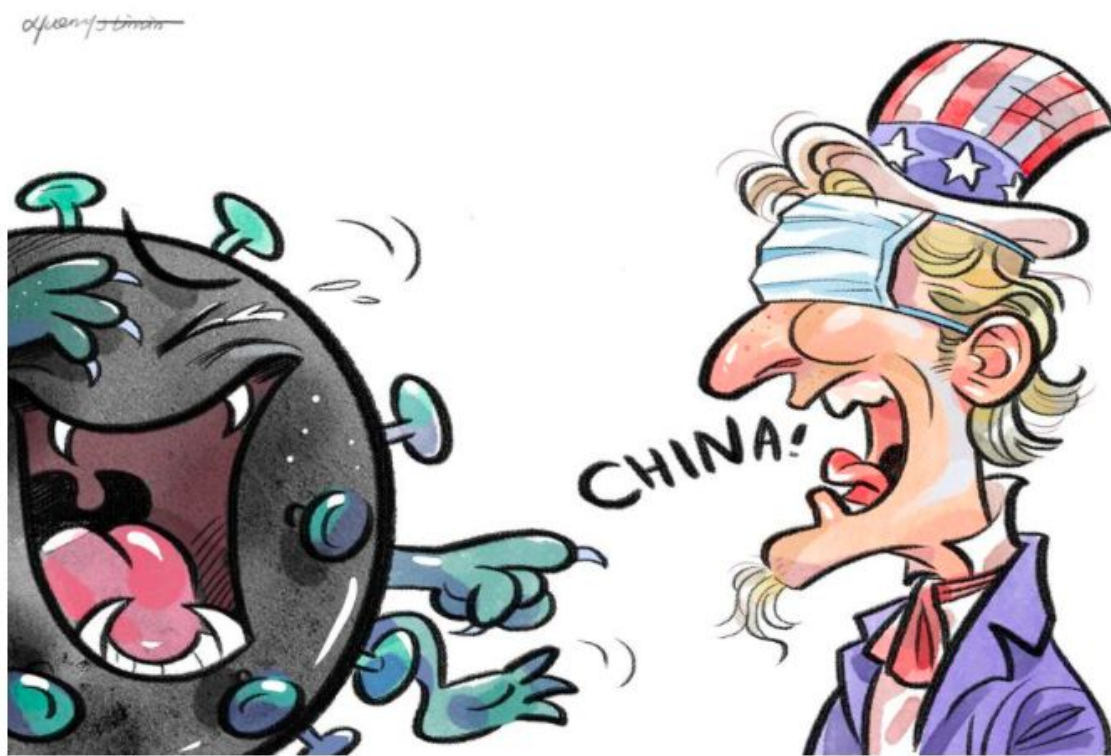


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Who has let the genie out of the bottle?

By Xin Ping | chinadaily.com.cn | Updated: 2021-08-13 22:05



[Li Min/ China Daily]

Who has Let the Genie out of the Bottle?

-- A Letter to Dr. Ralph Baric

Dear Dr. Ralph S. Baric,

I am impressed by the founding principles of the University of North Carolina (UNC) at Chapel Hill—*lux, libertas*—light and liberty, a motto that inspires the university to chart "a bold course of leading change to improve society and to help solve the world's greatest problems". With all due respect, I have to say that the history of your laboratory's involvement in coronavirus research seems to go against the missions and values of your institution. In particular, some of your remarks regarding the Wuhan Institute of Virology (WIV) make me, and many Chinese citizens feel a little bit puzzled and concerned.

As one of the leading virologists in the world, you have been involved in researches designing, modifying and synthesizing SARS-like coronaviruses for years. Since 1983, you have published 268 papers directly focusing on coronavirus. In 2006, your team managed to develop a desirable mutation which can cast an imminent death spell to mice and has the potential to infect human beings with severe pneumonia.

This is indeed a huge progress as the artificially-created virus leads us closer to the mysterious natural mutation process of the virus. Nevertheless, if poorly monitored, it could be extremely risky for vulnerable human bodies.

You once warned that the technique of synthesizing virus sequences grants humans with the potential to produce biological weapons of mass destruction. Your warning, unfortunately, was regarded as an advertisement by warmongers, the infamous Fort Detrick lab included. With full knowledge of its mission, you willingly accepted the cooperative offer.

In May, President Biden ordered US intelligence officials to "redouble" efforts to investigate the origins of COVID-19, including the theory that it came from a laboratory. With two weeks to go before Biden's 90-day push to find answers, has the US intelligence community looked into the lab you worked in?

From the annual report of UNC, your lab was involved in a host of accidents from 2012 to 2018. In 2017 and 2018, 42 and 43 accidents were reported respectively. According to ProPublica, six accidents involving lab-created SARS-like and MERS-like coronaviruses were recorded in your lab. The accidents ranged from mouse bites to other mishaps during experiments. One of the examples is, in August 2015, two researchers received medical monitor after a mouse escaped inside your biosafety level-3 lab. "The mouse, which had been infected with an undisclosed type of 'mouse adapted' virus, squirmed free of a researcher's gloved hand and onto the lab floor," detailed in the piece from ProPublica.

Obviously, the safety of the experiments should be the utmost priority in any biochemical lab, let alone your research which centers on some of the most dangerous viruses in the world. Why such errors and accidents happened again and again in your lab?

Ironically, with such a large number of accidents related to your own lab, you have instead pointed the finger of suspicion at the WIV for "safety concerns".

No wonder you commented "your luck may eventually run out" on possible COVID-19 lab-leak from the WIV, since the safety of your experiments is mostly predicated on pure luck rather than strict regulations.

Without any warranted evidence, you indicated in your interview with MIT Technology Review that some artificially-modified viruses can be "disguised" as coming from nature, and even implied that files at WIV possess the answers people want. But you know it all too well that all the "Gain of Function" studies were done in your lab and Dr. Shi Zhengli from WIV did nothing but offer viral sequences.

Dr. Baric, as one of the leading coronavirus experts, you know what virus origin-tracing means in scientific sense. In the global effort to discover the origins of coronavirus, whoever first cracks the scientific puzzle is certainly laudable. Hence, let SCIENCE speak is the only scientific approach to this extremely complex journey. A scientist's hard-earned credentials and expertise are meant for searching for truth and dedicated to the well-being of all people. It is dangerous when science degenerates into servant of politics. Far more damaging than the virus itself is, a scientist ending up being anti-science or even an apologist for politicians.

The genie, no matter wherever it may escape from, will be found in the end.

Sincerely,

Xin Ping

(Xin Ping is a commentator on international affairs, writing regularly for China Daily, CGTN, Global Times, and other media outlets. He can be reached at xinping604@gmail.com.)

Conspiracy theory or reasonable skepticism? Why we should demand an investigation into US labs for origins of COVID-19

By GT staff reporters

Published: Aug 15, 2021 07:48 PM Updated: Aug 15, 2021 09:28 PM



Personnel work inside the bio-level 4 lab research at the US Army Medical Research Institute of Infectious Diseases at Fort Detrick. Photo: AFP

Baric and his virus modification technique

Let's start with Ralph Baric, an American scientist who is called the "Coronavirus Hunter."

WUNC 91.5
NORTH CAROLINA PUBLIC RADIO

91.5 Chapel Hill 88.9 Manteo 90.9 Rocky Mount
91.1 Welcome 91.9 Fayetteville 90.5 Buxton

WUNC
BBC World News

NEXT UP: 5:00

Meet 'The Coronavirus Hunter' Ralph Baric

North Carolina Public Radio | By Frank Stasio
Published May 25, 2020 at 11:55 AM EDT

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in

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Screenshot on North Carolina-based media WUNC

Ralph S. Baric, PhD

WILLIAM R. KENAN, JR. DISTINGUISHED PROFESSOR
DEPARTMENT OF EPIDEMIOLOGY

PROFESSOR
DEPARTMENT OF MICROBIOLOGY AND IMMUNOLOGY

MEMBER
LINEBERGER COMPREHENSIVE CANCER CENTER

Screenshot on UNC

According to [a report](#) from the school-run media of the University of North Carolina, where Baric works, "Baric has been tracking coronaviruses for decades and working on medications to treat coronavirus-caused infections."

According to the [MIT Technology Review](#), with the support of this technique, Baric can not only cultivate a living virus based on the gene fragments of the coronaviruses, but also modify the genes of the coronaviruses and create new ones to explore the harm of the viruses to humans.

Baric had developed a way around that problem—a technique for “reverse genetics” in coronaviruses. Not only did it allow him to bring an actual virus to life from its genetic code, but he could mix and match parts of multiple viruses. He wanted to take the “spike” gene from SHC014 and move it into a genetic copy of the SARS virus he already had in his lab. The spike molecule is what lets a coronavirus open a cell and get inside it. The resulting chimera would demonstrate whether the spike of SHC014 would attach to human cells.

Screenshot on MIT

In 2003, [a published paper](#), of which Baric was a co-author, showcased the power of this technique. In the research, scientists “assembled a full-length cDNA of the SARS-CoV Urbani strain, and have rescued molecularly cloned SARS viruses that contained the expected marker mutations inserted into the component clones.”



RESEARCH ARTICLE

Reverse genetics with a full-length infectious cDNA of severe acute respiratory syndrome coronavirus

Boyd Yount, Kristopher M. Curtis, Elizabeth A. Fritz, Lisa E. Hensley, Peter B. Jahrling, Erik Prentice, Mark R. Denison, Thomas W. Geisbert, and Ralph S. Baric

^{*}Departments of Epidemiology and Microbiology and Immunology and [†]Carolina Vaccine Institute, University of North Carolina, Chapel Hill, NC 27599-7435; [‡]U.S. Army Medical Research Institute of Infectious Diseases, Fort Detrick, MD 21702; and [§]Departments of

Screenshot on PNAS Journal

Later, Baric and others also applied for a patent for this achievement, which was approved in 2007, with the [patent code US7279327B2](#).

RESEARCH ARTICLE

Reverse genetics with a full-length infectious cDNA of severe acute respiratory syndrome coronavirus

Boyd Yount, Kristopher M. Curtis, Elizabeth A. Fritz, Lisa E. Hensley, Peter B. Jahrling, Erik Prentice, Mark R. Denison, Thomas W. Geisbert, and Ralph S. Baric

^{*}Departments of Epidemiology and Microbiology and Immunology and [†]Carolina Vaccine Institute, University of North Carolina, Chapel Hill, NC 27599-7435; [‡]U.S. Army Medical Research Institute of Infectious Diseases, Fort Detrick, MD 21702; and [§]Departments of

Screenshot on PNAS Journal

Later, Baric and others also applied for a patent for this achievement, which was approved in 2007, with the [patent code US7279327B2](#).

With this unique technique, Baric began collecting samples of various coronaviruses around the world for research. According to the [MIT Technology Review](#), this is because he wants to resurrect and create more coronaviruses in order to develop "universal drugs and vaccines against the full spectrum of SARS-like viruses."

<https://www.globaltimes.cn/page/202108/1231519.shtml>

Controversial virus research seen both as groundbreaking, reckless

Tags: [coronavirus](#), [UNC](#), [research](#), [WRAL Investigates](#)

Posted May 14, 2020 6:00 p.m. EDT

Updated May 14, 2020 7:32 p.m. EDT

Screenshot on WRAL

Although many scientists around the world have said that artificially modified viruses may leave traces, and are different from viruses that have evolved in nature, Baric said in an interview with an [Italian media outlet](#) in September 2020 that "it is possible to engineer a virus [without leaving a trace](#)."



Ralph Baric takes interview with Italian media in September, 2020. Photo: Screenshot on Huffington Post

Ralph S. Baric

From Wikipedia, the free encyclopedia

Ralph Steven Baric (born 1954) is [William R. Kenan Jr.](#) Distinguished Professor in the Department of Epidemiology, and Professor in the Department of Microbiology and Immunology at the [The University of North Carolina at Chapel Hill](#).

Baric's work involves coronaviruses, including gain of function research aimed at devising effective vaccines against coronaviruses.^[1] Baric has warned of emerging coronaviruses presenting as a significant threat to global health, due to zoonosis.^{[2][3]}

Ralph S. Baric	
Born	1954 (age 66–67)
Nationality	American
Alma mater	North Carolina State University
Scientific career	
Fields	Epidemiology
Institutions	University of North Carolina at Chapel Hill
Thesis	<i>Inhibitors of host transcription block Sindbis virus replication</i> ↗ (1982)
Doctoral advisor	Robert E. Johnston



Methods and compositions for infectious cDNA of SARS coronavirus

Publication number: 20060240530

Abstract: The present invention provides a cDNA of a severe acute respiratory syndrome (SARS) coronavirus, recombinant SARS coronavirus vectors, and SARS coronavirus replicon particles.

Also provided are methods of making the compositions of this invention and methods of using the compositions as immunogens and/or vaccines and/or to express heterologous nucleic acids.

Type: Application

Filed: January 19, 2006

Publication date: October 26, 2006

Inventors: Ralph Baric, Rhonda Roberts, Boyd Yount, Kristopher Curtis

STATEMENT OF FEDERAL SUPPORT

This invention was supported by government funding under grant numbers A123946 and GM 63228 from the National Institute of Health, Allergy and Infectious Diseases. The United States Government has certain rights to this invention.

The present invention further provides a method of introducing a heterologous RNA into a subject, comprising administering to the subject an effective amount of the particles or populations and/or compositions of this invention.

Also provided herein is a method of inducing an immune response and/or treating and/or preventing a SARS coronavirus infection in a subject, comprising administering to the subject an effective amount of the viruses, vectors, particles or populations and/or compositions of this invention.

cDNA of SARS Coronavirus

Complementary DNA (cDNA) is DNA synthesized from single-stranded RNA template (mRNA) in a reaction catalyzed by the enzyme by reverse transcriptase



Compositions of Chimeric Spike Proteins

Methods and compositions for chimeric coronavirus spike proteins

Patent number: 9884895

Abstract: The present invention provides compositions and methods comprising a chimeric coronavirus spike protein.

Type: Grant

Filed: March 20, 2015

Date of Patent: February 6, 2018

Assignee: The University of North Carolina at Chapel Hill

Inventors: Ralph Baric, Sudhakar Agnihothram, Boyd Yount

STATEMENT OF FEDERAL SUPPORT

This invention was made with government support under Grant No. U54AI057157 awarded by the National Institutes of Health. The government has certain rights in the invention.

FIELD OF THE INVENTION

The present invention relates to methods and compositions comprising a chimeric coronavirus spike protein for treating and/or preventing a disease or disorder caused by a coronavirus infection.

SUMMARY OF THE INVENTION

In one aspect, the present invention provides a chimeric coronavirus spike protein comprising, in orientation from amino to carboxy terminus: a) a first region comprising a portion of a coronavirus spike protein ectodomain that precedes the receptor binding domain (RBD) as located in a nonchimeric coronavirus spike protein, of a first coronavirus; b) a second region comprising a coronavirus spike protein receptor binding domain (RBD) of a second coronavirus that is different from said first coronavirus; c) a third region comprising a portion of a coronavirus spike protein S1 domain as located in a nonchimeric coronavirus spike protein immediately downstream of the RBD, contiguous with a portion comprising a coronavirus spike protein S2 domain as located immediately upstream of a fusion protein domain in a nonchimeric coronavirus spike protein, wherein said third region is of said first coronavirus; and d) a fourth region comprising a portion of a coronavirus spike protein from the start of the fusion protein domain through the carboxy terminal end as located in a nonchimeric coronavirus spike protein of a third coronavirus that is different from said first coronavirus and said second coronavirus.

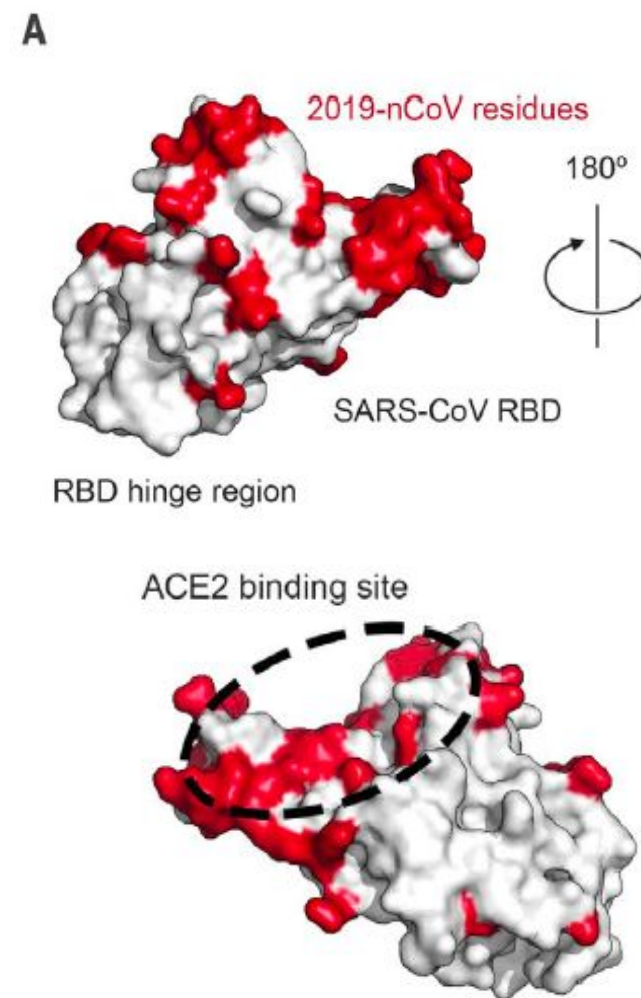


Cryo-EM structure of the 2019-nCoV spike in the prefusion conformation

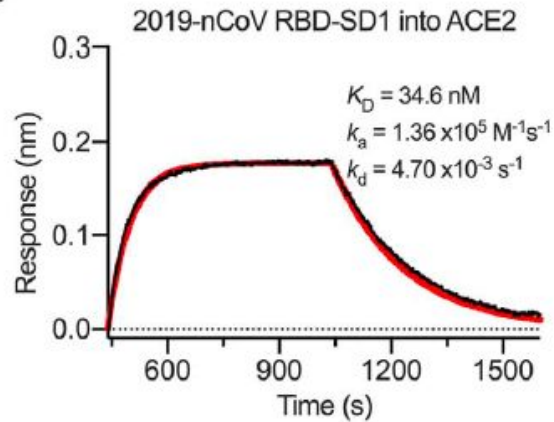
Daniel Wrapp^{1*}, Nianshuang Wang^{1*}, Kizzmekia S. Corbett², Jory A. Goldsmith¹, Ching-Lin Hsieh¹, Olubukola Abiona², Barney S. Graham², Jason S. McLellan^{1†}

The outbreak of a novel coronavirus (2019-nCoV) represents a pandemic threat that has been declared a public health emergency of international concern. The CoV spike (S) glycoprotein is a key target for vaccines, therapeutic antibodies, and diagnostics. To facilitate medical countermeasure development, we determined a 3.5-angstrom-resolution cryo-electron microscopy structure of the 2019-nCoV S trimer in the prefusion conformation. The predominant state of the trimer has one of the three receptor-binding domains (RBDs) rotated up in a receptor-accessible conformation. We also provide biophysical and structural evidence that the 2019-nCoV S protein binds angiotensin-converting enzyme 2 (ACE2) with higher affinity than does severe acute respiratory syndrome (SARS)-CoV S. Additionally, we tested several published SARS-CoV RBD-specific monoclonal antibodies and found that they do not have appreciable binding to 2019-nCoV S, suggesting that antibody cross-reactivity may be limited between the two RBDs. The structure of 2019-nCoV S should enable the rapid development and evaluation of medical countermeasures to address the ongoing public health crisis.

Fig. 4. Antigenicity of the 2019-nCoV RBD. (A) SARS-CoV RBD shown as a white molecular surface (PDB ID: 2AJF), with residues that vary in the 2019-nCoV RBD colored red. The ACE2-binding site is outlined with a black dashed line. (B) Biolayer interferometry sensorgram showing binding to ACE2 by the 2019-nCoV RBD-SD1. Binding data are shown as a black



B

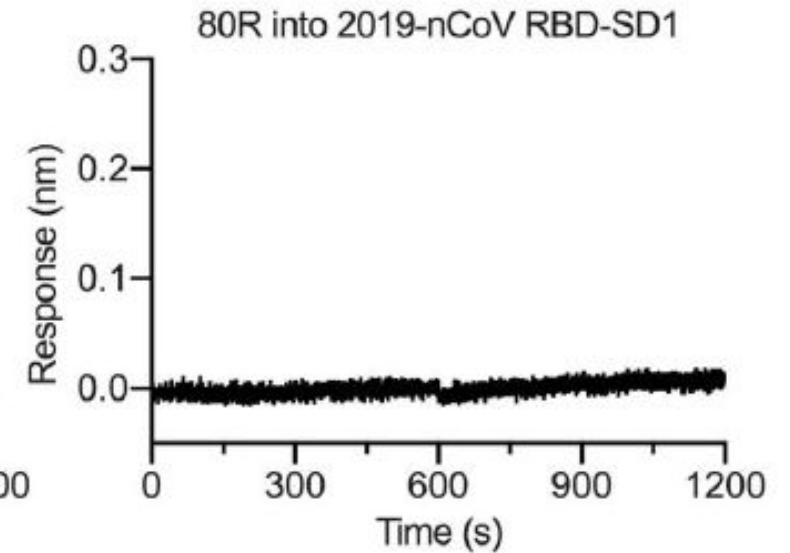
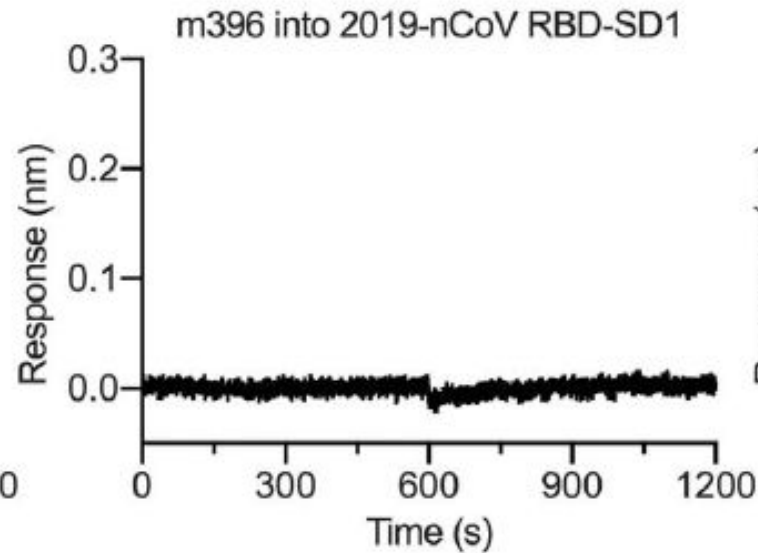
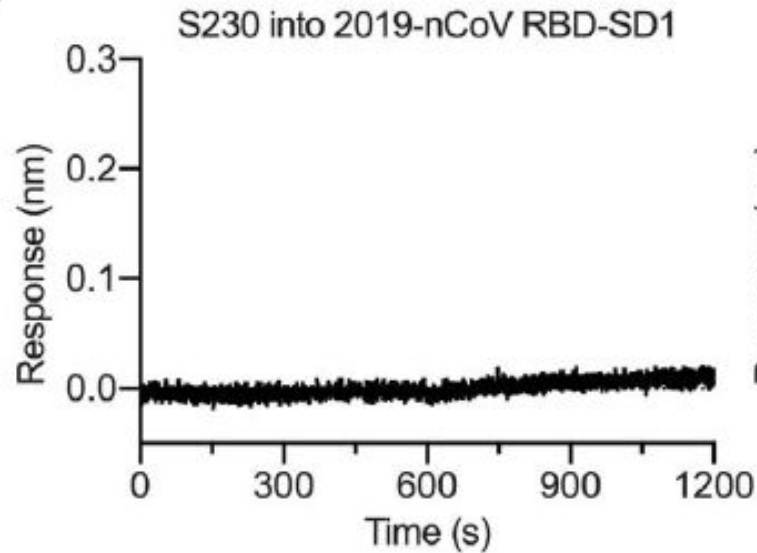


ACE-2 Receptor
Binding of 2019
SARS-CoV-2(S) and

line, and the best fit of the data to a 1:1 binding model is shown in red. (C) Biolayer interferometry to measure cross-reactivity of the SARS-CoV RBD-directed antibodies S230, m396, and 80R. Sensor tips with immobilized antibodies were dipped into wells containing 2019-nCoV RBD-SD1, and the resulting data are shown as a black line.

MONOCLONAL - ANTIBODIES from the EUA COVID-19 'VACCINE' mRNA
RECOMBINANT (S) PROTEIN **DO NOT BIND** TO SARS-CoV-2 PROTEIN

C



STATEMENT OF FEDERAL SUPPORT

This invention was made with government support under Grant Nos. AI085524, AI057157, and U19 AI107810 awarded by the National Institutes of Health. The government has certain rights in the invention.

FIELD OF THE INVENTION

The present invention relates to methods and compositions for detecting and identifying a coronavirus by subgroup as well as treating and/or preventing a disease or disorder caused by a coronavirus infection.

SUMMARY OF THE INVENTION

The present invention provides a method of detecting the presence of a coronavirus in a sample and identifying the subgroup of the coronavirus in the sample, comprising: a) contacting a sample with a panel of proteins comprising: 1) one or more nucleocapsid proteins from a subgroup 2c coronavirus, 2) one or more nucleocapsid proteins from a subgroup 2b coronavirus, 3) one or more nucleocapsid proteins from a subgroup 2a coronavirus, 4) one or more

The present invention also provides a method of identifying a coronavirus spike protein for administration to elicit an immune response to coronavirus in a subject infected by a coronavirus; and/or a subject at risk of coronavirus infection and/or to a subject for whom eliciting an immune response to a coronavirus is needed or desired, comprising: a) contacting a sample obtained from a subject infected with a coronavirus with a panel of proteins comprising: 1) one or more spike proteins from a subgroup 2c coronavirus, 2) one or more spike proteins from a subgroup 2b coronavirus, 3) one or more spike proteins from a subgroup 2a coronavirus, 4) one

Methods and Compositions for Coronavirus Diagnostics and Therapeutics

METHODS AND COMPOSITIONS FOR CORONAVIRUS DIAGNOSTICS AND THERAPEUTICS

Publication number: 20160238601

Abstract: The present invention provides methods and compositions for detecting a coronavirus in a sample and identifying the subgroup of the coronavirus in the sample.

Type: Application

Filed: October 14, 2014

Publication date: August 18, 2016

Inventors: Ralph Baric Sudhakar Agnihothram, Boyd Yount



Ralph Baric's research lab leads the ...
newsobserver.com

METHODS AND COMPOSITIONS FOR RECOMBINANT DENGUE VIRUSES FOR VACCINE AND DIAGNOSTIC DEVELOPMENT

Publication number: 20200230224

Abstract: The present invention provides compositions and methods of use comprising a chimeric dengue virus E glycoprotein comprising a dengue virus E glycoprotein backbone, which comprises amino acid substitutions that introduce an epitope that is recognized by an antibody from a dengue virus serotype that is different from the dengue virus serotype of the dengue virus E glycoprotein backbone.

Type: Application

Filed: August 29, 2019

Publication date: July 23, 2020

Inventors: Ralph Baric, Douglas Widman, Boyd Yount, Emily Gallichotte, Scott Royal, Aravinda Desilva, Jessica Swanstrom

Also provided herein is a method of protecting a subject from the effects of dengue virus infection, comprising administering to the subject an effective amount of the E glycoprotein of this invention, the flavivirus particle of this invention, the VLP of this invention, the nucleic acid molecule of this invention, the population of this invention, and/or the composition of this invention and any combination thereof.

The present invention also provides various diagnostic methods, including, for example, a method of identifying the presence of a neutralizing antibody to dengue virus serotype 3 and/or 4 in a biological sample from a subject, comprising: a) administering a composition comprising an E glycoprotein comprising an E glycoprotein backbone of serotype 4 comprising amino acid substitutions that introduce an epitope that is recognized by an antibody that is reactive with

Recombinant Chimeric Dengue Viruses for Vaccines & Diagnostic Development

Dengue viruses are mosquito-borne viral disease, naturally occurring in tropical and subtropical areas with an estimated 400 million cases worldwide each year.

According to WebMD, risk is increasing for those living along the Texas-Mexico Border. People with weakened immune systems and second (subsequent) infection have greater risk of more severe disease symptoms.

1. <https://patents.justia.com/patent/20200230224>

2. <https://www.webmd.com/a-to-z-guides/dengue-fever-reference>

Search Results

7 patents

Search Text: Baric

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- > Org Names
-
- > Agencies
-
- > States
-
- > Countries
-
- > Principal Investigators

Core NIH Project Number	Patent Number	Patent Title	Patent Owner ?	Primary Agency ?
U19AI109761	10829823	Compositions and methods for the rapid differential detection of Zika virus	COLUMBIA UNIV NEW YORK MORNINGSIDE	NIH
R01AI023946	7618802	Compositions of coronaviruses with a recombination-resistant genome	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH
R01GM063228	7618802	Compositions of coronaviruses with a recombination-resistant genome	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH
R01AI023946	6593111	Directional assembly of large viral genomes and chromosomes	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH
R01AI056351	9975923	Methods and compositions for norovirus blockade epitopes	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH
R01AI023946	7279327	Methods for producing recombinant coronavirus	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH
R01GM063228	7279327	Methods for producing recombinant coronavirus	UNIV OF NORTH CAROLINA CHAPEL HILL	NIH

2020 - 2021
NIH/NIAID GRANTS

RALPH BARIC
(UNC/TEXAS/VANDERBILT)

T Act Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC	Similar Projects
Respiratory Virus Vaccine and Adjuvant Exploration									
5U01AI149644-03			BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2021	NIAID	NIAID	\$1,000,000	
Task A24: Establishment of Chronic Pseudomonas aeruginosa and Staphylococcus aureus Infection in Mouse Models of Cystic Fibrosis									
272201700036I-P00008-759301900131-1			BARIC, RALPH	UNIV OF NORTH CAROLINA CHAPEL HILL	2021	NIAID	NIAID	\$279,707	
Determinants of Coronavirus Fidelity in Replication and Pathogenesis									
5R01AI108197-09			DENISON, MARK R. BARIC, RALPH S.	VANDERBILT UNIVERSITY MEDICAL CENTER	2021	NIAID	NIAID	\$672,084	
Antibody Landscape following Human Norovirus Infection and Vaccination									
5R01AI148260-02			BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2021	NIAID	NIAID	\$739,405	
Determinants of Coronavirus Fidelity in Replication and Pathogenesis									
3R01AI108197-09S1			DENISON, MARK R. BARIC, RALPH S.	VANDERBILT UNIVERSITY MEDICAL CENTER	2021	NIAID	NIAID	\$447,335	
Project 3: SARS CoV-2 Lung Organoid Interactions in Replication and Pathogenesis									
2U19AI116484-06	6912		BARIC, RALPH S.	STANFORD UNIVERSITY	2021	NIAID		\$340,241	
Human antibody-based countermeasures against the Wuhan Coronavirus SARS-CoV-2									
1R01AI157155-01			DIAMOND, MICHAEL S. BARIC, RALPH S. CROWE, JAMES E.	WASHINGTON UNIVERSITY	2020	NIAID	NIAID	\$1,193,309	

\$	279,707
\$	1,000,000
\$	672,084
\$	739,405
\$	447,335
\$	340,241
\$	1,193,309
\$	4,672,081

Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross						
3U19AI100625-09S26276	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID		\$412,634
Core A: Administrative Core						
1U54CA260543-01	8131 BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NCI		\$517,617
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
5U19AI100625-09	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$2,332,322
Respiratory Virus Vaccine and Adjuvant Exploration - Equipment Supplement						
3U01AI149644-02S1	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$1,088,512
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
3U19AI100625-09S3	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$91,160
Project 1: Serological Correlates of SARS CoV2 Immunity and Disease						
1U54CA260543-01	8134 BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NCI		\$658,139
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
3U19AI100625-09S2	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$412,634
Administrative Core						
5U19AI100625-09	7724 BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID		\$220,411
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
3U19AI100625-09S1	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$564,671
Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross						
3U19AI100625-09S38833	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID		\$91,160

2020 CONTINUED
NIH/NIAID GRANTS

RALPH BARIC
(UNC/TEXAS/VANDERBILT)

\$	412,634
\$	517,617
\$	2,332,322
\$	1,088,512
\$	91,160
\$	658,139
\$	412,634
\$	220,411
\$	564,671
\$	6,298,100

Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross

5U19AI100625-09	6276	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID		\$428,666
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Genetic Analysis of COVID-19 Susceptibility and Resistance Determinants in the Collaborative Cross

1R01AI157253-01		HEISE, MARK T BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$748,384
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Cell entry, cross-species transmission and pathogenesis of novel coronavirus from Wuhan

2R01AI110700-06		BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$766,414
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North Carolina Seronet Center for Excellence

1U54CA260543-01		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NCI	NCI	\$3,974,612
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Respiratory Virus Vaccine and Adjuvant Exploration - Equipment Supplement

3U01AI149644-02S1		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$1,088,512
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Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross

3U19AI100625-09S2		BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$412,634
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Project 1: Serological Correlates of SARS CoV2 Immunity and Disease

1U54CA260543-01	8134	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NCI		\$658,139
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Human antibody-based countermeasures against the Wuhan Coronavirus SARS-CoV-2

1R01AI157155-01		DIAMOND, MICHAEL S BARIC, RALPH S CROWE, JAMES E	WASHINGTON UNIVERSITY	2020	NIAID	NIAID	\$1,193,309
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Cell entry, cross-species transmission and pathogenesis of novel coronavirus from Wuhan

2R01AI110700-06		BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$766,414
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Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV

3R01AI132178-04S1		BARIC, RALPH S SHEAHAN, TIMOTHY PATRICK	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$458,053
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Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV

3R01AI132178-03S1		BARIC, RALPH S SHEAHAN, TIMOTHY PATRICK	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$450,462
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Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV

5R01AI132178-04		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$1,166,670
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2020 CONTINUED NIH/NIAID GRANTS

RALPH BARIC
(UNC/TEXAS/VANDERBILT)

\$	428,666
\$	748,384
\$	766,414
\$	3,974,612
\$	1,088,512
\$	412,634
\$	658,139
\$	1,193,309
\$	766,414
\$	458,053
\$	450,462
\$	1,166,670
\$	12,112,269



Determinants of Coronavirus Fidelity in Replication and Pathogenesis

3 R01AI108197-08S1	DENISON, MARK R.  BARIC, RALPH S. 	VANDERBILT UNIVERSITY MEDICAL CENTER	2020	NIAID	NIAID	\$318,794
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Determinants of Coronavirus Fidelity in Replication and Pathogenesis

5 R01AI108197-08	DENISON, MARK R.  BARIC, RALPH S. 	VANDERBILT UNIVERSITY MEDICAL CENTER	2020	NIAID	NIAID	\$672,084
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Genetic Analysis of COVID-19 Susceptibility and Resistance Determinants in the Collaborative Cross

1 R01AI157253-01	HEISE, MARK T.  BARIC, RALPH S. 	UNIV OF NORTH CAROLINA CHAPEL HILL	2020	NIAID	NIAID	\$748,384
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\$	318,794
\$	672,084
\$	748,384
\$	1,739,262

2020 – 2021 TOTAL
NIH/NIAID GRANTS
RALPH BARIC

\$	4,672,081
\$	6,298,100
\$	12,112,269
\$	23,082,450

<https://reporter.nih.gov/search/709S4FzbS0ScdpKxNlg6Q/projects>

https://en.wikipedia.org/wiki/Ralph_S._Baric

Ralph S. Baric

From Wikipedia, the free encyclopedia

Ralph Steven Baric (born 1954) is [William R. Kenan Jr.](#) Distinguished Professor in the Department of Epidemiology, and Professor in the Department of Microbiology and Immunology at the [The University of North Carolina at Chapel Hill](#).

Baric's work involves coronaviruses, including gain of function research aimed at devising effective vaccines against coronaviruses.^[1] Baric has warned of emerging coronaviruses presenting as a significant threat to global health, due to zoonosis.^{[2][3]}

Ralph S. Baric	
Born	1954 (age 66–67)
Nationality	American
Alma mater	North Carolina State University
Scientific career	
Fields	Epidemiology
Institutions	University of North Carolina at Chapel Hill
Thesis	<i>Inhibitors of host transcription block Sindbis virus replication</i>  (1982)
Doctoral advisor	Robert E. Johnston

ESTABLISHMENT OF CHRONIC BACTERIAL INFECTION MODELS IN MOUSE MODELS OF CYSTICs WITH PSEUDOMONAS AERUGINOSA AND STAPHYL						
272201700036I-0-75 9301900131-1	BARIC, RALPH	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID	NIAID	\$442,129
Respiratory Virus Vaccine and Adjuvant Exploration						
1 U01AI149644-01	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID	NIAID	\$1,000,000
Administrative Core						
5 U19AI100625-08 7724	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID		\$818,006
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
5 U19AI100625-08	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID	NIAID	\$2,769,729
Mechanisms of MERS-CoV Entry, Cross-species Transmission and Pathogenesis						
5 R01AI110700-05	BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID	NIAID	\$721,207
Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV						
5 R01AI132178-03	BARIC, RALPH S SHEAHAN, TIMOTHY PATRICK	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID	NIAID	\$1,166,670
Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross						
5 U19AI100625-08 7727	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2019	NIAID		\$428,666
Determinants of Coronavirus Fidelity in Replication and Pathogenesis						
5 R01AI108197-07	DENISON, MARK R BARIC, RALPH S	VANDERBILT UNIVERSITY MEDICAL CENTER	2019	NIAID	NIAID	\$672,084
Molecular Analysis of Serum Antibody Constituents in Zika Virus Infection						
5 R21AI135682-02	GEORGIOU, GEORGE BARIC, RALPH S	UNIVERSITY OF TEXAS, AUSTIN	2019	NIAID	NIAID	\$181,149
Systems Immunogenetics of Biodefense and Emerging Pathogens in the Collaborative Cross						
5 U19AI100625-07	BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2018	NIAID	NIAID	\$2,727,484
Administrative Core						
5 U19AI100625-07 7724	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2018	NIAID		\$342,898
Mechanisms of MERS-CoV Entry, Cross-species Transmission and Pathogenesis						
5 R01AI110700-04	BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2018	NIAID	NIAID	\$727,370
Diagnostic and Prognostic Biomarkers for Viral Severe Lung Disease						
5 U19AI109761-05 8481	BARIC, RALPH S	COLUMBIA UNIVERSITY HEALTH SCIENCES	2018	NIAID		\$889,074
Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV						
5 R01AI132178-02	BARIC, RALPH S SHEAHAN, TIMOTHY PATRICK	UNIV OF NORTH CAROLINA CHAPEL HILL	2018	NIAID	NIAID	\$1,166,670
Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross						
5 U19AI100625-07 7727	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2018	NIAID		\$419,667
Molecular Analysis of Serum Antibody Constituents in Zika Virus Infection						
1 R21AI135682-01	GEORGIOU, GEORGE BARIC, RALPH S	UNIVERSITY OF TEXAS, AUSTIN	2018	NIAID	NIAID	\$233,638
Determinants of Coronavirus Fidelity in Replication and Pathogenesis						
2 R01AI108197-06	DENISON, MARK R BARIC, RALPH S	VANDERBILT UNIVERSITY MEDICAL CENTER	2018	NIAID	NIAID	\$686,584

miFiGHT	
2018 -2019 NIH/NIAID GRANTS	
RALPH BARIC (UNC/TEXAS/VANDERBILT)	
\$	442,129
\$	1,000,000
\$	818,006
\$	2,769,729
\$	721,207
\$	1,166,670
\$	428,666
\$	672,084
\$	181,149
\$	2,727,484
\$	342,898
\$	727,370
\$	889,074
\$	419,667
\$	233,638
\$	686,584
\$	14,226,355
https://reporter.nih.gov/search/70954Fzb50ScdpKlxNlg6Q/projects	

2016-2017
CORONAVIRUS, GOF, ET AL.
NIH/NIAID GRANTS

RALPH BARIC
(UNC/Columbia/Vanderbilt)

\$	211,216
\$	733,354
\$	1,131,261
\$	1,455,240
\$	418,635
\$	396,549
\$	2,021,134
\$	649,146
\$	252,483
\$	135,110
\$	739,947
\$	739,162
\$	899,034
\$	4,853,040
\$	2,322,470
\$	135,110
\$	704,107
\$	548,803
\$	18,345,801

T	Act	Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC
Administrative Core										
5	U19AI107810	-05	8688		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID		\$211,216
Mechanisms of MERS-CoV Entry, Cross-species Transmission and Pathogenesis										
5	R01AI110700	-03			BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID	NIAID	\$733,354
Diagnostic and Prognostic Biomarkers for Viral Severe Lung Disease										
5	U19AI109761	-04	8481		BARIC, RALPH S	COLUMBIA UNIVERSITY HEALTH SCIENCES	2017	NIAID		\$1,131,261
Broad-spectrum antiviral GS-5734 to treat MERS-CoV and related emerging CoV										
1	R01AI132178	-01			BARIC, RALPH S SHEAHAN, TIMOTHY PATRICK	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID	NIAID	\$1,455,240
Systems Immunogenetics of Emerging Coronavirus Infections in the Collaborative Cross										
2	U19AI100625	-06	7727		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID		\$418,635
Role of Uncharacterized Genes in High Pathogenic Human Coronavirus Infection										
5	U19AI107810	-05	8683		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID		\$396,549
Characterization of novel genes encoded ty RNA and DNA viruses										
5	U19AI107810	-05			BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2017	NIAID	NIAID	\$2,021,134
Administrative and Education Core										
4	U19AI100625	-05	6159		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID		\$649,146
Administrative Core										
4	U19AI107810	-04	8688		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID		\$252,483
System Immunogenetics of SARS-CoV Infection										
3	U19AI100625	-04S1	5232		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID		\$135,110
System Immunogenetics of SARS-CoV Infection										
4	U19AI100625	-05	6155		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID		\$739,947
Mechanisms of MERS-CoV Entry, Cross-species Transmission and Pathogenesis										
5	R01AI110700	-02			BARIC, RALPH S LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID	NIAID	\$739,162
Diagnostic and Prognostic Biomarkers for Viral Severe Lung Disease										
5	U19AI109761	-03	8481		BARIC, RALPH S	COLUMBIA UNIVERSITY HEALTH SCIENCES	2016	NIAID		\$889,034
Systems Immunogenetics of Biodefense Pathogens in the Collaborative Cross										
4	U19AI100625	-05			BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID	NIAID	\$4,853,040
Characterization of novel genes encoded ty RNA and DNA viruses										
4	U19AI107810	-04			BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID	NIAID	\$2,322,470
Unlocking Zika Virus Immune Control and Pathogenesis with the Collaborative Cross										
3	U19AI100625	-04S1			BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID	NIAID	\$135,110
Role of Uncharacterized Genes in High Pathogenic Human Coronavirus Infection										
4	U19AI107810	-04	8683		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2016	NIAID		\$704,107
Determinants of Coronavirus Fidelity in Replication and Pathogenesis										
5	R01AI108197	-05			DENISON, MARK R BARIC, RALPH S	VANDERBILT UNIVERSITY MEDICAL CENTER	2016	NIAID	NIAID	\$548,803

2014-2015 CORONAVIRUS, GOF, ET AL.

RALPH BARIC
(UNC/Vanderbilt/Columbia/)

Act	Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC
Determinants of Coronavirus Fidelity in Replication and Pathogenesis									
	R01AI108197-04			DENISON, MARK R. BARIC, RALPH S.	VANDERBILT UNIVERSITY MEDICAL CENTER	2015	NIAID	NIAID	\$306,487
Administrative Core									
	U19AI107810-03		8688	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID		\$188,494
Administrative and Education Core									
	U19AI100625-04		6159	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID		\$279,874
Diagnostic and Prognostic Biomarkers for Viral Severe Lung Disease									
	U19AI109761-02		8481	BARIC, RALPH S.	COLUMBIA UNIVERSITY HEALTH SCIENCES	2015	NIAID		\$1,137,211
Mechanisms of MERS-CoV Entry, Cross-species Transmission and Pathogenesis									
	R01AI110700-01A1			BARIC, RALPH S. LI, FANG	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID	NIAID	\$754,420
System Immunogenetics of SARS-CoV Infection									
	U19AI100625-04		6155	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID		\$412,965
Systems Immunogenetics of Biodefense Pathogens in the Collaborative Cross									
	U19AI100625-04			BARIC, RALPH S. HEISE, MARK T.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID	NIAID	\$4,144,540
Characterization of novel genes encoded ty RNA and DNA viruses									
	U19AI107810-03			BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID	NIAID	\$2,115,911
Role of Uncharacterized Genes in High Pathogenic Human Coronavirus Infection									
	U19AI107810-03		8683	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2015	NIAID		\$370,377
Determinants of Coronavirus Fidelity in Replication and Pathogenesis									
	R01AI108197-03			DENISON, MARK R. BARIC, RALPH S.	VANDERBILT UNIVERSITY	2015	NIAID	NIAID	\$242,311
Administrative Core									
	U19AI107810-02		8688	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID		\$223,265
Administrative and Education Core									
	U19AI100625-03		6159	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID		\$832,756
Mechanisms of Norovirus Protective Immunity									
	R56AI106006-01A1			BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID	NIAID	\$759,938
System Immunogenetics of SARS-CoV Infection									
	U19AI100625-03		6155	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID		\$334,030
Diagnostic and Prognostic Biomarkers for Viral Severe Lung Disease									
	U19AI109761-01		8481	BARIC, RALPH S.	COLUMBIA UNIVERSITY HEALTH SCIENCES	2014	NIAID		\$1,184,414
Systems Immunogenetics of Biodefense Pathogens in the Collaborative Cross									
	U19AI100625-03			BARIC, RALPH S. HEISE, MARK T.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID	NIAID	\$4,148,261
Role of Uncharacterized Genes in High Pathogenic Human Coronavirus Infection									
	U19AI107810-02		8683	BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID		\$420,831
Characterization of novel genes encoded ty RNA and DNA viruses									
	U19AI107810-02			BARIC, RALPH S.	UNIV OF NORTH CAROLINA CHAPEL HILL	2014	NIAID	NIAID	\$2,102,641
Determinants of Coronavirus Fidelity in Replication and Pathogenesis									
	R01AI108197-02			DENISON, MARK R. BARIC, RALPH S.	VANDERBILT UNIVERSITY	2014	NIAID	NIAID	\$547,101

\$	306,487
\$	188,494
\$	279,874
\$	1,137,211
\$	754,420
\$	412,965
\$	4,144,540
\$	2,115,911
\$	370,377
\$	242,311
\$	233,265
\$	832,756
\$	759,938
\$	334,030
\$	1,184,414
\$	4,148,261
\$	420,831
\$	2,102,641
\$	547,101
\$	20,515,827

T	Act	Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC
Administrative Core										
1	U19AI107810-01		8688		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID		\$196,909
Administrative and Education Core										
5	U19AI100625-02		6159		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID		\$206,337
System Immunogenetics of SARS-CoV Infection										
5	U19AI100625-02		6155		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID		\$367,995
Systems Immunogenetics of Biodefense Pathogens in the Collaborative Cross										
5	U19AI100625-02				BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID	NIAID	\$4,030,980
Systems Pathogenomics of Severe Acute Respiratory Virus Infection										
5	U54AI081680-05		6278		BARIC, RALPH S	OREGON HEALTH & SCIENCE UNIVERSITY	2013	NIAID		\$818,118
Role of Uncharacterized Genes in High Pathogenic Human Coronavirus Infection										
1	U19AI107810-01		8683		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID		\$371,153
Characterization of novel genes encoded ty RNA and DNA viruses										
1	U19AI107810-01				BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID	NIAID	\$2,027,645
Susceptibility and Protective Immunity to Noroviruses										
5	R01AI056351-10				BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2013	NIAID	NIAID	\$409,466
Determinants of Coronavirus Fidelity in Replication and Pathogenesis										
1	R01AI108197-01				DENISON, MARK R. BARIC, RALPH S	VANDERBILT UNIVERSITY	2013	NIAID	NIAID	\$560,000
Administrative and Education Core										
1	U19AI100625-01		6159		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2012	NIAID		\$174,333
SARS-CoV Pathogenic Mechanisms in Senescent Mice										
5	R01AI075297-05				BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2012	NIAID	NIAID	\$484,651
System Immunogenetics of SARS-CoV Infection										
1	U19AI100625-01		6155		BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2012	NIAID		\$435,565
Systems Immunogenetics of Biodefense Pathogens in the Collaborative Cross										
1	U19AI100625-01				BARIC, RALPH S HEISE, MARK T	UNIV OF NORTH CAROLINA CHAPEL HILL	2012	NIAID	NIAID	\$4,594,721
Systems Pathogenomics of Severe Acute Respiratory Virus Infection										
5	U54AI081680-04		6278		BARIC, RALPH S	OREGON HEALTH & SCIENCE UNIVERSITY	2012	NIAID		\$746,369
Susceptibility and Protective Immunity to Noroviruses										
5	R01AI056351-09				BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2012	NIAID	NIAID	\$435,605

2012-2013
CORONAVIRUS, GOF, ET AL.

RALPH BARIC (UNC/Vanderbilt/OR)

\$ 196,909

\$ 206,337

\$ 367,995

\$ 4,030,980

\$ 818,118

\$ 371,153

\$ 2,027,645

\$ 409,466

\$ 560,000

\$ 174,333

\$ 484,651

\$ 435,565

\$ 4,594,721

\$ 746,369

\$ 435,605

\$ 15,859,847

Systems Pathogenomics of Severe Acute Respiratory Virus Infection						
U54AI081680-03 6278	BARIC, RALPH S	OREGON HEALTH & SCIENCE UNIVERSITY	2011	NIAID		\$758,603
Susceptibility and Protective Immunity to Noroviruses						
R01AI056351-08	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2011	NIAID	NIAID	\$457,050
SARS-CoV Pathogenic Mechanisms in Senescent Mice						
R01AI075297-03	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2010	NIAID	NIAID	\$492,533
Systems Pathogenomics of Severe Acute Respiratory Virus Infection						
U54AI081680-02 6278	BARIC, RALPH S	OREGON HEALTH & SCIENCE UNIVERSITY	2010	NIAID		\$739,767
Susceptibility and Protective Immunity to Noroviruses						
R01AI056351-07	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2010	NIAID	NIAID	\$606,993
High-Throughput Based Assays for Chemical Screens Against SARS-CoV Targets						
R21NS063854-01S1	ENGEL, DANIEL A BARIC, RALPH S	UNIVERSITY OF VIRGINIA	2010	NIHNS	OD	\$38,500
Developing Vaccine Candidates for the SARS Coronavirus						
P01AI059443-05	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2009	NIAID	NIAID	\$1,907,553
Core--SARS Coronavirus Clone and Reagent						
P01AI059443-05 9002	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2009	NIAID		\$202,418
Rewiring SARS-CoV Genome Organization and Pathogenesis						
P01AI059443-05 0001	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2009	NIAID		\$264,982
Systems Pathogenomics of Severe Acute Respiratory Virus Infection						
U54AI081680-01 6278	BARIC, RALPH S	OREGON HEALTH & SCIENCE UNIVERSITY	2009	NIAID		\$747,616
SARS-CoV Pathogenic Mechanisms in Senescent Mice						
R01AI075297-02	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2009	NIAID	NIAID	\$490,616
Susceptibility and Protective Immunity to Noroviruses						
R01AI056351-06A1	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2009	NIAID	NIAID	\$660,673
Developing Vaccine Candidates for the SARS Coronavirus						
P01AI059443-04	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2008	NIAID	NIAID	\$1,854,836
Core--SARS Coronavirus Clone and Reagent						
P01AI059443-04 9002	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2008	NIAID		\$196,600
Rewiring SARS-CoV Genome Organization and Pathogenesis						
P01AI059443-04 0001	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2008	NIAID		\$257,344
SARS Reverse Genetics						
R01AI059136-05	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2008	NIAID	NIAID	\$271,608
SARS-CoV Pathogenic Mechanisms in Senescent Mice						
R01AI075297-01A1	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2008	NIAID	NIAID	\$496,151
High-Throughput Based Assays for Chemical Screens Against SARS-CoV Targets						
R21NS063854-01	ENGEL, DANIEL A BARIC, RALPH S	UNIVERSITY OF VIRGINIA	2008	NIHNS	OD	\$157,259
SARS Reverse Genetics						
R01AI059136-04	BARIC, RALPH S	UNIV OF NORTH CAROLINA CHAPEL HILL	2007	NIAID	NIAID	\$276,869

\$ 758,603

\$ 457,050

\$ 492,533

\$ 739,767

\$ 606,993

\$ 38,500

\$ 1,907,553

\$ 202,418

\$ 264,418

\$ 264,982

\$ 747,616

\$ 490,616

\$ 660,673

\$ 1,854,836

\$ 196,600

\$ 257,344

\$ 271,608

\$ 496,151

\$ 157,259

\$ 276,869

\$ 11,142,389

2003 – 2006
SARS-CORONAVIRU,
NOROVIRUS, Et al.
NIH/NIAID GRANTS

RALPH BARIC (UNC)

\$	743
\$	1,840,683
\$	181,677
\$	239,406
\$	337,090
\$	285,138
\$	377,578
\$	146,495
\$	1,676,513
\$	174,680
\$	218,242
\$	395,202
\$	290,054
\$	367,042
\$	345,202
\$	278,647
\$	367,042
\$	384,133
\$	519,733
\$	129,665
\$	8,554,965

T	Act	Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC
5	M01RR000046-	46	1437		BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NCRR		\$743
Developing Vaccine Candidates for the SARS Coronavirus										
5	P01AI059443-	02			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID	NIAID	\$1,840,683
Core--SARS Coronavirus Clone and Reagent										
5	P01AI059443-	02	9002		BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID		\$181,677
Rewiring SARS-CoV Genome Organization and Pathogenesis										
5	P01AI059443-	02	0001		BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID		\$239,406
Studies into the Mechanisms for MHV Replication										
5	R01AI023946-	17			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID	NIAID	\$337,090
SARS Reverse Genetics										
5	R01AI059136-	03			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID	NIAID	\$285,138
Susceptibility and Protective Immunity to Noroviruses										
5	R01AI056351-	04			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2006	NIAID	NIAID	\$377,578
Administrative Core										
1	P01AI059443-	01A17958			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID		\$146,495
Developing Vaccine Candidates for the SARS Coronavirus										
1	P01AI059443-	01A1			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID	NIAID	\$1,676,513
Core--SARS Coronavirus Clone and Reagent										
1	P01AI059443-	01A19002			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID		\$174,680
Rewiring SARS-CoV Genome Organization and Pathogenesis										
1	P01AI059443-	01A10001			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID		\$218,242
Studies into the Mechanisms for MHV Replication										
5	R01AI023946-	16			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID	NIAID	\$395,202
SARS Reverse Genetics										
5	R01AI059136-	02			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID	NIAID	\$290,054
Susceptibility and Protective Immunity to Noroviruses										
5	R01AI056351-	03			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2005	NIAID	NIAID	\$376,502
REVERSE GENETICS WITH A CORONAVIRUS INFECTIOUS CONSTRUCT										
5	R01AI059228-	04			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2004	NIAID	NIAID	\$290,054
Studies into the Mechanisms for MHV Replication										
5	R01AI023946-	15			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2004	NIAID	NIAID	\$345,202
SARS Reverse Genetics										
1	R01AI059136-	01			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2004	NIAID	NIAID	\$278,647
Remodeling SARS Coronavirus Genome Regulatory Networks										
1	R01AI061819-	01			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2004	NIAID	NIAID	\$367,042
Susceptibility and Protective Immunity to Noroviruses										
5	R01AI056351-	02			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2004	NIAID	NIAID	\$384,133
REVERSE GENETICS WITH A CORONAVIRUS INFECTIOUS CONSTRUCT										
5	R01AI059228-	03			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2003	NIAID	NIAID	\$290,054
Studies into the Mechanisms for MHV Replication										
2	R01AI023946-	14A1			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2003	NIAID	NIAID	\$519,733
Susceptibility and Protective Immunity to Noroviruses										
1	R01AI056351-	01			BARIC, RALPH S	UNIVERSITY OF NORTH CAROLINA CHAPEL HILL	2003	NIAID	NIAID	\$129,665

2003 – 2021 TOTAL NIH/NIAID GRANTS RALPH BARIC

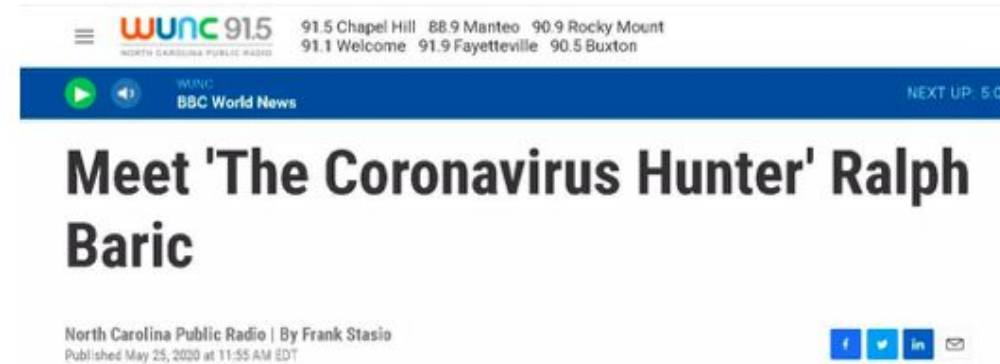
\$	23,082,450
\$	14,226,335
\$	18,345,801
\$	20,515,827
\$	15,859,847
\$	11,142,389
\$	8,554,965
\$	111,727,614



US TAX DOLLARS

Baric and his virus modification technique

Let's start with Ralph Baric, an American scientist who is called the "Coronavirus Hunter."



Screenshot on North Carolina-based media WUNC



Screenshot on UNC

According to [a report](#) from the school-run media of the University of North Carolina, where Baric works, "Baric has been tracking coronaviruses for decades and working on medications to treat coronavirus-caused infections."



UNIVERSITY NEWS

Leading COVID experts to deliver Carolina's Spring Commencement address

Drs. Anthony Fauci and Kizzmekia Corbett have been pivotal in understanding and combatting the virus.

By University Communications, Thursday, March 4th, 2021



T	Act	Project	Year	Sub	Principal Investigator(s)/ Project Leader(s)	Organization	Fiscal Year	Admin IC	Funding IC	FY Total Cost by IC
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-20			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2020	NIAID	NIAID	\$1,001,661
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-20			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2020	NIAID	NIAID	\$1,001,661
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-19			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2019	NIAID	NIAID	\$1,069,830
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-19			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2019	NIAID	NIAID	\$1,069,830
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-18			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2018	NIAID	NIAID	\$984,045
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-18			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2018	NIAID	NIAID	\$984,045
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-17			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2017	NIAID	NIAID	\$1,117,501
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-17			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2017	NIAID	NIAID	\$921,475
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-16			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2016	NIAID	NIAID	\$744,742
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-16			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2016	NIAID	NIAID	\$744,742
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-15			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2015	NIAID	NIAID	\$853,062
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-15			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2015	NIAID	NIAID	\$853,062
		Role of Viral Reservoirs in the Pathogenesis of HIV Disease								
1		ZIAAI000851-15			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2015	NIAID	NIAID	\$1,384,173
		Therapeutic Strategies for the Management of HCV/HIV co- infection								
1		ZIAAI000390-31			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2014	NIAID	NIAID	\$689,091
		Interaction of HIV envelope with cell surface receptors								
1		ZIAAI000883-14			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2014	NIAID	NIAID	\$612,656
		Role of HIV Envelope Proteins In Viral Replication and HIV Pathogenesis								
1		ZIAAI000887-14			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2014	NIAID	NIAID	\$501,264
		Role of Viral Reservoirs in the Pathogenesis of HIV Disease								
1		ZIAAI000851-14			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2014	NIAID	NIAID	\$966,899
		Role of B Lymphocytes In HIV Infection And Pathogenesis								
1		ZIAAI000825-17			FAUCI, ANTHONY S.	NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES	2014	NIAID	NIAID	\$772,682

2014 -2020
NIAID GRANTS
HIV ENVELOPE (GP120)

[ANTHONY FAUCI](#)

\$	1,001,661
\$	1,069,830
\$	984,045
\$	1,175,501
\$	921,475
\$	744,742
\$	853,062
\$	1,384,173
\$	689,091
\$	612,656
\$	501,264
\$	966,899
\$	772,682
\$	11,677,081

https://reporter.nih.gov/search/3_Ch3YpiTEum9wv2qeurbA/projects?PI=2403678

ANTHONY FAUCI - US PATENTS – HIV GP120

Search Results						Export	Share
6 patents							
Search Text: FAUCI, ANTHONY							
Projects Publications Patents Clinical Studies News & More							
Filters							
<input type="checkbox"/> Active Projects							
> Fiscal Years							
> Org Names							
> Agencies							
> Countries							
> Principal Investigators							
Core NIH Project Number		Patent Number	Patent Title	Patent Owner	Primary Agency		
Z01AI000887		9896509	Use of antagonists of the interaction between HIV GP120 and .alpha.4.beta.7 integrin		NIH		
Z01AI000887		9441041	Use of antagonists of the interaction between HIV GP120 and .alpha.4.beta.7 integrin		NIH		
ZIAAI000887		9441041	Use of antagonists of the interaction between HIV GP120 and .alpha.4.beta.7 integrin		NIH		
ZIAAI000887		9896509	Use of antagonists of the interaction between HIV GP120 and .alpha.4.beta.7 integrin		NIH		
Z01AI000887		9193790	Use of antagonists of the interaction between HIV GP120 and A4B7 integrin		NIH		
ZIAAI000887		9193790	Use of antagonists of the interaction between HIV GP120 and A4B7 integrin		NIH		