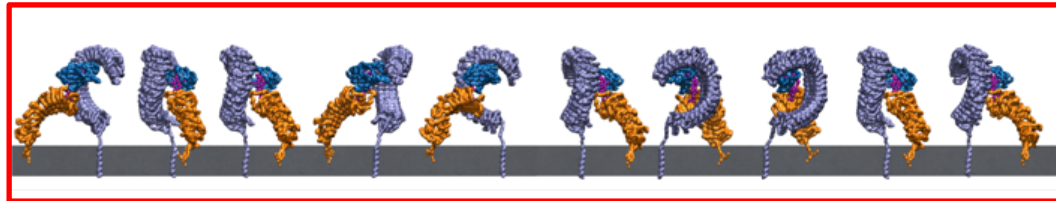
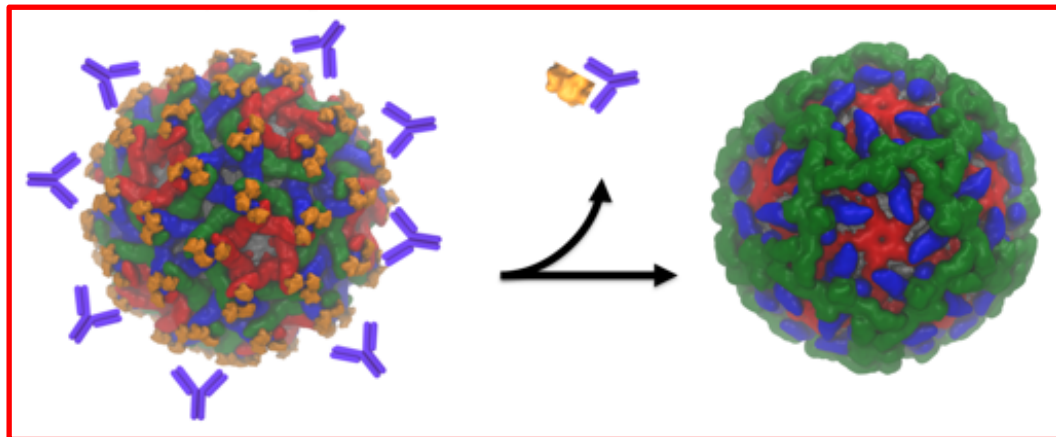
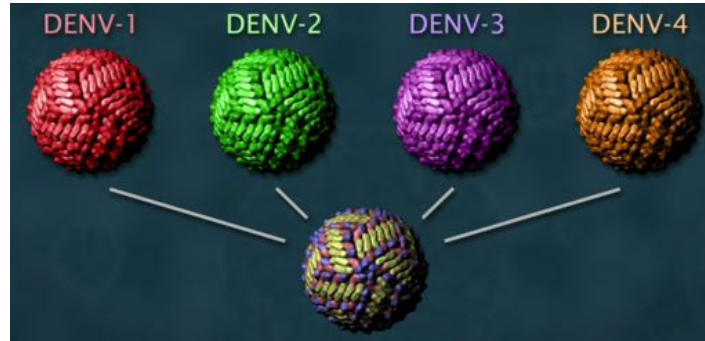
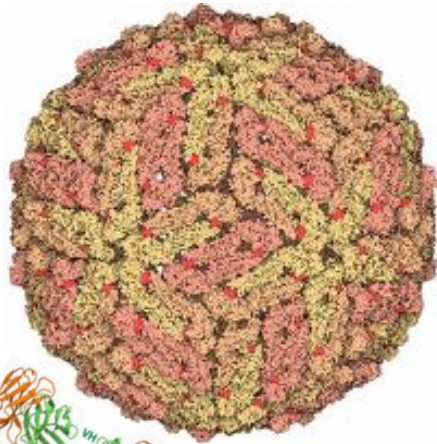


Multiscale Simulation, Modelling & Design: Infectious Diseases and the Host Response

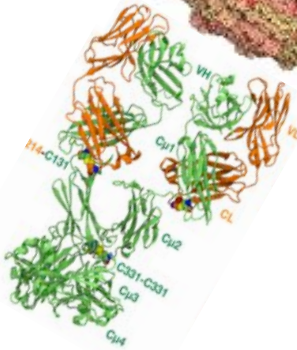
Peter J. Bond



Research Focus: Host-Pathogen Interactions



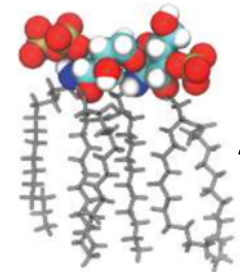
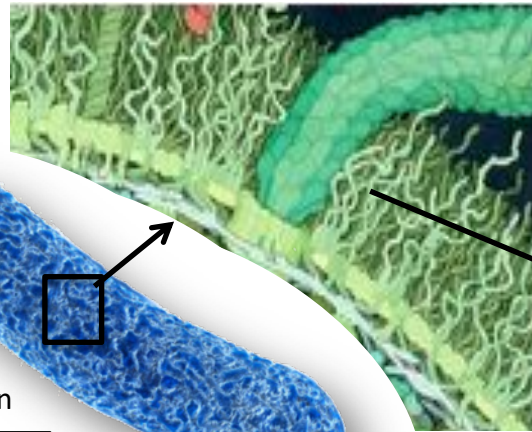
Dengue: flavivirus with lipid bilayer & envelope proteins



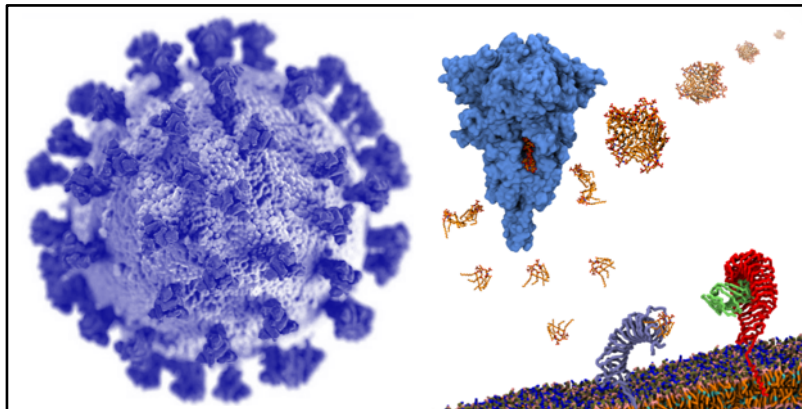
Viral envelope dynamics, antivirals & antibodies / vaccines



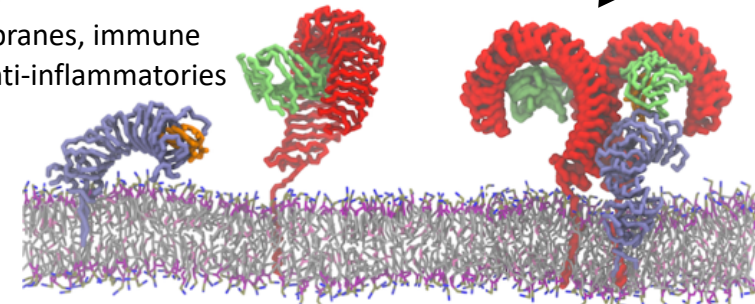
(Gram-negative) bacterial cell envelopes, membranes (lipopolysaccharide), & antibiotics



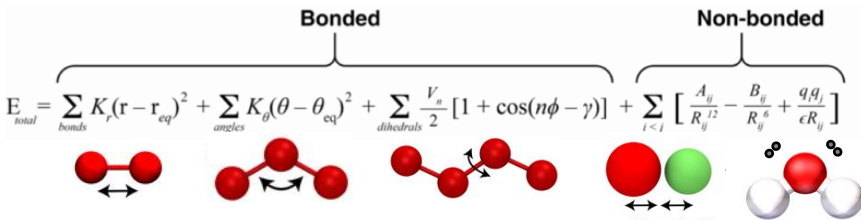
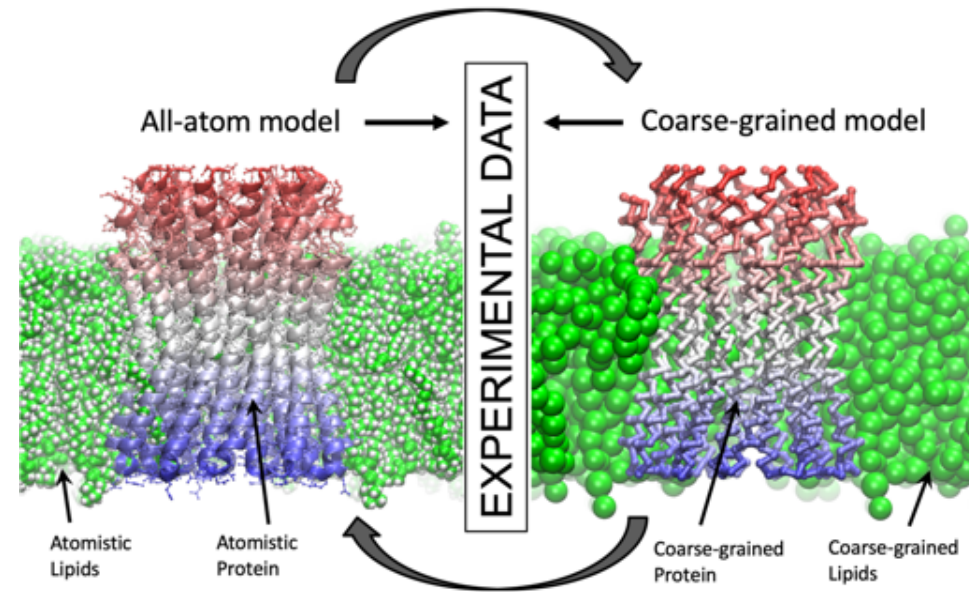
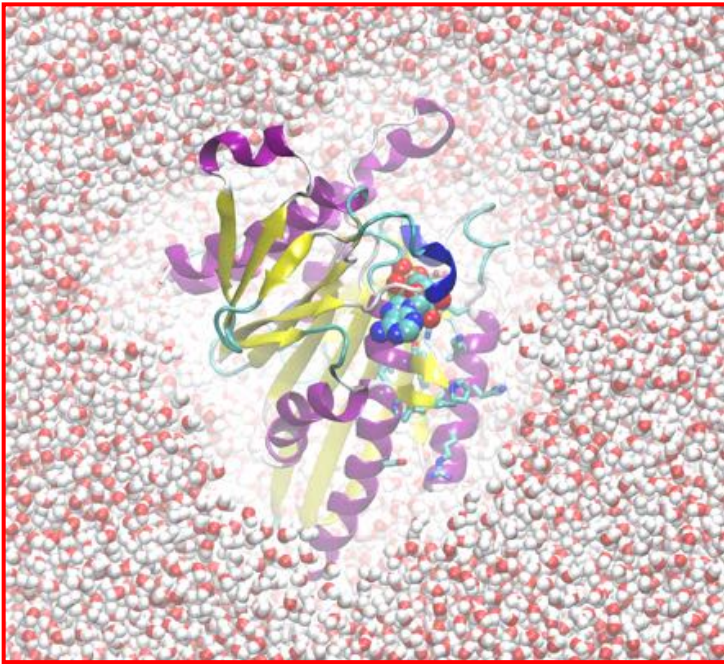
Coronaviruses: spike proteins & LPS – a novel interaction



Host cell membranes, immune receptors, & anti-inflammatories



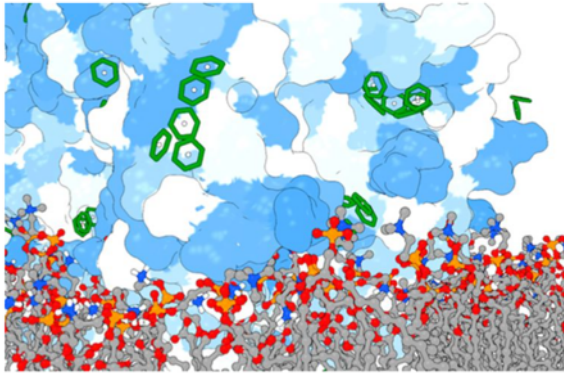
Biomolecular Simulations & Multiscale Approaches



- Methods of choice:
- (1) Molecular simulation.
 - (2) Multiscale approaches.
 - (3) Integrative modelling.

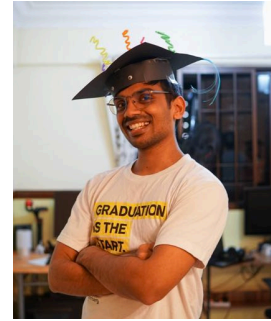
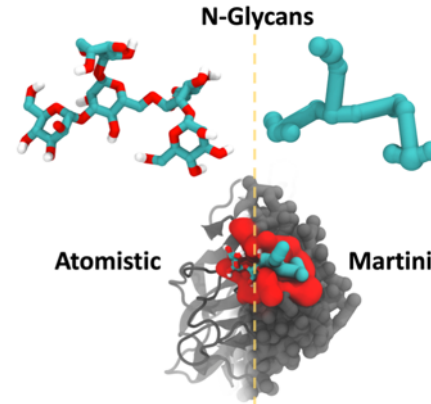
Method Development & Applications in Biology

Cryptic pockets in membrane targets



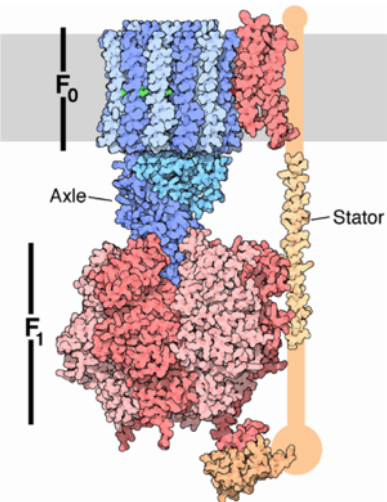
- Benzene mapping for lipid systems – novel druggable pockets discovered in dengue E protein.
J. Chem. Theory Comput. 16:5948-59 (2020)

Multiscale library for N-glycans in glycoproteins

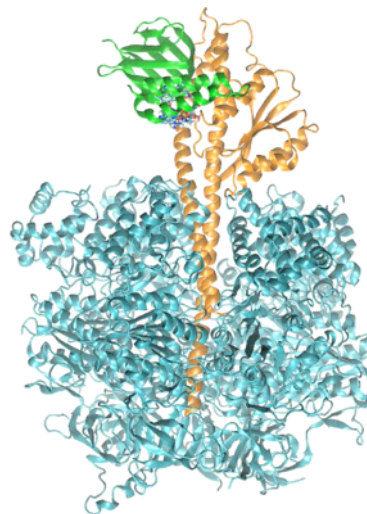


- Multiscale N-glycan library for glycoproteins (with Chandra - BII, NUS, Uni. Manchester)
J. Chem. Inf. Model. 60:3864-83 (2020)

Membrane-embedded ATP-driven enzymes – sensors & targets



Actin-filament on stalk tagged with fluorescent dye (Noji et al)



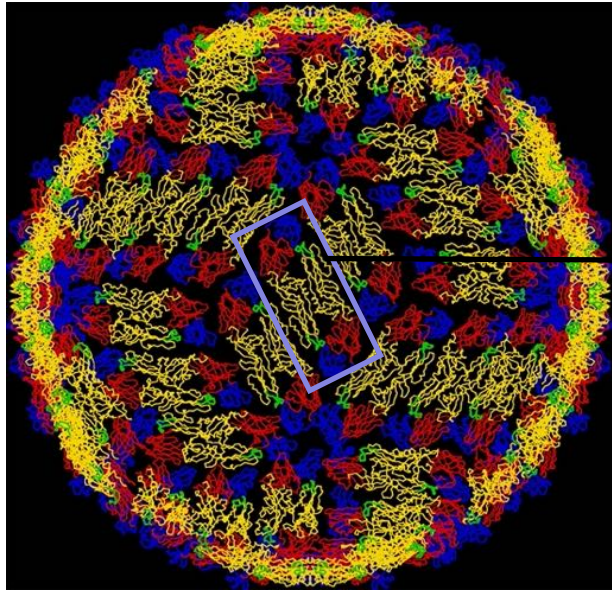
Diagnostic sensors (NUS, Delft Uni.) & antimicrobials (NTU, Dundee) / drugs targeting membrane transporters (Duke-NUS, Dundee), HIV enzymes (with Sam - BII), & flagellar motor machinery (Uni. Warwick).

ChemPhysChem. 21(9):916-26 (2020); *ChemBioChem.* 21:3249 (2020); *J. Phys. Chem. B.* 124:7176-83 (2020); *Biochim. Biophys. Acta. Gen.* 1865:129766 (2020); *Biochim. Biophys. Acta. Biomembr.* 1862:183137 (2020); *Molecules* 25:5902 (2020)

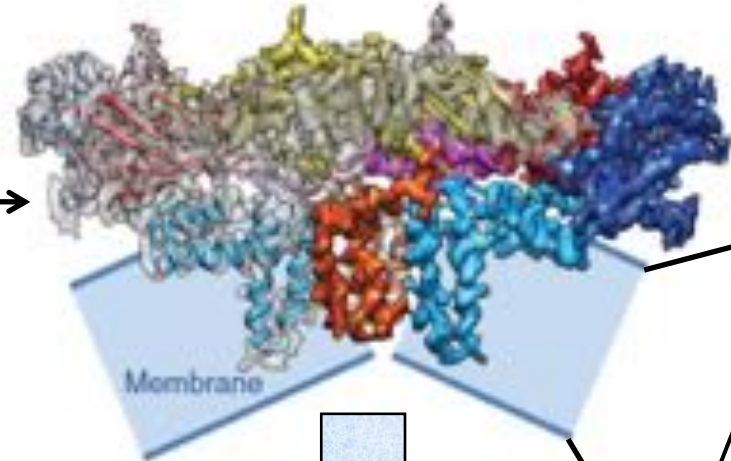
Integrative Modelling of Dengue Viral Envelope



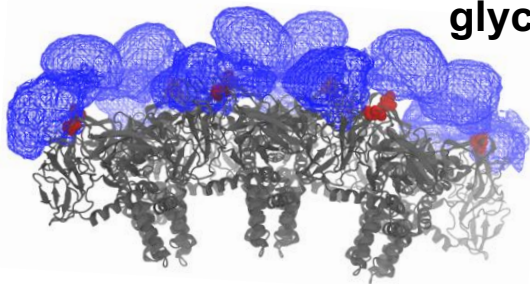
cryo-EM & other structural data



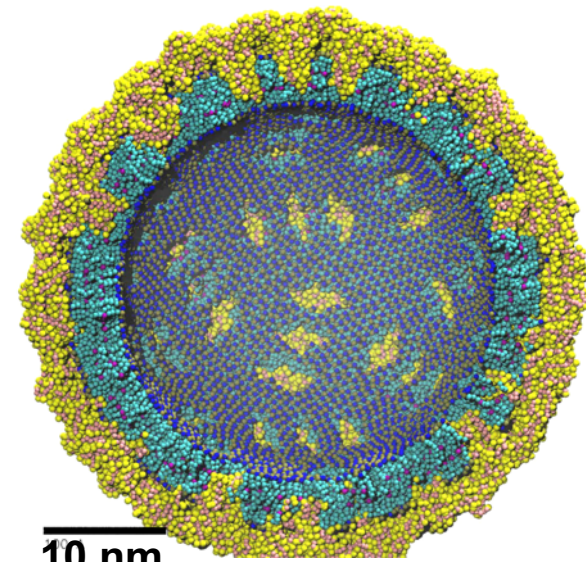
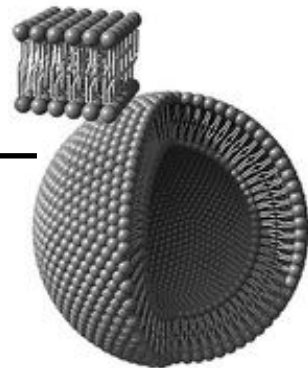
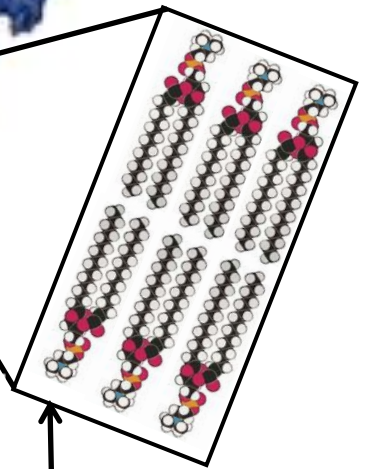
biophysics, e.g. HDX-MS



glycomics data



lipidomics data

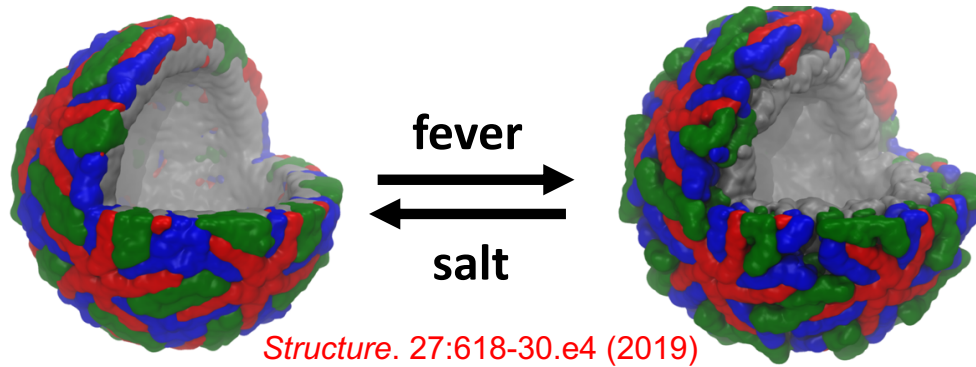


10 nm

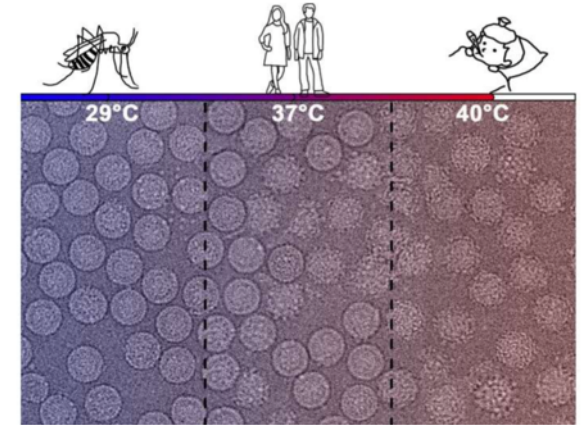
Structure (2016) 24:1410;
Structure (2019) 27:253;
Structure (2019) 27:618;
Curr Opin Struct Biol (2020) 61:146

Dengue – “Shape-Shifting” in the “Arms Race”

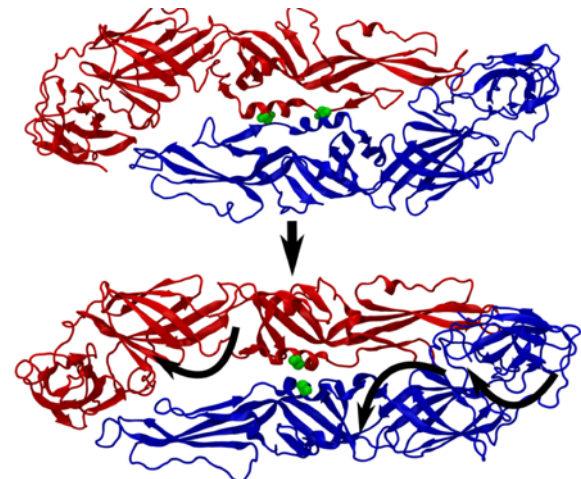
With Duke-NUS, NUS, SigN, SGH (NRF CRP)



(2) Modelling of clinical vs. lab strains: envelope mutations alter virus morphology at different temperature (e.g. fever). Altered epitopes = resistance to antibodies/vaccines.

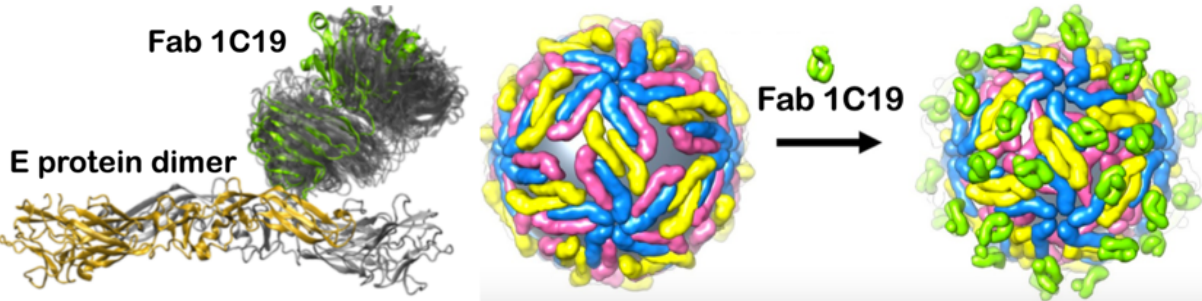


PLoS Pathog. 15:e1007996 (2019)



(1) Modelling & biophysics show that host environment leads to viral envelope “shape-shifting” between “smooth” and “bumpy” morphologies.

PLoS Pathog. 17:e1009331 (2021)

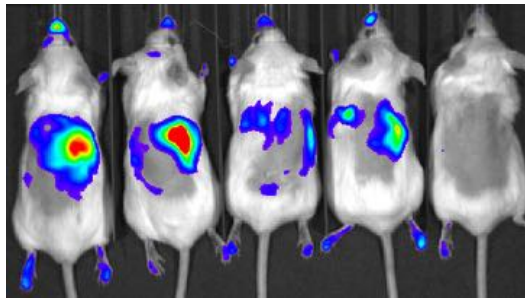
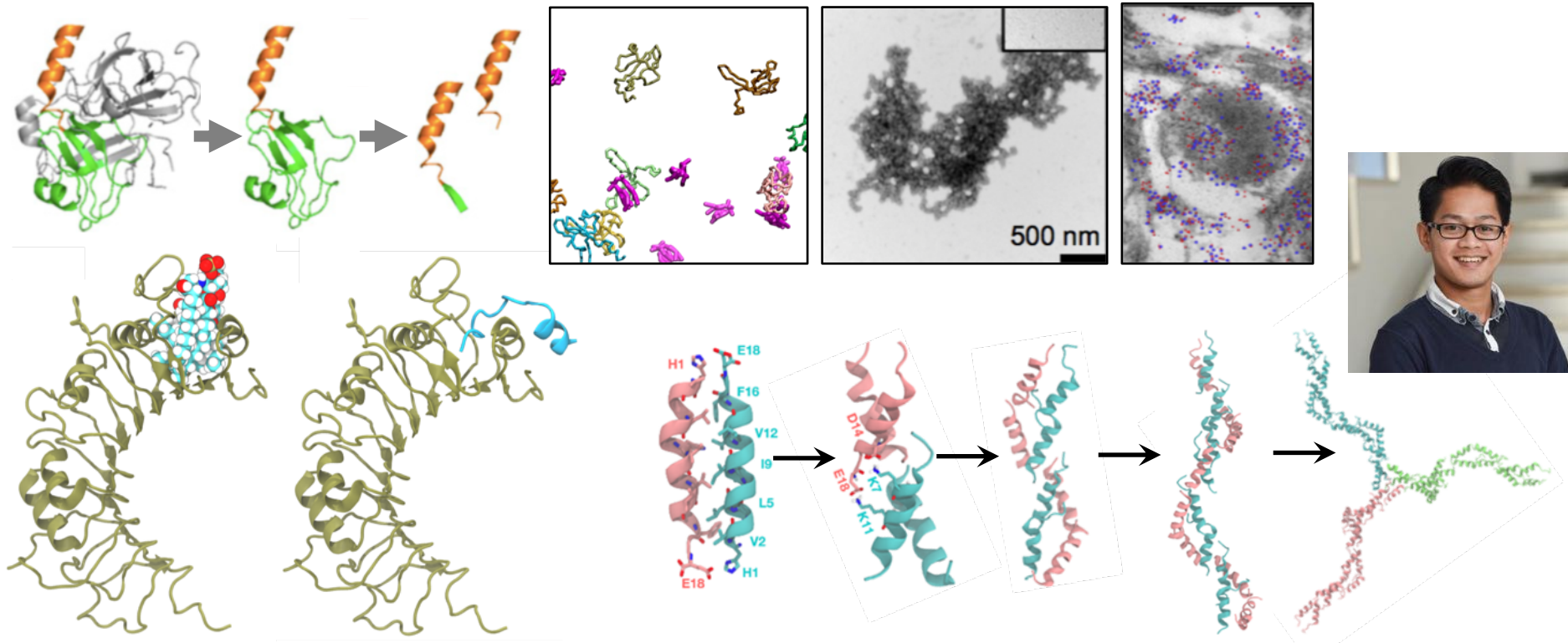


(3) Modelling & biophysics reveal antibody epitope hidden below viral surface – but a sufficiently strong antibody can force the virus to “shape-shift”, exposing it for neutralization.

→ Next-gen therapeutics: (a) high-affinity antibodies that (b) recognize diverse morphologies.

Anti-inflammatory & Antibacterial Peptides (1)

- Growing evidence that “membrane-busting” AMPs / HDPs may in fact be multi-faceted, and/or may target cell-surface receptors & enzymes.
- Thrombin-derived C-terminal peptides – anti-inflammatory potential.



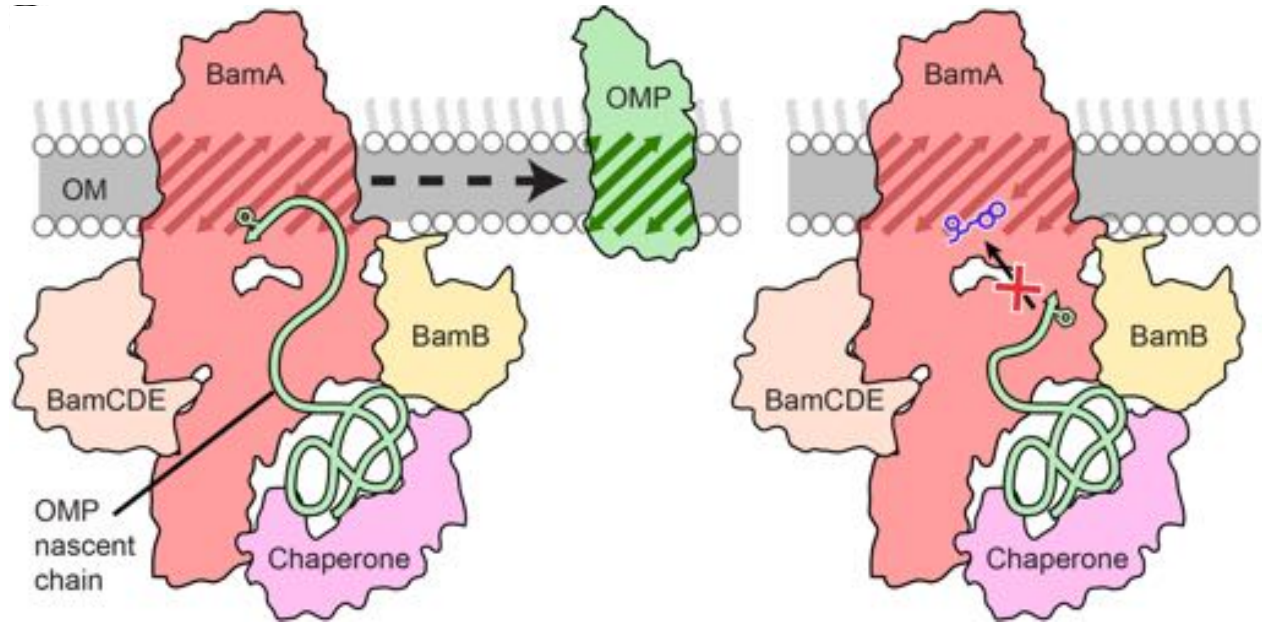
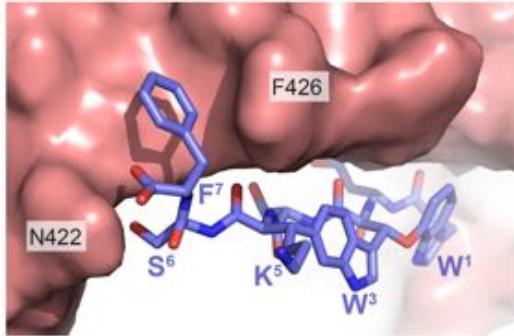
LPS LPS+rTCP96 rTCP96

- Receptor binding mode, oligomerization, pH dependence, etc...
- Preclinical development for wound healing with Artur Schmidtchen, Jitka Petrova, Anna Petruk (Lund University / In2cure), Rathi Saravanan (NTU / LKCMed)

J Biol Chem (2020) 295:3417; *Biomolecules* (2020) 10:1572; *Pharmacol Res* (2019) 147:104372; *Nat Commun* (2018) 9:2762; *Structure* (2018) 26:1151; *BBA Biomembr* (2018) 1860:2374; *PNAS* (2017) 114:E4213

Anti-inflammatory & Antibacterial Peptides (2)

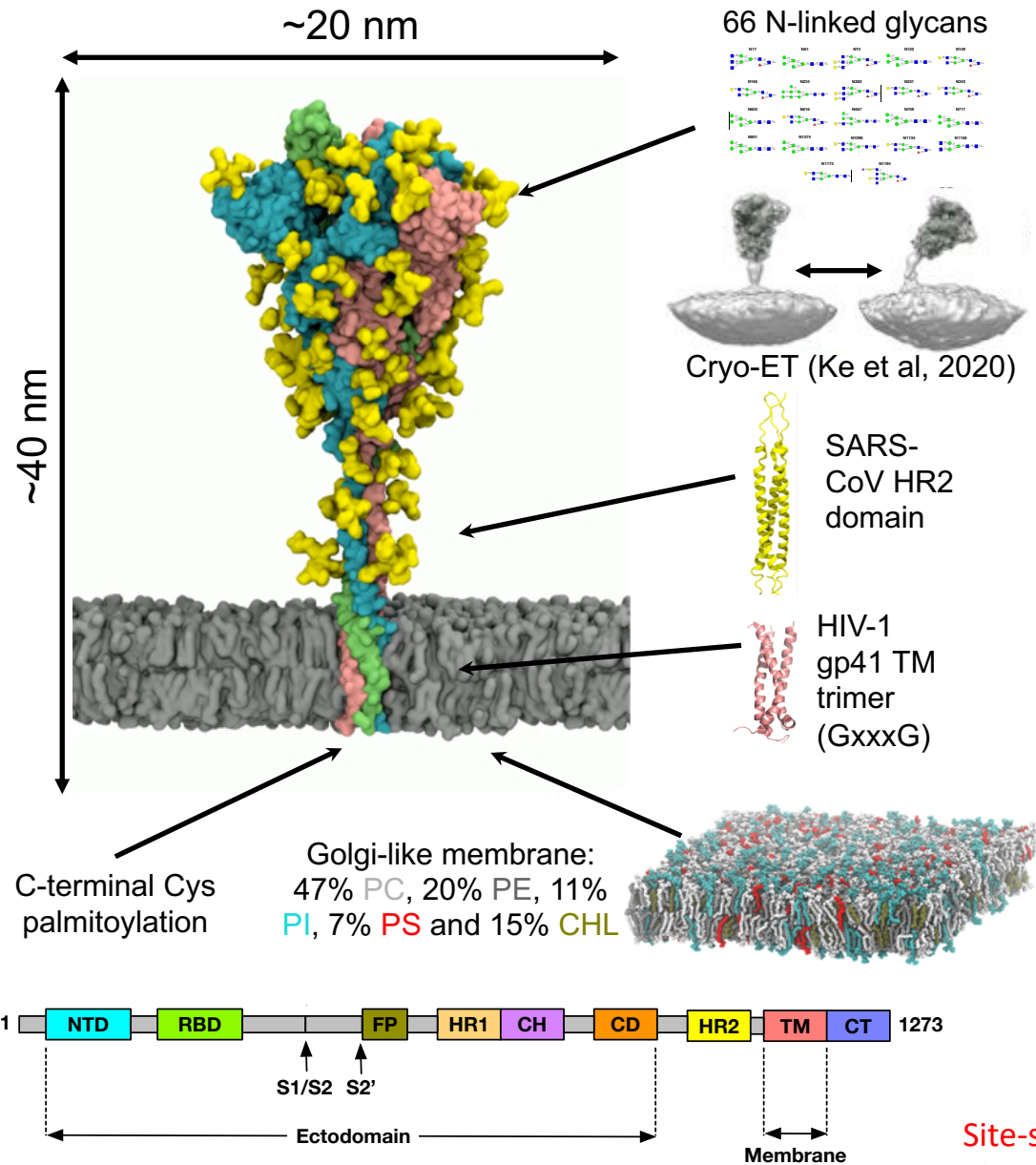
- Growing evidence that “membrane-busting” AMPs / HDPs may in fact be multi-faceted, and/or may target cell-surface receptors & enzymes.
- Antibiotics targeting Gram-negative bacteria: (last new class, quinolones >50 years ago).



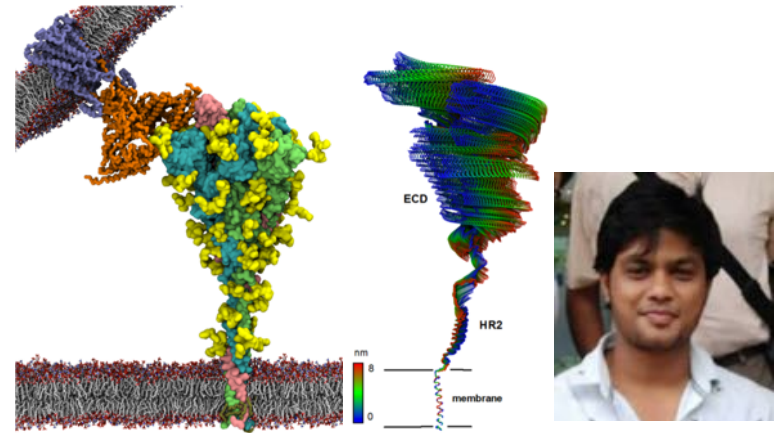
The antibiotic darobactin mimics a β -strand to inhibit outer membrane insertase. *Nature*. Epub, April 2021

- Darobactin β -hairpin peptide binds to OMP insertase (BAM complex), blocking native substrates.
- Replaces cardiolipin molecule: membrane environment is an unusual “extended binding pocket”.
- Interaction mediated via backbone: uniquely robust against potential resistance mutations.
- Jan, with Sebastian Hiller (Uni. Basel) & Polyphor; other scaffolds under investigation...

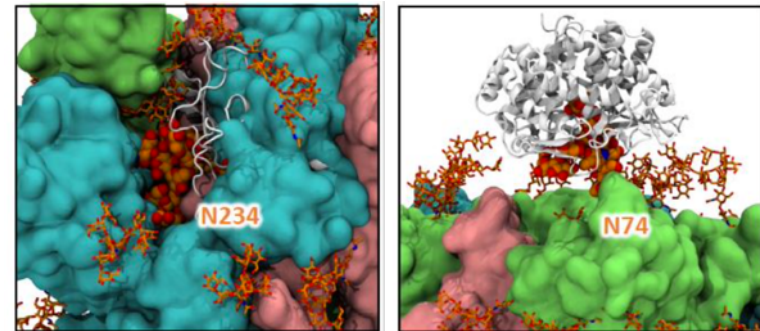
Understanding Structure-Function of SARS-CoV-2 Spike



- Firdaus, Lorena, Raghu, Aishwary, Jan, Alex
- Ganesh Anand (NUS, Penn State), Artur Schmidtchen (Lund), Max Crispin (Soton)
- BII collab's - Igor, Sebastian



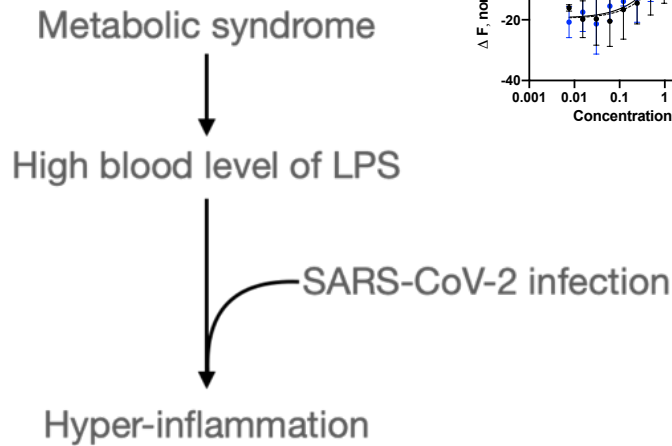
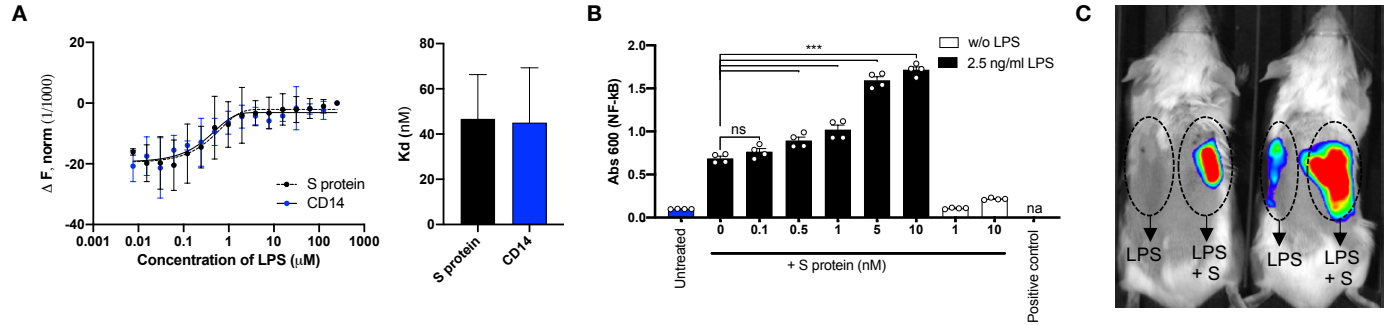
SARS-CoV-2 S protein:ACE2 interaction reveals novel allosteric targets. *Elife*. 17:e1009331 (2021)



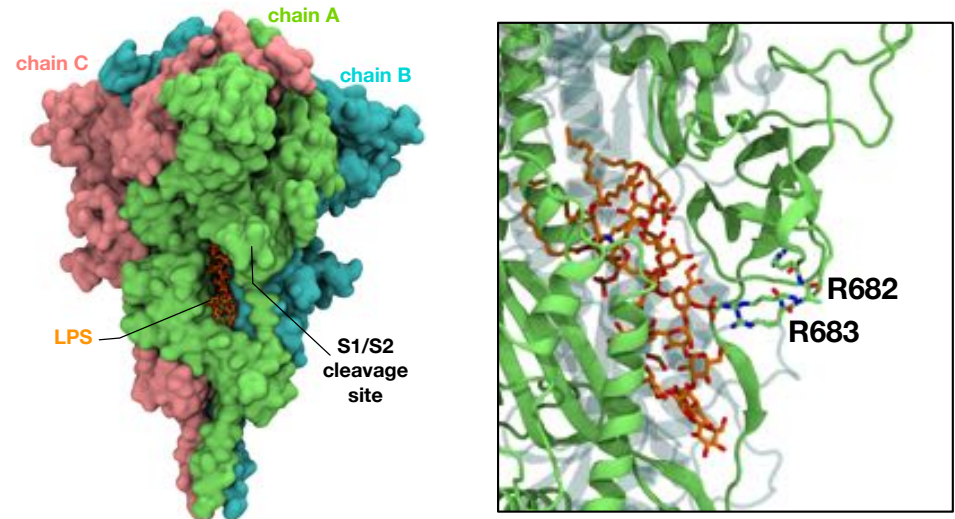
Site-specific steric control of SARS-CoV-2 spike glycosylation. *bioRxiv* (doi: 10.1101/2021.03.08.433764)

Defining a Novel Interaction: SARS-CoV-2 Spike / LPS

- (A) Spike protein:LPS affinity \equiv CD14 (microscale thermophoresis).
 (B) Spike protein boosts NF- κ B response to LPS.
 (C) Inflammation in NF- κ B reporter mice.



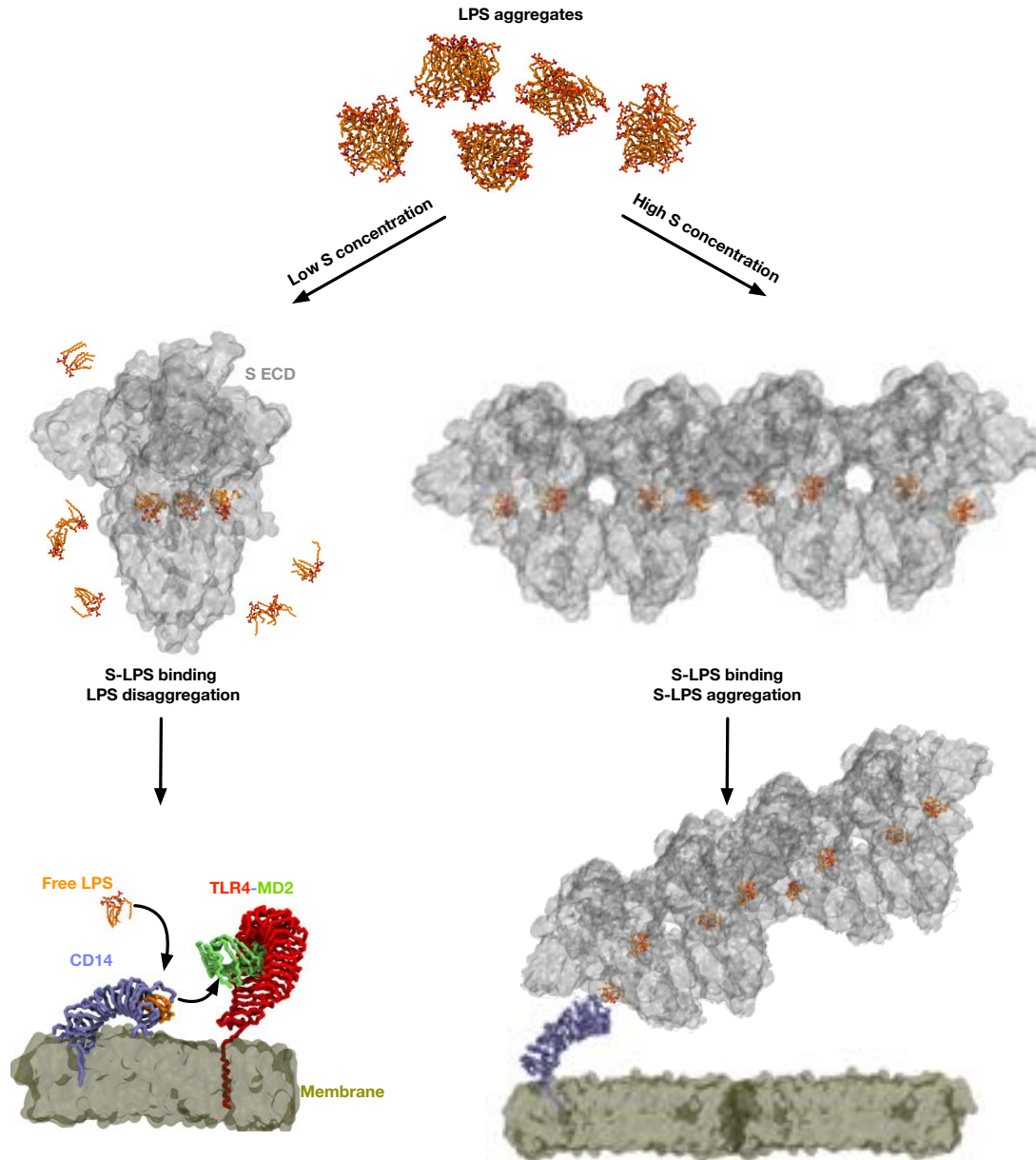
SARS-CoV-2 spike protein binds to bacterial lipopolysaccharide and boosts proinflammatory activity. *J Mol Cell Biol* (2021) 12:916



- Severe COVID19 common in those with metabolic syndrome (diabetes, obesity etc.) linked with raised LPS levels
- Metabolic syndrome predisposes patients to severe COVID19: Hyper-inflammation in lungs \rightarrow respiratory failure, sepsis & death.
- **Multiple recent papers: (i) raised LPS/intestinal permeability & sepsis; (ii) “direct” activation of TLR4 by spike...**

- Conserved LPS site detected via docking, solvent mapping.
- **Validation and follow-up studies underway.**

Viruses vs. Bacteria vs. Host...



Multiscale Simulation, Modelling & Design Group

Singapore, viruses

Sheemei Lok
Ganesh Anand
Sylvie Alonso
Thorsten Wohland
Paul Macary
Eng-Eong Ooi
Subhash Vasudevan
Yue Wan
Terry Nguyen-Khuong
Paul Matsudaira

Singapore, general

Gerhard Gruber
Shu Sin Chng
Xue Li Guan
Rachel Ee Pui Lai
Koji Itahana
Markus Wenk

BII

Chandra Verma
Igor Berezovsky
Sebastian Maurer-Stroh
Roland Huber
Samuel Gan

Computing

BII, NSCC

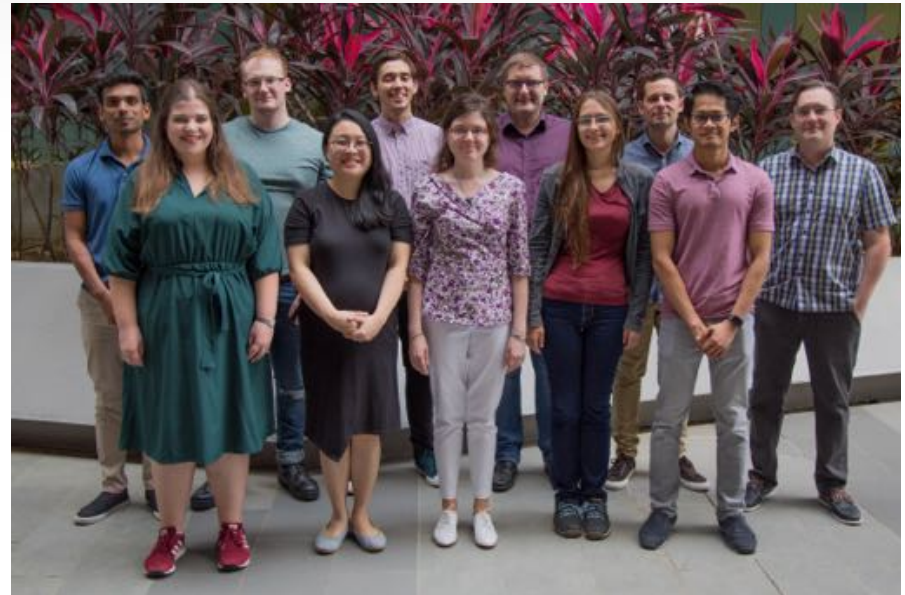


International

Sebastian Hiller
Max Crispin
Olivera Francetic
Jane Allison
Jim Warwicker
Tom Piggot
Syma Khalid
Luning Liu
Martin Ulmschneider
Ivo Martins
Slawomir Boncel
Duncan McMillan

International, "LPS network"

Artur Schmidtchen
Jitka Petrlova
Anna Petruk
Rathi Saravanan
(Lund, Copenhagen,
LKCMed, NTU)
Graeme Lancaster
Mark Febbraio
(BakerIDI, Melbourne)
Clare Bryant
Nick Gay
(Cambridge)



Jan Marzinek (**YIRG**)

Firdaus Samsudin (**YIRG**)

Alexander Krahl

Lorena Zuzic

Aishwary Shivgan

Dagnija Tupina

Sam Weetman

Priscilla Boon

Jack Copping

Alister Boags

Eilish McBurnie

