

More Evidence COVID Was Tinkered within a Lab? Now Scientists Find Virus Contains Tiny Chunk of DNA that Matches Sequence Patented by Moderna Three Years Before Pandemic Began

By <u>Connor Boyd</u> Global Research, February 24, 2022 Daily Mail Online 23 February 2022 Region: <u>Europe</u> Theme: <u>Intelligence</u>, <u>Science and Medicine</u>

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Fresh suspicion that COVID may have been tinkered with in a lab emerged today after scientists found genetic material owned by <u>Moderna</u> in the virus's spike protein.

They identified a tiny snippet of code that is identical to part of a gene patented by the vaccine maker three years before the pandemic.

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(54)	MODIFIED POLYNUCLEOTIDES FOR THE PRODUCTION OF ONCOLOGY-RELATED PROTEINS AND PEPTIDES				61/618,873, filed on Apr. 2, 2012, provisional application No. 61/681,650, filed on Aug. 10, 2012 provisional application No. 61/737,147, filed on Dec 14, 2012, provisional application No. 61/618,878 filed on Apr. 2, 2012, provisional application No 61/681,654, filed on Aug. 10, 2012, provisiona application No. 61/737,152, filed on Dec. 14, 2012				
(71)	Applicant: Moderna Therapeutles, Inc., Cambridge, MA (US)								
(72)	Inventors: Stephane Bancel, Cambridge, MA (US); Tirtha Chakraborty, Medford, MA (US); Antonin de Fougerolles, Waterloo (BE); Sayda M. Elbashir, Cambridge, MA (US); Matthias John, Cambridge, MA (US); Matthias John, Cambridge, MA (US); Susan Whoriskey, Belmont, MA (US); Kristy M. Wood, Wellesley, MA (US); Paul Hatala, Charlestown, MA (US); Jason P. Schrum, Philadelphia, PA (US); Kencehi Ejebe, New York, NY (US); Jeff Lynn Ellsworth, Lexington, MA (US); Justin Guild, Framingham, MA (US)			n, sty n	provisional application No. 61/618,885, filed on Apr. 2, 2012, provisional application No. 61/681,658, filed on Aug. 10, 2012, provisional application No. 61/737,155, filed on Dec. 14, 2012, provisional application No. 61/618,896, filed on Apr. 2, 2012, provisional application No. 61/668,157, filed on Jul 5, 2012, provisional application No. 61/668,1661, filed on Aug. 10, 2012, provisional application No. 61/737,160, filed on Dec. 14, 2012, provisional application No. 61/618,911, filed on Apr. 2, 2012, provisional application No. 61/681,667, filed on Aug. 10, 2012, provisional application No. 61/737,168, filed on Dec. 14, 2012, provisional application No. 61/618,922, filed on Apr. 2, 2012, provisional application No. 61/681,675, filed on Aug. 10, 2012, provisional application No. 61/737,174, filed on Dec. 14, 2012, provisional application No. 61/618,967, 104 on Dec. 14, 2012, provisional application No. 61/618,967, 104 on Dec.				
(73)	Assignce: ModernaTX, Inc., Cambridge, MA (US)				filed on Apr. 2, 2012, provisional application No 61/681,687, filed on Aug. 10, 2012, provisional				
(*)	Notice:	patent is ext	ny diselaimer, the term of t tended or adjusted under b) by 0 days.		Int. Cl. C07K 14/535 C07K 14/505		(2006.0 (2006.0	- A	
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(65)		Prior Pul	blication Data		See application	on file fo	r comple	ete search history.	
	US 2016/0152678 A1 Jun. 2, 2016		(56)		Referen	ces Cite	d		
	Palated U.S. Ameliantian Data			U.S.	PATENT	DOCUN	MENTS		
(63)	Continuati Dec. 16, 2	Related U.S. Application Data antinuation of application No. 14/106,988, filed on res. 16, 2013, now Pat. No. 9,301,993, which is a			2,008,526 A 3,552,394 A	1/1971	Wrapple Horn et tinued)		
	continuation of application No. PCT/US2013/030070, filed on Mar. 9, 2013.		NO.	FOREIGN PATENT DOCUMENTS					
(60)	10, 2012,	provisional	No. 61/681,742, filed on A application No. 61/737,2 , provisional application)	24. CA		6634 AI 3135 C (Con	12/200 6/200		

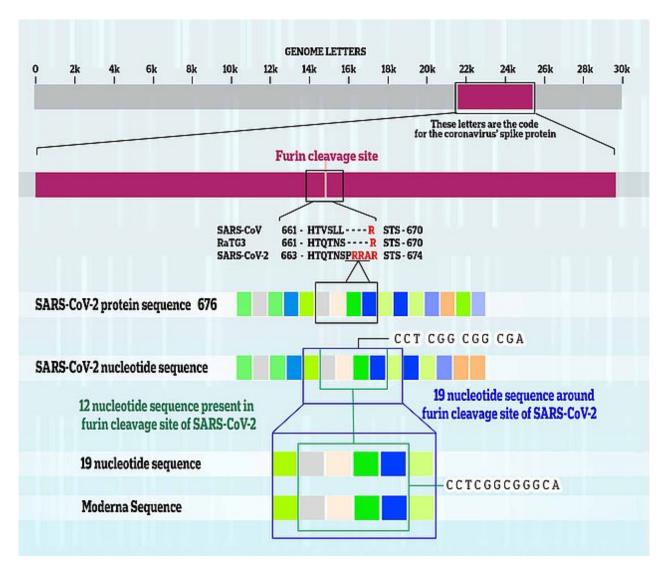
It was discovered in SARS-CoV-2's unique furin cleavage site, the part that makes it so good at infecting people and separates it from other coronaviruses.

The structure has been one of the focal points of debate about the virus's origin, with some scientists claiming it could not have been acquired naturally.

The international team of researchers suggest the virus may have mutated to have a furin cleavage site during experiments on human cells in a lab.

They claim there is a one-in-three-trillion chance Moderna's sequence randomly appeared through natural evolution.

But there is some debate about whether the match is as rare as the study claims, with other experts describing it as a 'quirky' coincidence rather than a 'smoking gun'.



SARS-CoV-2, which causes Covid, carries all the information needed for it to spread in around 30,000 letters of genetic code, known as RNA. The virus shares a sequence of 19 specific letters with a genetic section owned by Moderna. Twelve of the shared letters make up the structure of Covid's furin cleavage site, with the rest being a match with nucleotides on a nearby part of the genome. Source: Daily Mail Online

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