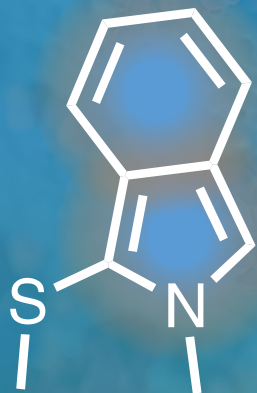


CPPC
2021

First Canadian Peptide and Protein Community Virtual Symposium

27-28 MAY 2021 | ONLINE



Canadian
Peptide
Protein
Community



biomedicines



FIICKed ACE2 mimics inhibit SARS-CoV-2 spike protein-ACE2 interaction

Mihajlo Todorovic¹, Antonio A. W. L. Wong¹, Naysilla L. Dayanara, Taylor Navi, David M. Perrin²

¹ These presenters contributed equally

² Corresponding authors contact: dperrin@chem.ubc.ca | 2036 Main Mall, Vancouver, BC CANADA V6T 1Z1



Coronavirus Disease 2019
COVID-19

CPPC
2021

Global Cases

151.397.857

Cases by Country/Region/Sovereignty

32.345.329 US

19.164.969 India

14.659.011 Brazil

5.677.835 France

4.820.591 Turkey

4.750.755 Russia

4.432.246 United Kingdom

4.022.653 Italy

3.524.077 Spain

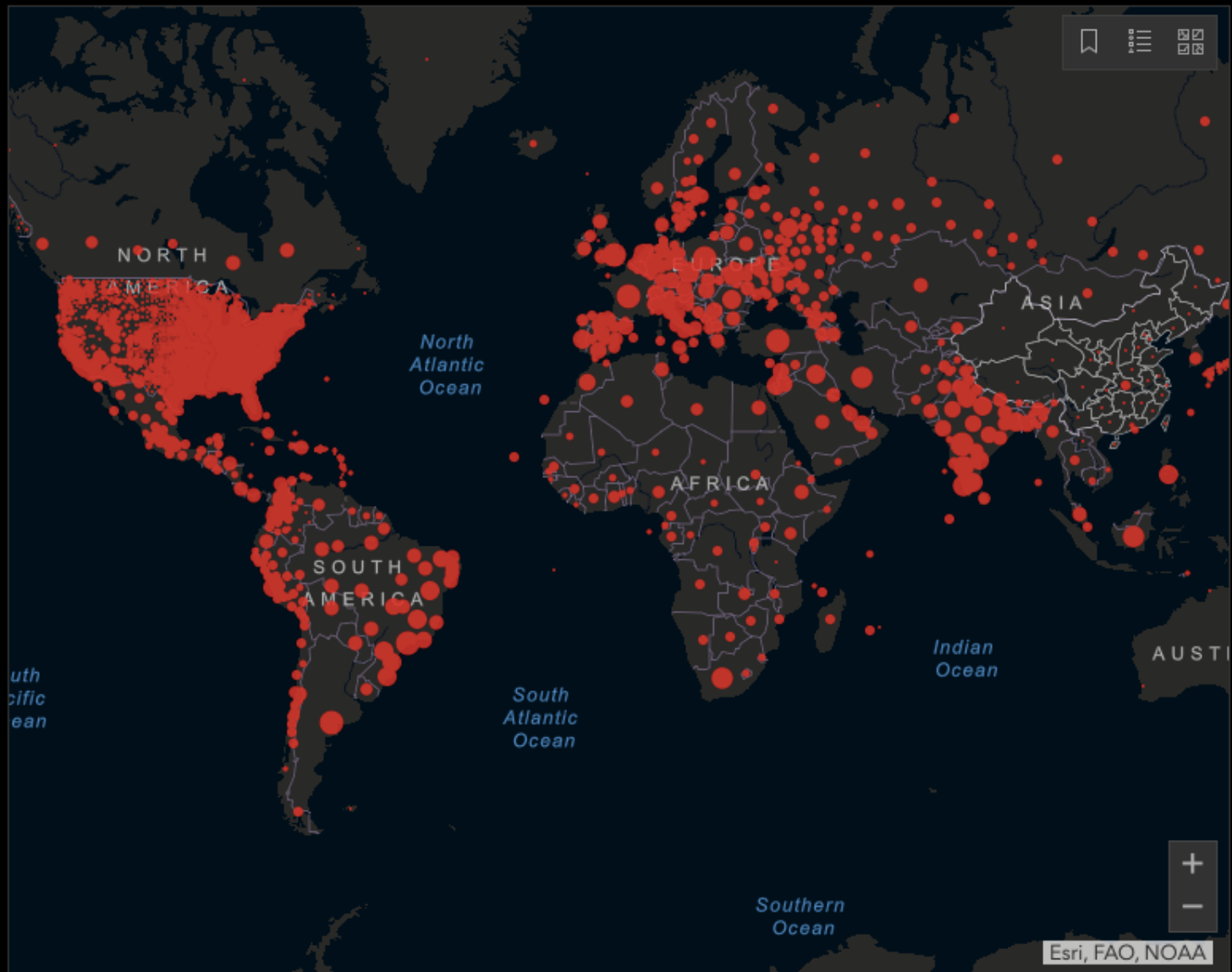
3.405.365 Germany

2.977.363 Argentina

2.859.724 Colombia

2.792.142 Poland

2.499.077 Iran



Cumulative Cases

Incidence Rate

Case-Fatality Ratio

Testing Rate

192

countries/region

3.179.927

Global Deaths

CPPC 2021

Last Updated at (M/D/YYYY)

4/30/2021 10:20 p. m.



Admin0



Global Cases

151.397.857

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32.345.329 US

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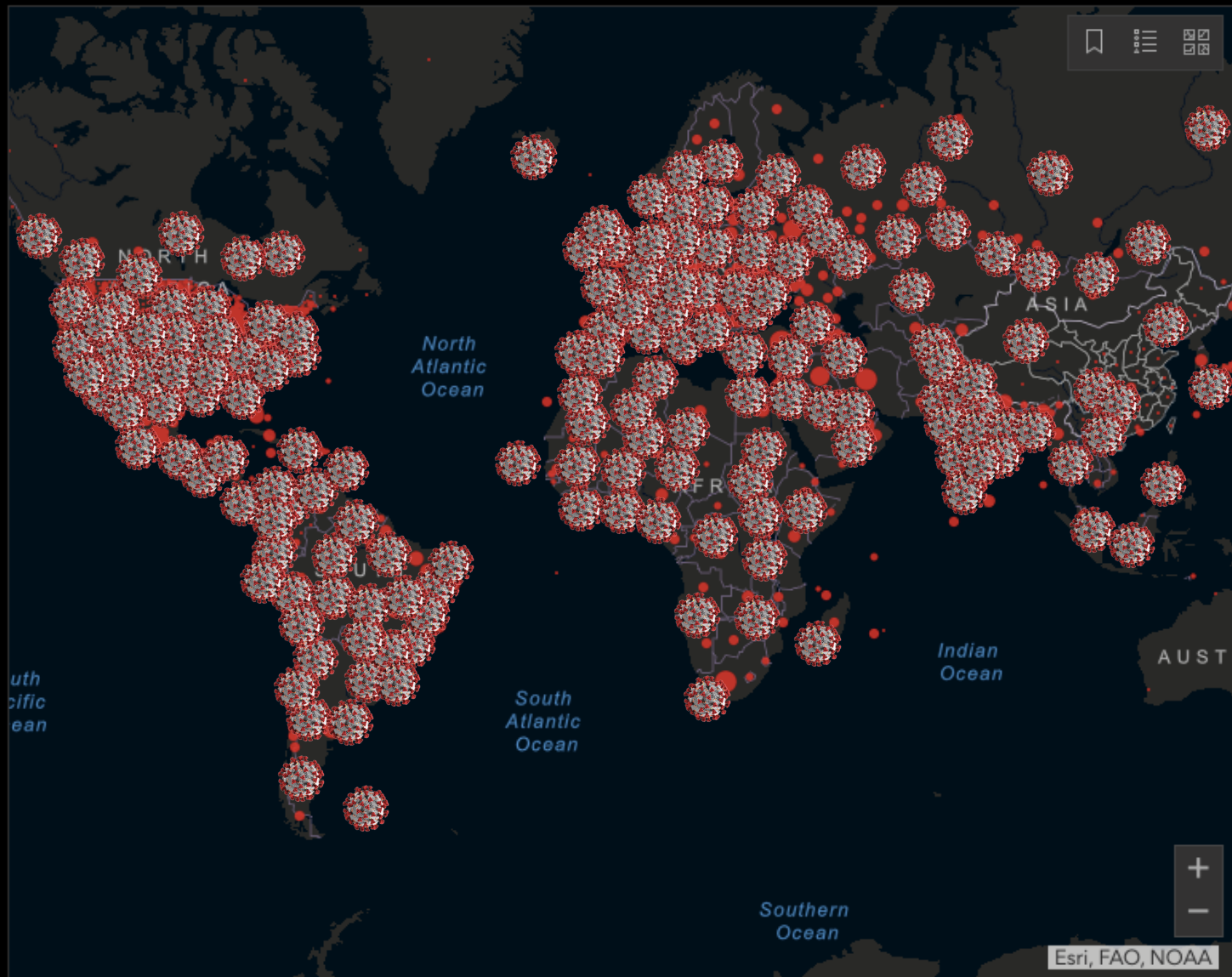
3.405.365 Germany

2.977.363 Argentina

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2.792.142 Poland

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Cumulative Cases

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Case-Fatality Ratio

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Admin0



Last Updated at (M/D/YYYY)
4/30/2021 10:20 p. m.

192

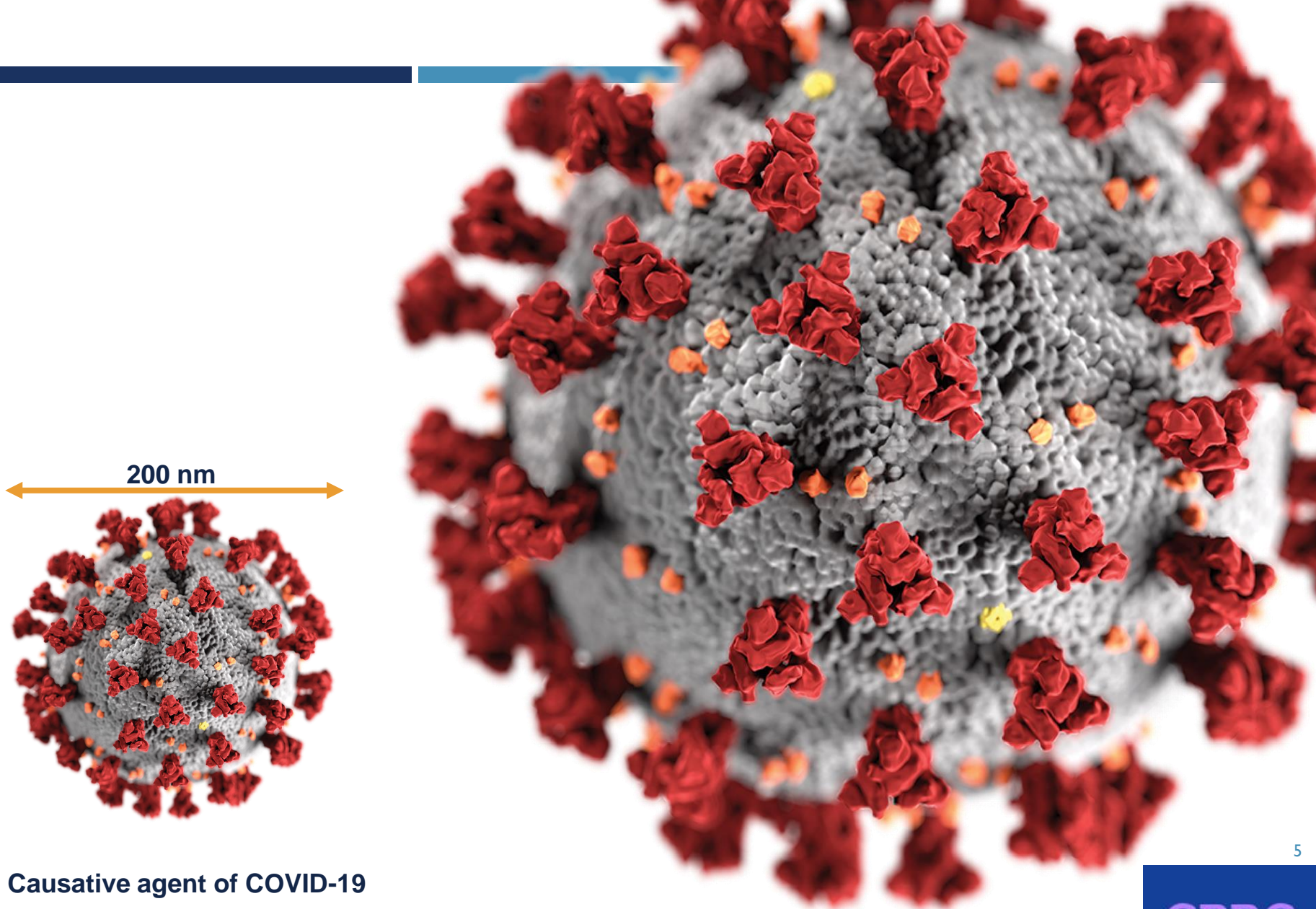
countries/region

3.179.927

Global Deaths

CPPC 2021

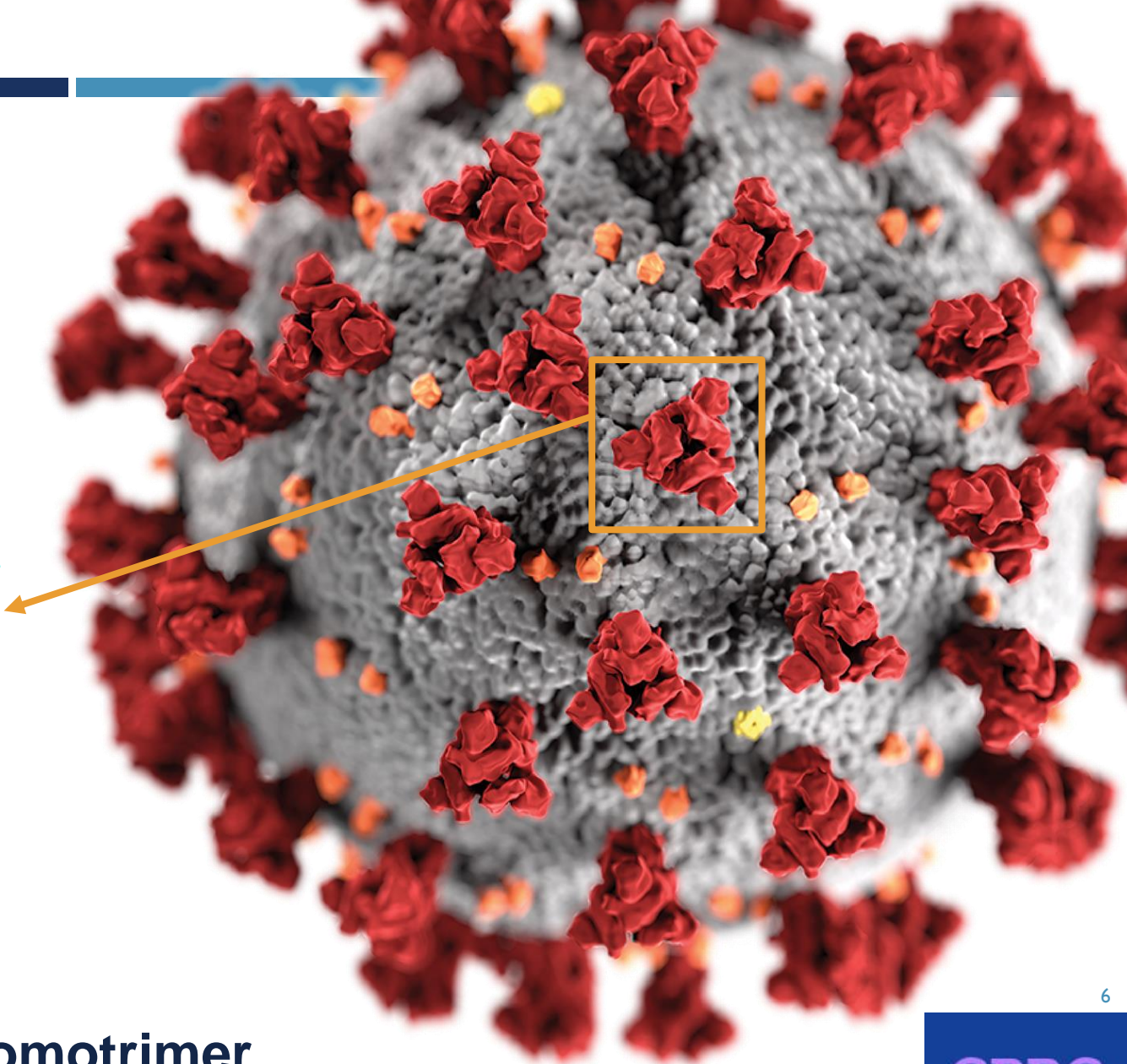
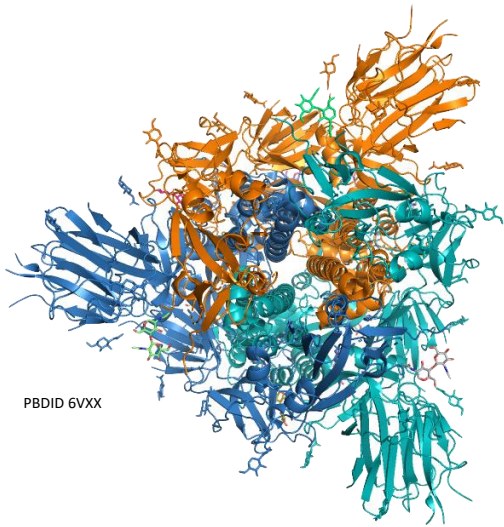
Esri, FAO, NOAA



Causative agent of COVID-19

SARS-CoV-2

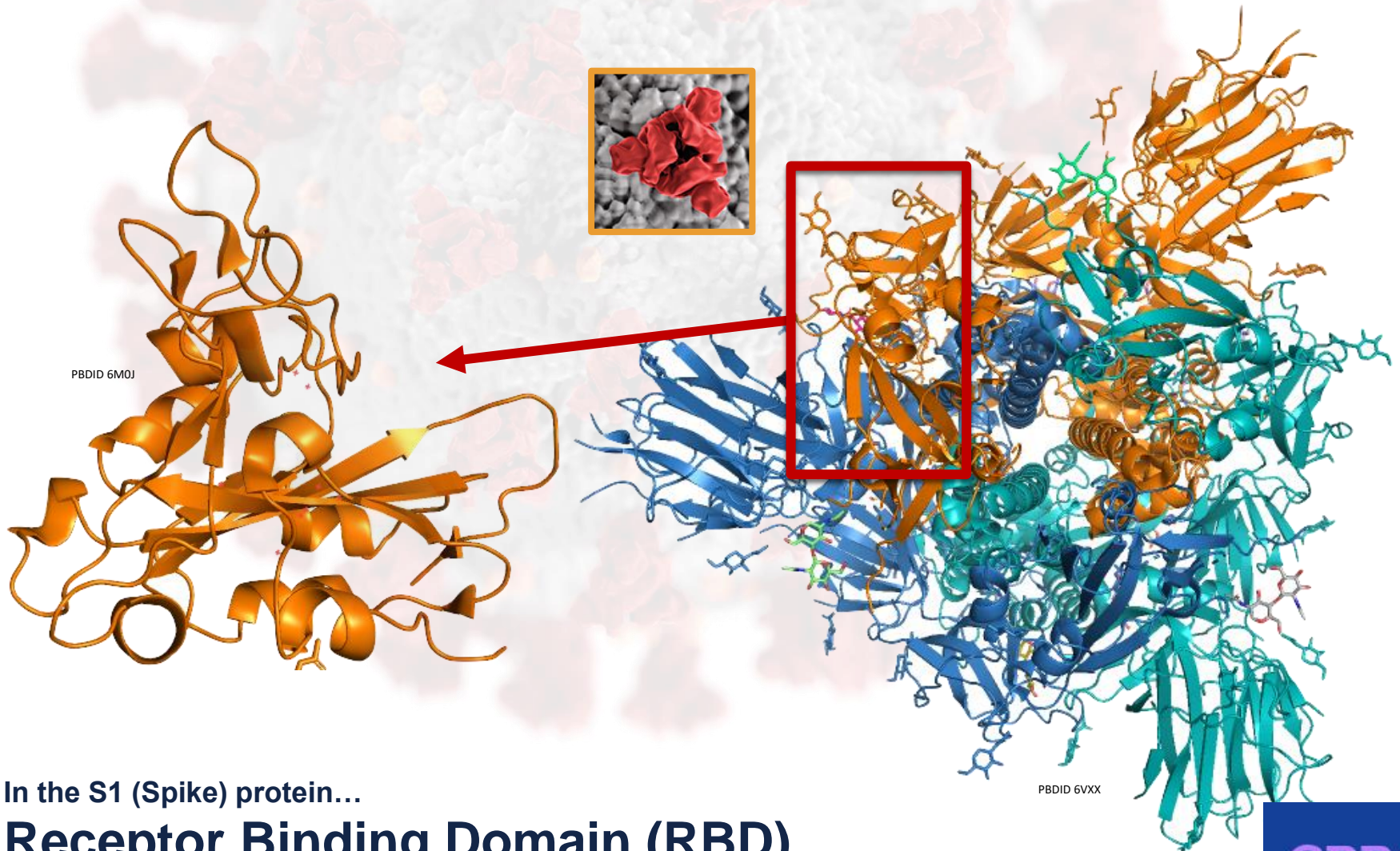
Severe Acute Respiratory Syndrome Coronavirus 2



On the surface...

Spike protein homotrimer

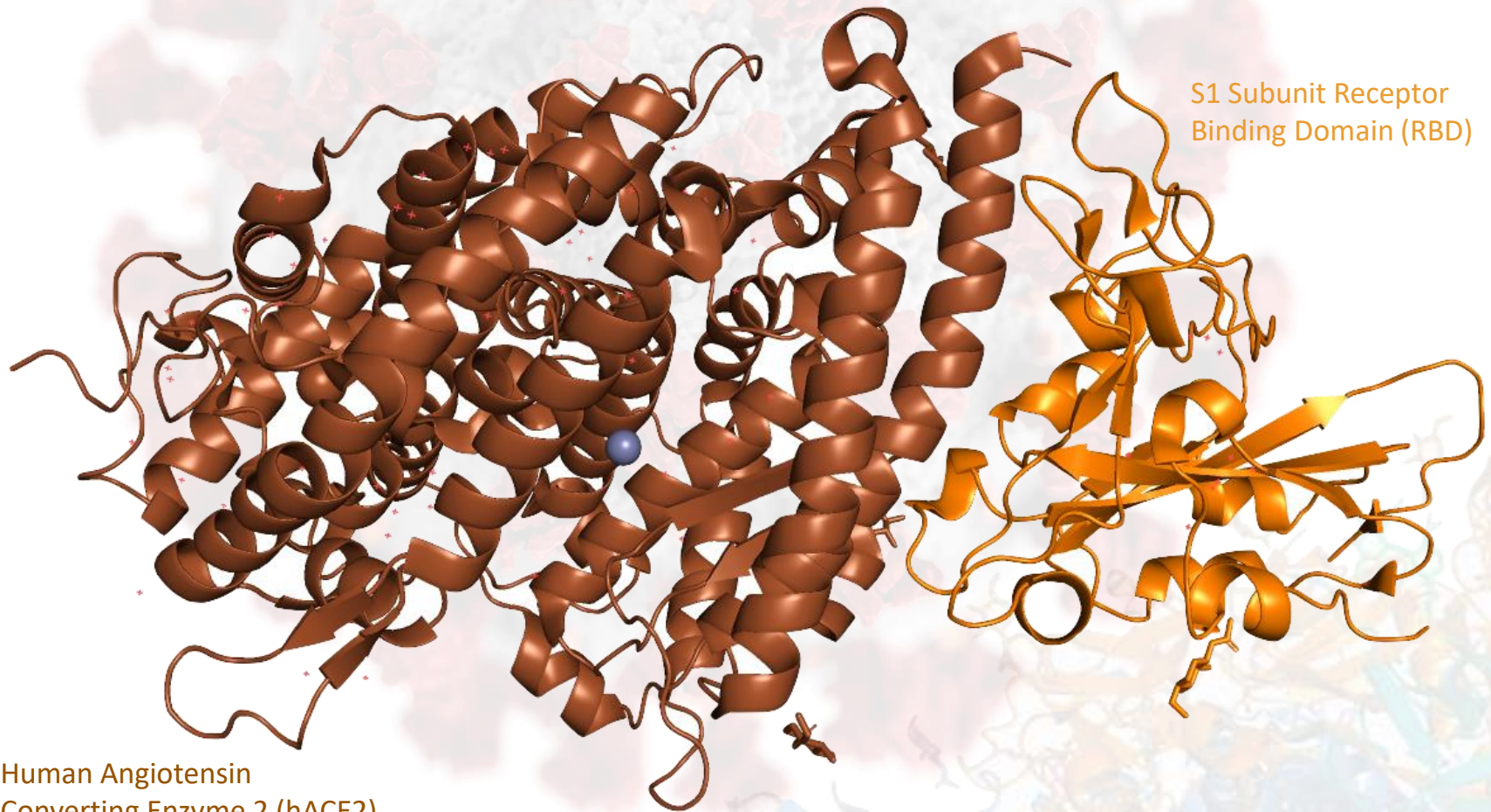
Allow penetration and subsequent entry



In the S1 (Spike) protein...

Receptor Binding Domain (RBD)

Residue number (resn) 319-541



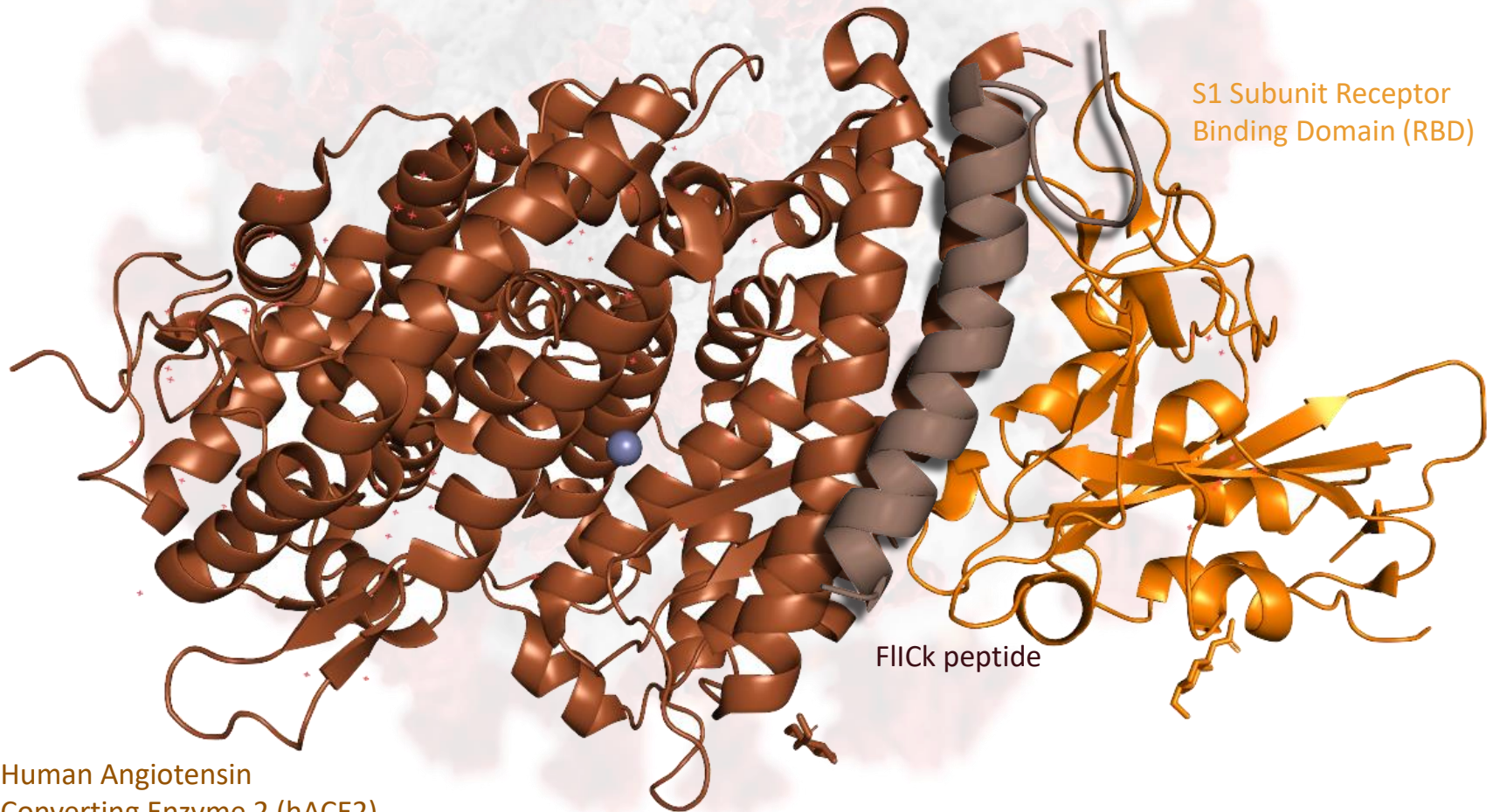
S1 Subunit Receptor
Binding Domain (RBD)

Human Angiotensin
Converting Enzyme 2 (hACE2)

In the S1 (Spike) protein...

Substrate bound X-ray structure

PDBID 6M0J



Human Angiotensin
Converting Enzyme 2 (hACE2)

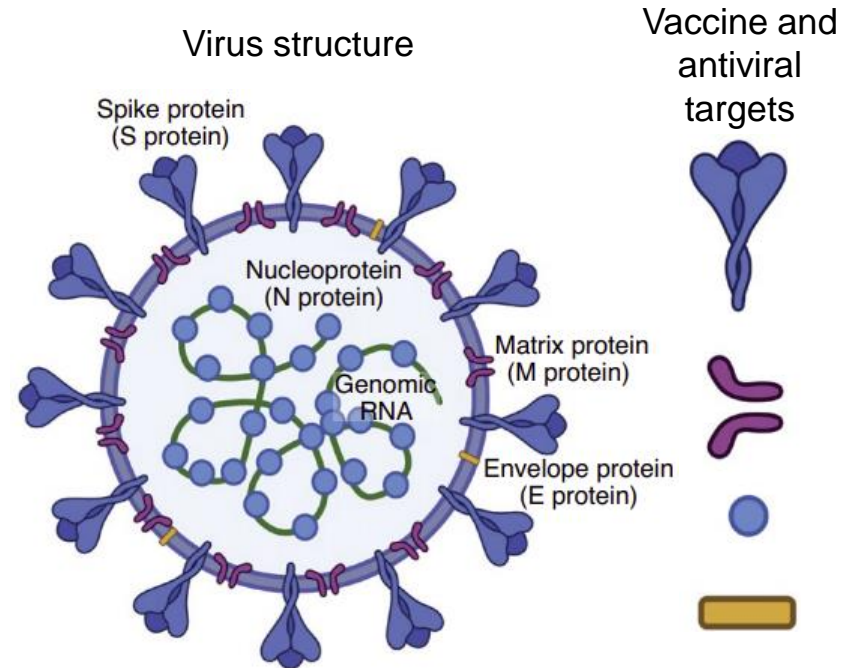
In the S1 (Spike) protein...

Substrate bound X-ray structure

PDBID 6M0J

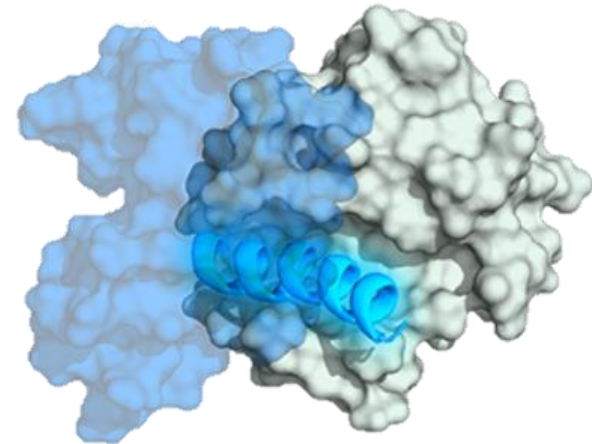
(Peptide Introduction): Challenges associated with mass vaccination

- Production and distribution of vaccines
- Variable vaccine acceptance
- Imperfect vaccine efficacy
- Development of vaccination-resistant strains

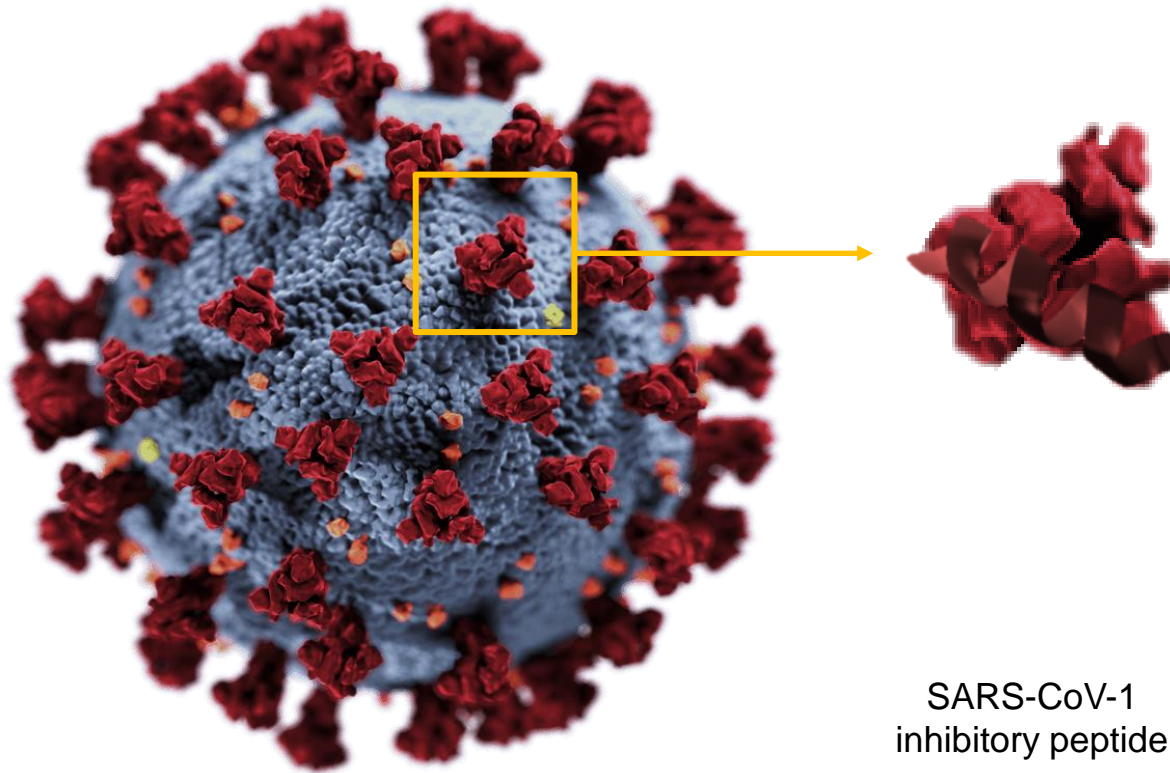


(Peptide Introduction): Peptides as drugs

- Better at targeting protein-protein interactions
- Effective for extracellular targets
- Less prone to resistance
- Stapling increases protease resistance



(Peptide Introduction): SARS-CoV-1 inhibitory peptides

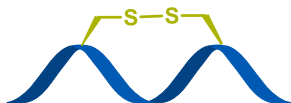


Ho, T. Y.; Wu, S. L.; Chen, J. C.; Wei, Y. C.; Cheng, S. E.; Chang, Y. H.; Liu, H. J.; Hsiang, C. Y.
Antiviral Res. **2006**, *69* (2), 70–76. <https://doi.org/10.1016/j.antiviral.2005.10.005>.

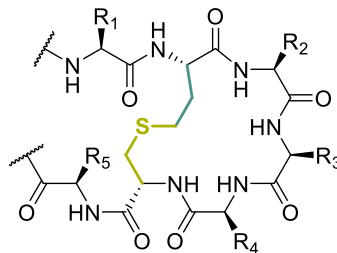
(Peptide Introduction): Peptide Stapling

Natural Peptide Crosslinks

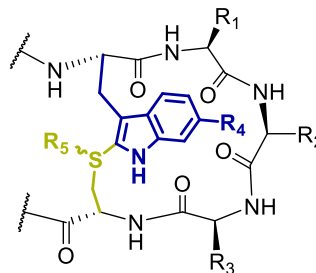
Disulfide Linkage



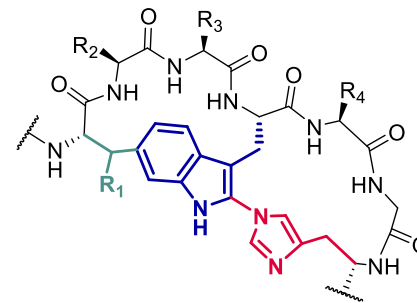
Thioether Bridge



Tryptathionine Bridge

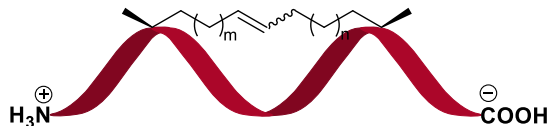


Tryptophan-Histidine Crosslink

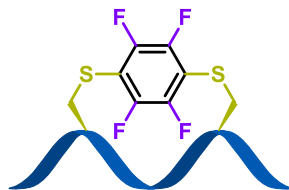


Synthetic Peptide Crosslinks

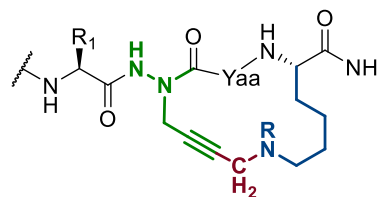
Olefin Metathesis



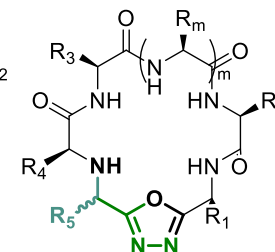
Perfluoroaryl Stapling



A³ Cyclization



Oxadiazole Grafting



Blackwell, Helen E., & Grubbs, Robert H. (1998). *Angewandte Chemie International Edition*, 37(23), 3281–3284.

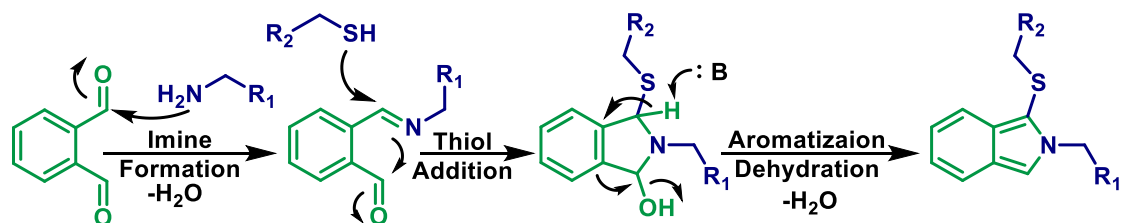
Spokoyny, A. M., Zou, Y., Ling, J. J., Yu, H., Lin, Y. S., & Pentelute, B. L. (2013). *Journal of the American Chemical Society*, 135(16), 5946–5949.

Zhang, J., Mulumba, M., Ong, H., & Lubell, W. D. (2017). *Angewandte Chemie – International Edition*, 56(22), 6284–6288.

Frost, J. R., Scully, C. C. G., & Yudin, A. K. (2016). *Nature Chemistry*, 8(12), 1105–1111.

<https://doi.org/10.1038/nchem.2636>

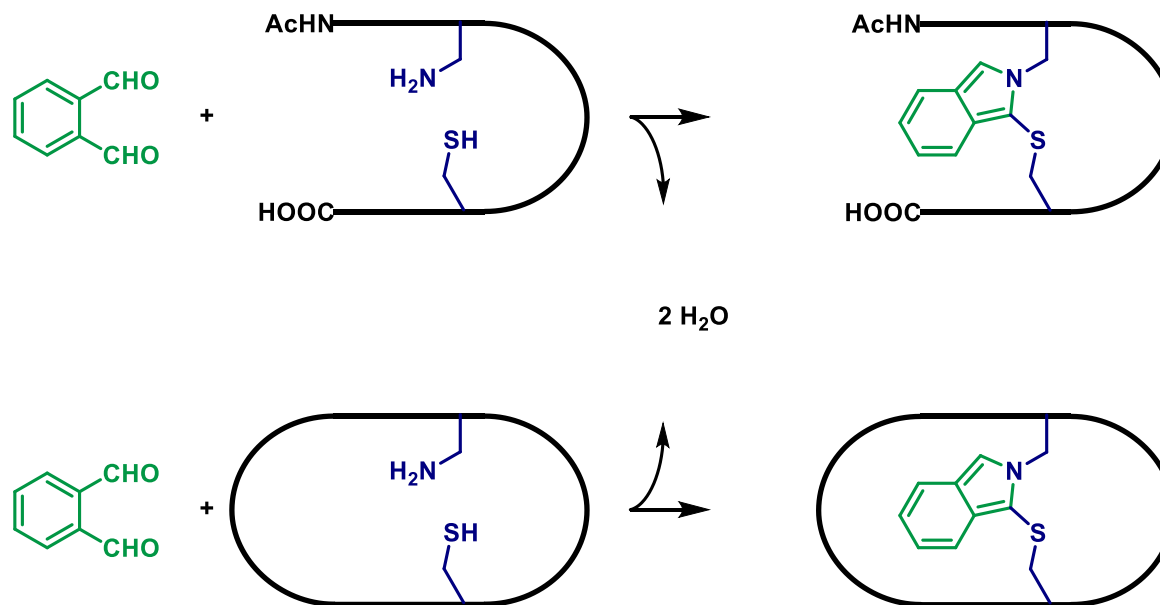
(FIICk): Isoindole Formation



- **Isoindole Formation** is a classic analytical reaction
- Condensation of ortho-phthalaldehyde with an amine and a thiol results in a fluorescent heterocycle
- Used to quantify amino acids and peptides and as an active site titrant

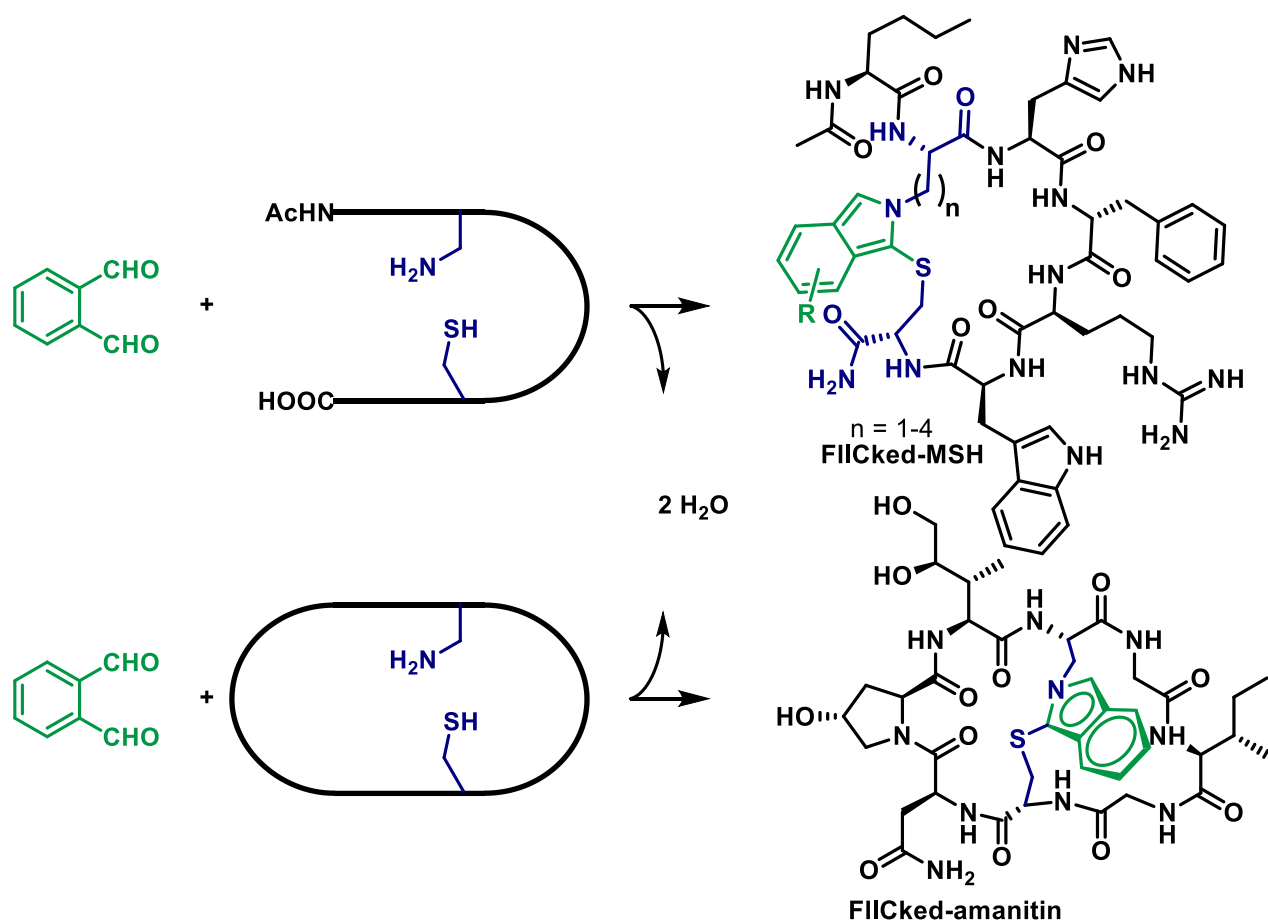
Todorovic, M.; Schwab, K. D.; Zeisler, J.; Zhang, C.; Bénard, F.; Perrin, D. M. *Angew. Chemie - Int. Ed.* **2019**, 58 (40), 14120–14124. <https://doi.org/10.1002/anie.201906514>.

(FI)Ck: Monocycles and Bicycles



Todorovic, M.; Schwab, K. D.; Zeisler, J.; Zhang, C.; Bénard, F.; Perrin, D. M. *Angew. Chemie - Int. Ed.* **2019**, 58 (40), 14120–14124. <https://doi.org/10.1002/anie.201906514>.

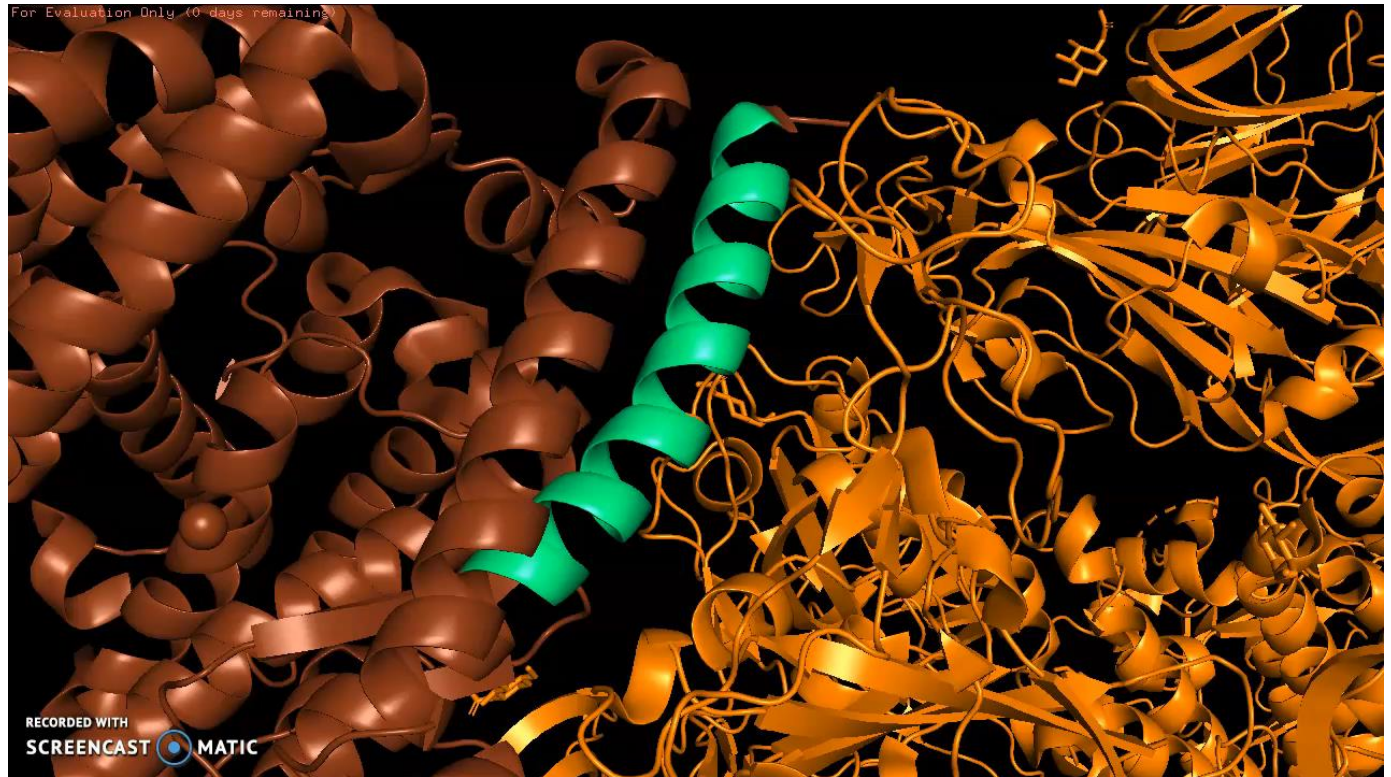
(FIICK): Proof of Concept Peptides



Todorovic, M.; Schwab, K. D.; Zeisler, J.; Zhang, C.; Bénard, F.; Perrin, D. M. *Angew. Chemie - Int. Ed.* **2019**, 58 (40), 14120–14124. <https://doi.org/10.1002/anie.201906514>.

(FIICK): ACE2 Mimicking Peptide Literature Precedence

- Direct cut-outs of N-terminal fragment of hACE2 were made by Pentelute¹ and Altseens²
- Binding to Spike protein tested by Biolayer Interferometry (BLI) or Atomic Force Microscopy
- Little binding in the high micromolar range was reported and no inhibition of the ACE2-Spike protein interaction was observed in our hands
- Doubled stapled (olefin metathesis) versions of ACE2 helix made by Debnath³
- Although binding measured by BLI, activity in in-vitro antiviral show

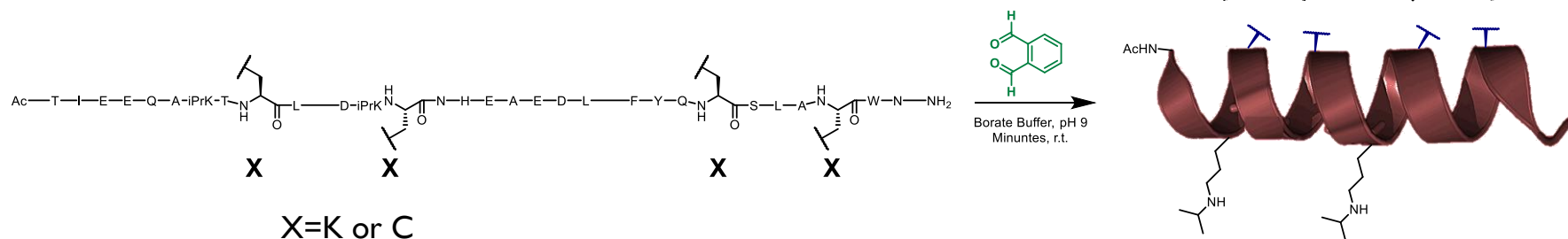


1) Zhang, G.; Pomplun, S.; Loftis, A. R.; Loas, A.; Pentelute, B. L. *bioRxiv* **2020**, 2020.03.19.999318. <https://doi.org/10.1101/2020.03.19.999318>.

2) Yang, J.; Petitjean, S. J. L.; Koehler, M.; Zhang, Q.; Dumitru, A. C.; Chen, W.; Derclaye, S.; Vincent, S. P.; Soumillon, P.; Alsteens, D. *Nat. Commun.* **2020**, // <https://doi.org/10.1038/s41467-020-18319-6>.

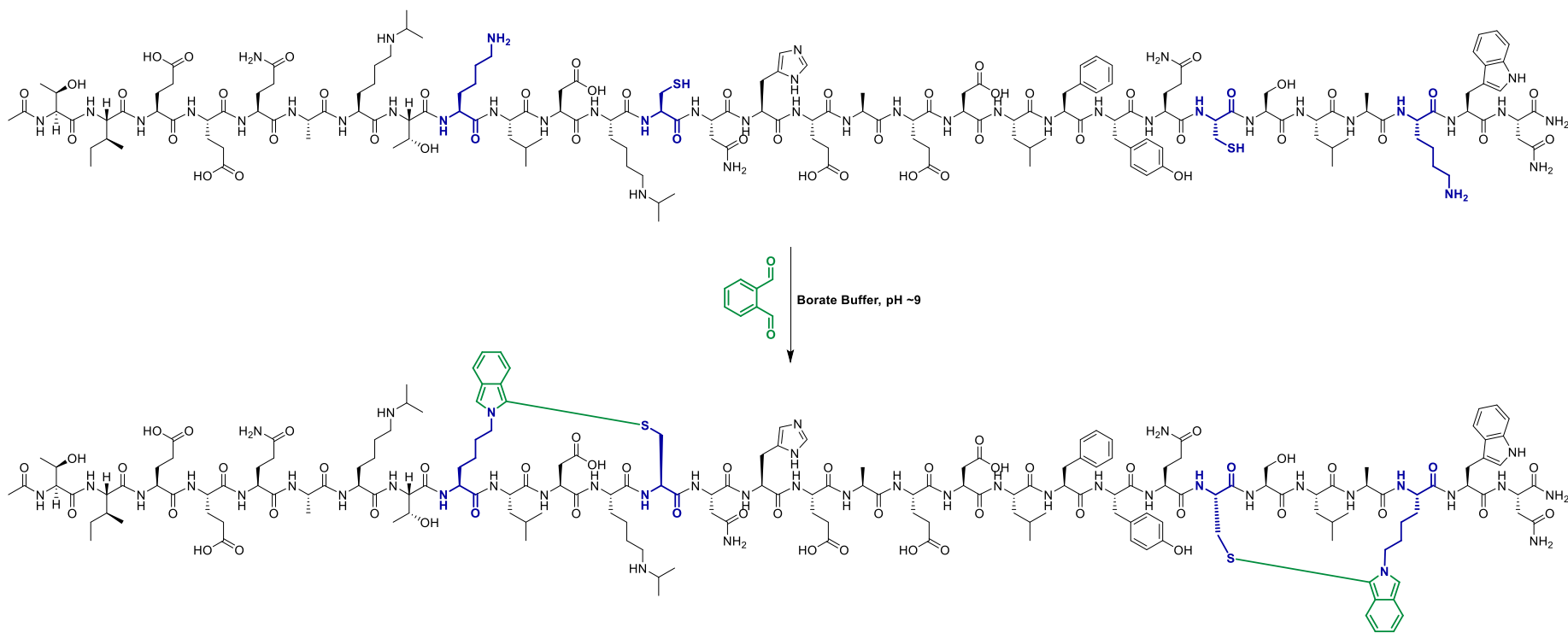
3) Curreli, F.; Victor, S. M. B.; Ahmed, S.; Drelich, A.; Tong, X.; Tseng, C.-T. K.; Hillyer, C. D.; Debnath, A. K. *MBio* **2020**, // (6), e02451-20.

(FIICK): Double FIICKed ACE2 Peptides

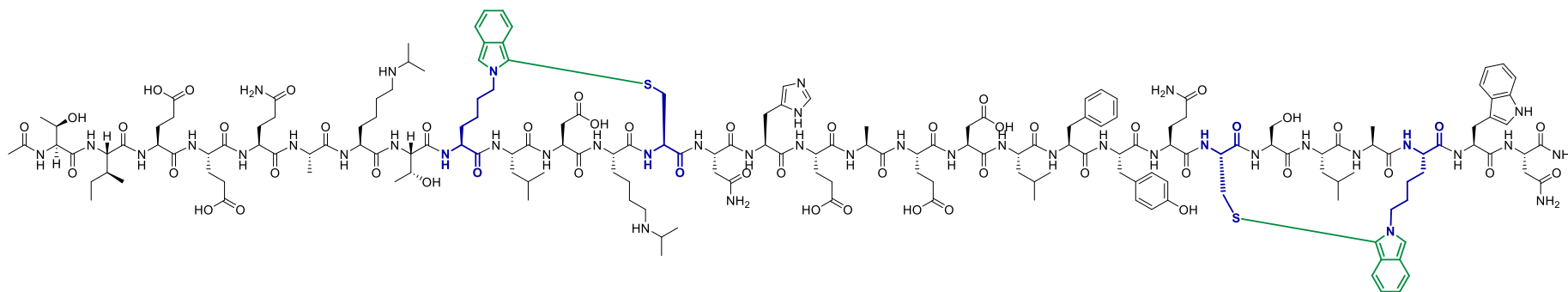


Peptide	Sequence
1	CKCK
2	CKKC
3	KCCK
4	KCKC

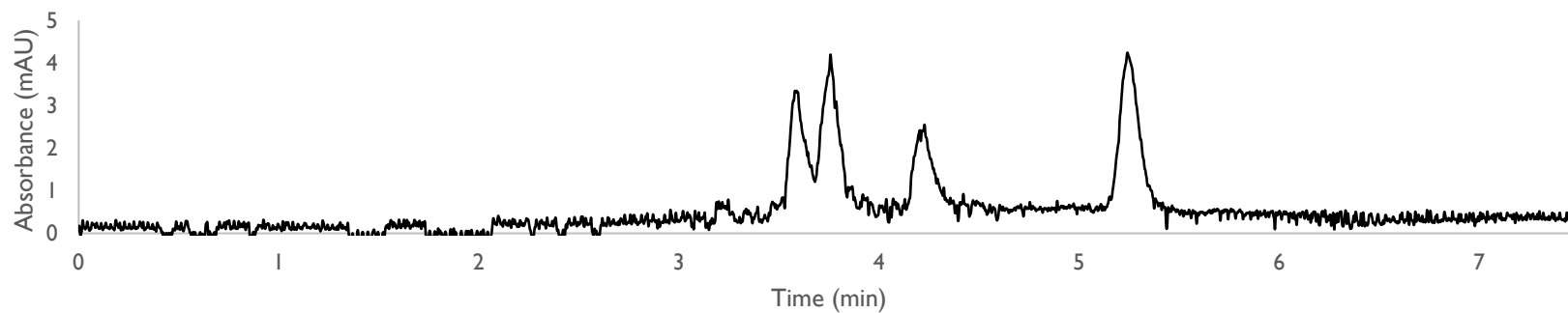
(FIICk): Peptide 3 Double Flicking



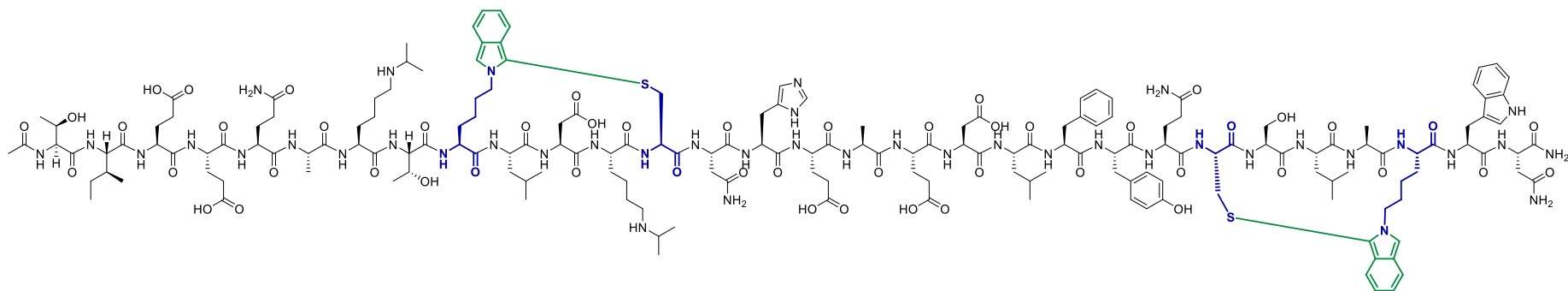
(FIICK): Peptide 3 Double Flicking



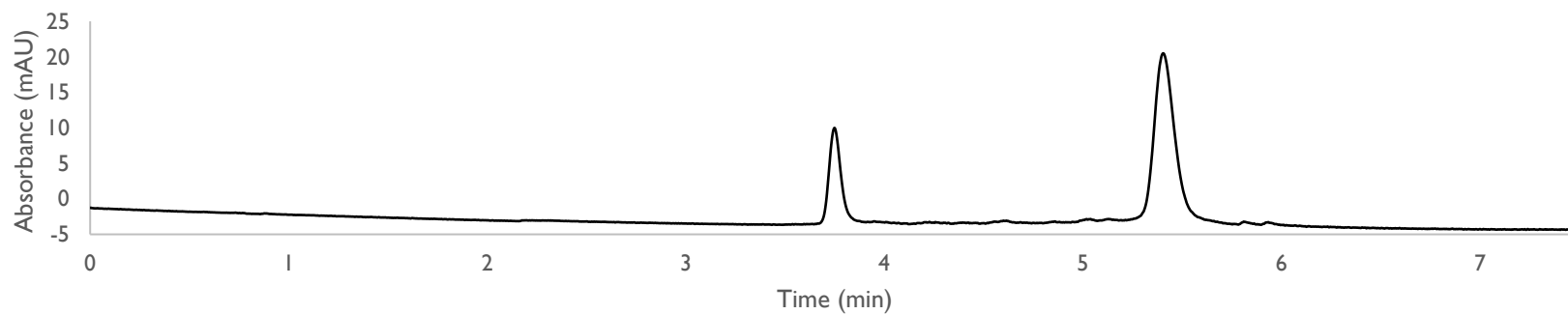
Peptide 3 Double Flicked Crude (335nm) 2eq OPA



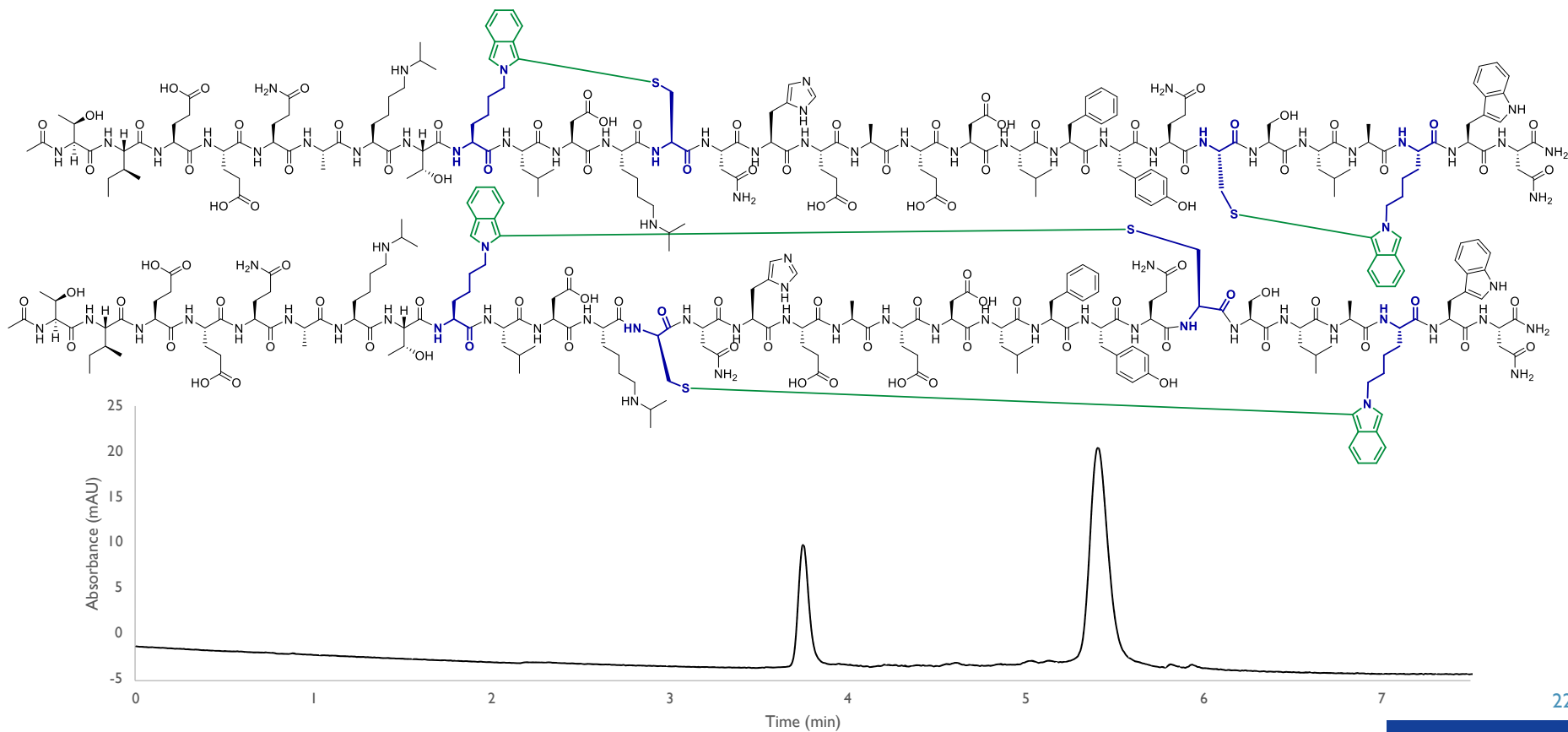
(FIICK): Peptide 3 Double Flicking



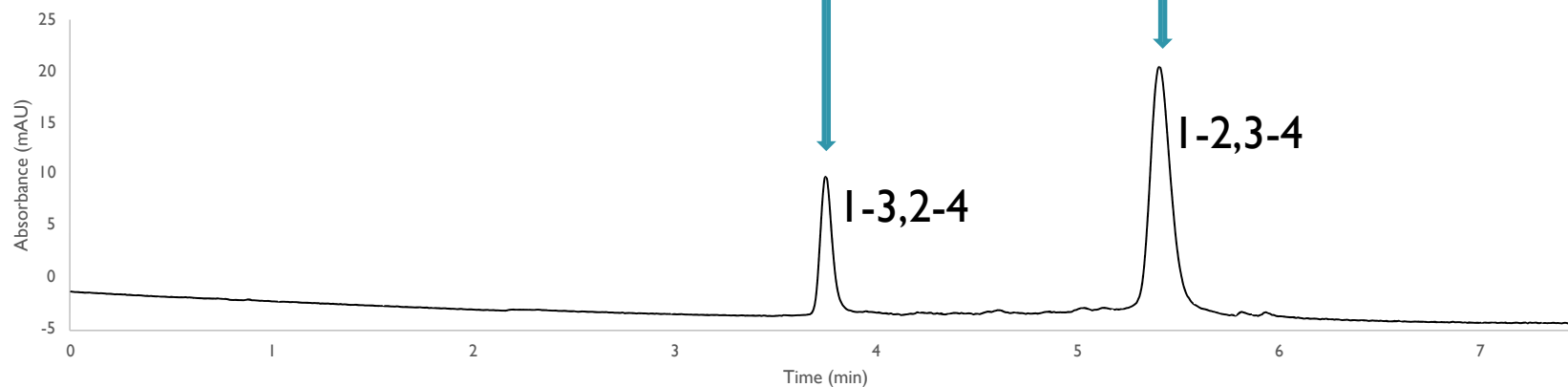
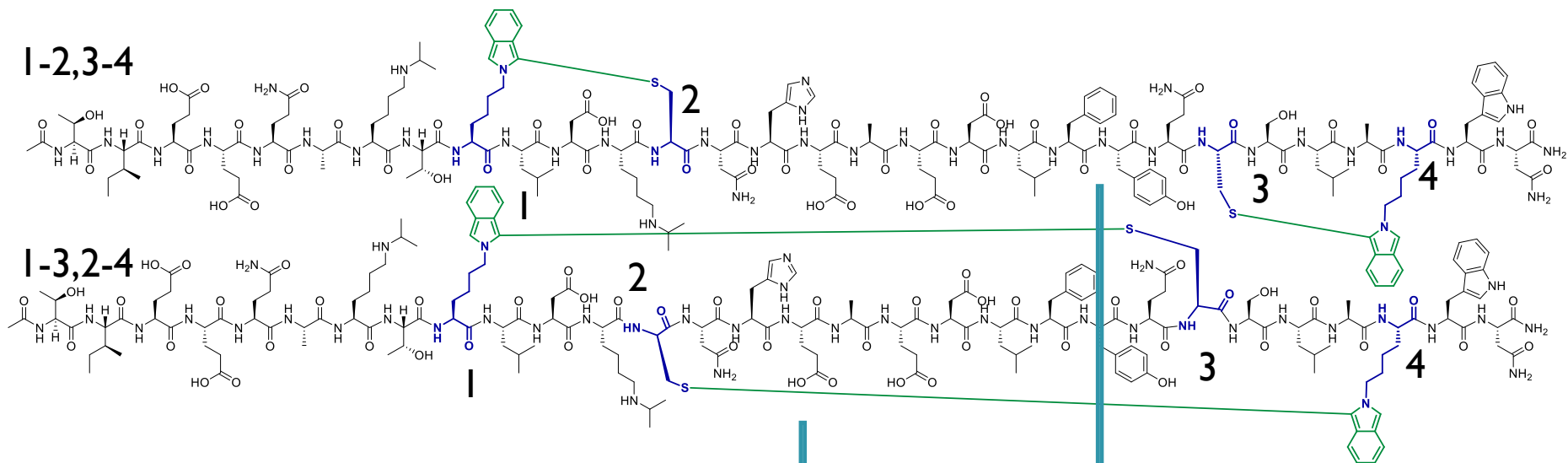
Peptide 3 Double Flicked Crude (335nm) 10eq OPA



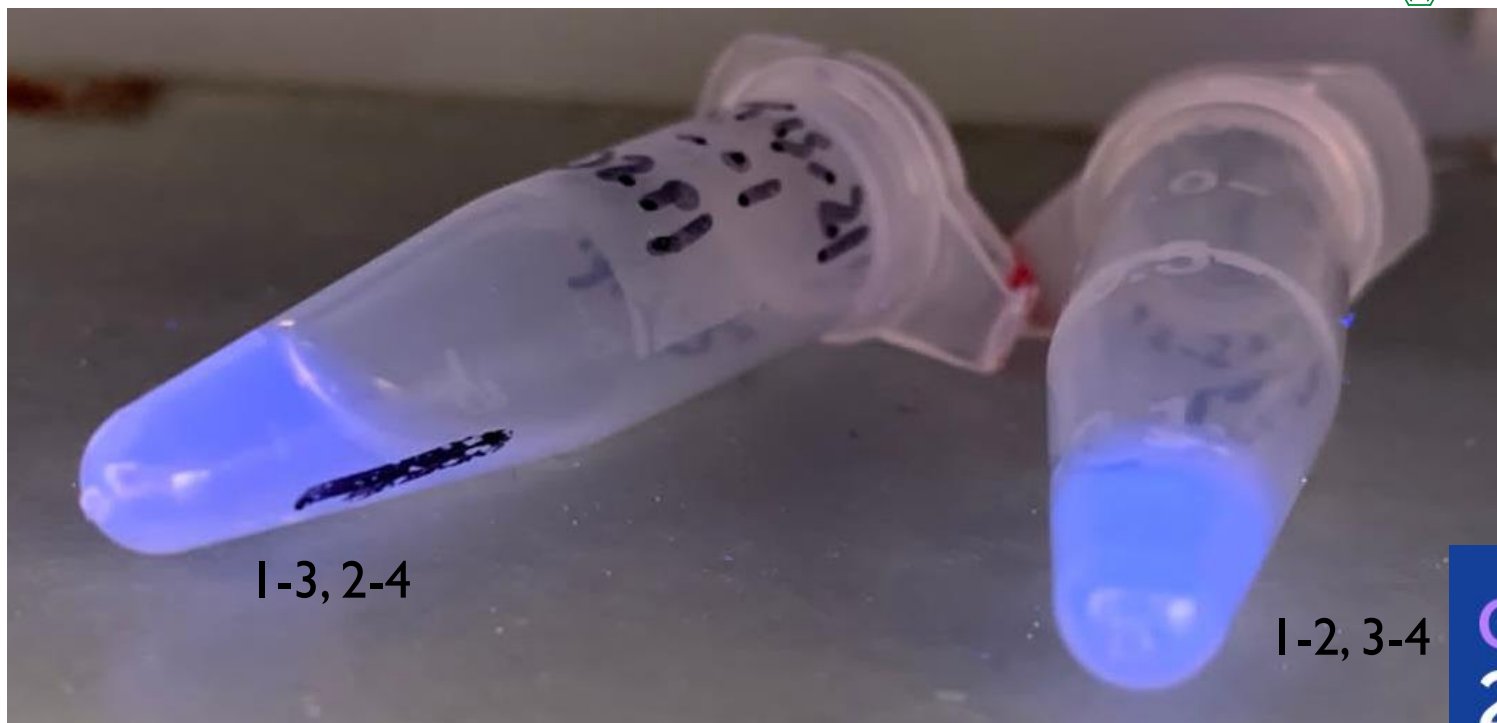
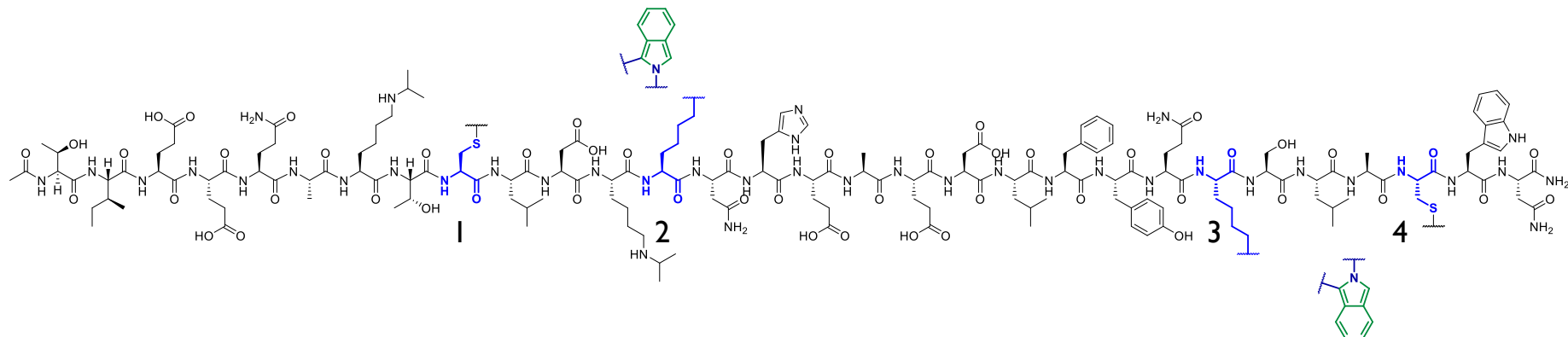
(FIICK): Peptide 3 Double Flicking

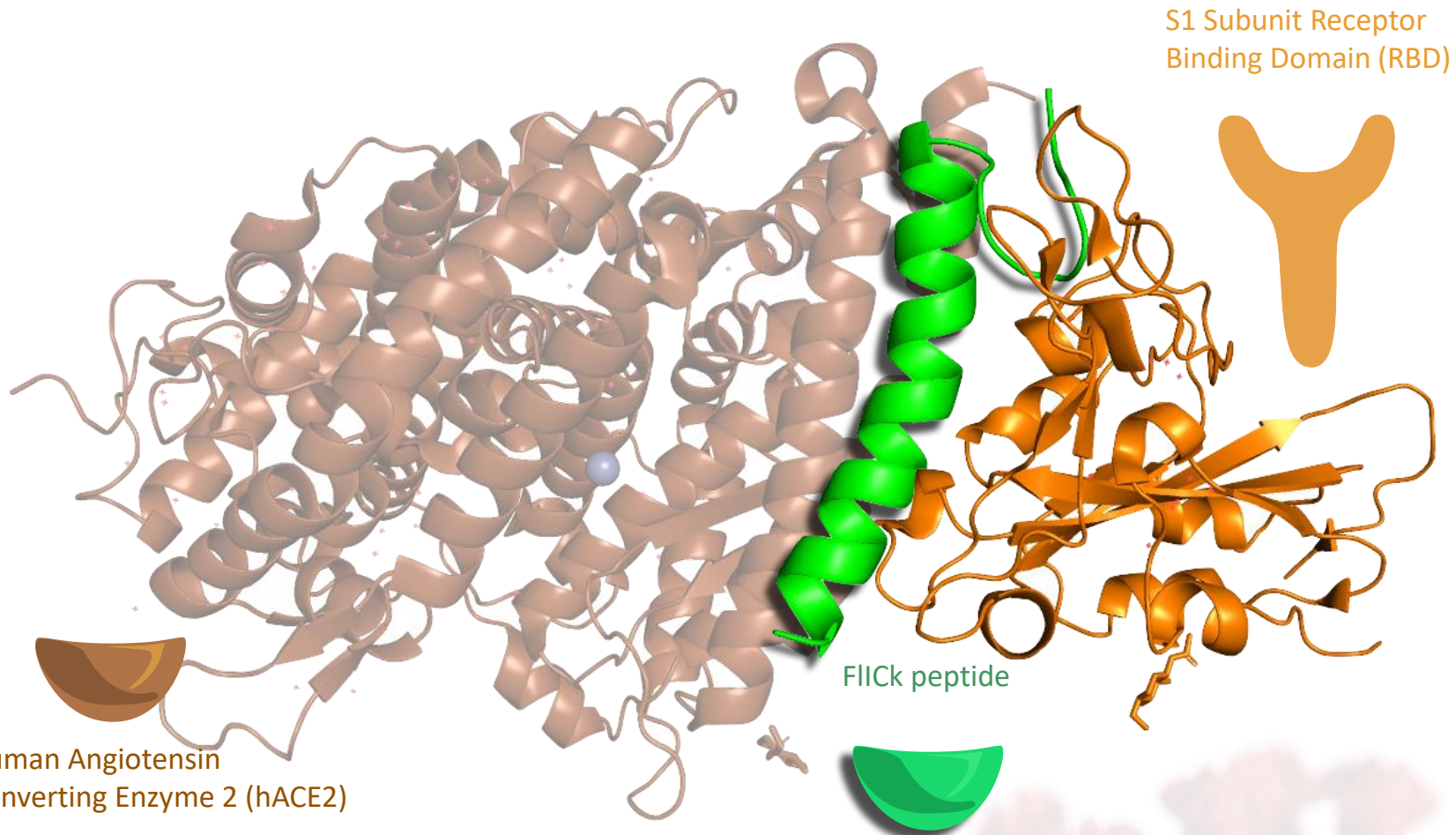


(FI)Ck: Peptide 3 Double Flicking



(FIICk): Double Flicked Peptide Fluorescence





Enzyme ImmunoSorbent Assay
ELISA

ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

Positive results/ good binder



ELISA: Enzyme ImmunoSorbent Assay

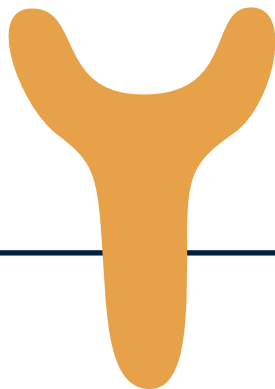
Negative results/ poor binder



hACE2



FIICk Peptide
(Poor)



SARS-CoV-2
Spike RBD

ELISA Plate

Positive results/ good binder



FIICk Peptide
(Good)



hACE

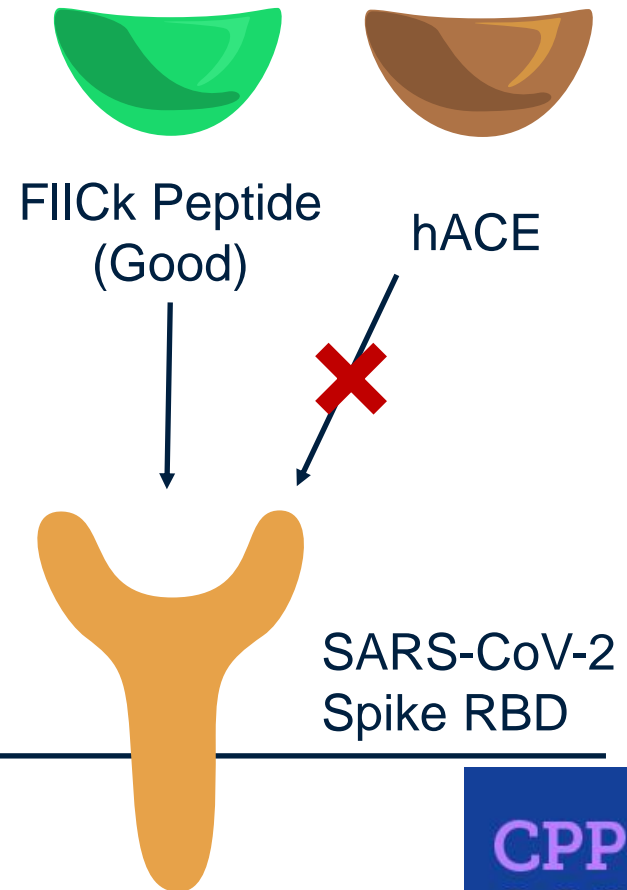
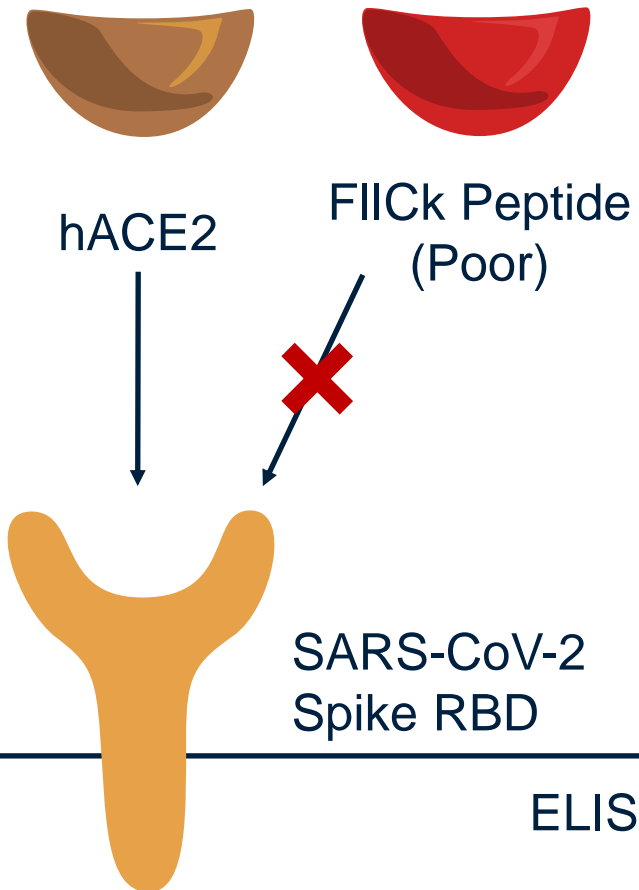


SARS-CoV-2
Spike RBD

ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

Positive results/ good binder

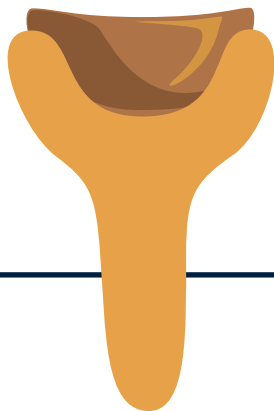


ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder



FIICk Peptide
(Poor)



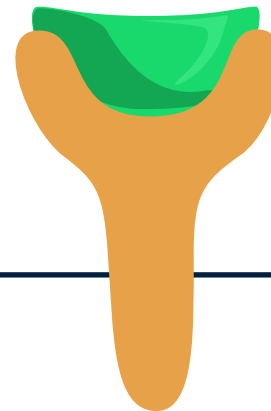
hACE2

SARS-CoV-2
Spike RBD

Positive results/ good binder



hACE



FIICk Peptide
(Good)

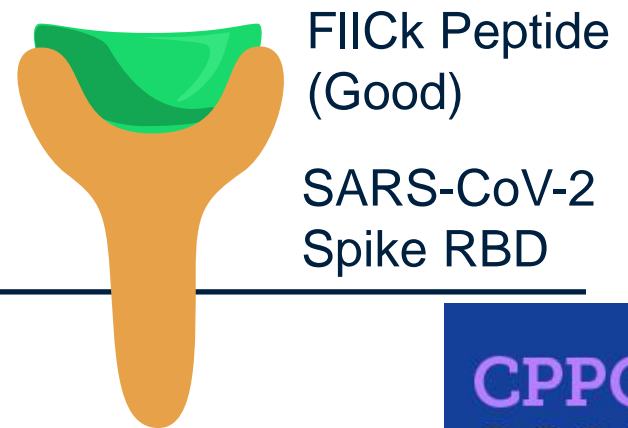
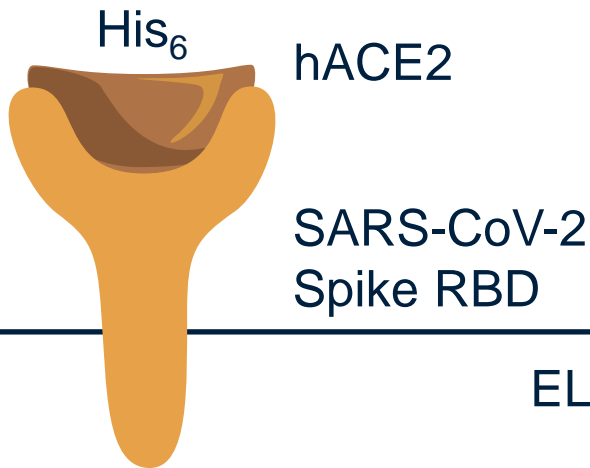
SARS-CoV-2
Spike RBD

ELISA Plate

ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

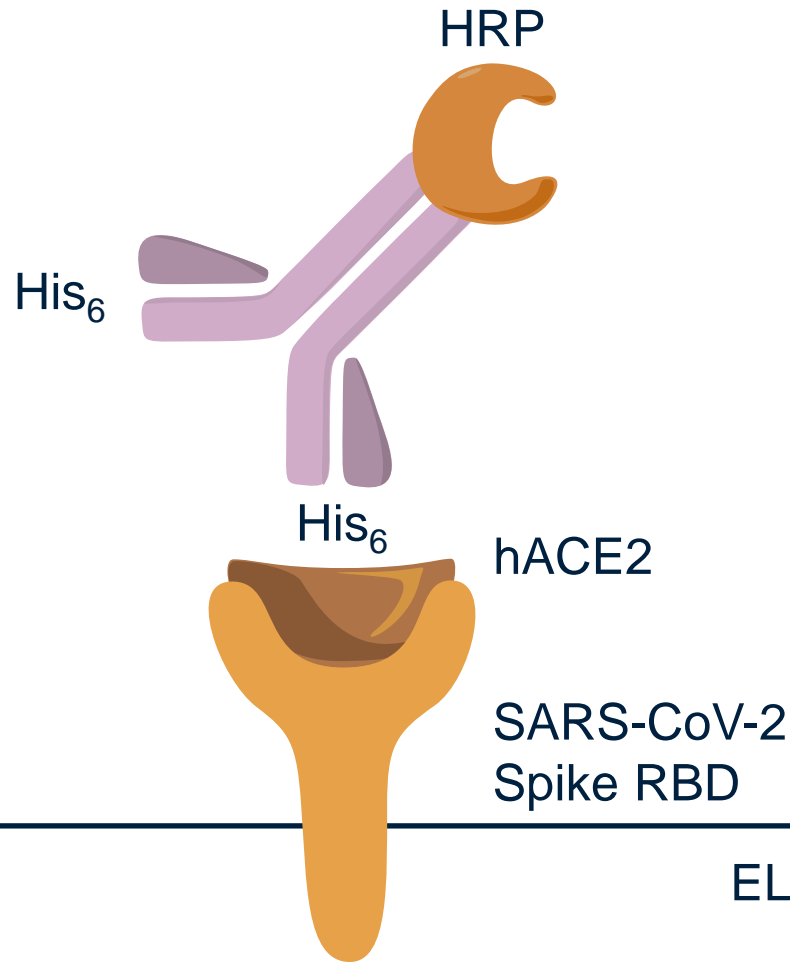
Positive results/ good binder



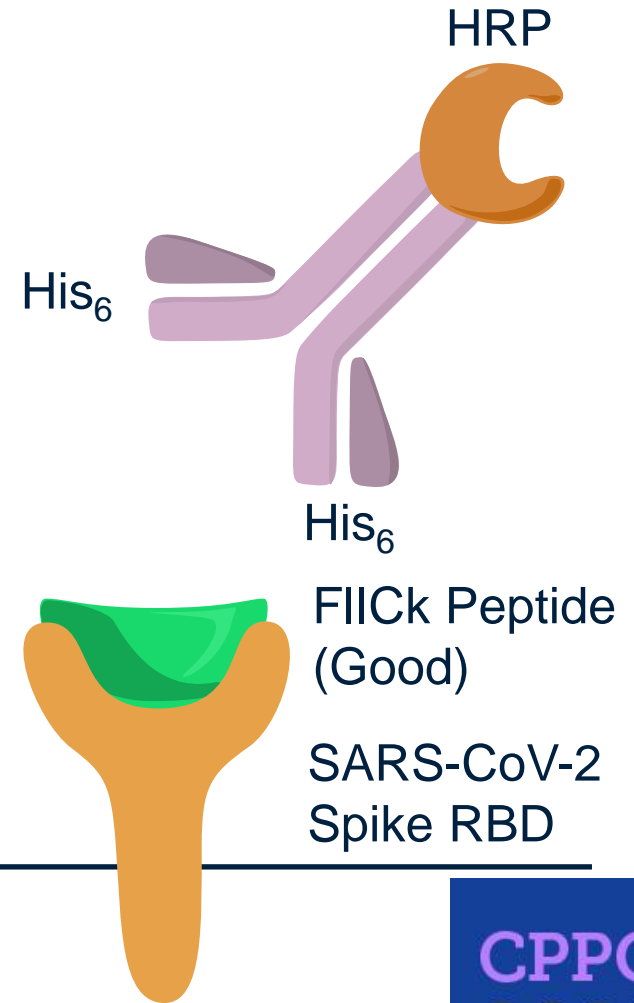
ELISA Plate

ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder



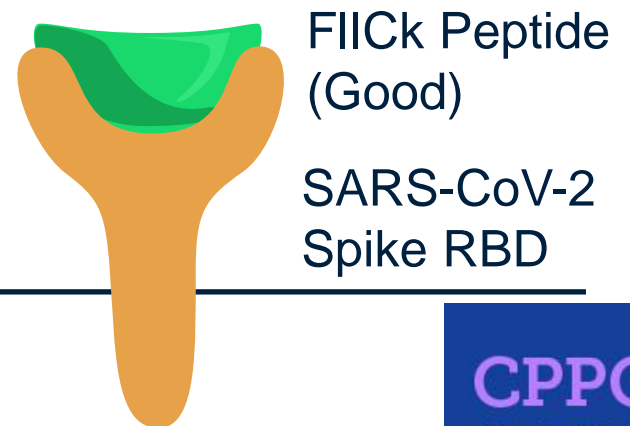
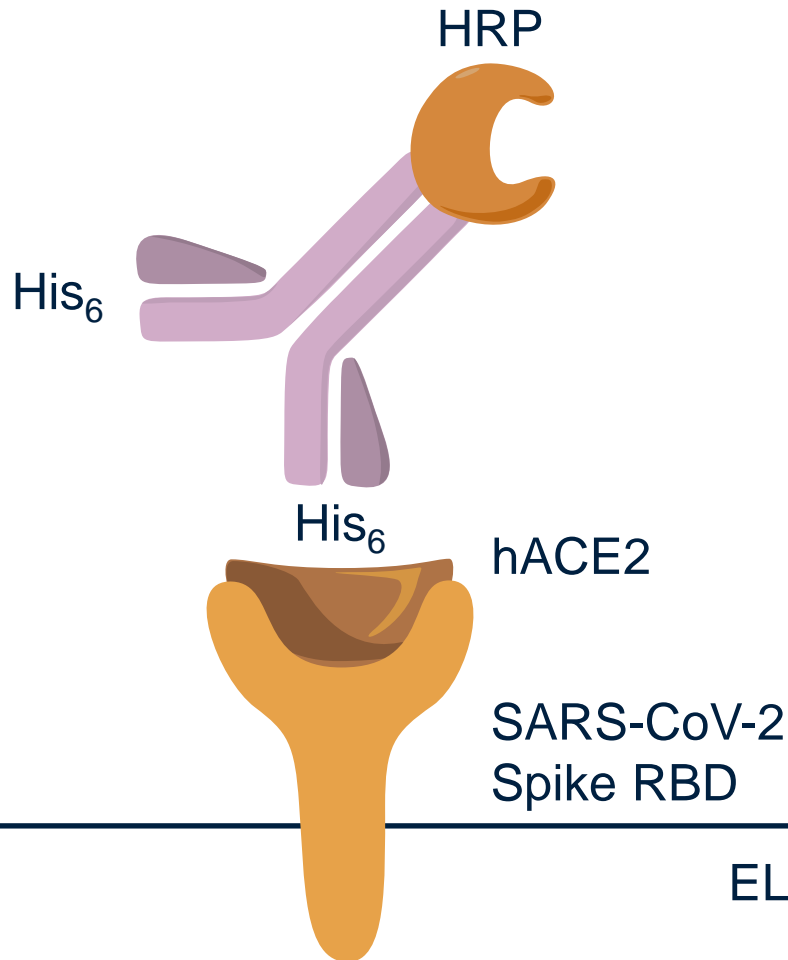
Positive results/ good binder



ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

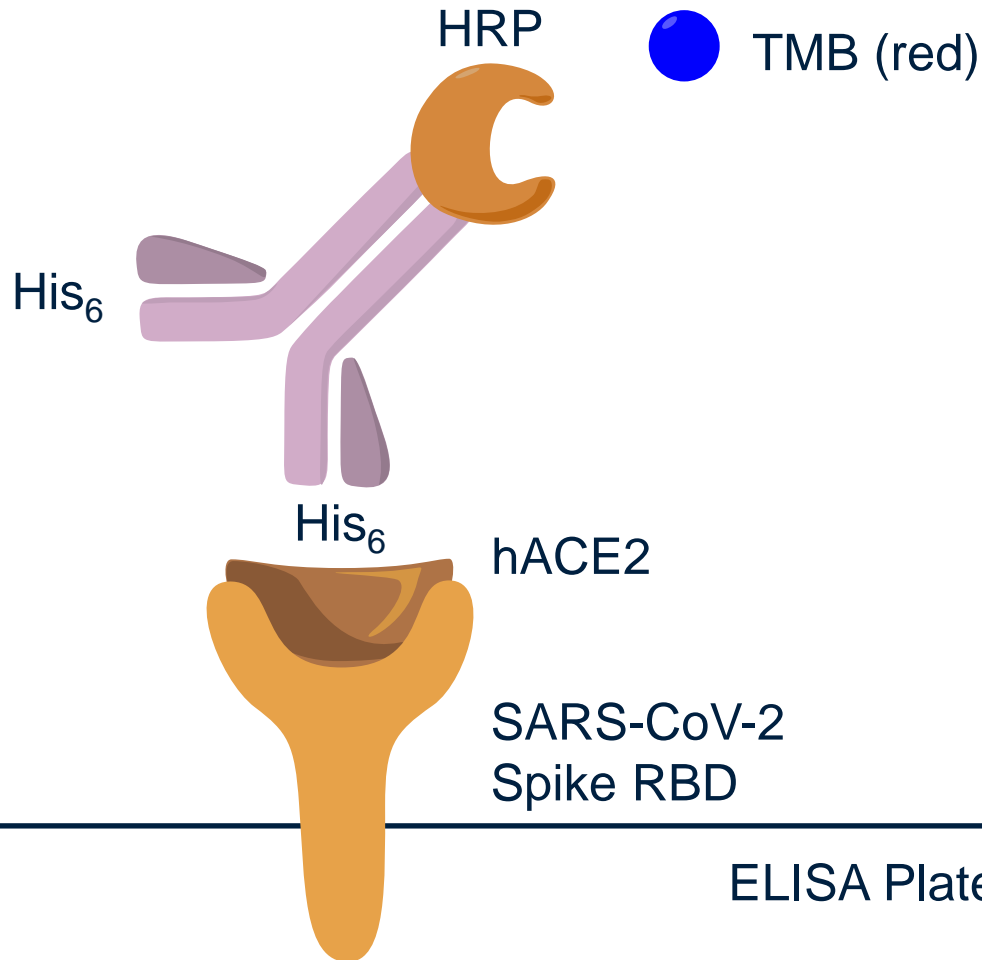
Positive results/ good binder



ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

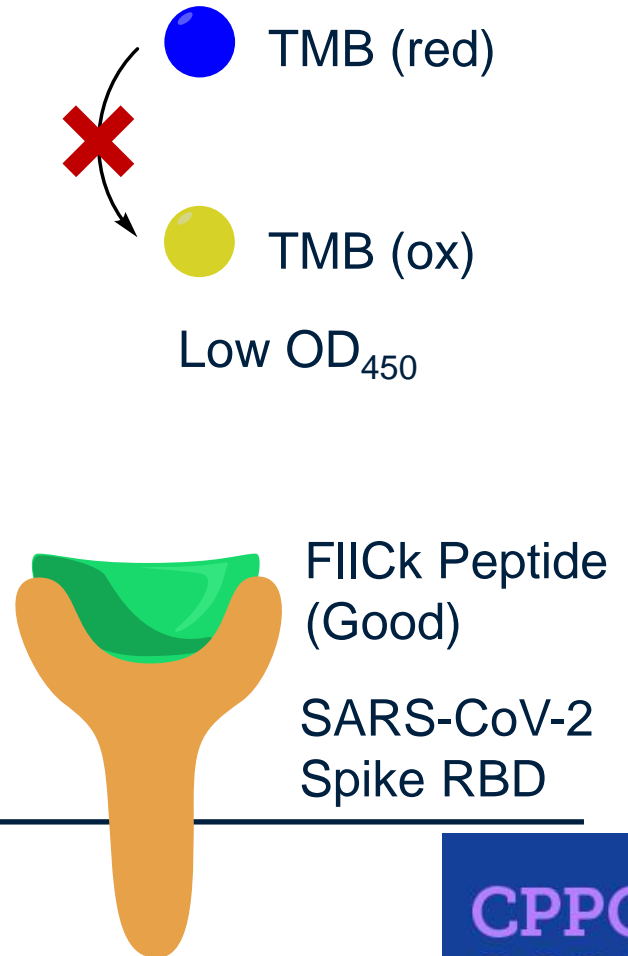
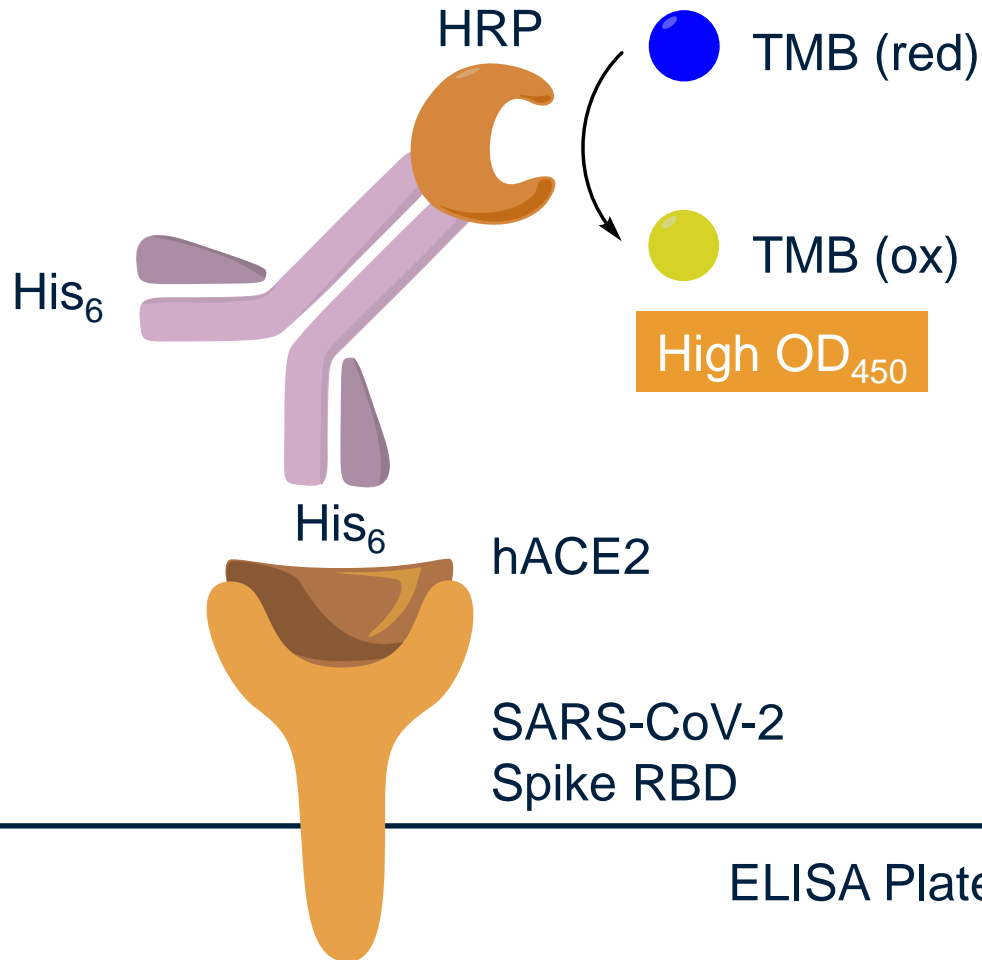
Positive results/ good binder



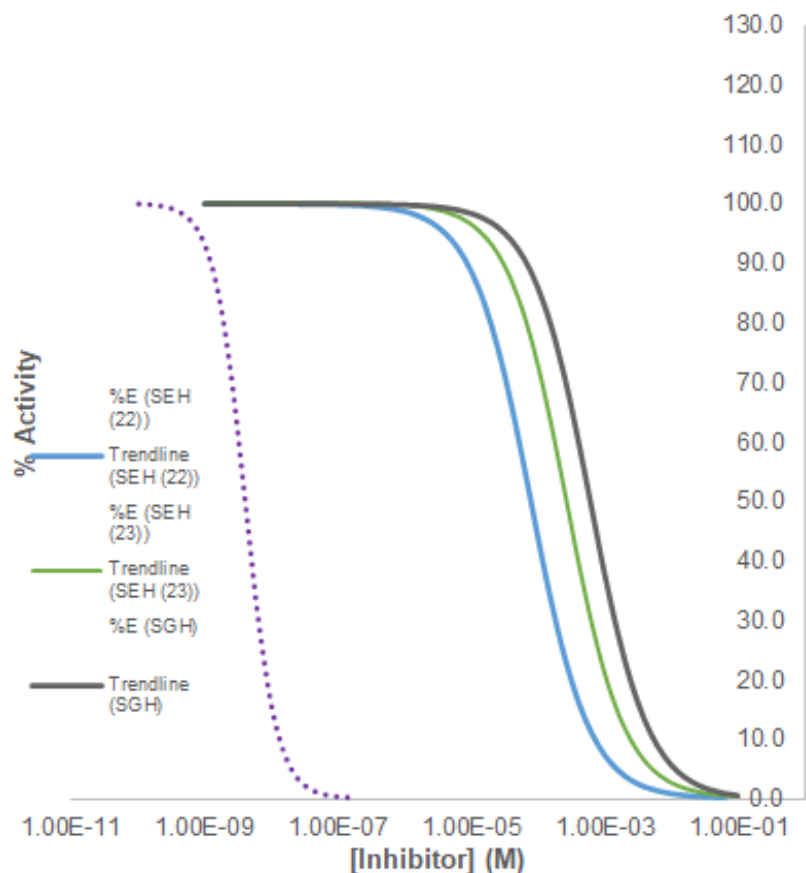
ELISA: Enzyme ImmunoSorbent Assay

Negative results/ poor binder

Positive results/ good binder



(FIICK): Double Flicked Peptide IC50s

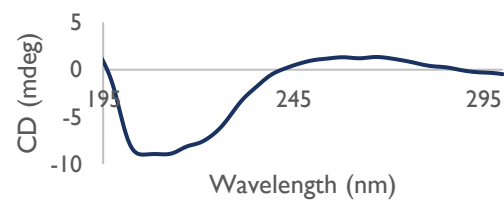


Entry	I234	Topology	R.T. (min)	IC50(μM)
1	CKCK	1-4,2-3	4.71	Not active (>900)
2	CKCK	1-2,3-4	5.21	Not active (>900)
3	CKKC	1-3,2-4	3.57	77
4	CKKC	1-2,3-4	5.22	260
5	KCCK	1-3,2-4	3.75	Not active (>500)
6	KCCK	1-2,3-4	5.39	Not active (>900)
7	KCKC	1-4,2-3	5.00	Not active (>900)
8	KCKC	1-2,3-4	5.36	620

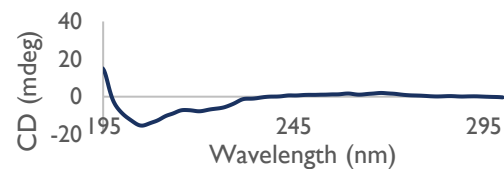
(FIICK): Double Flicked Peptide IC50s

Entry	XXXX 1234	Topology	R.T. (min)	IC50(μ M)
1	CKCK	1-4,2-3	4.71	Not active (>900)
2	CKCK	1-2,3-4	5.21	Not active (>900)
3	CKKC	1-3,2-4	3.57	77
4	CKKC	1-2,3-4	5.22	260
5	KCCK	1-3,2-4	3.75	Not active (>500)
6	KCCK	1-2,3-4	5.39	Not active (>900)
7	KCKC	1-4,2-3	5.00	Not active (>900)
8	KCKC	1-2,3-4	5.36	620

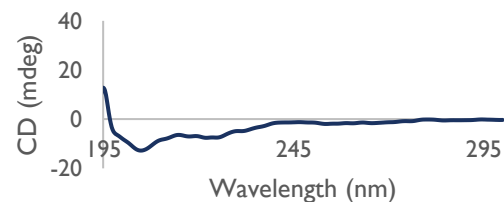
Entry 3 CD Spectrum



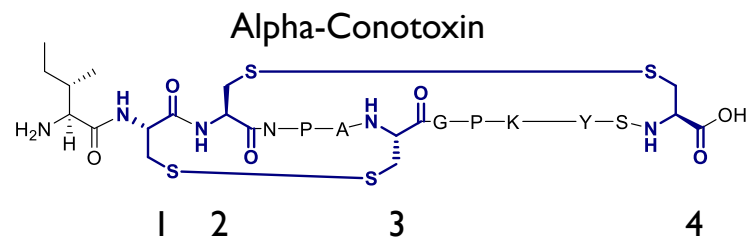
Entry 4 CD Spectrum



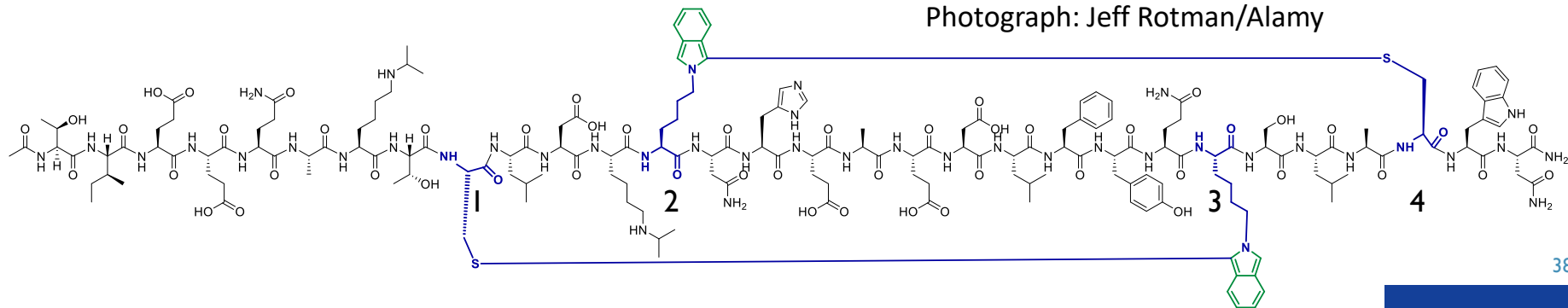
Entry 8 CD Spectrum



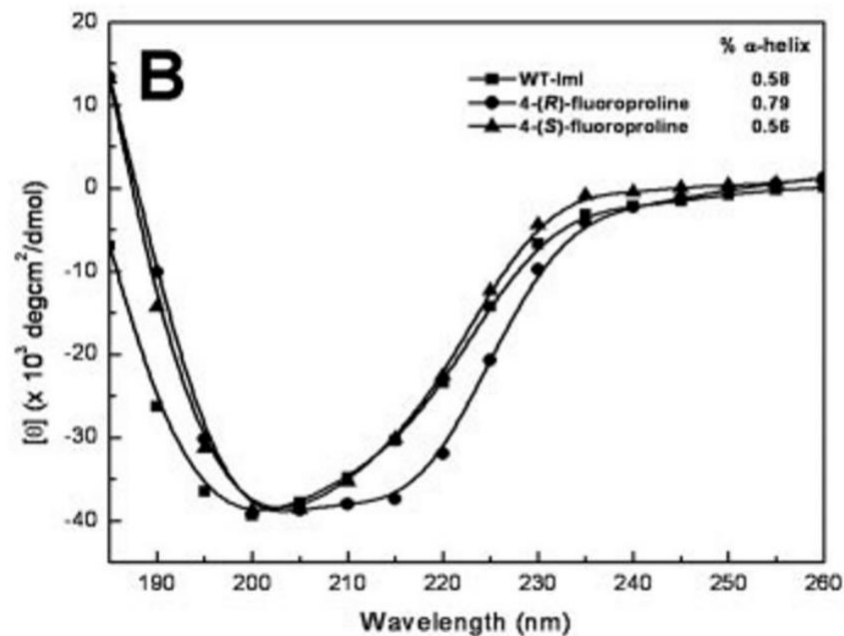
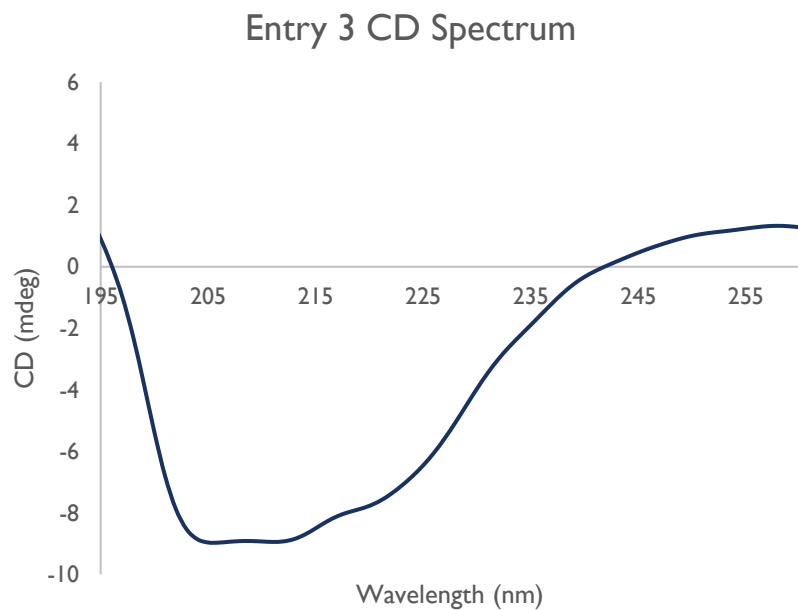
(FIICk): Alpha Conotoxin Topology



Photograph: Jeff Rotman/Alamy











(FIICK): Alpha Conotoxin Topology



Armishaw, C.; Jensen, A. A.; Balle, T.; Clark, R. J.; Harpsøe, K.; Skonberg, C.; Liljefors, T.; Strømgaard, K. *J. Biol. Chem.* **2009**, *284* (14), 9498–9512.
<https://doi.org/10.1074/jbc.M806136200>.

(FIICK): Effect of chaotropes and salts on Peptide 3 crosslinking selectivity

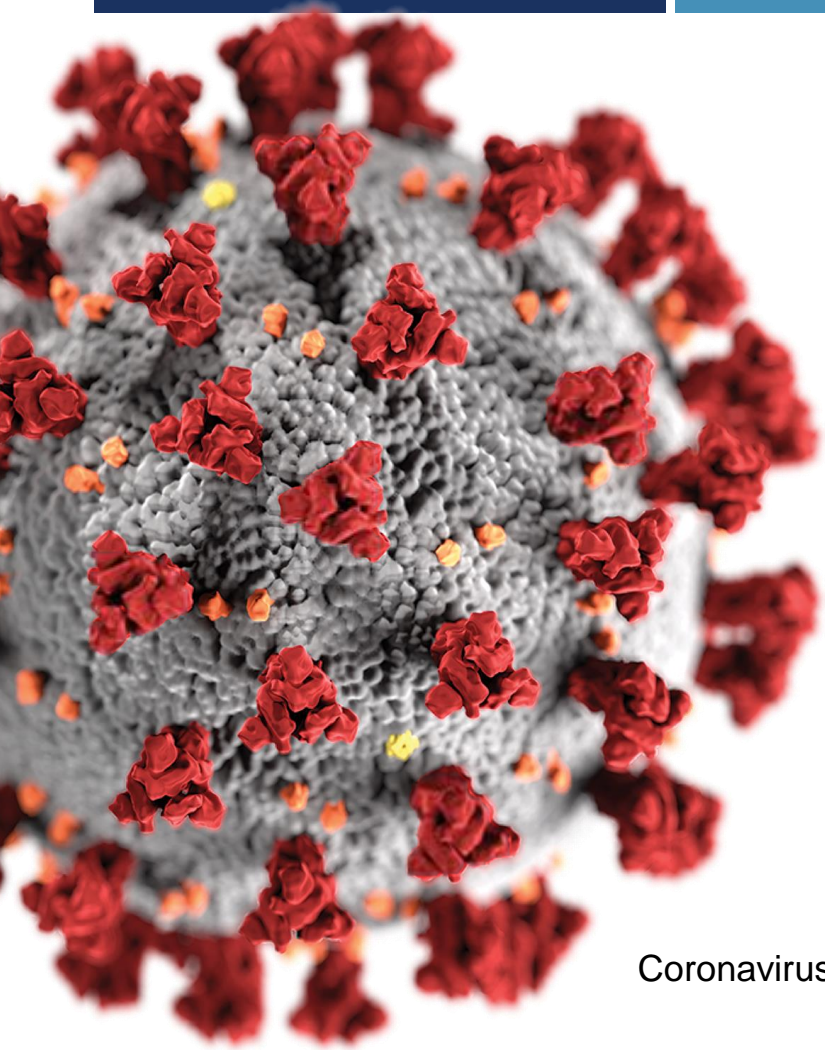
Chaotrope/salt (concentration)	1-2,3-4 : 1-3,2-4 ratio	HPLC chromatogram
none	2.5	
Guanidinium chloride (2% w/w)	2.5	
Urea (2% w/w)	2.4	
Polysorbate 20 (2% w/w)	3.2	
Polysorbate 80 (2% w/w)	2.7	
Triton X-100 (2% w/w)	3.5	
Mr. Clean® dish detergent (2% w/w)	2.9	
Magnesium chloride (450 mM)	2.3	

(FIICK): Effect of temperature and solvent on Peptide 3 crosslinking selectivity

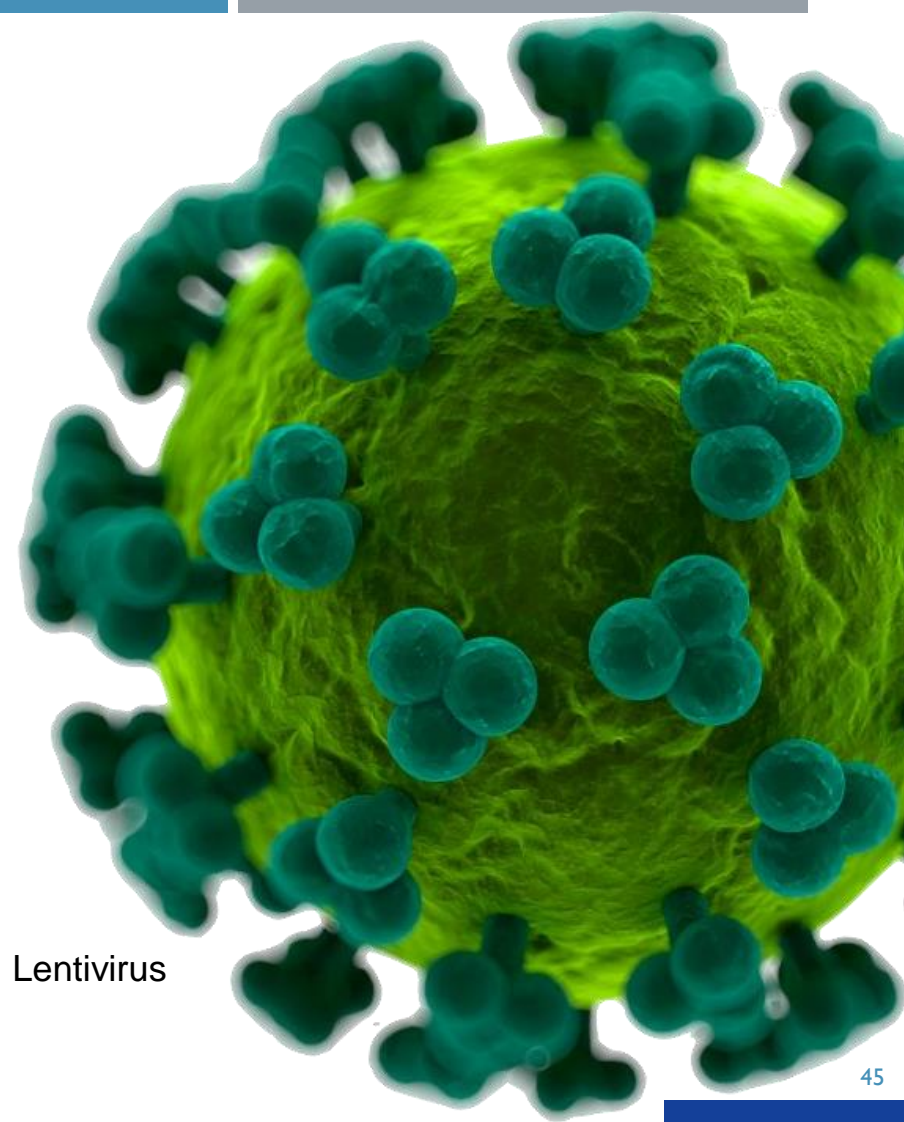
Solvent	Temperature (°C)	1-2,3-4 : 1-3,2-4 ratio	HPLC chromatogram
Sodium borate buffer, pH 9	4	1.90	
Sodium borate buffer, pH 9	20	2.52	
Sodium borate buffer, pH 9	37	2.34	
Sodium borate buffer, pH 9/EtOH (1:1)	-20	7.78	
Sodium borate buffer, pH 9/EtOH (1:1)	20	7.46	

(FIICK): Effect of solvent on Peptide 3 crosslinking selectivity

Peptide	Solvent	1-2,3-4 : 1-3,2-4 ratio	HPLC chromatogram
3	Sodium borate buffer, pH 9	2.52	
3	Sodium borate buffer, pH 9/EtOH (1:1)	7.46	
4	Sodium borate buffer, pH 9	2.26	
4	Sodium borate buffer, pH 9/DMF (1:1)	3.95	
4	Sodium borate buffer, pH 9/DMA (1:1)	3.39	
4	Sodium borate buffer, pH 9/MeCN (1:1)	2.46	
4	Sodium borate buffer, pH 9/THF (1:1)	6.98	



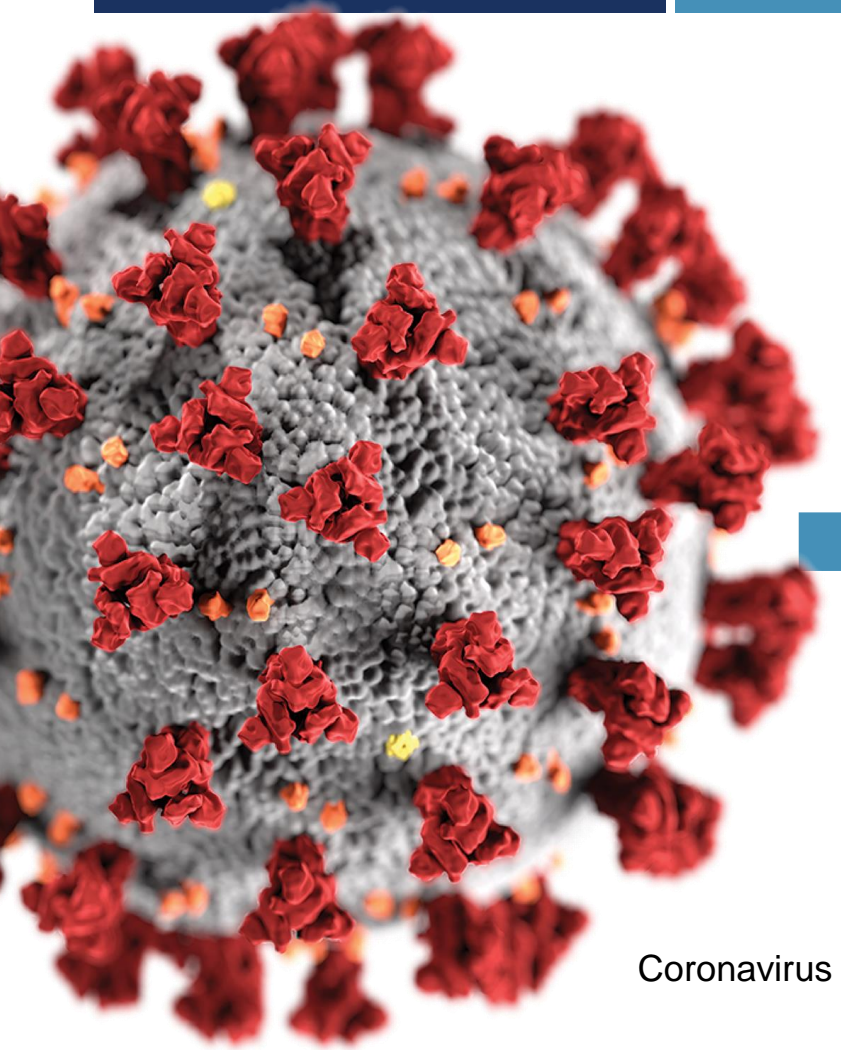
Coronavirus



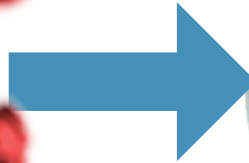
Lentivirus

Future direction

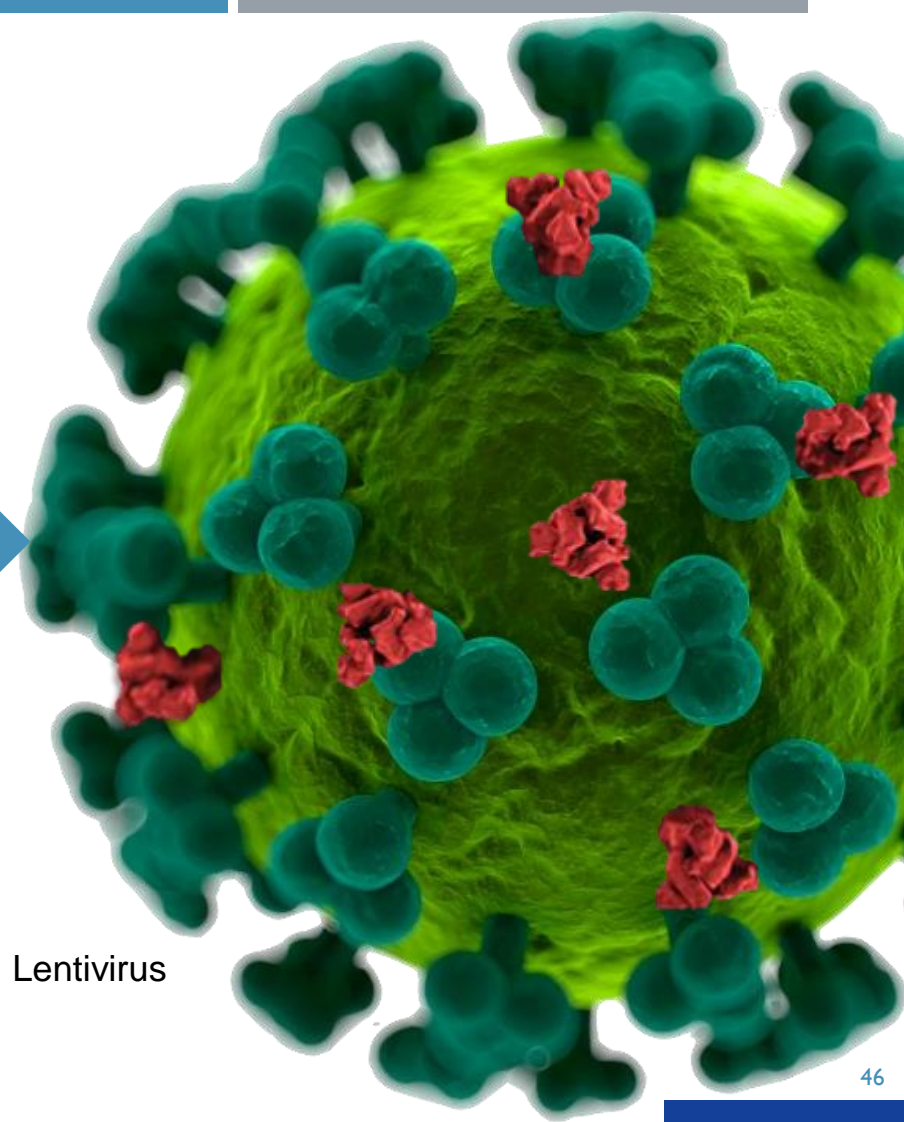
Pseudovirus Neutralization Assay (PNA)



Coronavirus

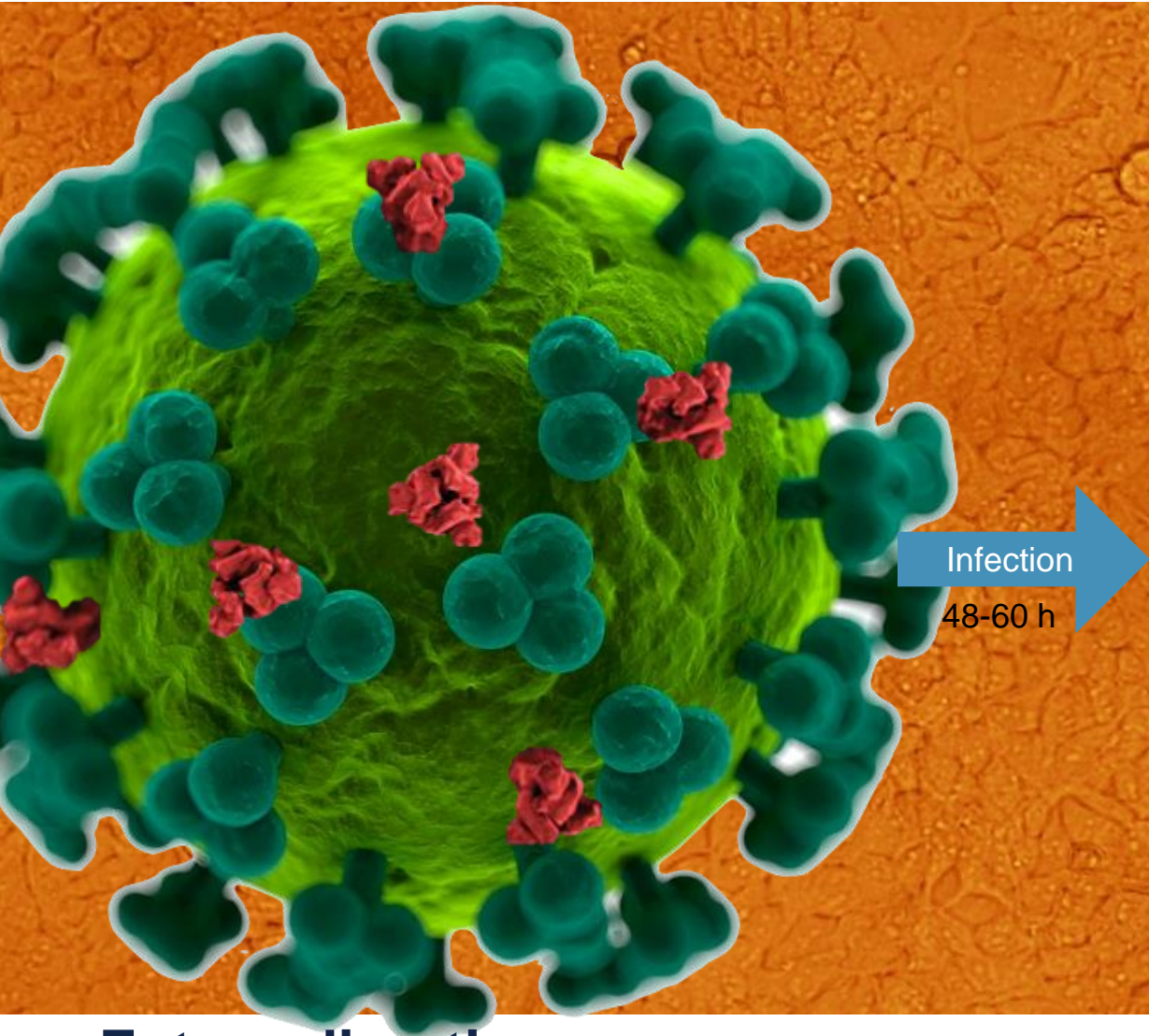


Lentivirus



Future direction

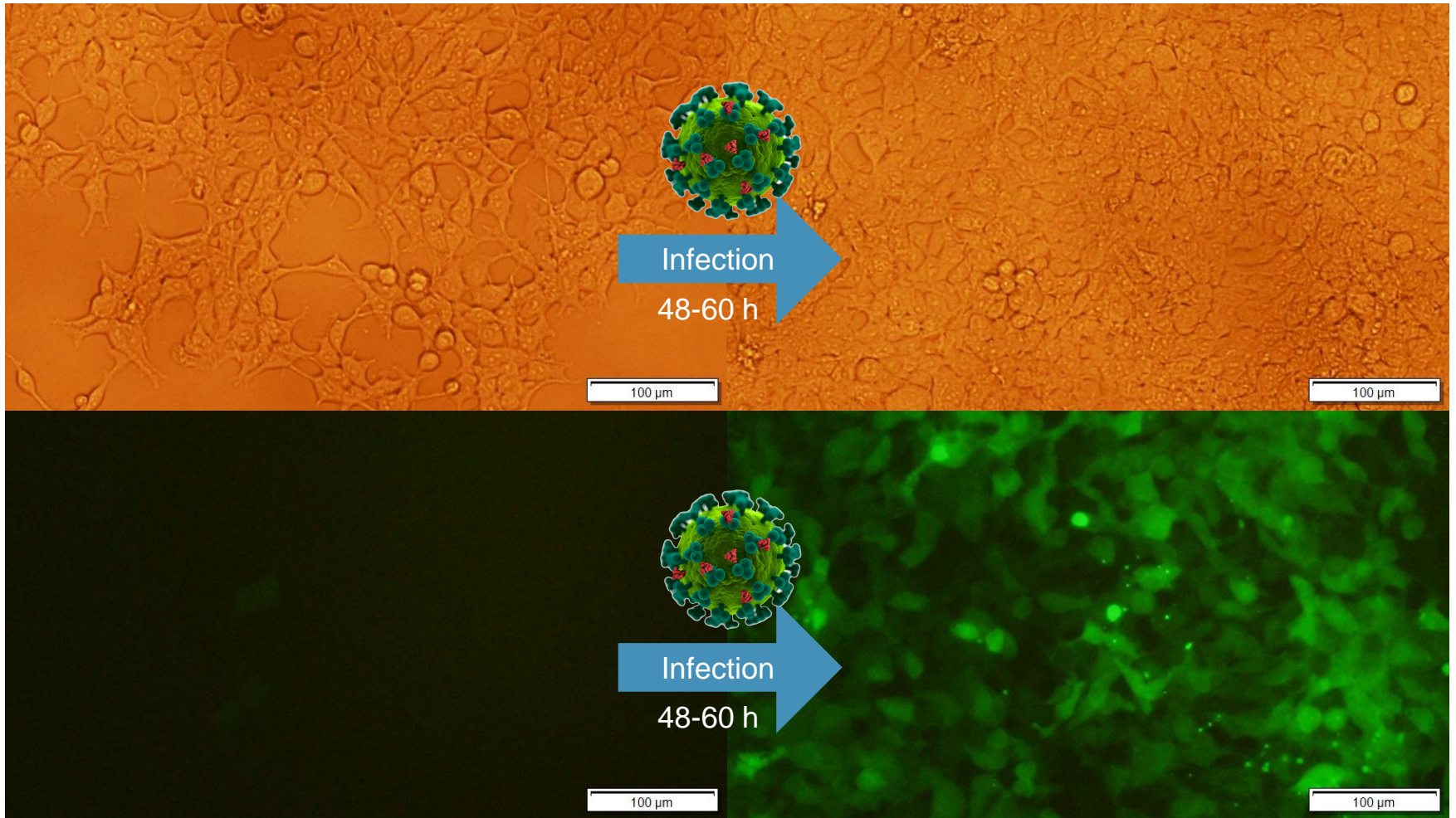
Pseudovirus Neutralization Assay (PNA)



Infection
48-60 h

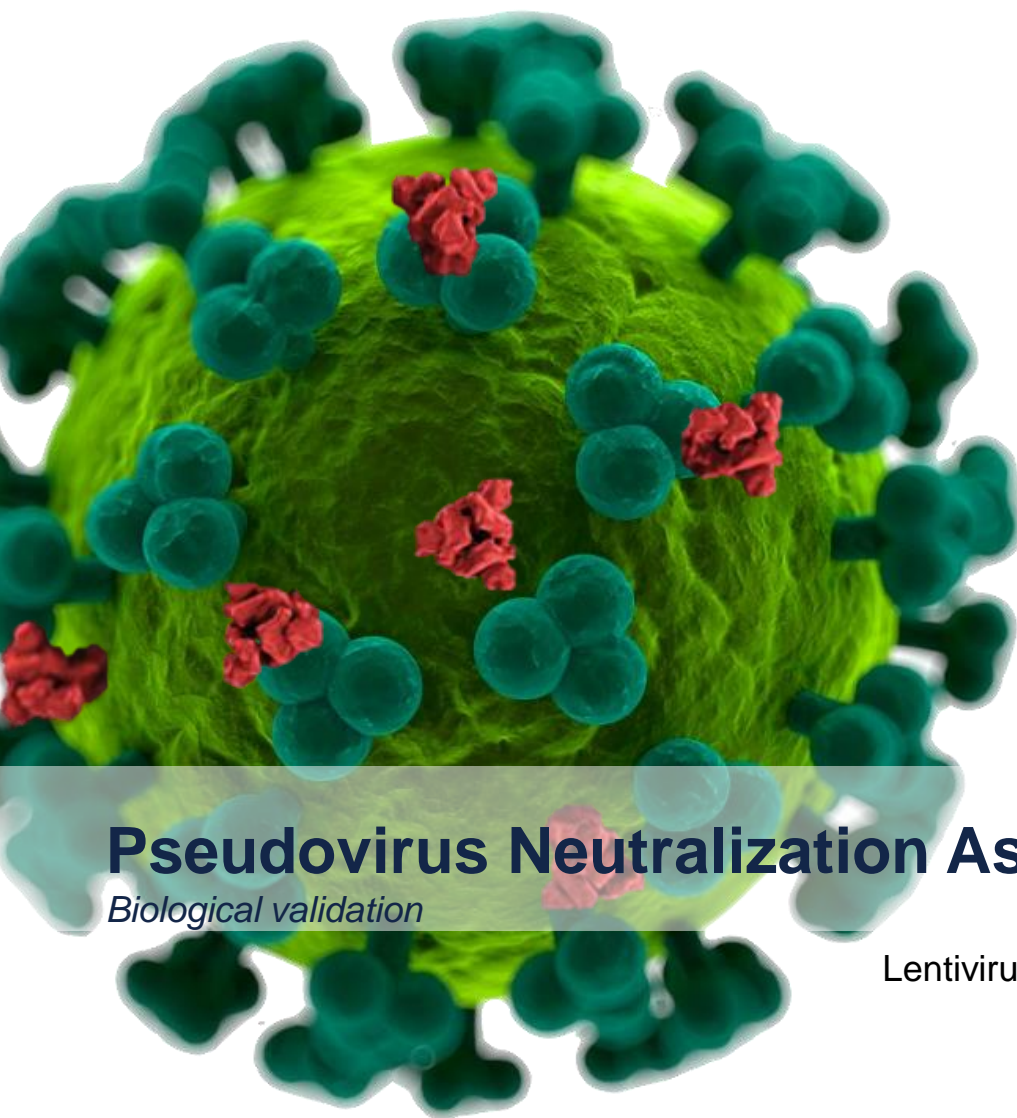
Future direction

Pseudovirus Neutralization Assay (PNA)



Future direction

Pseudovirus Neutralization Assay (PNA)



→
+ ACE2⁺ cells

✓ Infection

C1=CC=C2C(=C1)N(C2)S
+ inhibitor

→
+ ACE2⁺ cells

✗ Infection

Pseudovirus Neutralization Assay

Biological validation

Lentivirus

Acknowledgements

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THE UNIVERSITY OF BRITISH COLUMBIA

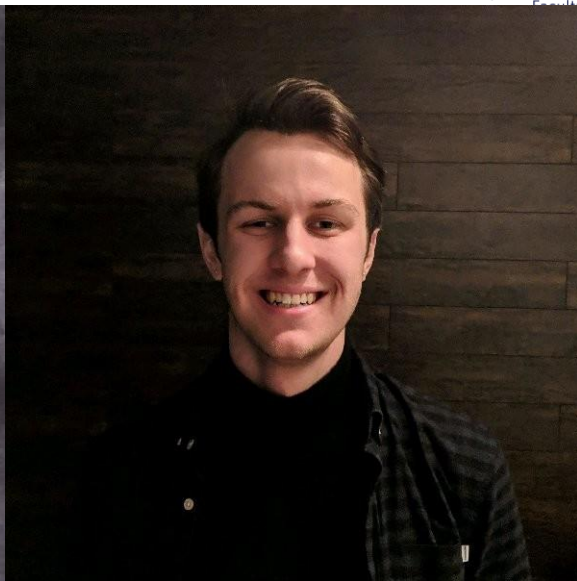
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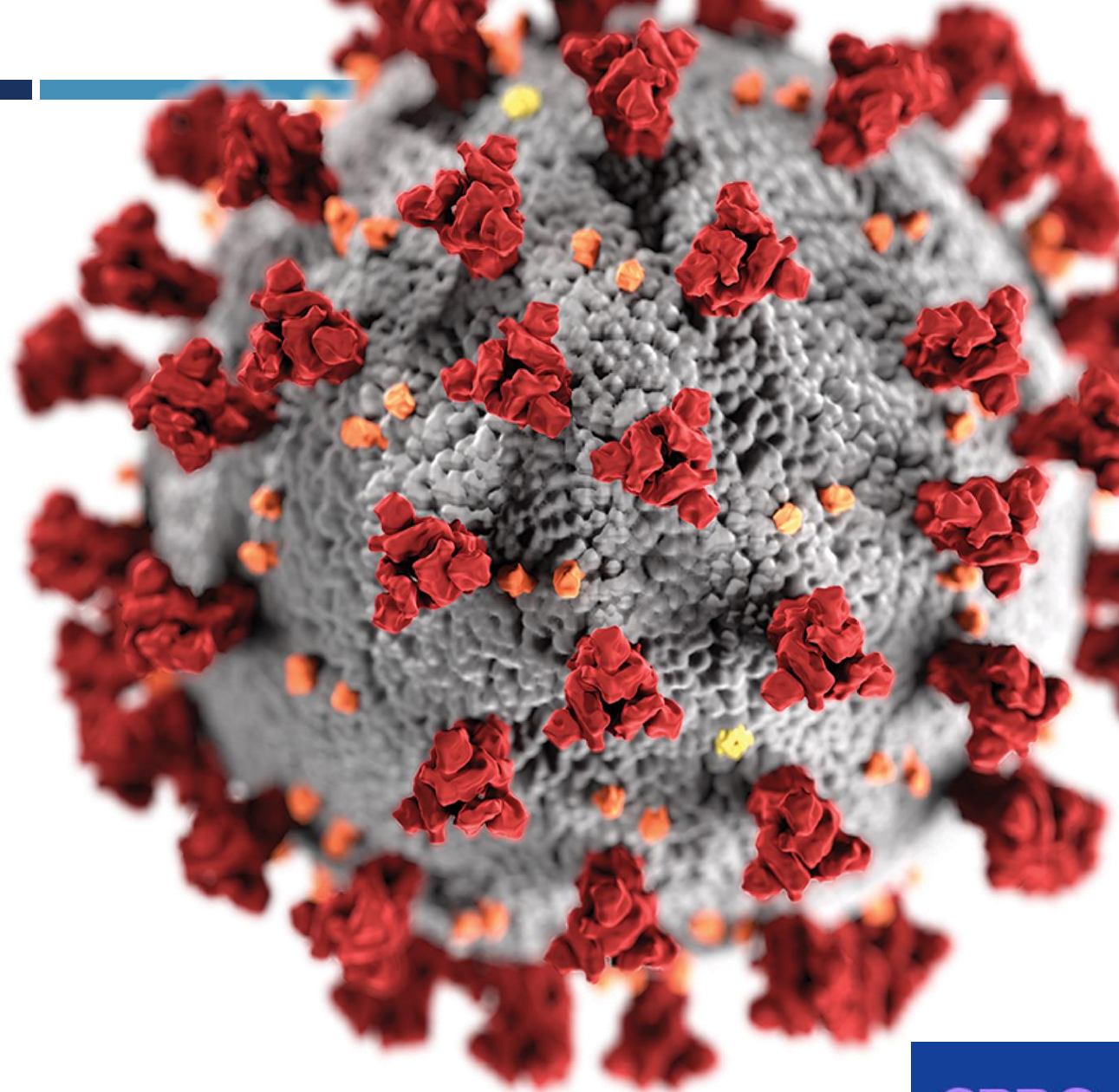
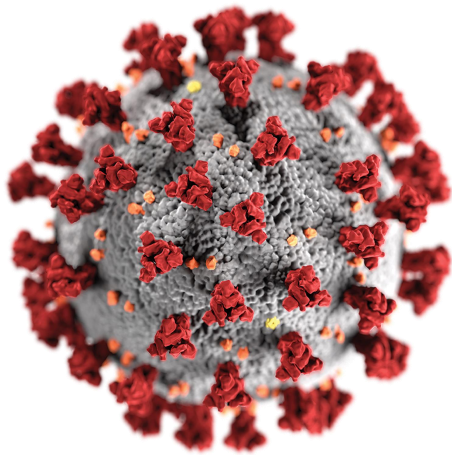
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Questions?