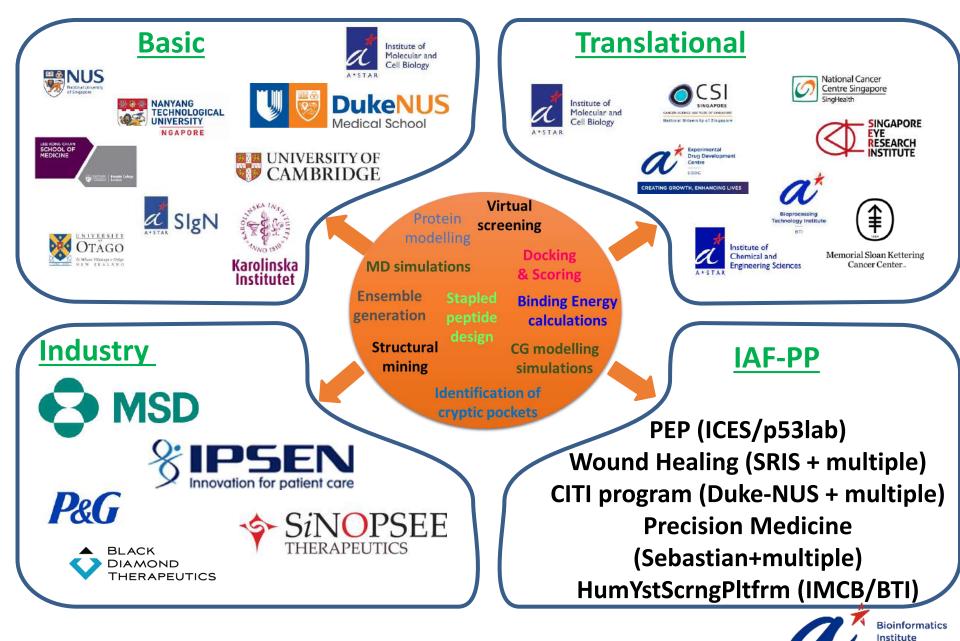
Atomistic Simulations & Design



BII PI seminars April 2021



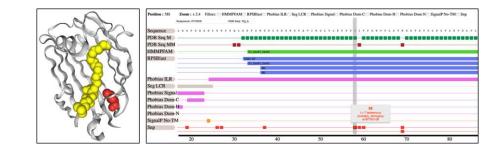
Understanding mechanism, designing molecules as probes/inhibitors/therapeutics



BII

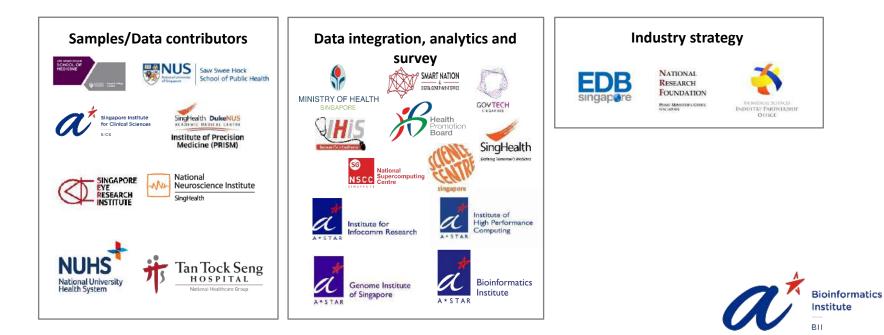


National Precision Medicine: SNPDrug3D



Phase I

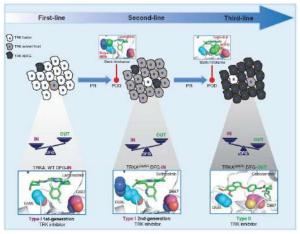
SNP database to help clinicians to make more informed decision.



Translation into clinic

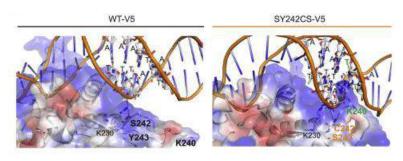


xDFG Mutations Trigger a Sensitivity Switch from Type I to II Kinase Inhibitors



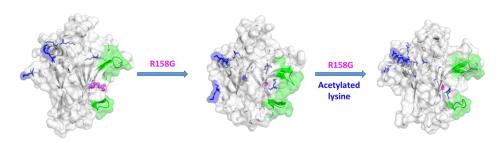
Cancer Discovery, 11, 126 (2021)

FOXA1 Mutations Influence Therapeutic Response in Breast Cancer

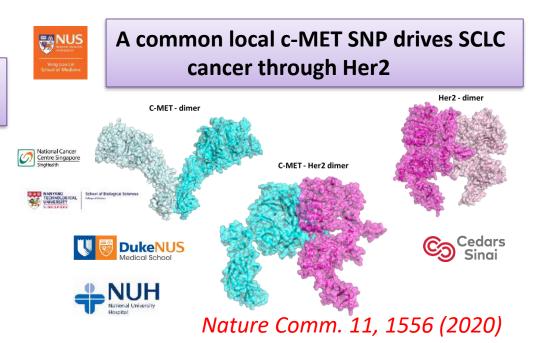


Cancer Cell. 38,534 (2020)

Acetylation reverses a locally observed mutant p53 and restores its activity

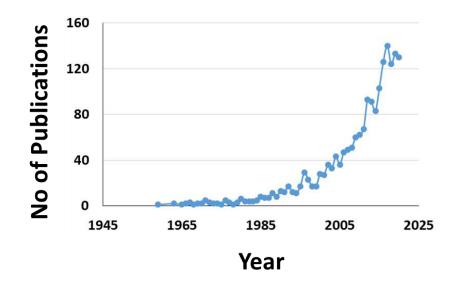


Nature Comm. 11, 2086 (2020)



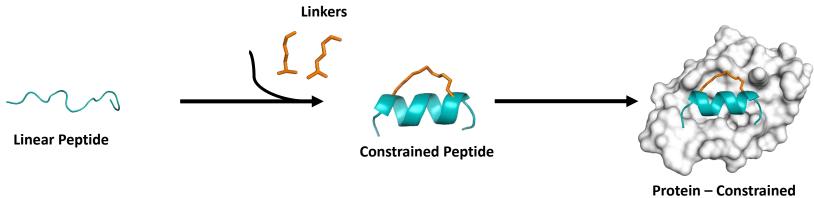
Peptide (stapled) Design – Basic & Industry

A new kid in town



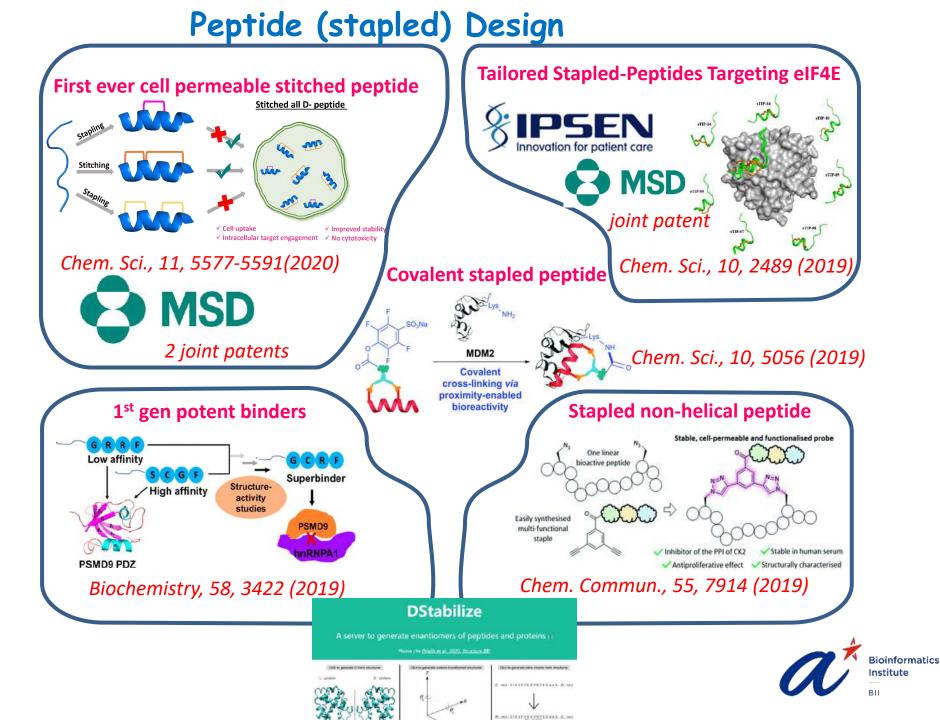
Raghav, Pietro, Shilpa, Jianguo, Minh + Yaw sing, Sonia, Shruti, Minh, Aishwary, Ashar



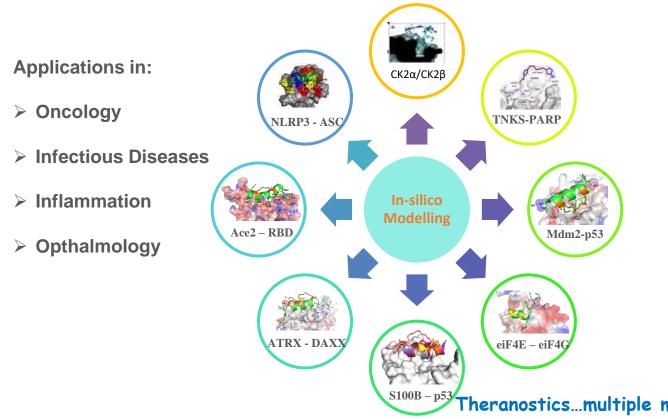


Peptide complex

	Small	Peptides			Biologics
	Molecules	Linear	Cyclic	Stapled	
Potency/specificity/toxicity		++	+++	++++	++++
Metabolic stability		+	++	++++	++++
Acquired Resistance		++++	++++	++++	+++
Inhibition of PPI*		+++	++++	++++	++++
Tissue penetration	++++	+	+	++++	
Immunogenicity	++++	++	++++	++++	
Patient compliance	++++			++++	
FDA approved drugs	2500	49	11	0	250



Some examples....



> >60 publications (JACS,PNAS, ACS Chem Biol, Chem. Sci, Angew. Chem Int Ed., NRDD, DDT)

- > 10 patents filed/granted/process
- Pharma collaborations (lpsen, MSD)
- Developed several first in class molecules

*Theranostics...multiple myeloma novel approach..NUH

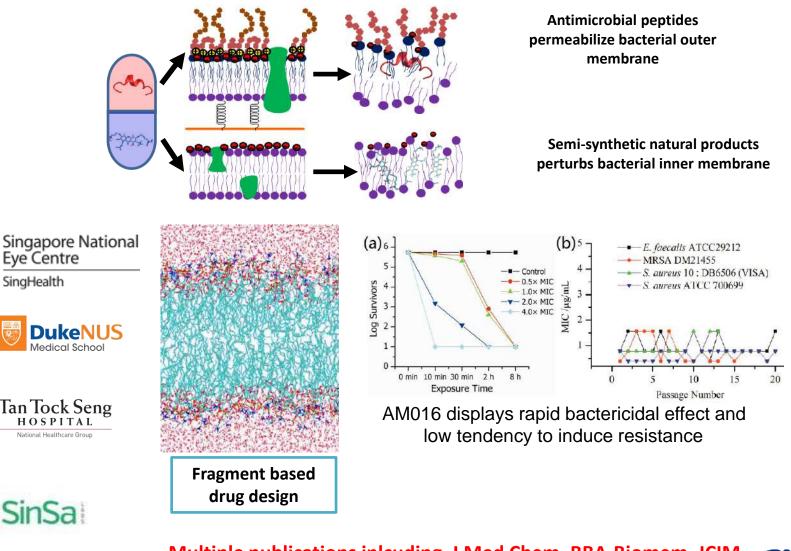
Stratification of immunotherapy patients: CITI program pan Singapore

Shruti, Raghav



Jianguo

Combating antimicrobial resistance: Superbugs succumb to combination therapy

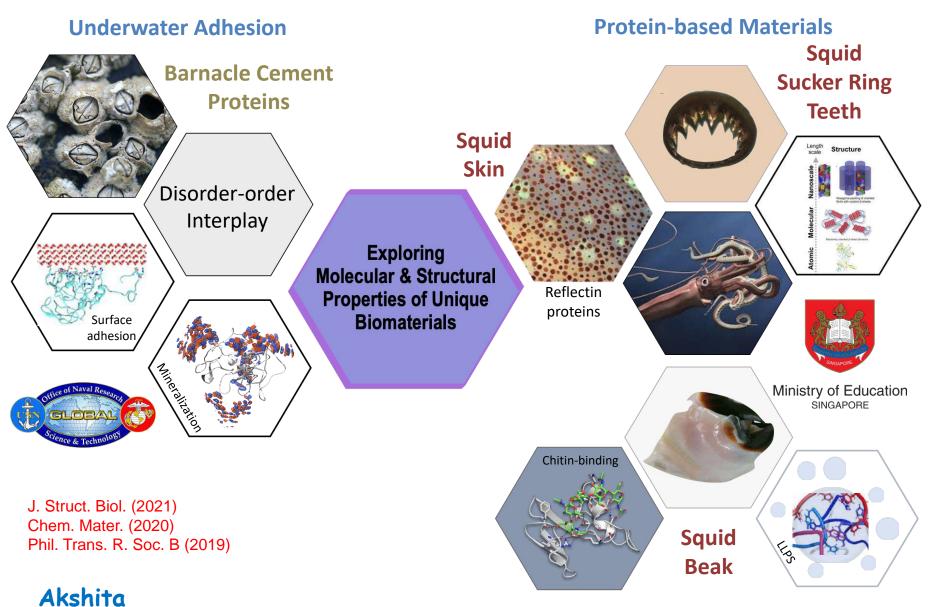


A patents A patent A patent



Computational Biomaterial Science

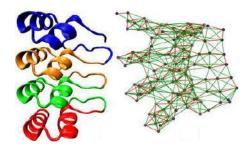




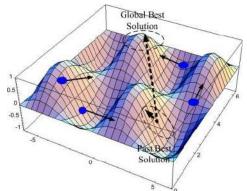




Exploring Applications of Quantum Computing to understand Biomolecular Mechanisms

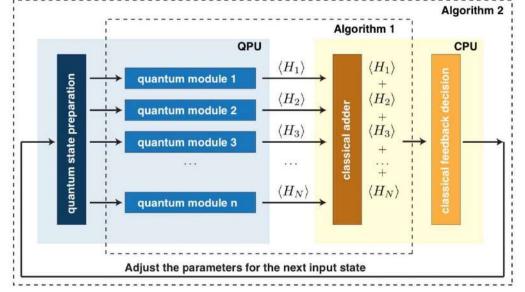


Protein Structure depicted as a network, comprising nodes and edges.



Protein-ligand interaction visualized as a optimization problem.

By recasting certain biological problems existing hybrid Quantum-Classical computing methods can be used on near-term noisy quantum computers to enhance our understanding of



Use of hybrid Quantum and Classical computing, in VQE to address problems solvable by near-term noisy quantum computers.

Thank You





