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New Paradigm for Host and Viral Gene Regulation by MERS Coronavirus nsp1

Project Number Contact PI/Project Leader 5R01AI114657-05 MAKINO, SHINJI

Awardee Organization
UNIVERSITY OF TEXAS
MED BR GALVESTON



Abstract Text

DESCRIPTION (provided by applicant): Middle East respiratory syndrome coronavirus (MCoV) emerged in Saudi Arabia in 2012 and has been disseminated into other countries in the Middle East, North Africa and Europe. Although MERS is most probably a zoonosis, MCoV can spread person-toperson. MCoV infection causes fever, cough and pneumonia leading to respiratory failure, and the reported case fatality is ~50%. Little is known about the mechanisms of MCoV's high virulence and pathogenesis. Coronaviruses (CoVs) carry a single-stranded, positive-sense RNA genome of ~30 kb. Immediately after infection, the translation of the viral genome produces two large polyproteins that are processed into 15 or 16 mature nonstructural proteins (nsp1-nsp16), most of which are involved in viral RNA synthesis, while some have other biological functions. Nsp1 protein of various CoVs inhibits host gene expression. Viral proteins that inhibit host gene expression are often major virulence factors, and, hence, CoV nsp1 proteins most probably play a critical role in CoV pathogenesis. Indeed, mouse hepatitis virus nsp1 is a major virulence factor, and SARS-CoV (SCoV) nsp1 inhibits the production of type I interferon and interferon-stimulated genes in infected cells. Nsp1 proteins of different CoVs use divergent strategies to exert host gene expression suppression; by binding to the 40S ribosomal subunit, SCoV nsp1 inhibits mRNA translation and induces endonucleolytic mRNA cleavage, while nsp1 of transmissible gastroenteritis virus does not induce mRNA cleavage, yet suppresses translation without binding to the 40S ribosomal subunits. Our data that MCoV nsp1 inhibited translation of host mRNAs and promoted mRNA cleavage without binding to the 40S subunits and other experimental results led us to hypothesize that MCoV nsp1 suppresses host gene expression by using mechanisms that have not been described in any viral proteins. This application aims to delineate the mechanisms of MCoV nsp1-induced translation inhibition and mRNA cleavage. We will also clarify the strategy that allows robust viral gene expression in MCoVinfected cells under conditions of nsp1- induced translation inhibition. The proposed studies will provide a foundation for understanding the modulation of host gene expression by MCoV and expand our knowledge of viral pathogenicity at the molecular level.

Public Health Relevance Statement

PUBLIC HEALTH RELEVANCE: Middle East respiratory syndrome coronavirus (MCoV) emerged in Saudi Arabia in 2012 and has been disseminated into other countries in the Middle East, North Africa and Europe. Because MCoV can spread person-to-person, it is an area of international public health concern. To understand MCoV pathogenicity at the molecular level, we will study how nsp1 protein, one of MCoV proteins, inhibits host gene expression and exerts robust viral gene expression in infected cells, where nsp1 efficiently suppresses host gene expression.

NIH Spending Category

Biodefense Biotechnology Emerging Infectious Diseases Genetics

Infectious Diseases Lung Pneumonia Pneumonia & Influenza Rare Diseases

Project Terms

5' Untranslated Regions **Acute Kidney Failure Animals** Binding Area **Biological Process Camels Case Fatality Rates** Case Study **Cell Nucleus** Cells Chiroptera Chronic Complex Coronavirus Coughing Country 11/27/21, 5:14 AM RePORT) RePORTER

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Project Number Contact PI/Project Leader 5R01AI114657-05 MAKINO, SHINJI

Awardee Organization UNIVERSITY OF TEXAS MED BR GALVESTON

Congressional District

Project End

Budget Start

Date

Date

30-April-

01-May-

2021

2021

Read More

Details

Contact PI/ Project Other PIs Program Official

Leader Not Applicable Name

Name STEMMY, ERIK J

MAKINO, SHINJI

Contact

erik.stemmy@nih.gov

PROFESSOR Contact

shmakino@utmb.edu

Organization

Name Department Type State Code

UNIVERSITY OF TEXAS MED MICROBIOLOGY/IMMUN/VIROL(TX

BR GALVESTON Organization Type

City SCHOOLS OF MEDICINE 14
GALVESTON

Country
UNITED STATES (US)

Other Information

FOA Administering Institutes or Project Start 01-MayPA-13-302 Centers Date 2015

Study Section

Virology - A Study
Section[VIRA]

NATIONAL INSTITUTE OF
ALLERGY AND INFECTIOUS
DISEASES

Award Notice DUNS Number CFDA Code

Fiscal Year Date 800771149 855 Date 2019

2019 12-April-2019 Budget End 30-April-

Project Funding Information for 2019

Total Funding Direct Costs Indirect Costs \$387,500 \$250,000 \$137,500

Year Funding IC

2019 NATIONAL INSTITUTE OF ALLERGY AND INFECTIOUS DISEASES \$387,500

NIH Categorical Spending Click here for more information on NIH Categorical Spending

Funding IC FY Total Cost by IC NIH Spending Category

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Project Number Contact PI/Project Leader 5R01AI114657-05 **MAKINO, SHINJI**

Awardee Organization **UNIVERSITY OF TEXAS MED BR GALVESTON**

> intectious Diseases; Lung; Pneumonia; Pneumonia & Influenza; Rare Diseases;

品 Sub Projects

No Sub Projects information available for 5R01Al1114657-05



Export

Simila

Public

Publication

Year

Characterization of the Molecular Interactions That Govern the Packaging of Viral RNA **Segments into Rift Valley Fever Phlebovirus Particles.**

Authors

2021

Journal of virology 2021 06 24; 95 (14) e0042921

Journal (Link to PubMed abstract)

Tercero, Breanna;

Narayanan,

Krishna; Terasaki, Kaori; Makino,

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Mechanisms of Coronavirus Nsp1-Mediated Control of Host and Viral Gene Expression.

Cells 2021 02 02; 10 (2)

Nakagawa, Keisuke; Makino, Shinji

Shinji

2021

2020

2020

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An Infectious cDNA Clone of SARS-CoV-2.

Cell host & microbe 2020 05 13; 27 Xie, Xuping; (5) 841-848.e3

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iCite 44.63

Reverse genetics approaches for the development of bunyavirus vaccines.

ΑII

Current opinion in virology 2020

<u>10; 44 16-25</u>

Tercero, Breanna; Makino,

Shinji

MG

<u>i</u> G i Cite 1.04

Severe Acute Respiratory Syndrome Coronavirus 2 from Patient with Coronavirus Disease, **United States.**

Emerging infectious diseases 2020 Harcourt, <u>06; 26 (6) 1266-1273</u>

Jennifer: View

All

2020

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iCite 56.15

A nanoluciferase SARS-CoV-2 for rapid neutralization testing and screening of antiinfective drugs for COVID-19.

Thank you for your feedback!

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Project Number

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Contact PI/Project Leader MAKINO, SHINJI

Awardee Organization
UNIVERSITY OF TEXAS
MED BR GALVESTON



The Project Outcomes shown here are displayed verbatim as submitted by the Principal Investigator (PI) for this award. Any opinions, findings, and conclusions or recommendations expressed are those of the PI and do not necessarily reflect the views of the National Institutes of Health. NIH has not endorsed the content below.

No Outcomes available for 5R01Al1114657-05

Clinical Studies

No Clinical Studies information available for 5R01Al1114657-05

News and More

Related News Releases

No news release information available for 5R01AI114657-05

History

No Historical information available for 5R01Al1114657-05

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No Similar Projects information available for 5R01Al114657-05