



DINESH KUMAR SRIRAMULU

Postdoctoral Researcher

PROFILE

Hardworking Postdoctoral Research fellow with demonstrated record of research success in. Self-directed and motivated professional with a highly ethical, meticulous and thorough approach.

CONTACT

PHONE:
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EDUCATION

Pusan National University

2017 – 2021 Ph.D. Chemical Engineering

- Thesis: Study on the Effect of Protein-Ligand properties on the Prediction Accuracy of a Molecular Docking Tool, AutoDock.
- GPA 4.17/4.5

Pondicherry University

2014 – 2016 M. Tech Computational Biology

- Thesis: Identification of potential BACE1 inhibitor – an *in-silico* analysis
- GPA 8.67/10

SASTRA University

2009 – 2013 B. Tech Bioinformatics

- Thesis: Analysis of temperature factor distribution in ultra-high-resolution protein structures
GPA 7.27/10

WORK EXPERIENCE

POSTDOCTORAL RESEARCHER, KOREA INSTITUTE OF SCIENCE AND TECHNOLOGY

OCTOBER 2024 – PRESENT

POSTDOCTORAL RESEARCHER, KOREA INSTITUTE OF SCIENCE AND TECHNOLOGY

MARCH 2023 – FEBRUARY 2024

- Devised and carried out research initiatives to investigate challenges in the context of Target Screening.
- Collaborated with multiple research groups to work on interdisciplinary projects.

POSTDOCTORAL RESEARCHER, PUSAN NATIONAL UNIVERSITY

MARCH 2021 – FEBRUARY 2023

- Conducted independent computational research and development to attain short- and long-term objectives.
- Contributed to and actively participated in research conception, design, and execution to address defined problems.

PUBLICATIONS

- G Munussami, S Sokalingam, **DK Sriramulu**, SG Lee., Identification of common and distinct features of ligand-binding sites in kernel and outlier lipocalins. *Journal of Industrial and Engineering Chemistry*, **2019**, 78, 344-351.
- **Sriramulu, D. K.**; Wu, S.; Lee, S.-G., Effect of ligand torsion number on the AutoDock mediated prediction of protein-ligand binding affinity. *Journal of Industrial and Engineering Chemistry* **2020**, 83, 359-365
- **Sriramulu, D. K.**; Lee, S.-G., Combinatorial Effect of Ligand and Ligand-Binding Site Hydrophobicities on Binding Affinity. *Journal of Chemical Information and Modeling* **2020**, 60 (3), 1678-1684.
- P Jebamani, S Sokalingam, **DK Sriramulu**, ST Jung, SG Lee., Assessment of Computational Modeling of Fc-Fc Receptor Binding Through Protein-protein Docking Tool. *Biotechnology and Bioprocess Engineering*, **2020**, 25 (5), 734-741
- **Sriramulu, D. K.**; Lee, S.-G., Effect of molecular properties of the protein-ligand complex on the prediction accuracy of AutoDock. *Journal of Molecular Graphics and Modeling* **2021**, 106.

- Jebamani, P., **Sriramulu, D.K.**, Jung, S.T. *et al.* Structural Study on the Impact of S239D/I332E Mutations in the Binding of Fc and FcγRIIIa. *Biotechnol Bioproc E* **26**, 985–992 (2021).
- Hyeona Kang, Dinesh Kumar Sriramulu, Sun-Gu, Lee In silico Study on Binding Specificities of Cellular Retinol Binding Protein and Its Q108R Mutant. *Biotechnol Bioproc E* **27**, 126-134 (2022).
- Jebamani, P., **Sriramulu, D.K.**, Lee, S.-G., Residue interaction network and molecular dynamics simulation study on the binding of S239D/I332E Fc variant with enhanced affinity to FcγRIIIa receptor. *Journal of Molecular Graphics and Modeling* **2023**, 118
- **Dinesh Kumar Sriramulu**, Sun-Gu Lee. Combinatorial Effect of Ligand Aromaticity, Torsion Number and Hydrophobicity on the Predictability of AutoDock for Protein-Ligand Binding Conformation. *KSBB Journal*, 38 (1), 70-75 (2023).
- **Dinesh Kumar Sriramulu**, Sun-Gu Lee, Analysis of protein-protein interface with incorporating low-frequency molecular interactions in molecular dynamics simulation, *Journal of Molecular Graphics and Modelling*, Volume 122, 2023,108461
- InWha Park, Keunwan Park, Hyeon-Seong Lee, Seong-Min Hong, **Dinesh Kumar Sriramulu**, Hoseong Hwang, Sungmin Song, Jong Gwon Baek, Dong Hoon Kim, Sun Yeou Kim, Jaeyoung Kwon, Hak Cheol Kwon. Ecdysteroids from the Korean Endemic Species *Ajuga spectabilis* with Activities against Glucocorticoid Receptors and 11β-Hydroxysteroid Dehydrogenase Type 1, *ACS Omega* 2023 8 (29), 26191-26200.

SKILLS

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|-------------------------------------|----------------------|
| • Biomolecular interaction analysis | • Perl |
| • Computer-Aided drug design | • C++ |
| • Drug-Target prediction | • R programming |
| • Python | • MATLAB |
| | • Scientific Writing |

REFERENCES

1. Dr. Pavlo Polishchuk,
Senior Researcher, Institute of Molecular and Translational
Medicine, Faculty of Medicine and Dentistry,
Palacky University, Olomouc,
Czech Republic
E-mail: pavlo.polishchuk@upol.cz

2. Dr. Sun-Gu, Lee

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Engineering, Pusan National University, Busan,

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