

## Rapid detection of viruses of concern with Blazar<sup>®</sup> Platform - In line with ICH Q5A(R2)

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The life science business of Merck operates as MilliporeSigma in the U.S. and Canada.

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## Viral safety assurance requires risk mitigation at a number of stages



# Traditional cell line characterization tests for ~20 specific viruses and MERCK takes 45+ days



- Mice, hamsters and/or rats inoculated
- Animal observations for ~28 days
- LCMV challenge
- Animals euthanized and samples taken
- Antibody production measured immunologically – e.g. ELISA

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# Adventitious agent testing is often a rate-limiting step in downstream processing



For CHO-based processes, the industry shift towards intensified and continuous manufacturing increases the need for rapid testing methods to detect:

- Bacteria/fungi
- Mycoplasma
- Adventitious viruses

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Endogenous
retrovirus-like
particles

#### Alternatives to culture-based testing approaches are required

- Traditionally, most testing at the bulk harvest stage is performed using culture-based methods
- Alternative methods should provide breadth of detection and sensitivity, as well as speed

### Molecular methods can accelerate virus testing



# The Blazar<sup>®</sup> platform workflow uses a 3-step method for virus detection







#### Extraction

- Nucleic acid extraction from total bulk harvest material (cells and supernatant)
- Samples are pre-spiked with control viruses to confirm that extraction and PCR perform as expected

#### **Degenerate PCR**

- Two rounds of PCR (first round and nested PCR) to increase sensitivity and specificity
- Separate assays for DNA and RNA viruses (with preceding reverse transcriptase step for RNA viruses)

#### **Fragment analysis**

 Separation by capillary electrophoresis and automated analysis of peaks in relation to expected size ranges

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## Use of degenerate primers provides broad specificity





#### **1997 ICH Q5(A) guidelines**

MAP	HAP	RAP
Ectromelia Virus <sup>2,3</sup>	Lymphocytic Choriomeningitis Virus (LCM) <sup>1,3</sup>	Hantaan Virus <sup>1,3</sup>
Hantaan Virus <sup>1,3</sup>	Pneumonia Virus of Mice (PVM) <sup>2,3</sup>	Kilham Rat Virus (KRV) <sup>2,3</sup>
K Virus <sup>2</sup>	Reovirus Type 3 (Reo3) <sup>1,3</sup>	Mouse Encephalomyelitis Virus (Theilers, GDVII) <sup>2</sup>
Lactic Dehydrogenase Virus (LDM) <sup>1,3</sup>	Sendai Virus <sup>1,3</sup>	Pneumonia Virus of Mice (PVM) <sup>2,3</sup>
Lymphocytic Choriomeningitis Virus (LCM) <sup>1,3</sup>	SV5	Rat Coronavirus (RCV) <sup>2</sup>
Minute Virus of Mice <sup>2,3</sup>		Reovirus Type 3 (Reo3) <sup>1,3</sup>
Mouse Adenovirus (MAV) <sup>2,3</sup>		Sendai Virus <sup>1,3</sup>
Mouse Cytomegalovirus (MCMV) <sup>2,3</sup>		Sialoacryoadenitis Virus (SDAV) <sup>2</sup>
Mouse Encephalomyelitis Virus (Theilers, GDVII) <sup>2</sup>		Toolan Virus (HI) <sup>2,3</sup>
Mouse Hepatitis Virus (MHV) <sup>2</sup>		
Mouse Rotavirus (EDIM) <sup>2,3</sup>		
Pneumonia Virus of Mice (PVM) <sup>2,3</sup>		
Polyoma Virus <sup>2</sup>		
Reovirus Type 3 (Reo3) <sup>1,3</sup> Sendai Virus <sup>1,3</sup>		
Thymic Virus <sup>2</sup>		

#### **Blazar<sup>®</sup> Rodent virus panel**



\* DNA Viruses from ICH List \* RNA Viruses from ICH List

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# Accelerating CHO bulk harvest testing requires a full set of rapid methods



# Accelerating CHO bulk harvest testing requires a full set of rapid methods



Timelines are shown as minimum total assay turnaround time

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## **Blazar® CHO AOF panel targets 15 virus families**



Internal experts, industry consultants

Consideration of historical CHO cell contamination events

Focus on viruses for which CHO cells are permissive, and/or which may have potential to infect human cells

Emerging viruses included to cover future risks

Virus family	Туре	Example viruses covered by CHO AOF panel	
Adenoviridae	dsDNA	human adenovirus A/B/C/D/E/F/G	
Anelloviridae	ssDNA	rat torque teno virus-1/2	
Circoviridae	ssDNA	rat circovirus, porcine circovirus-1/2/3	
Parvoviridae	ssDNA	minute virus of mice, hamster parvovirus, rodent protoparvovirus, rat parvovirus 2, rat bocavirus	
Polyomaviridae	dsDNA	rat polyomavirus 2	
Bornaviridae	ssRNA	borna virus	
Caliciviridae	ssRNA	vesivirus 2117, rat norovirus, mouse norovirus, Calicivirus-Allston-2008	
Coronaviridae	ssRNA	mouse hepatitis virus	
Hepeviridae	ssRNA	hepatitis E virus	
Orthomyxoviridae	ssRNA	influenza virus A/B	
Paramyxoviridae	ssRNA	beilong virus, Human respirovirus 1 strain, Human parainfluenza virus types 1/2/3/4/5, Measles virus, Mumps	
Picornaviridae	ssRNA	coxsackie virus B3, Encephalomyocarditis virus	
Reoviridae	dsRNA	reovirus-1/2/3, Mammalian orthoreovirus, Epizootic hemorrhagic disease virus	
Rhabdoviridae	ssRNA	vesicular stomatitis virus	
Togaviridae	ssRNA	semliki forest virus, Chikungunya virus strain, Eastern equine encephalitis virus	



## Degenerate PCR enables detection of emerging viruses

#### **Multiple variants detected**

- For each virus family, degenerate PCR primers are designed against genomic regions representing conserved protein motifs
- This provides broad specificity, allowing multiple family members and related variants to be detected
- The Blazar<sup>®</sup> CHO AOF RNA panel is predicted to detect >26,900 viral target sequences

#### **Proof of concept**

- A coronavirus primer design has the ability to detect the emergent human SARS-CoV-2 virus
- The Blazar<sup>®</sup> CHO AOF primers were designed in 2018, more than one year before SARS-CoV-2 was described

#### Nature article



https://doi.org/10.1038/s41586-020-2008-3 Received: 7 January 2020 Accepted: 28 January 2020 Published online: 3 February 2020 Open access

8-3 Fan Wu<sup>17</sup>, Su Zhao<sup>27</sup>, Bin Yu<sup>37</sup>, Yan-Mei Chen<sup>17</sup>, Wen Wang<sup>47</sup>, Zhi-Gang Song<sup>17</sup>, Yi Hu<sup>27</sup>,
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Yi Liu<sup>1</sup>, Qi-Min Wang<sup>1</sup>, Jiao-Jiao Zheng<sup>1</sup>, Lin Xu<sup>1</sup>, Edward C. Holmes<sup>1.5</sup> & Yong-Zhen Zhang<sup>1,4,6</sup>

Emerging infectious diseases, such as severe acute respiratory syndrome (SARS) and Zika virus disease, present a maior threat to public health<sup>1-3</sup>. Despite intense research





The Blazar<sup>®</sup> platform mitigates the risk of emerging viruses, which pose a major threat to biologics manufacture

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## **Blazar® CHO AOF panel specifications**

Assay code	399003GMP.BSV (Rockville, US) Available 399003GMP.BUK (Glasgow, UK) Coming soon!	
Sample format	2x1ml of test article	
Total turnaround time	12 days	
Virus coverage	DNA and RNA viruses from 15 virus families	
Internal controls	Spike recovery: DNA and RNA virus each spiked at detection limit in the test article prior to extraction, to demonstrate extraction and PCR efficiency	
Sensitivity	10 genomic copies per reaction*	
False positive rate	<1%	
True positive rate	>99%	
Specificity (pass criteria)	No target peaks observed in 3/3 test article wells Spike recovery control is detected in $\geq 1/3$ wells	
System suitability	No Template Control (NTC) signal detected in 0/3 wells Spike recovery must fall within specified sizing window	

\* 100 genomic copies per reaction for polyomaviruses

Suitable for CHO bulk harvest from animal origin-free processes, where the MCB has been fully characterized



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## Building on our success...

Our established Blazar<sup>®</sup> rodent panel, which uses the same award-winning technology as the CHO AOF panel, is now familiar to regulators through IND submissions and is used by the majority of our clients for cell line characterization

#### ... in line with updated regulatory guidance

The ICH Q5A revision will promote molecular methods for virus detection



### ICH Q5A Revision Addresses Important regulatory and industry Advances



#### Emerging product types

Virus-like particles (VLPs), subunit proteins, and viralvectored vaccines and gene therapies using novel mammalian and insectbased vector/cell expression systems



## Analytical technologies

Nucleic acid-based assays such as PCR and NGS may provide rapid and sensitive detection of adventitious and endogenous viruses in the starting and harvest materials



Virus clearance

validation strategies

Flexibility in validation

approaches should be

allowed in order to

effectively leverage

knowledge gained during

development



#### Manufacturing

Emerging or advanced manufacturing approaches beyond traditional unit and batch process operations



### General guidance on the use of molecular methods ICH Q5A R2 draft (Oct 2022)



#### 3Rs

Virus-specific PCR, targeted molecular methods or NGS can be used as replacement assays for animal-based methods, No headto-head comparison

#### **Infectious agents**

Positive results should be investigated to determine infectivity



Assays should be appropriately qualified or validated for their intended use.

#### **Types**

NGS and Nucleic Acid Amplification Techniques such as PCR may be appropriate for broad and specific virus detection, respectively

#### Comparability

These tests may be introduced without a systematic head-tohead comparison with the currently recommended in vivo assays



### **Summary: Blazar® CHO AOF panel**



Rapid and sensitive virus detection for effective risk mitigation in CHO bulk harvest and other applications



#### Shorten timelines to release

- **12-day** turnaround for CHO bulk harvest samples to replace and/or supplement *in vitro* assay results
- Combine with other rapid methods for an accelerated testing package for animal origin-free processes

#### **Meet regulatory compliance**

- Molecular assays are acceptable to regulators and reflect industry trends towards more robust and sensitive methods
- Established Blazar® technology provides the expected sensitivity and breadth of detection
- Reduce need for animal-derived assay components

#### **Reduce sample requirements**

Assay requires only 2 x 1 ml bulk harvest

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