

SARS-CoV-2: A unicorn?

In October 2020, [Dr Tom Cowan](#) wrote a piece, '[Only Poisoned Monkey Cells 'Grew' the 'Virus'](#)', after analysing a key article posted by the CDC, in their journal, Emerging Infectious Diseases: "[Severe Acute Respiratory Syndrome Coronavirus 2 from Patient with Coronavirus Disease, United States](#)", published in June 2020.

Excerpts from Dr. Cowan's piece:

The purpose of the [CDC] article was for a group of about 20 virologists to describe the state of the science of the isolation, purification and biological characteristics of the new SARS-CoV-2 virus, and to share this information with other scientists for their own research.

In the section titled 'Whole Genome Sequencing,' we find that rather than having isolated the virus and sequencing the genome from end to end, they found 37 base pairs from unpurified samples using PCR probes. This means **they actually looked at 37 out of the approximately 30,000 of the base pairs that are claimed to be the genome of the intact virus. They then took these 37 segments and put them into a computer program, which filled in the rest of the base pairs.**

To me, this computer-generation step constitutes scientific fraud. Here is an equivalency: A group of researchers claim to have found a unicorn because they found a piece of a hoof, a hair from a tail, and a snippet of a horn. They then add that information into a computer and program it to re-create the unicorn, and they then claim this computer re-creation is the real unicorn. Of course, they had never actually seen a unicorn so could not possibly have examined its genetic makeup to compare their samples with the actual unicorn's hair, hooves and horn.

The researchers claim they decided which is the real genome of SARS-CoV-2 by 'consensus,' sort of like a vote. Again, different computer programs will come up with different versions of the imaginary 'unicorn,' so they come together as a group and decide which is the real imaginary unicorn.