# Covid-19 is A Bioweapon

## Part 1

As Pfizer and Moderna vaccines are being given throughout the world, we are seeing a growing pushback, as side effects, and deaths continue to mount up. It has been a year since I first reported that Corona or Covid-19 was a bioweapon. Recently, one or two voices have expressed words to this effect.

At least once a day someone will tell me that the Vaccine does not change our DNA. We are being told that our DNA cannot be changed, by people who are ignorant to the facts. Most people believe our DNA is fixed, meaning it does not change. That was what most of us were taught in high school decade ago; however, the science of genetics has changed since the late 1980s. An initial rough draft of the human genome was available in June 2000 and by February 2001 a working draft had been completed and published followed by the final sequencing mapping of the human genome on April 14, 2003.

The main goals of the Human Genome Project were first articulated in 1988 by a special committee of the U.S. National Academy of Sciences, and later adopted through a detailed series of five-year plans jointly written by the National Institutes of Health and the Department of Energy.

Congress funded both the NIH and the DOE to embark on further exploration of this concept, and the two government agencies formalized an agreement by signing a Memorandum of Understanding to "coordinate research and technical activities related to the human genome."

James Watson was appointed to lead the NIH component, which was dubbed the Office of Human Genome Research. The following year, the Office of Human Genome Research evolved into the National Center for Human Genome Research.

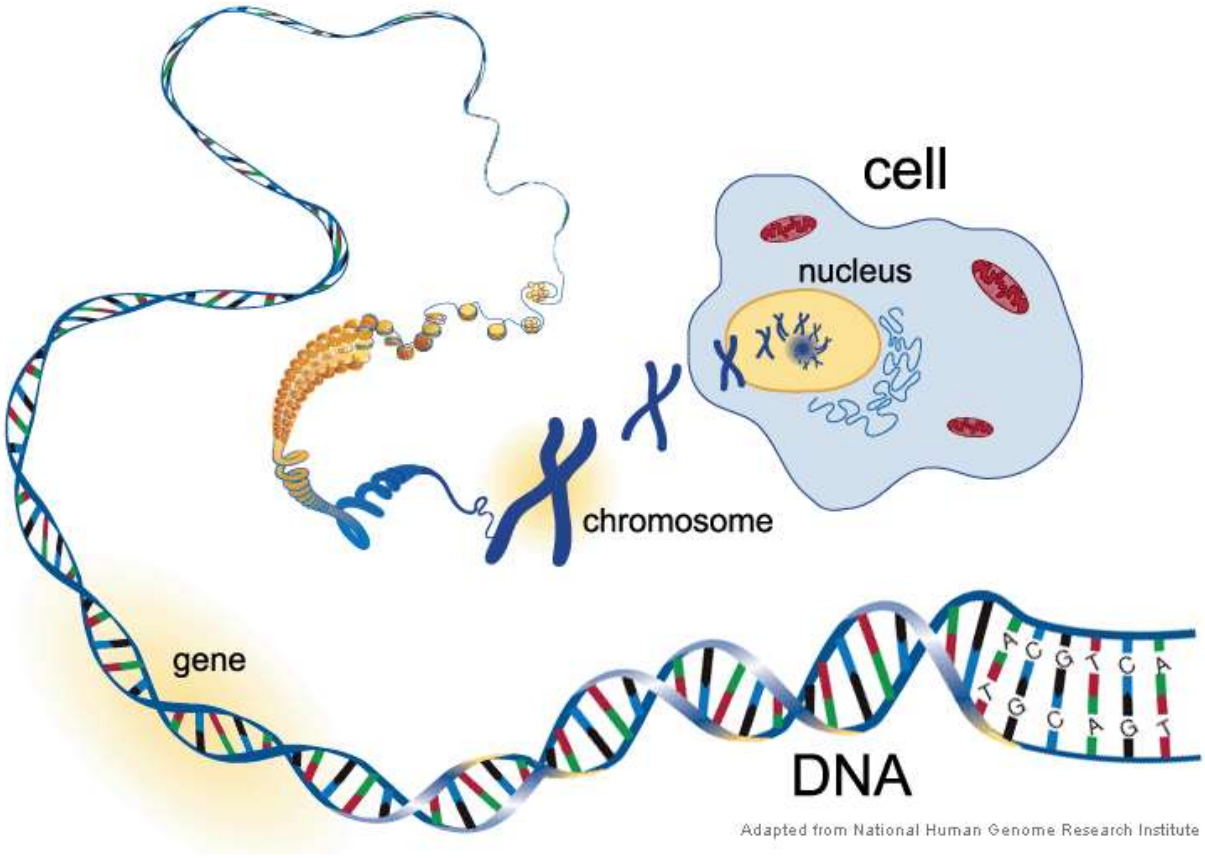
In 1990, the initial planning stage was completed with the publication of a joint research plan, "Understanding Our Genetic Inheritance: The Human Genome Project, The First Five Years, FY 1991-1995." This initial research plan set out specific goals for the first five years of what was then projected to be a 15-year research effort.

HGP researchers deciphered the human genome in three major ways: determining the order, or "sequence," of all the bases in our genome's DNA; making maps that show the locations of genes for major sections of all our chromosomes; and producing what are called linkage maps, through which inherited traits (such as those for genetic disease) can be tracked over generations.

The HGP has revealed that there are probably about 20,500 human genes. This ultimate product of the HGP has given the world a resource of detailed information

about the structure, organization and function of the complete set of human genes. This information can be thought of as the basic set of inheritable "instructions" for the development and function of a human being.

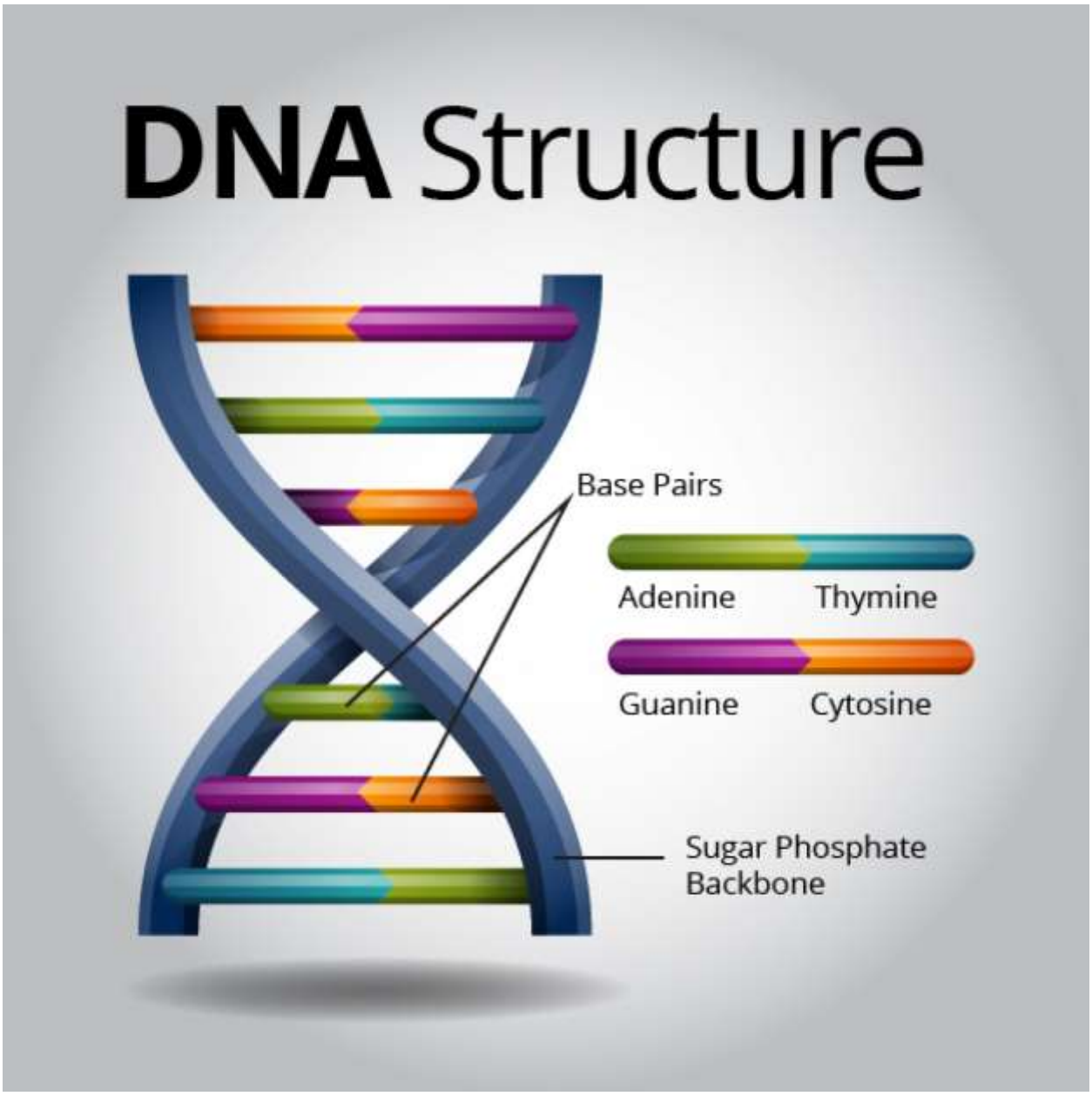
The International Human Genome Sequencing Consortium published the first draft of the human genome in the journal Nature in February 2001 with the sequence of the entire genome's three billion base pairs some 90 percent complete. More than 2,800 researchers who took part in the consortium shared authorship. A startling finding of this first draft was that the number of human genes appeared to be significantly fewer than previous estimates, which ranged from 50,000 genes to as many as 140,000. The full sequence was completed and published in April 2003.




Human beings have roughly 20,500 genes, all coiled up in DNA, housed in each and every one of the trillions of cells that make you who you are. That's 20,500 places where the machinery of human life can be altered. Many of these alterations would make life impossible. Some scientists have said that the mapping of the human Genome has opened a Pandora's box issue, meaning it could prove problematic over time because of what it could unleash. For example, designer babies are within limitless possibilities.

Upon publication of the majority of the genome in February 2001, Francis Collins, then director of the National Human Genome Research Institute, noted that the genome

could be thought of in terms of a book with multiple uses: "It's a history book - a narrative of the journey of our species through time. It's a shop manual, with an incredibly detailed blueprint for building every human cell. And it's a transformative textbook of medicine, with insights that will give health care providers immense new powers to treat, prevent and cure disease."

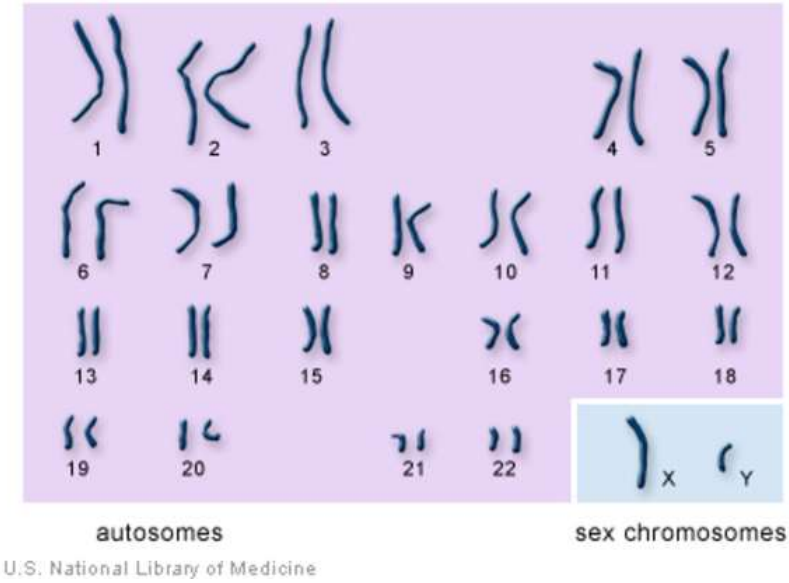



The Building Block of Life

In humans, each cell normally contains 23 pairs of chromosomes, for a total of 46. Twenty-two of these pairs, called autosomes, look the same in both males and females. The 23rd pair, the sex chromosomes, differs between males and females. Females have two copies of the X chromosome, while males have one X and one Y

chromosome. Genetics exposes the fact that there can only be two genders at birth, the science settles that despite all the efforts to surgically alter one's gender.

**The 22 autosomes are numbered by size. The other two chromosomes, X and Y, are the sex chromosomes. This picture of the human chromosomes lined up in pairs is called a karyotype.**




Credit: U.S. National Library of Medicine

The tools created through the HGP also continue to inform efforts to characterize the entire genomes of several other organisms used extensively in biological research, such as mice, fruit flies and flatworms. These efforts support each other, because most organisms have many similar, or "homologous," genes with similar functions. Therefore, the identification of the sequence or function of a gene in a model organism, for example, the roundworm C. elegans, has the potential to explain a homologous gene in human beings, or in one of the other model organisms.

Of course, information is only as good as the ability to use it. Therefore, advanced methods for widely disseminating the information generated by the HGP to scientists, physicians and others, is necessary in order to ensure the most rapid application of research results for the benefit of humanity. Biomedical technology and research are particular beneficiaries of the HGP.

However, the momentous implications for individuals and society for possessing the detailed genetic information made possible by the HGP were recognized from the outset. Another major component of the HGP - and an ongoing component of NHGRI - is therefore devoted to the analysis of the ethical, legal and social implications (ELSI) of our newfound knowledge, and the subsequent development of policy options for public consideration.

We still have no idea what 20 per cent of protein-coding genes are for. What's more, we have stopped making progress, according to a study looking at what we know about yeast and human proteins.

"Basically we really don't have a clue," says team leader Valerie Wood at the University of Cambridge.

Her team started by defining what is known or unknown. For instance, we might be able to tell that a protein is an enzyme from its sequence, but if we don't know what reaction it catalyzes, its function cannot be said to be known. Wood compares it to taking a car to pieces – recognizing that one piece is, say, a wire isn't much help in understanding what it is for. It is this specific concern that has raised ethical questions when it comes to the CRISPR technology in gene therapy. The CRISPR-like tool for RNA editing could temporarily alter your proteins.

A gene is a sequence of DNA base pairs that codes for a particular protein. These sequences account for only about 2% of the human genome. The rest is referred to as noncoding DNA, some of which serves various functions and some of which is apparently nonfunctional. The number of functional genes in the human genome is actually quite low, but the number of genes is not an indicator of the complexity of an organism, and many creatures have more genes than humans do. Although the human genome has been mapped for about twenty years, there is much that is unknown to the mystery of God's design.

My interest in genetics goes back to 1967, when my wife became pregnant with our first child. She began to experience epileptic grand mal and peti mal seizures. Throughout her pregnancy we were soon to learn that she had inherited the genetic disorder known as Huntingdon's disease. She inherited the disease from her mother's genetic disease. My interest was related to the fact that we had three children with the risk of inheriting the gene for Huntingdon's. My oldest son and my youngest daughter were at risk. We lost our second son as the result of a grand mal seizure that led to my wife falling out of bed that led to a still-birth.

As the result of the genome mapping, gene therapy has made huge advancements in the matter of gene therapy. The process of CRISPR today has enabled medical science to basically cut and paste, or to remove defective genes and correct issues that formerly could not be resolved or meant a life of suffering with various genetic problems. Persons contracting the Covid-19 virus, do undergo an alteration to the DNA of the infected cells, the so-called "host DNA". This is due to the inclusion of retrovial HIV-1 fragments. They do not however contract HIV itself, as the complete genomic sequence of this retrovirus is not present.

Retroviruses are a type of virus that uses RNA as their genetic material and a special enzyme called reverse transcriptase to translate the virus' genetic information into DNA. That DNA can then integrate into the host (your) cell's DNA. At this point, the retrovirus can replicate itself using your cells resources. It "goes viral". The HIV-1 fragments act

in the normal biological manner as the entire HIV virus itself. More details will be explained later.

The mRNA-based vaccine that Pfizer and Moderna are using does contain genomic fragments of the retrovirus HIV-1, and a modified RNA genomic sequence of Covid-19. Likewise, persons inoculated will neither contract HIV, nor Covid-19.

However, and this is the major difference between the virus and the vaccine, the latter does permanently alter a person's entire DNA. Anyone telling you something other, does not know what they are talking about or are ignorant of genetics.

The difference lays within the additional components embedded within the vaccine itself. These include quantum dots, luciferase, hydrogel and nanomaterials such as gold and diamond (functioning as biosensors inside the human body). Working in tandem with the Covid-19 virus, and the fragments of the retrovirus HIV-1, the entire host" DNA is permanently changed. This exceeds the natural biological change brought about by a "common" retrovirus.

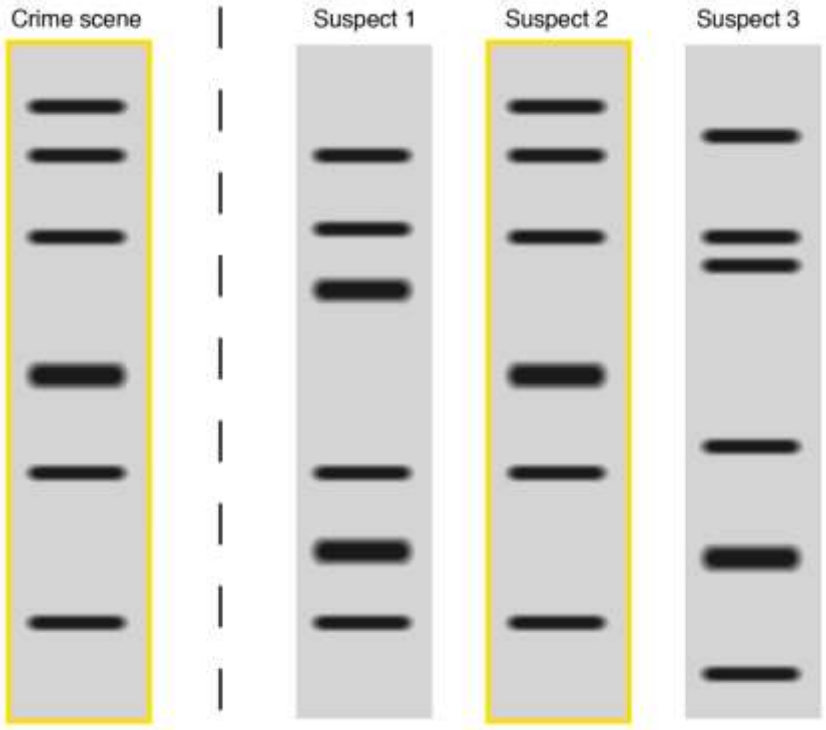
As a RNA virus, Covid-19 functions as a vector to transmit the HIV-1 fragments into a cell's cytosol. At this point the roles are reversed with the retroviral HIV-1 fragments now acting as a vector for the delivery of the Covid-19 virus into the cell's nucleus.

Thee fragments of HIV-1 improve the "gain of function" of Covid-19 as a virus, increasing its mobidity, without an increase in mortality. Likewise, this process will be detailed later.

The grand delusion of 2019-nCOV/COVID-19 and its attendant mainstream narrative proclaiming the virus being of natural origin, has obfuscated verified scientific evidence to the contrary. Specifically, not 4, or 6, but 18 genomic fragments (inserts) of HIV-1 (Human Immunodeficiency Virus) have deliberately and artificially been placed within the overall genome of Covid-19, a single-stranded RNA (Ribonucleic acid) virus. A retrovirus.

As a retrovirus, these HIV-1 fragments while not forming the entire genome of the HIV virus itself, is directly relevant to every human. As many scientists and their associates prove with their published papers, these fragments serve the designed biological objective known as "gain of function" as scientists from France, China, and India have discovered. As will be explained later, the function of the HIV-1 fragments is to promote viral entry into a host cell. Not only entry of HIV-1, but the 2019-nCoV/COVID-19 virus itself! A retrovirus. It became a bioweapon at most-likely a BSL4 lab. BSL4 labs are the most secured labs in the world for dealing with pathogens and can only be found in a military facility setting.

Most of you are familiar with crime scene investigations in which gathering DNA material is of critical value. The illustration below is a simplified picture of how DNA does not lie in identifying the person or persons connected to solving a crime.




Why this is so extremely critical point is that in viruses there is what I refer to as an audit trail or a finger print. This basically means scientists can confirm that the virus was modified in a BSL4 lab. These inserts confirm a non-natural appearance of 18 fragments of HIV-1 have been found in the Coronavirus. There were also 4 fragments of the SARS-2. In simple terms someone somewhere jacked up the virus to make it more potent.

The wording below of the 9 or 10 Indian scientists, who first discovered this fact early on, I believe it was March, 2020, and I quote from the Abstract of their findings.

**"We are currently witnessing a major epidemic caused by the 2019 novel corona-virus (2019-nCoV. The evolution of 2019-nCoV remains elusive. We found 4 insertions in the spike glycoprotein (S) which are unique to the 2019-nCoV and are not present in other coronaviruses. Importantly, amino acid residues in all the 4 inserts have identity or similarity to those in the HIV-1 gp120 or HIV-1 Gag. Interestingly, despite the inserts being discontinuous on the primary amino acid sequence, 3D-modelling of the 2019-nCoV suggests that they converge to constitute the receptor binding site. <u>The finding of 4 unique inserts in the 2019-nCoV, all of which have identity/similarity to amino acid residues in key structural proteins of HIV-1 to unlikely to be fortuitous in nature</u>. This work provides yet unknown insights on 2019-nCoV and sheds light on the evaluation and pathogenicity of this virus with important implications for diagnosis of this virus.**

The Indian scientists submitted their paper for publishing only to later withdraw it as a result of pressure from likely the WHO, World Health Organization and others. It is

important to know that since research in the area of genetics and viruses, requires money, lots of it, one's career can be seriously damaged when one chooses to go against the flow of information being made public by health agencies. That said, their studies were examined by the 2008 Nobel Laureate in Physiology Dr. Luc Antoine Montagnier. Dr. Montagnier is recognized for his discovery of the human immunno-deficiency virus (HIV). Later a Chinese scientist, Huang Yan Ling confirmed ***"THIS VIRUS WAS MANIPULATED METICUOUSLY, IT'S NOT NATURAL".***

Although I cannot share with the reader the graphic illustrations and their lesson that explain the process involved to modify the genetic structure, we have scientists from Australia reporting this finding just a month ago, in addition to three scientific teams from India, France, and China. Australia cancelled their $billion dollar vaccine program, after finding HIV-1 in individuals who had been vaccinated with the Covid-19 vaccine. Interestingly, the media has not publicized the fact that Australia cancelled its virus vaccine program over what was found in the vaccine.

We have now reached the point confirming that the Covid-19 vaccine is an engineered Bioweapon, and did not suddenly spread from China around the world all by itself. A Bioweapon is created to do one of two objectives if not both: **1. Debilitate** or **2. Kill.**

Moving on we need to understand what a retrovirus is and does. The National Human Genome Research Institute describes a retrovirus including HIV-1: *a retrovirus is a virus that uses RNA as its genetic material. When a retrovirus infects a cell, it makes a DNA copy of its genome that is inserted into the DNA of the host cell*. There are a variety of different retroviruses that cause human diseases such as some forms of cancer and AIDS.

In this case, the DNA resides within the nucleus of human cells. Once fused, the DNA copy of HIV-1 and 2019-nCoV/COVID-19 permanently alters the existing DNA of the human host. I regret that I can't find visual slides that explain the process of what is taking place in your body. My point is that the Covid-19 vaccine, containing the inserts will change your DNA!

There is a biological symmetry between 2019-nCoV/COVID-19 and HIV-1, with one requiring the natural characteristics of the other. They work in concert with one another, each acting as a vector for the other. A vector is an organism that does not cause disease itself but which spreads infection by conveying pathogens from one host to another.

Within their respective papers, Indian researchers, and in Dr. Montagnier and his co-researcher Jean Claude Perez noted within the complete genomic DNA sequence of 2019-nCoV/COVID-19 the fragments (inserts) of HIV-1. Meaning, the 2019-cCoV/COVID-19 virus itself exists due the presence of HIV-1. The spread, or virility of 2019-nCoV/COVID-19 is dependent upon these fragments of HIV. This is the aforementioned "gain of function".

Once infected, the DNA of the host cell produces a new single strand of RNA. One consisting of both the original DNA of the host, and the DNA of 2019-nCoV/COVID-19, and HIV-1.

Initially, as a single RNA strand, 2019-nCoV/COVID-19 and HIV-1 move into the fluid of a cell, the cytoplasm. The RNA then produces a copy, a complimentary single strand of itself, referred to as cDNA. It is a single strand of DNA. Next, a second complimentary strand is produced, matching the first. The two intertwine into the familiar double helix of common DNA. This is still referred to as cDNA, or dscDNA (double stranded complimentary DNA). This cDNA consists of the combined genome of 2019-nCoV/COVID-19 and HIV-1.

The new cDNA (as a familiar double helix) enters the nucleus of the cell and binds to the existing DNA of the host (human) cell. This permanently changes the host DNA of this specific cell into a retrovirus: 2019-nCoV/COVID-19 and HIV-1.

The virus, like all viruses, wants to reproduce. It wants to "go viral". It does so by producing a single strand of messenger RNA (mRNA), itself a copy of the viral cDNA (the newly formed, altered host DNA). This single strand mRNA exits the nucleus, then the cell itself, moving on to the next cell, thus "going viral".

If you have read my articles on Mycoplasma, you recognize that this is exactly what I dubbed the "sin" pathogen does. It goes viral consuming dead or dying cells, and eventually the healthy cells succumb to the process of the Mycoplasma pathogen.

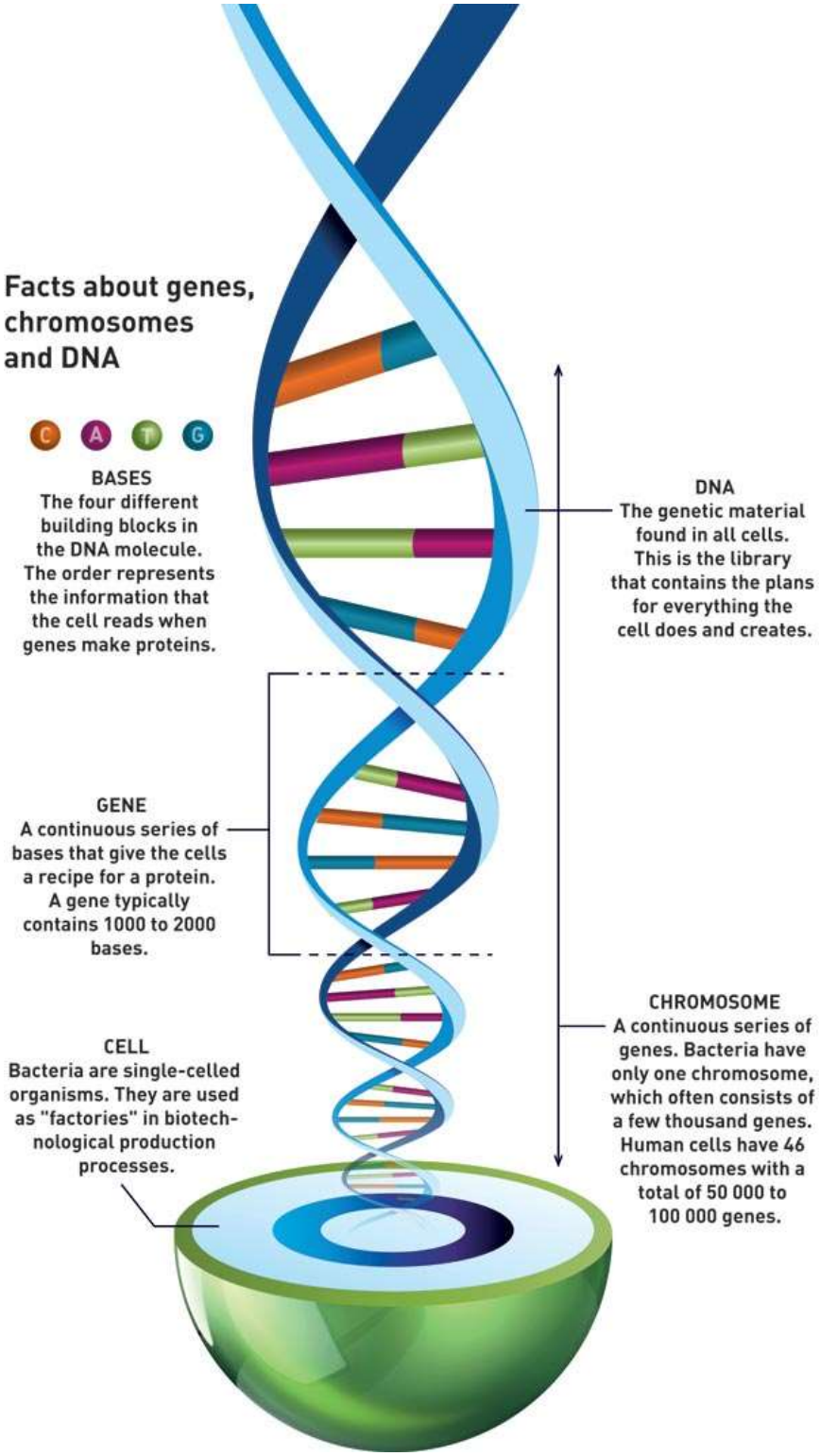
"18 RNA fragments of homology equal or more than 80% with human or simian retroviruses have been found in the COVID-19 genome."

"HIV-1 EIE with a crucial Spike mutation."

Through the facts relating to this article, everything converges toward possible laboratory manipulations which contributed to modifications of the genome of COVID-19, but also, very probably much older SARS, with perhaps double objective of vaccine design and "gain of function" in terms of penetration of this virus into the cell."

The conclusions of those scientists and others cited are: this genome of 2019-nCoV/COVID-19 was man-made in a laboratory and contains not simply 4 inserts as cited by the Indian scientists, but 18 human or simian retroviruses (HIV-1). The word "simian" here refers to animal or monkeys. The earliest polio vaccines were tainted with the cancer agent known as Simian-40.

With respect to the "Gain of Function" objective, this is for the purpose of increasing the rate of infection, the pathogenic transmissibility (mobidity) of a virus within a host population. Not necessarily leading to an increase of mortalities. This is leveraging the natural processes of a virus in increasing the quantity of infected hosts, without killing them off.


Facts about genes, chromosomes and DNA
C A T G
BASES
The four different building blocks in the DNA molecule. The order represents the information that the cell reads when genes make proteins.
DNA
The genetic material found in all cells. This is the library that contains the plans for everything the cell does and creates.
GENE
A continuous series of bases that give the cells a recipe for a protein. A gene typically contains 1000 to 2000 bases.
CHROMOSOME
A continuous series of genes. Bacteria have only one chromosome, which often consists of a few thousand genes. Human cells have 46 chromosomes with a total of 50 000 to 100 000 genes.
CELL
Bacteria are single-celled organisms. They are used as "factories" in biotechnological production processes.

This is accomplished by enhancing the ability of a virus to penetrate into a host cell. The presence of HIV-1 genetic material, the 18 RNA fragments, indicates the targeting of host cell RNA. The natural process of all retroviruses like HIV-1 is to fuse its genetic material to that of the host DNA, thus permanently changing it. From there, this modified DNA naturally produces a single strand of RNA containing the virus and moves on to infect the next host cell. It "goes viral".

Why have you not heard this fact in the media? If you have your radio or television on, all day long, all you hear is the vaccines are here or the mask will protect you. It's all a huge scam to change your DNA with an engineered bioweapon.

The science is being repressed in the media, and when you experience this kind of censorship, you can be sure it is true and they do not want you to hear or read it.

The Covid-19 vaccine is a bioweapon and it will permanently change your DNA. There are lots of questions you may be asking about what the world has been put through by the fear-mongering people into believing this was a pandemic. It was a "Plandemic" to bring the world into submission to the Great Reset planned leading to the cashless society and a global world government. The New World Order elite are meeting in Davos, Switzerland, this very week to plan their next attack on humanity. I saw yesterday that a hot dog in Davos, Switzerland costs $24 during the world elite meet to plan the next phase.

DO NOT TAKE A VACCINE, AND DO NOT ALLOW ANYONE TO PUT A SWAB ON A STICK UP YOUR NOSE. IT JUST MIGHT INFECT YOU WITH SOMETHING YOU MAY REGRET. SHARE THIS IF YOU WISH!

Blessings,

Pastor Bob, EvanTeachr@aol.com
www.pastorbobreid.com
http://jesusisthewaythetruththelife.com/node/22