 SciKit Learn, TensorFlow, Keras, PyTorch.
Cheminformatics	 RDKit.

## Miscellaneous Experience







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### Awards and Achievements

- 2021 – 2022     **Cross the border**, Tokyo-Tech pioneering doctoral research program by Tokyo Institute of Technology.
- 2019 – 2021     **Tokyo Tech Tsubame Scholarship**, Tokyo Institute of Technology.
-  **Ph.D. Scholarship**, The Egypt-Japan Education Partnership (EJEP).
- 2010             **Medal of Superiority**, Faculty of Science, Ain Shams University, Cairo, Egypt.

### Workshops, Training and Online Courses

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- June, 2024         **IUPAB 2024 Hands-on Training Program**, CHARMM-GUI/GENESIS MD Tutorial, RIKEN, Kobe, Japan.
- January, 2024     **Deep Learning Specialization**, Stanford ONLINE & DeepLearning.AI, Coursera Platform (Prof. Andrew Ng).
- November, 2023     **Machine Learning Specialization**, Stanford ONLINE & DeepLearning.AI, Coursera Platform (Prof. Andrew Ng).
- February, 2022     **Fundamentals of Deep Learning**, NVIDIA Deep Learning Institute.
- June, 2021         **BioExcel Summer School on Biomolecular Simulations**, Centre of Excellence for Computational Biomolecular Research.
- March, 2021        **Introduction to Simulation Environments for Life Sciences**, Online PATC@BSC Training Course.