

BOMBSHELL: WHO Coronavirus PCR Test Primer Sequence is Found in All Human DNA

Primers and probes

Name	Sequences (5'-3')	Length (bases)	PCR product size	Ref.
RdRp gene / nCoV_IP2				
nCoV_IP2-12669Fw	ATGAGCTTAGTCCTGTTG	17	108 bp	1
nCoV_IP2-12759Rv	CTCCCTTGTGTGTTGT	18		
nCoV_IP2-12696bProbe(+)	AGATGCTTGTGCTGCCGGTA [5']Hex [3']BHQ-1	21		
RdRp gene / nCoV_IP4				
nCoV_IP4-14059Fw	GGTAACTGGTATGATTCG	19	107 bp	1
nCoV_IP4-14146Rv	CTGGTCAAGTTAATATAGG	20		
nCoV_IP4-14084Probe(+)	TCATACAAACCACGCCAGG [5']Fam [3']BHQ-1	19		
E gene / E_Sarbeco				
E_Sarbeco_F1	ACAGGTACGTTAATAGTTAATAGCGT	18	125 bp	2
E_Sarbeco_R2	ATATTGCAGCAGTACGCACACA	20		
E_Sarbeco_P1	ACACTAGCCATCCTTACTGCGCTTCG [5']Fam [3']BHQ-1	20		

1/ National Reference Center for Respiratory Viruses, Institut Pasteur, Paris.

2/ Corman et al. Eurosurveillance¹

This was important enough that I wanted to get it out immediately. My research into the NCBI database for nucleotide sequences has led to a stunning discovery. One of the WHO primer sequences in the PCR test for SARS-CoV-2 is found in all human DNA!

The sequence "CTCCCTTGTGTGTTGT" is an 18-character primer sequence found in the **WHO coronavirus PCR testing protocol document** (https://www.who.int/docs/default-source/coronaviruse/real-time-rt-pcr-assays-for-the-detection-of-sars-cov-2-institut-pasteur-paris.pdf?sfvrsn=3662fcb6_2). The primer sequences are what get amplified by the PCR process in order to be detected and designated a "positive" test result. It just so happens this *exact same* 18-character sequence, verbatim, is also found on Homo sapiens chromosome 8! As far as I can tell, this means that the WHO test kits should find a positive result in all humans. Can anyone explain this otherwise?

I really cannot overstate the significance of this finding. At minimum, it should have a notable impact on test results.

[Download](#) [GenBank](#) [Graphics](#) Sort by:

Homo sapiens chromosome 8, GRCh38.p12 Primary Assembly
Sequence ID: [NC_000008.11](#) Length: 145138636 Number of Matches: 3

Range 1: 63648346 to 63648363 [GenBank](#) [Graphics](#) [Next Match](#) [Previous Match](#)

Score	Expect	Identities	Gaps	Strand
36.2 bits(18)	0.098	18/18(100%)	0/18(0%)	Plus/Plus

Features: [438640 bp at 5' side: YTH domain-containing family protein 3 isoform X2](#)
[932428 bp at 3' side: class E basic helix-loop-helix protein 22](#)

Query 1 CTCCCTTGTGTGTTGT 18
Sbjct 63648346 CTCCCTTGTGTGTTGT 63648363

Homo sapiens chromosome 8, GRCh38.p12 Primary Assembly
Sequence ID: NC_000008.11 Length: 145138636
Range 1: 63648346 to 63648363 is "CTCCCTTTGTTGTGTTGT"

Update: After some effort, I have finally discovered a way to display proof (beyond my screenshots) that human chromosome 8 has this exact same 18-character sequence. Please try the link below. The sequence is shown at the bottom of the page.

[https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log\\$=nuclalign&from=63648346&to=63648363](https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log$=nuclalign&from=63648346&to=63648363)
([https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log\\$=nuclalign&from=63648346&to=63648363](https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log$=nuclalign&from=63648346&to=63648363))

Posted on April 6, 2020 June 18, 2020 by Fauxlex Posted in Coronavirus, COVID-19

77 thoughts on "BOMBSHELL: WHO Coronavirus PCR Test Primer Sequence is Found in All Human DNA"

1. Pingback: [We all have Corona | Fakeologist.com](#)
2. MAARTEN ROSSAERT SAYS:

April 6, 2020 at 12:04 pm

How then do some people test negative for COVID-19?

FAUXLEX SAYS:

April 6, 2020 at 12:07 pm

This is the WHO protocol. I also came across several private companies (Sigma-Aldrich, Thermo Fisher) whose PCR primer sequences were longer and had no such match. So I am not saying that ALL test kits belong to WHO, but if we were to find shenanigans we would expect to find it with the WHO. This really has me shaken, and I am really starting to believe that the WHO designed a test where they would be able to find a positive anywhere they wanted to find one. Also, the PCR test process is inherently very error-prone and even the WHO test may occasionally have false negatives.

I'm stunned.

FAUXLEX SAYS:

April 6, 2020 at 12:15 pm

I mean seriously, what are the odds?

1. MARK TOKARSKI SAYS:

April 6, 2020 at 12:54 pm

Why does anyone assume the test itself is even being done, and if being done, that the results are being reported honestly? Has anyone every heard the six magic words in Climate Science: "They lie, they lie, they lie." Why not in virology as well?

Somebody please, follow this link ...

<https://thedoctorwithin.com/blog/2020/03/10/newsletter-march-2020/>, scroll down to #3, and then read on for a few minutes. This is written by a doctor in the Bay Area who decided to follow publicly reported cases of people who tested positive for Covid-19, starting with Santa Clara Health and then getting in touch with 50 pulmonologists/clinics.

Not one was equipped to be able to test for a virus or to distinguish between influenza and Covid. His conclusion: The 60 (at that time) cases reported in Santa Clara County did not exist and the County website was lying.

Someone follow the link. Please. Just this once.

FAUXLEX SAYS:

April 6, 2020 at 12:57 pm

C'mon Mark, this is a bombshell finding. And it even supports your ideas! Taken straight from the WHO testing protocol. I am not suggesting that all the testing they claim to be doing is actually being done, but the implications in this finding are huge.

Repeating my question...what are the odds?

1. **PETRA LIVERANI SAYS:**

April 12, 2020 at 8:13 am

Bombshell finding. It certainly seems so to me.

1. **PETRA LIVERANI SAYS:**

April 12, 2020 at 8:02 am

LOL. I've got it on my blog post too.

<https://occamsrazorterrevents.weebly.com/blog/coronavirus-hoax-jan-2020>

FAUXLEX SAYS:

April 6, 2020 at 1:02 pm

And I should be clear that it is NOT easy to find an 18-character match like this. In most of my searches, I was finding 14-15 character long matches in human DNA for 20-25 character long primers. Then suddenly BOOM the short 18-character WHO primer sequence has a perfect 18-character match. That is nuts. I really think that the odds are very high that this is a meaningful finding.

3. **LAWRENCE ROTHMAN SAYS:**

April 6, 2020 at 12:17 pm

So in the case of the CDC well documented and media covered faulty COVID-19 tests... was the CDC faking their incompetence or would we say they were really incompetent? BTW, Greetings to Appalachia!

4. **BLAIR SAYS:**

April 6, 2020 at 1:40 pm

Can you post a link to the Homo Sapiens Chromosome 8 18 character sequence data please?

FAUXLEX SAYS:

April 6, 2020 at 2:48 pm

Blair, I think I finally found a way to present this 18 character sequence on Homo sapiens chromosome 8 in a permanent way. If you go to the bottom of the page at the link below, it will show that the nucleotide sequence from place 63648346 thru 63648363 on human chromosome 8 is exactly CTCCTTTGTTGTGTTGT

[https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?](https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log$=nuclalign&blast_rank=2&from=63648346&to=63648363)

[report=genbank&log\\$=nuclalign&blast_rank=2&from=63648346&to=63648363](https://www.ncbi.nlm.nih.gov/nucleotide/NC_000008.11?report=genbank&log$=nuclalign&blast_rank=2&from=63648346&to=63648363)

5. **STEPHERS SAYS:**

April 7, 2020 at 6:41 am

I think you hit the nail on the head here. I had been searching and searching for this throughout the ncbi site and Genbank, but had not hit upon it. I had a hunch that the Human Genome Project was involved with this. I also suspected the data from 23&me and the Ancestry DNA database was being covertly utilized as well. I also watched videos of Kary Mullis discussing his invention of the PCR test, as well as the significance of SNPs (genomic polymorphisms). If you read this excerpt below by Kary Mullis (the man who INVENTED the PCR test being used to diagnose CV), you will see him talk about it detecting “proteins”. Chromosome 8 is a protein. If it is a perfect match, then that is precisely what the PCR test is intended for – to detect and amplify chromosome 8 (under the guise of a “novel” coronavirus). I am still researching this, but it seems Chromosome 8 has something to do with our individual susceptibility to illness. If we combine the research by Duessberg and others that claimed HIV did not cause AIDS, then we can apply the same logic here. I recently received Duessberg’s book on the AIDS fakery (thoughtfully sent and signed by him) and the forward is by Kary Mullis. I still do not trust Kary Mullis, given his background, and I highly suspect his involvement with this current fakery at the highest levels. He allegedly died from PNEUMONIA last summer. I think there is a “tell” in that. Just my hunch though. Nevertheless, this is what Mullis said (I will follow with a link to this in a subsequent post) — — — Kary Mullis ... is thoroughly convinced that HIV is not the cause of AIDS. With regard to the viral-load tests, which attempt to use PCR for counting viruses, Mullis has stated: “Quantitative PCR is an oxymoron.” PCR is intended to identify substances qualitatively, but by its very nature is unsuited for estimating numbers. Although there is a common misimpression that the viral-load tests actually count the number of viruses in the blood, these tests cannot detect free, infectious viruses at all; they can only detect proteins that are believed, in some cases wrongly, to be unique to HIV. The tests can detect genetic sequences of viruses, but not viruses themselves [(29), p. 3].

FAUXLEX SAYS:

April 7, 2020 at 7:23 am

Thanks for the links. Very interesting stuff! Granted, if chromosome 8 is being multiplied it is only this one teeny strip of code being multiplied. But yes, I really did not expect to find this. I hoped that I would not, because the implications would be remarkable. Still, I had to check and am amazed at finding this match.

1. **STEPHERS SAYS:**

April 7, 2020 at 2:09 pm

Agreed – I should have clarified – only one small component/strip of code being amplified. Regardless, the “virus” narrative is a complete diversion to obscure what is really being tested and treated (within a massive psyop and 4IR/digital disruption agenda layered on top – just read all the relevant white papers from the World Economic Forum). It seems to me there are highly classified clinical trials underway with current patients who have signed NDAs in return for treatment – most likely for various cancers, including leukemia and lymphoma, etc. I suspect the vitamin c treatments are simply placebo controls. I also posit that the plasma from “cured” patients (not from a virus, but rather a SNP/mutation contributing to illness) has some truth to it, as it could be “revelation of the method” with respect to de-activating/neutralizing these polymorphisms which are leading to potentially fatal disease. My guess, though, is the VIPs in Hollywood and government, etc, will keep this information (and relevant treatments) to themselves, for the most part. Hope this makes some sense. I am happy to elaborate or clarify at any time...

1. **SMJ SAYS:**

April 7, 2020 at 8:04 am

Funny how ole eu“kary”ote mullis got wrote out of the script before his nucleic chain reaction test came under corona scrutiny...

“PCR was born in the mind of Kary Mullis on a moonlit night in northern California in 1983. As he drove through the hills, Mullis was thinking about the precision of DNA replication, and a way to tap into it popped into his mind. He excitedly explained his idea to his girlfriend and then went home to think it through. “It was difficult for me to sleep with deoxyribonuclear bombs exploding in my brain,” he wrote much later. (A quote so common that I can’t find the original source, which I think was Scientific American.)”
blogs.plos.org/dnascience/2019...

...seeing as eu“kary”ote had a reputation for being a bit of a mullis...

Mullis

Name used to describe someone who often does very normal things in crazy, insane, or unpredictable ways. Sometimes these people may confuse you by doing something normal one second and then doing the very same thing in some wacked out way the next second. An easy way to spot a Mullis is thier unique way of playing poker which causes them to either loose horribly in the first few hands OR confuse the other players so as to cause them to fold everytime the Mullis bets due to the unpredictability of the hand being bet on.

While playing poker, Matt went all in as he has done many times before. The other players all fold their hands. Matt reveals that he had absolutely nothing of any value in his hand. One player yells “I should have known he didn’t have anything because he’s a fucking Mullis!”

1. STEPHERS SAYS:

[April 7, 2020 at 1:50 pm](#)

So, it seems Mullis was a Mullis in all regards indeed!

FAUXLEX SAYS:

[April 7, 2020 at 8:49 pm](#)

I just find it insane that Mullis died less than a year before this whole thing, and of pneumonia no less! Mark, he does sound like an interesting man and I definitely also could see myself getting along with him.

6. STEPHERS SAYS:

[April 7, 2020 at 6:43 am](#)

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4172096/>

Questioning the HIV-AIDS Hypothesis: 30 Years of Dissent
Patricia Goodson

7. STEPHERS SAYS:

[April 7, 2020 at 6:45 am](#)

This is a history of the Human Genome Project

<https://science.sciencemag.org/content/291/5507/1195.full>

8. STEPHERS SAYS:

[April 7, 2020 at 6:47 am](#)

<https://www.ncbi.nlm.nih.gov/pmc/articles/PMC4502642/>

Defining “mutation” and “polymorphism” in the era of personal genomics

Roshan Karki, Deep Pandya, Robert C. Elston, and Cristiano Ferlini

Hopefully this helps to explain further the research into SNPs...

9. STEPHERS SAYS:

[April 7, 2020 at 2:54 pm](#)

<https://www.ncbi.nlm.nih.gov/probe/docs/projhapmap/> Haplotypes are very pertinent to this, as it would explain the reason why the US “decided” to use a different PCR test than China (hint: they “needed” a different test – one that can detect SNPs in our population). Hope this link

provides enough background info to explain.

10. **STEPHERS SAYS:**

April 7, 2020 at 3:13 pm

One more link for background on the HapMap

Project. <https://www.genome.gov/10001688/international-hapmap-project>.

To reiterate, my theory only makes sense if you take the “virus” narrative out of the equation, since it was there to obscure all along. If the PCR test is testing for SNPs encoded in DNA/chromosomes (in this one case with chromosome 8, as caught by FAUXLEX – there are most likely others being tested and corrected for as well), then this is all an extension of a massive epigenomics R&D operation (albeit highly classified and involving those with high security clearances). Kary Mullis, himself, set forth that the PCR test should only ever be used in R&D, and NOT as a diagnostic tool. Hence, I am suggesting that this is precisely how it is being utilized – even though it appears to be for diagnosis of said viral infection. With that said, the entire lot of “frontline” doctors and nurses shown all over the mockingbird media are in reality part of a massive, global simulation (functional pandemic preparedness exercise). Even “Peekay’s” recent compilation of the spectacle of singing and dancing doctors and nurses should speak for itself – there is NO emergency at the present moment.

1. **TED SAYS:**

April 8, 2020 at 12:51 pm

Excellent work, thank you.

11. **BERND PAYSAN SAYS:**

April 11, 2020 at 5:23 am

People, the virus is an RNA virus, and chromosome 8 is DNA. It’s not going to be replicated by the RT-PCR, because this process starts with the Reverse Transcriptase (RT!), and therefore does not copy DNA.

The remaining question is whether this snippet can exist as RNA in human cells, which is still possible, but not necessarily likely. The overall test is positive if all three primers+probes are found, so one false positive primer does not kill the entire test.

Furthermore, you also need to have the probe in proximity to the primer. If the PCR doesn’t hit the probe, the test will stay negative. Is the probe found in chromosome 8?

The Chinese had some problems with their test in the first weeks, and changed things to improve the situation. This is normal in such a situation, where tests are developed in a hurry. The US CDC had severe quality issues in the beginning, even water tested positive.

FAUXLEX SAYS:

April 12, 2020 at 9:53 am

Thank you for the comment. This is honestly what I was looking for...hoping to be wrong. Although I disagree that the process could not possibly be thrown off by this error. Any instances where chromosome 8 had been unzipped for replication would leave the matching sequence available for the PCR primer to anneal. This might not cause full replication by the PCR process, but if the test is so touchy you would really think this could throw off their results. If as you say plain water originally tested positive. Thanks again for the comment.

You would think they would have some technology to pick PCR primers that selects against sequences already found within human DNA. They also should have gone with a primer longer than 18 characters if they want reliable results. These tests are going to have a huge rate

of false positives and/or negatives to the point where it's questionable how much we can rely upon it. Let alone the idea that 80% of people who test positive show no outward symptoms. We need to end these draconian lockdowns immediately.

12. Pingback: [Unexpected Heroes of Coronavirus Dissent: Tanzania President John Magufuli – Piece of Mindful](#)
13. Pingback: [PCR Test - A Shaky Foundation - The Covid-19 Reader](#)
14. Pingback: [כיצד עובדת הבדיקה ומה היא בעצם בודקת? - קורונה - האמת המוסתרת](#)
15. Pingback: [BOMBSHELL: WHO Coronavirus PCR Test Primer Sequence is Found in All Human DNA – Philosophers Stone](#)
16. Pingback: [REBLOG: George Floyd, The Stiff Who Came In From The Cold – Piece of Mindful](#)
17. Pingback: [THE PLAGUE OF FEAR 2020 – Part 4 – Philosophers Stone](#)
18. Pingback: [La secuencia del cebador de la prueba PCR de coronavirus de la OMS se encuentra en todo el ADN humano - agarcime.com](#)
19. Pingback: [BOMBAZO: La secuencia del cebador de la prueba PCR de coronavirus de la OMS se encuentra en todo el ADN humano - Soy Latino](#)
20. Pingback: [You are the virus! | Fakeologist.com](#)
21. Pingback: [BOMBSHELL: WHO Coronavirus PCR Test Primer Sequence is Found in All Human DNA – Piece of Mindful – Zero Vacunas](#)
22. Pingback: [Noticia Bomba: Los Test PCR, Lo Que Detectan es el Cromosoma 8 de Todos los Seres Humanos. – Neo Jebus](#)
23. Pingback: [WHO'S COVID-19 PCR test primer sequence is found in ALL human DNA – NEWS FROM UNDERGROUND](#)
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30. Pingback: [Bombshell Evidence: COVID RNA Base Pairs are Identical to Chromosome 8 Human DNA | Our Greater Destiny](#)
31. Pingback: [you are the virus – Covid 19 Information](#)
32. Pingback: [5G, MICROCHIP, CORONA VURUS = SANATISMO SANITARIO, LE CRONACHE MAI RIVELATE. Salmi 5 – ITALIA ALATA](#)
33. Pingback: [Compassion, "Collective Risk", Pets, Humans 2.0 and MRNA Vaccines, Jackboots in Oz? - Investigating The Alleged COVID-19 Pandemic](#)
34. Pingback: [Evidence ,COVID-19, RNA, 5G,Chromosome 8, Human DNA,Amanda Vollmer,WHO, | altnews.org](#)
35. Pingback: [BOMBSHELL: WHO Coronavirus PCR Test Primer Sequence is Found in All Human DNA | Anti Oligarch](#)
36. Pingback: [David Icke and Dr. Andrew Kaufman analyze the match between SARS-CoV-2 and Chromosome 8 sequences | Lys-d'Or](#)
37. Pingback: [La dichiarazione di indipendenza Al Congresso, 4 luglio 1776 = The declaration of independence In Congress, July 4, 1776 – ITALIA ALATA](#)
38. Pingback: [ARE THOSE COVID TESTS SEARCHING FOR "SOMEONE"?](#)
39. Pingback: [I Have a Dream = Ho un Sogno by Martin Luther King, Jr. 1963, Washington – ITALIA ALATA](#)

40. Pingback: [Biohacking PCR to detect covid19 at home - Ramon Puig - Sales and Marketing expertise](#)
41. Pingback: [Dit is waarom bijna iedereen positief test op corona | Wakkeren](#)
42. Pingback: [Wat Covid-19 wordt genoemd is geen virus, maar een stukje van ons DNA | JDreport.com](#)
43. Pingback: [You are the virus! | Fakeologist.com](#)
44. Pingback: [Killuminati](#)
45. Pingback: [Wielki przekręt WHO!! – WolniLudzie.net](#)
46. Pingback: [¡¡BOMBAZO!! La secuencia genética que la OMS da para identificar el virus está presente en el gen 8 de todos los humanos - La mentira esta ahí fuera La mentira esta ahí fuera](#)
47. Pingback: [ORVOSOK A TISZTÁNLTÁSÉRT - Szakrális Magyarország](#)
48. Pingback: [A virus walks into a bar .. \(updated!\) * Vuurvogel](#)
49. Pingback: [KÖZLEMÉNY A KONFERENCIÁNK UTÁN 2020.08.21. - Orvosok és Egészségügyi Dolgozók a Tisztánltásért](#)
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62. Pingback: [2020: Do your Research! | My 25 Year Journey to Health](#)

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