

Shocking study finds covid vaccines REWRITE your DNA... criminal CDC proven to have repeatedly LIED about this very issue to deceive and harm the public

Thursday, March 03, 2022 by: Mike Adams

Tags: badhealth, badmedicine, CDC, covid-19, DNA, immunization, mRNA, pandemic, Pfizer, real science, reverse transcriptase, spike protein, transhumanism, vaccines

This article may contain statements that reflect the opinion of the author



(Natural News) As The Epoch Times and other media outlets reported yesterday, a shocking new study conducted at Lund University in Sweden has confirmed that mRNA nanoparticles from Pfizer's covid-19 vaccine enter human cells and are reverse-transcribed into DNA, achieving a permanent alteration of the person's genetic code.

This is something that the CDC lied about for the last several years, claiming it was impossible.

The study is published in the journal *Molecular Biology*. It's entitled, "Intracellular Reverse Transcription of Pfizer BioNTech COVID-19 mRNA Vaccine BNT162b2 In Vitro in Human Liver Cell Line."

The study concludes that, “BNT162b2 mRNA is reverse transcribed intracellularly into DNA as fast as 6 [hours] after BNT162b2 exposure.”

This means the vaccine alters your genetic code in mere hours after being injected. This may help explain why cancer rates are exploding in people who take covid vaccines. The integrity of their DNA is being destroyed.

It is also yet more proof that the mRNA particles from the vaccine do, indeed, circulate throughout the body and don't stay restricted to the injection site. That's another lie told to the world by the CDC, Big Tech and the pharma-funded corporate media which has been caught in lie after lie when it comes to covid origins and vaccine safety.

The CDC continues to lie and deceive the public about mRNA vaccines altering your DNA

As pointed out by The Epoch Times, the CDC's web page entitled, “Myths and Facts about COVID-19 Vaccines” still asserts the lie about vaccines not altering your DNA, claiming:



The genetic material delivered by mRNA vaccines never enters the nucleus of your cells, which is where your DNA is kept. Viral vector COVID-19 vaccines deliver genetic material to the cell nucleus to allow our cells to build protection against COVID-19. However, the vector virus does not have the machinery needed to integrate its genetic material into our DNA, so it cannot alter our DNA.

It turns out the CDC was lying all along. No one is surprised to learn this, and it begs the obvious question: What else is the CDC lying about? What other truths are they hiding from the public?

The CDC lied about vaccine distribution throughout your body, and they lied about vaccines altering your DNA. The CDC, which owns thousands of patents on vaccines, is actually a vaccine criminal cartel and propaganda operation that deliberately misleads the public about vaccines in order to sell deadly poison to the masses.

The CDC, in other words, is a pharmaceutical crime cartel that carries out crimes against humanity in order to enrich Big Pharma (and themselves).

TheCovidWorld.com has also found that the mRNA code injected into people as part of the covid

vaccines contains a gp130 cancer-causing gene sequence. While more research is needed, it appears the mRNA vaccine may actually “program” human DNA to grow cancer tumors.

BLAST ^{2.10} | blastn suite | results for RID: 1R3ZDZJY018

Job Title: Nucleotide Sequence
 RID: 1R3ZDZJY018
 Program: BLASTN
 Database: nt
 Query ID: id|Query_20439

Filter Results
 Organism: only top 20 will appear
 Percent Identity: [] to []
 E value: [] to []
 Query Coverage: [] to []

Sequences producing significant alignments

| Description | Scientific Name | Max Score | Total Score | Query Covr | E value | Per. Ident | Ann. Len | Accession |
|---|--------------------|-----------|-------------|------------|---------|------------|----------|---------------|
| MusMus musculus strain M03452.6.852.6.H., complete genome | Mus musculus | 472 | 472 | 98% | 5e-113 | 66.17% | 16870 | MM020032.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 431 | 431 | 88% | 3e-107 | 103.00% | 3753 | OK120542.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 381 | 381 | 93% | 4e-101 | 62.78% | 3828 | OK120542.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 368 | 368 | 84% | 1e-106 | 103.00% | 3753 | AB771336.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 355 | 355 | 82% | 5e-87 | 62.67% | 3793 | OK120540.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 327 | 327 | 93% | 6e-45 | 49.53% | 4257 | NC_030368.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 327 | 327 | 93% | 6e-45 | 69.37% | 4218 | MG_083837.1 |
| Recombinant vector SARS-CoV-2/US1911, complete sequence | synthetic const... | 311 | 311 | 93% | 1e-41 | 60.70% | 7403 | MM481182.1 |
| Synthetic construct of HCoV11972 (Novel Influenza A (H1N1) (SARS-CoV-2) vaccine sequence) | synthetic const... | 300 | 300 | 90% | 1e-16 | 87.00% | 4215 | MG_083838.1 |
| Expression vector SARS-CoV-2/US1911, complete sequence | Expression vect... | 285 | 285 | 82% | 1e-40 | 88.58% | 11445 | MM580962.1 |
| Expression vector SARS-CoV-2/US1911, complete sequence | Expression vect... | 285 | 285 | 82% | 1e-40 | 88.58% | 10732 | MM580962.1 |
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| Human genome (hg38) (Ensembl) (human) (HOMO) (HOMO), complete genome | Homo sapiens | 243 | 243 | 81% | 2e-59 | 67.18% | 1317 | CP009716.1 |
| PREDICTED: Pan paniscus TLE family member 5, transcriptional modulator (TLE5), transcript variant 1 | Pan paniscus | 241 | 241 | 90% | 7e-03 | 68.58% | 1803 | XM_02404032.1 |
| PREDICTED: Pan paniscus TLE family member 5, transcriptional modulator (TLE5), transcript variant 2 | Pan paniscus | 241 | 241 | 90% | 7e-04 | 68.58% | 1805 | XM_03884536.1 |
| PREDICTED: Pan paniscus TLE family member 5, transcriptional modulator (TLE5), transcript variant 3 | Pan paniscus | 241 | 241 | 90% | 7e-09 | 68.58% | 1791 | XM_03884538.1 |
| PREDICTED: Pan paniscus TLE family member 5, transcriptional modulator (TLE5), transcript variant 4 | Pan paniscus | 241 | 241 | 90% | 7e-09 | 68.58% | 1891 | XM_03884536.1 |

Annotations:
 - Red circle around '1R3ZDZJY018' in the search parameters.
 - Red arrow pointing to 'gp130' in the 'Human genome' entry.
 - Red text: 'Ignore these vaccine, Sars-Cov2, chimeric Sequences' (with a large red arrow pointing to the top of the table).
 - Red text: 'This is the mutated Cancer Gene gp130' (with a red circle around 'gp130' in the table entry).

Get the full story in today's Situation Update podcast below. The vaccine discussion starts at 34:50. Before that, we cover Joe Biden's Slur of the Union Address and you get to hear me belt out lines from the song, "I Give You My Word" as performed by President Komacho (Terry Crews) in the film Idiocracy ... which turns out to be a documentary of the Biden administration.

Nobody can beat Terry Crews at this, of course, as he's the master of comedy. But I give it a shot. Listen and weep:

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