

Of the many mutations in omicron the most interesting two are in the SEB domain.

Staphylococcus Enterotoxin B is a peptide that is classified as a bioweapon.

When ingested it creates symptoms very similar to COVID and it's know to trigger cytokine storms



It's well described in Cheng et al.

This known toxin was warp sped into spike only Vaccines. SEB is directly adjacent to the FCS.

Let's have a look at what Ahanotu et al has to say about SEB exposure and it's capacity as a bioweapon.

nnas.org/content/117/41

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https://www.pnas.org/content/117/41/25254

Exposure to these peptides has been known to cause cytokine storms. They are superantigens.

liebertpub.com/doi/pdfplus/10...

Now look at Omicron spike protein 671-692.

Two amino acid changes and one is the infamous proline change near the FCS. Prolines are right angle brackets in proteins. When they change, they alter structure.

P681H

N679K

This may attenuate the SEB toxicity. Fewer Ckine storms.

Ahanotu et al considerations on SEB as a bioweapon as prescient.

Some of this is discussed in our preprint.

I was not aware of these SEB mutations in omicron when I wrote this.

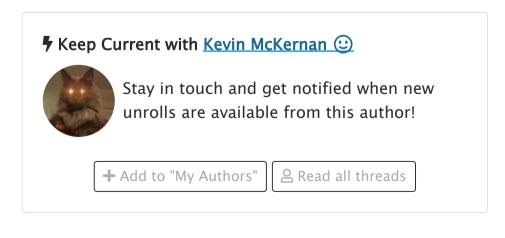
I was C19 +ve when I submitted this, ironically.

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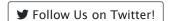
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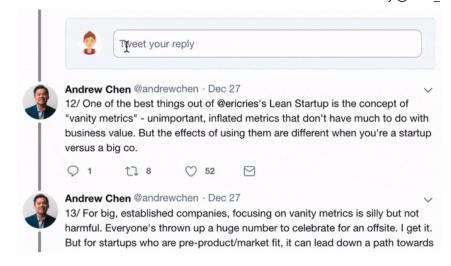




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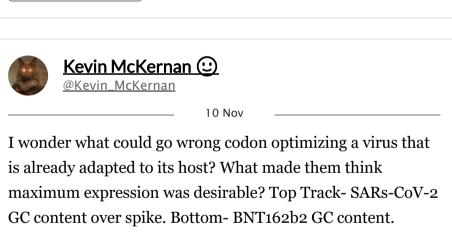




When you Warp Speed your codon optimizations and forget to look to at quadruplex G formation. These mRNA Vaccines should be considered Pro-Drugs, where proof of the final drug is measured, not assumed. osf.io/bcsa6/

So Molbio folks like to Jack the expression of genes sadly assuming that more is better. To do this they add strong beta globin promoters and swap the synonymous bases in the codons out with more human like codons. This is called codon optimization. There are consequences

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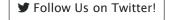


They certainly weren't thinking about secondary structure. This type of stuff matters to RNAses and other components of the innate immune response (Toll Like Receptors). Yes, their goal was to evade innate detection with pseudouridine

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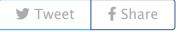
ALERT Is WIV Sloppy with BL4 viruses? What is Nipahvirus sequence doing in C19 patients sequenced by Wuhan Institute of Virology (WIV)? This is a BL4 virus far more deadly than C19. Why is it found in patients WIV sequenced





"We are never going to know how safe the vaccine is in children unless we start giving it". That's the Wisdom of the FDA review on stabbing your kids with experimental medicine.

26 Oct



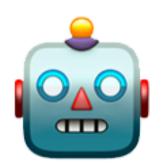
First described here- osf.io/s4td6/ How do we reproduce these claims? Lets walk through it.

1 J.L. D

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morons lips. This is after many parents showcased their wheel chair bound kids and that were vax injured & Pfizer refused to acknowledge the injury. You're never gonna know because your complicit is not looking at it.

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