Jordy Homing Lam

 ${\hbox{Ho-Ming-Lam}} \\ \hbox{University of Southern California, CA 90089}$

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High Performance Computing • Informatics • Geometric Deep Learning • Structural Biology

EDUCATION

University of Southern California

2019 - Present

Ph.D. Candidate Bioinformatics

Hong Kong University of Science and Technology

2017

M.Phil. Chemistry

The Chinese university of Hong Kong

2015

B.S.(Hons) Chemistry

SERVICE

Teaching | Methods in Computational Physics · Structural Bioinformatics · Medicinal Chemistry

Reviewer | Computer Physics Communications

AWARD

Dornsife Graduate Fellowship | University of Southern California

2019 - Present

• Ph.D. Funding

Sir David Trench Scholarship | The Chinese University of Hong Kong (1 per whole student body)

2015

• A scholarship "awarded to a final-year student in recognition of their all-round leadership".

Grace Chiu Po Yuen Memorial Prize | The Chinese University of Hong Kong (1 per class)

2015

 $\bullet\,$ Academic award for Biochemistry or Chemistry student.

Ho Man Sum Distinguished College Service Award | The Chinese University of Hong Kong (1 per college)

• Awarded to an achiever of "a balanced development in academic pursuit and contributions to the United College".

Diploma, Violin | The Royal Academy of Music, United Kingdom

2008

2012

• Passed with Distinction.

MISCELLANY

Languages: English \cdot Cantonese \cdot Mandarin \cdot French \cdot Beginning Japanese

Programming: Python · Cython · C · CUDA · Fortran90 · Matlab · Bash · Slurm · I 4 T_FX

Music: Instrumental · Polyphony

Publication

† indicates co-first authorship.

Scalable anisotropic vibrations of megascale macromolecules

Lam JH, Nakano A, Katritch V

Nature Commun. 2024 In Review, Revised.

Nimrodd: Enhancing multi-resolution geometric neural hierarchy with an expectation maximization kernel

IN PREPARATION 2024

Lam JH, Ferrari T, Liu Y, Nakano A, Katritch V

Structural Insights into Inverse Agonism of an Orphan Receptor

SUBMITTED 2024
In Review.

Barekatain M^{\dagger} , Johansson L^{\dagger} , Lam JH^{\dagger} , Chang H, Sadybekov A, Han GW, Russo J, Bliesath J, Brice N, Carlton M, Saikatendu K, Murphy S, Monenschein H, Schiffer H, Popov P, Lutomski C, Robinson CV, Liu ZJ, Hua T, Katritch V, Cherezov V

Constitutive activation mechanism of a class C GPCR NA	T. STRUCT. MOL. BIOL. 2024
Shin J, Park J, Jeong J, <u>Lam JH</u> , Qiu X, Wu D, Kim K, Lee J, Robinson CV, Hyun J, Katritch V, Kim KP, Cho Y	In Press.
Structure of the dopamine D3 receptor bound to a bitopic agonist reveals a	Nat. Commun. 2024
new specificity site in an expanded allosteric pocket Arroyo-Urea S, Nazarova AL, Carrión-Antolí A, Bonifazi A, Battiti FO, <u>Lam JH</u> ,	$In \ Review.$
Newman AH, Katritch V, García-Nafría J	
Ligand and G-protein selectivity in the κ -opioid receptor	Nature 2023
Han J, Zhang J, Nazarova AL, Bernhard SM, Krumm BE, Zhao L, <u>Lam JH</u> , Rangari VA, Majumdar S, Nichols DE, Katritch V, Yuan P, Fay JF, Che T	[DOI]
Structural details of a Class B GPCR-arrestin complex revealed by	Nat. Commun. 2023
genetically encoded crosslinkers in living cells	
Aydin Y [†] , Böttke T [†] , <u>Lam JH</u> [†] , Ernicke S, Fortmann A, Tretbar M, Zarzycka B, Gurevich VV, Katritch V, Coin I	[DOI]
Structural basis of GABA reuptake inhibition	Nature 2022
Motiwala Z, Aduri NG, Shaye H, Han GW, Lam JH, Katritch V, Cherezov V, Gati C	[DOI]
Structure of the full-length human Pannexin1 channel and insights into its	Cell Discov. 2021
role in pyroptosis	(= 0 =1
Zhang S, Yuan B, <u>Lam JH</u> , Zhou J, Zhou X, Ramos-Mandujano G, Tian X, Liu Y, Han R, Li Y, Gao X, Li M, Yang M	[DOI]
Structural basis of the activation of a metabotropic GABA receptor	Nature 2020
Shaye H, Ishchenko A, <u>Lam JH</u> , Han GW, Xue L, Rondard P, Pin JP, Katritch V,	[DOI]
Gati C, Cherezov V	
Self-assembling tetrameric peptides allow in situ 3D bioprinting under	J. Mater. Chem. B. 2019
physiological conditions	
Rauf S, Susapto HH, Kahin K, Alshehri S, Abdelrahman S, <u>Lam JH</u> , Asad S, Jadhav S, Sundaramurthi D, Gao X, Hauser CAE	[DOI]
A deep learning framework to predict binding preference of RNA	Nat. Commun. 2019
constituents on protein surface	TVAT. COMMON. 2013
Lam JH, Li Y, Zhu L, Umarov R, Jiang H, Héliou A, Sheong FK, Liu T, Long Y, Li Y, Fang L, Altman RB, Chen W, Huang X, Gao X	[DOI]
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