Technical Journal Club

30 June, 2020

Anna Henzi, MD PhD student



Synthesizing viruses

 Viral isolates for development of diagnostics, therapeutics and vaccines, or in vivo models

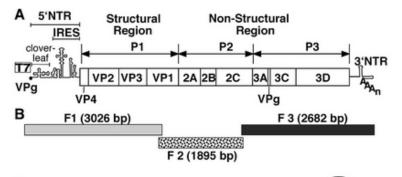
Synthesizing viruses

Chemical Synthesis of Poliovirus cDNA: Generation of Infectious Virus in the Absence of Natural template

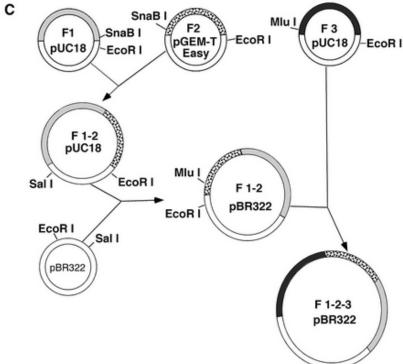
Poliovirus: small, nonenveloped, + ssRNA genome (7.5 kb)

One of the simplest genetic systems known for proliferation

Synthesizing viruses



 Use of known sequence as the only instruction for engineering the genome



- Phage T7 promoter for RNA polymerase > transcription into infectious RNA
- Generation of virus in cellfree cytoplasmic extracts

How to make recombinant Coronaviruses

- Thiel, J Gen Virol, 2001
- Coley, J Virol, 2005
- Thao, Nature, 2020

Coronaviruses

- enveloped, + stranded RNA viruses
- large genomes: 27-31 kb
- Gene sequences toxic to E.coli
- > not possible to clone into plasmid vectors or passage in procaryotic systems
- Bacterial artificial chromosomes or long range RT-PCR
- Alternative: optimization of in vitro DNA ligation, vaccinia virus eukaryotic cloning vector, cytoplasmic expression of transfected RNA

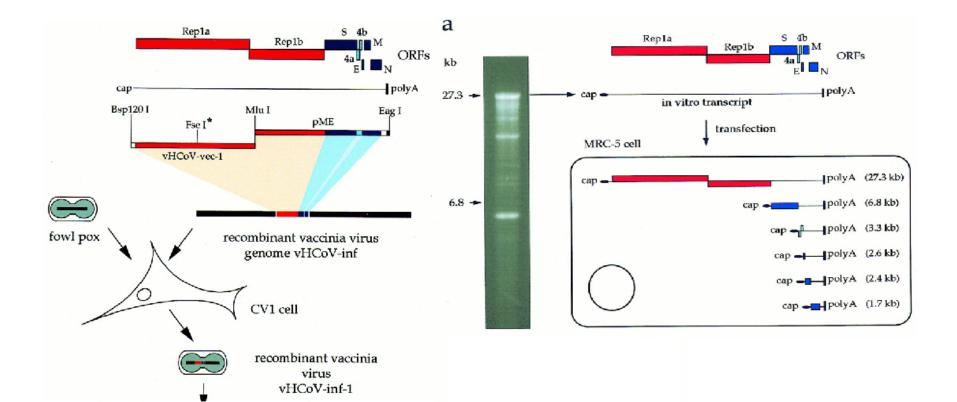
Infectious RNA transcribed in vitro from a cDNA copy of the human coronavirus genome cloned in vaccinia virus

Volker Thiel, Jens Herold,† Barbara Schelle and Stuart G. Siddell

Institute of Virology and Immunology, University of Würzburg, Versbacher Straße 7, 97078 Würzburg, Germany

- HCoV229E
- Vaccinia virus (poxvirus)
 - eukaryotic cloning vector

- High stability of cloned insert cDNA
- Retain infectivity, growth kinetics,
 stability regardless of insert size
- Large cDNA fragments can be cloned efficiently into vaccinia virus genome



Recombinant Mouse Hepatitis Virus Strain A59 from Cloned, Full-Length cDNA Replicates to High Titers In Vitro and Is Fully Pathogenic In Vivo

Scott E. Coley, Ehud Lavi, Stanley G. Sawicki, Li Fu, Barbara Schelle, Nadja Karl, Stuart G. Siddell, Volker Thiel

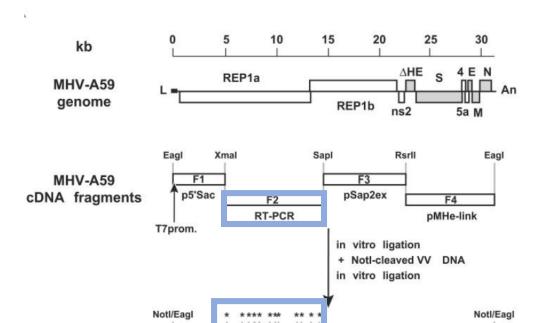
DOI: 10.1128/JVI.79.5.3097-3106.2005



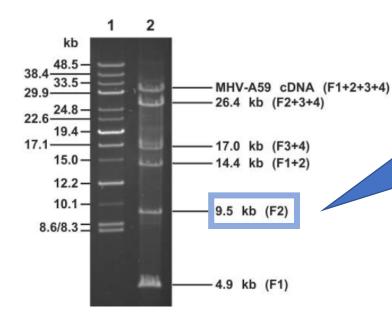
Vaccinia virus

MHV strain A59

- Versatile, universal tool to study all aspects of MHV molecular biology and pathogenesis
- Facilitation of mutagenesis
- Rescue of recombinant coronaviruses by nucleocapsid protein-expressing cell line



full-length MHV-A59 cDNA



vMHV-E6

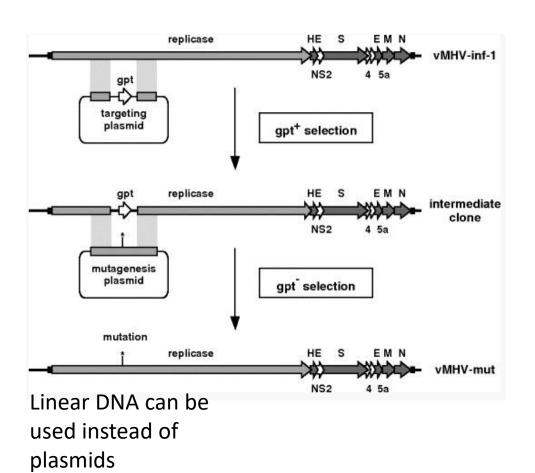
VV 1

T7prom.

Replaced with correct sequence in 4 rounds of virus-mediated recombination (with + and – selection)

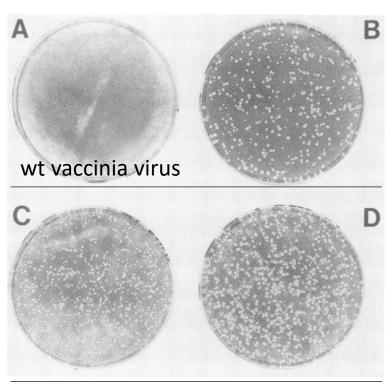
VV

Modification of Coronavirus cDNA by Vaccinia Virus-Mediated Homologous Recombination



SARS and other Coronaviruses, Eriksson, 2008;

Falkner, J Virol, 1988



Recombinant vaccinia virus (gpt+)

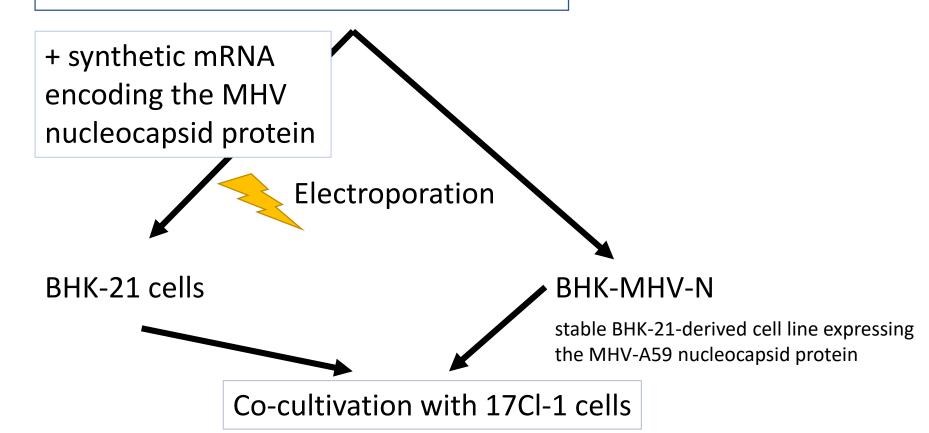
+MPA +(hypo) xanthine No treatment

Rescue of recombinant MHV-inf-1

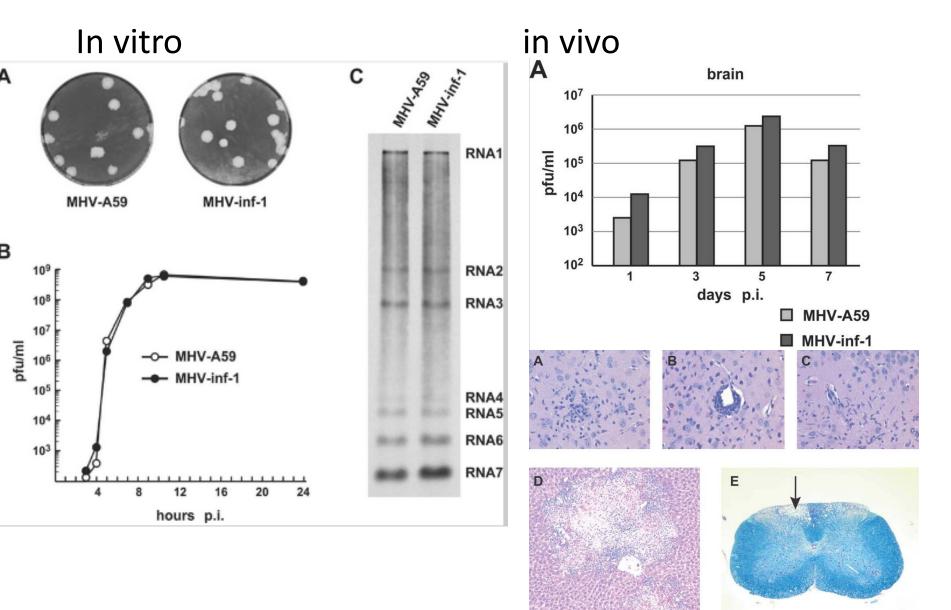
 «rescue of recombinant coronaviruses is greatly facilitated under conditions that allow N protein expression"

Rescue of recombinant MHV-inf-1

- Full length MHV RNA
 - in vitro transcription of DNA from
 - purified virions
 - Infected cells



Replication of recombinant MHV-inf-1



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DOI: 10.1128/JVI.79.5.3097-3106.2005



- Plasmid DNA and RT-PCR derived products for vacciniavirus mediated homologous recombination
- Infectious virus produced from 6-well dish
- Cell line expressing N protein (BHK-MHV-N) facilitates rescue
- MHV system to study similar coronaviruses like SARS-CoV

Reverse genetics platform

- Feline coronaviruses (Tekes, J Virol, 2008, 2012)
- SARS-CoV (van den Worm, PloS One, 2012)

Accelerated Article Preview

Rapid reconstruction of SARS-CoV-2 using a synthetic genomics platform

Received: 20 February 2020

Accepted: 24 April 2020

Accelerated Article Preview Published online 4 May 2020

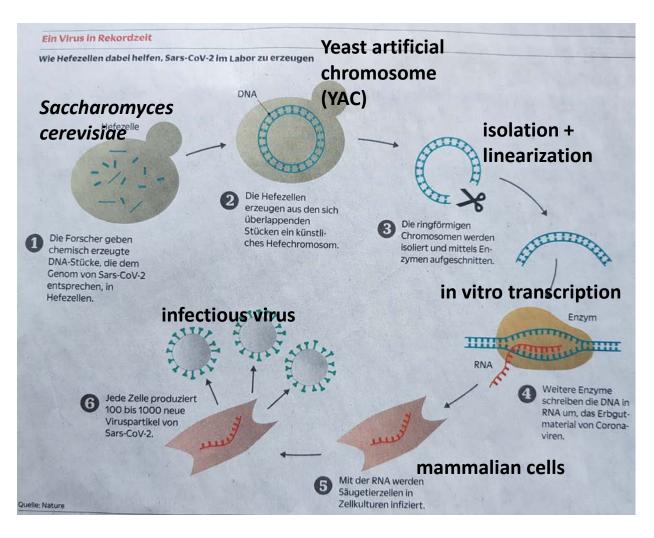
Cite this article as: Thao, T. T. N. et al. Rapid reconstruction of SARS-CoV-2 using a synthetic genomics platform. *Nature* https://doi.org/10.1038/s41586-020-2294-9 (2020).

Tran Thi Nhu Thao, Fabien Labroussaa, Nadine Ebert, Philip V'kovski, Hanspeter Stalder, Jasmine Portmann, Jenna Kelly, Silvio Steiner, Melle Holwerda, Annika Kratzel, Mitra Gultom, Kimberly Schmied, Laura Laloli, Linda Hüsser, Manon Wider, Stephanie Pfaender, Dagny Hirt, Valentina Cippà, Silvia Crespo-Pomar, Simon Schröder, Doreen Muth, Daniela Niemeyer, Victor Corman, Marcel A. Müller, Christian Drosten, Ronald Dijkman, Joerg Jores & Volker Thiel

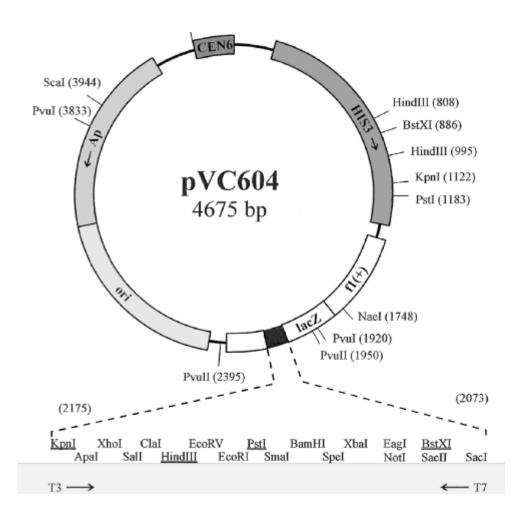
This is a PDF file of a peer-reviewed paper that has been accepted for publication.

Although unedited, the content has been subjected to preliminary formatting. Nature is providing this early version of the type set penetres a service to currenthers and

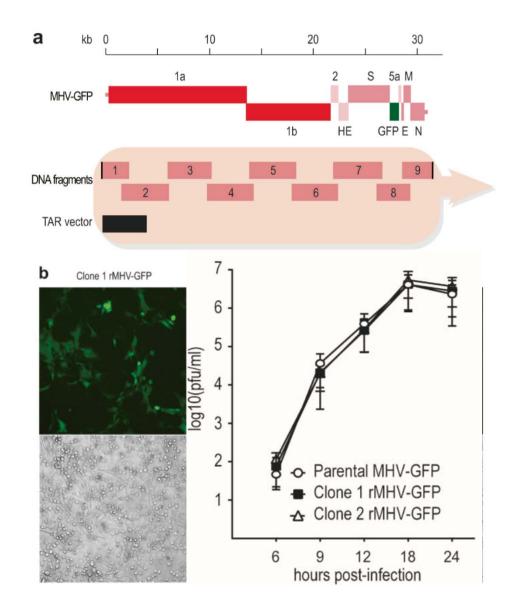
Transformation-associated recombination (TAR) cloning



TAR vector (Kouprina et al.)



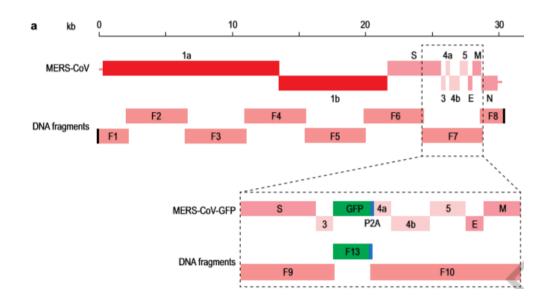
Test platform with MHV-GFP

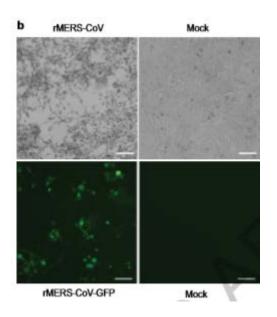


Virus rescue:

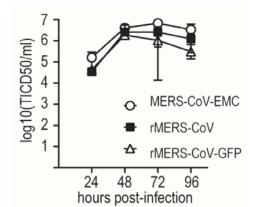
- 1) electroporation of BHK-MHV-N cells
- 2) co-cultivation with susceptible cells(17Cl-1)

Application to other coronaviruses + mutagenesis





Synthetic genomics platform suitable to genetically modify coronavirus genomes.

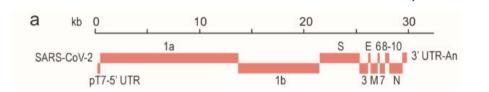


Application to other viruses, families and sources

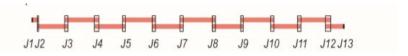
- Several other coronaviruses
- ZIKA virus (Flaviviridae)
- hRSV
- Cloning of nRSV-B from clinical sample (nasopharyngeal aspirate) without any prior information about the virus genotype using RSV consensus primers
- Synthetic genomics platform proviced the technical advance to rapidly generate molecular clones of diverse RNA viruses by using virus isolates, cloned DNA, synthetic DNA or clinical samples as starting material.

SARS-CoV-2

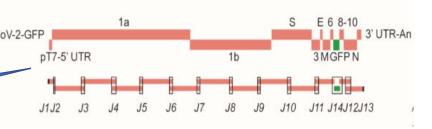
 Genome sequences released 10.01.2020



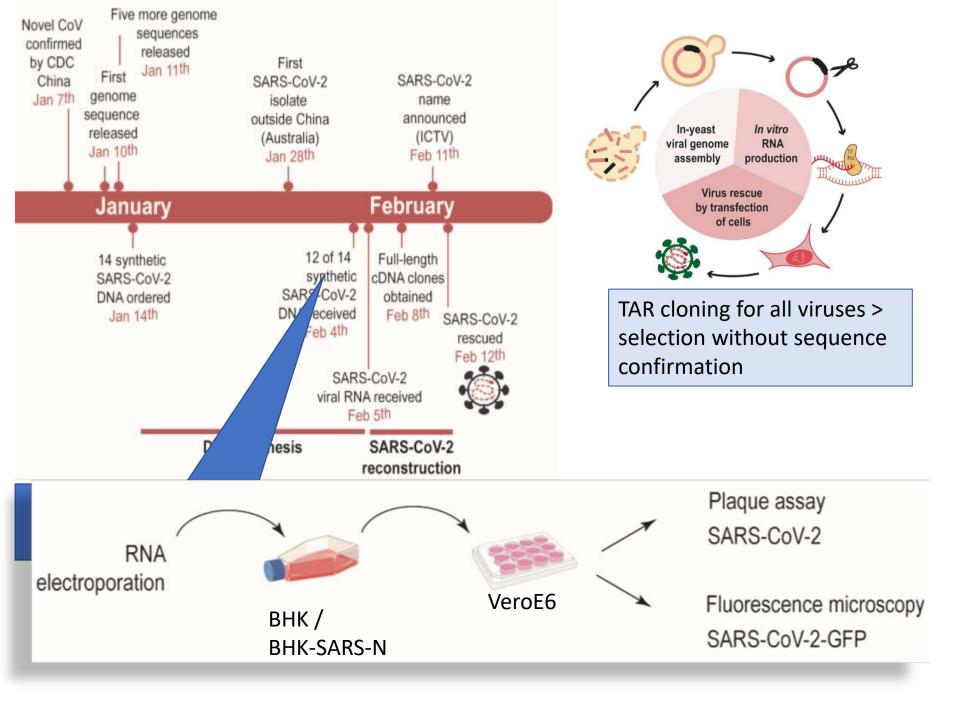
12 overlapping DNA fragments



In parallel: SARS-CoV-2-GFP SARS-CoV-2-GFP



Variations in nt 3-5 > generate 3 versions +/- GFP



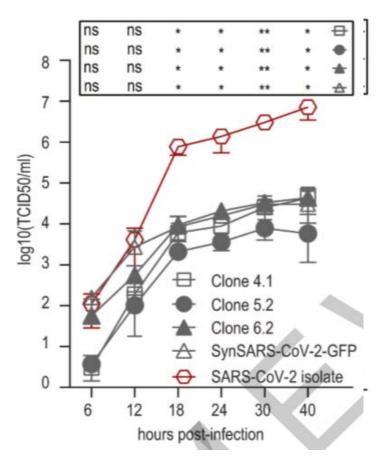
Analysis of rSARS-CoV-2

- All cloned 5' ends retained
- similar replication kinetics

Replication affected by GFP and/or partial deletion of ORF

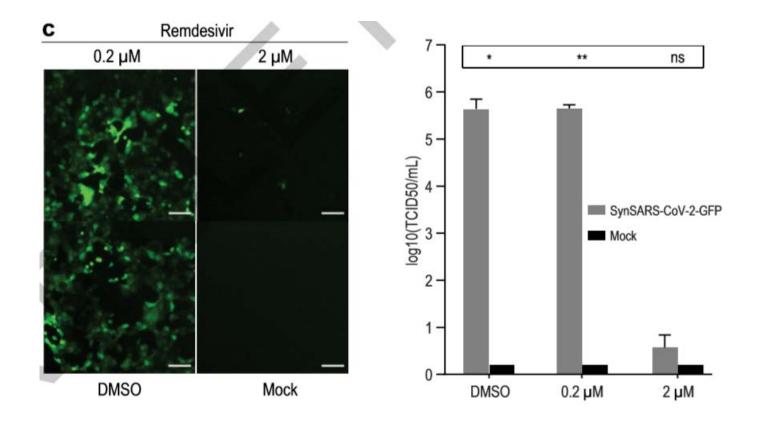
rSARS-CoV-2-GFP





Analysis of rSARS-CoV-2

Simple GFP readout: Remdesivir



Analysis of rSARS-CoV-2

Simple GFP readout: serum

Mock

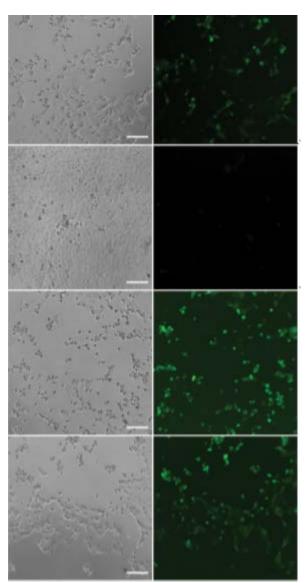
No serum

Convalescent Human anti-SARS-CoV-2

Ctrl

ctrl

Convalescent Human anti-SARS-CoV



Synthetic genomics platform

- Provide infectious virus to health authorities and diagnostic laboratories without the need of having access to clinical samples
- Rapid introduction of emerging sequence variations into infectious clones > functional characterization

Ethical concerns

- Synthesis of viruses
- Synthetic life

«If the ability to replicate is an attribute of life, then poliovirus

is a chemical $(C_{332,652}H_{492,388}O_{131,196}-P_{7501}S_{2340})$ with a life cycle.»

Cello, Science, 2002

Thank you for your attention!

