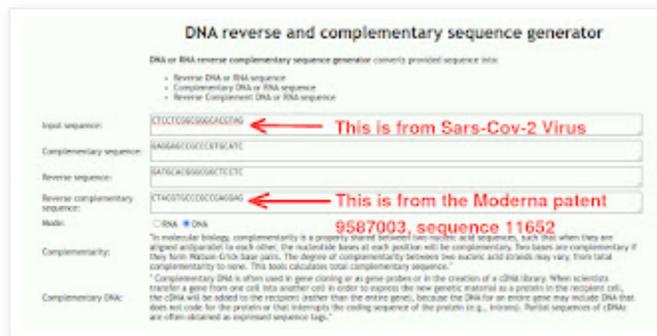




Go to the “Reverse Complement Calculator”:

https://bugaco.com/calculators/dna_reverse_complement.php

Now type in the genes from Sars-Cov-2 (which is CTCCTCGGCGGGCACGTAG):



So, this shows that the Sars-Cov-2 sequence CTCCTCGGCGGGCACGTAG, matches the Moderna patent sequence CTACGTGCCCGCCGAGGAG as a “reverse complement”.

This is the explanation of “reverse complement”. Remember that Sars-Cov-2 is a RNA virus and Moderna patents are DNA

(DNA STRUCTURE AND THE REVERSE COMPLEMENT

OPERATION)

None of this tells us whether this sequence is “good”, “bad”, “pure chance”, etc. All we know is:

- CTCCTCGGCGGGCACGTAG is the most crucial part of Sars-Cov-2 RNA, containing the code for the “furin cleavage site”
- No natural coronaviruses contain a “furin cleavage site”
- Furin cleavage site greatly enhances infectivity of Sars-Cov-2 and its ability to infect humans and other species
- This code only appears in a Moderna **oncology (cancer)** patent 9587003, and not in any multicellular organisms or viruses known before 2020
- The mutation of MSH3 gene related to this patent, is known to break DNA recombination and lead to cancers (see my previous article)
- Spike protein is also known to penetrate cell nuclei and interfere with V(D)J DNA recombination and BRCA1 and p53 repair, for a reason that MAY OR MAY NOT be related to CTCCTCGGCGGGCACGTAG
- BRCA mutation is responsible for breast and ovarian cancer