Design and validation of highthroughput real-time PCR systems for detection of porcine and bovine respiratory and enteric pathogens using the BioMark HD (Fluidigm) platform

The 4th AVDC-China

Nicole B. Goecke Postdoc



UNIVERSITY OF COPENHAGEN



The idea behind the use of high-throughput real-time PCR

- The Danish National Veterinary Institute
 - Large diagnostic unit
 - Main focus on diseases of domestic animals
 - Identification of cause of disease and pathogens
 - Surveillance of avian and swine influenza A viruses











The diagnostic unit – the flow



The used low-throughput real-time PCR platform





Rotor-Gene Q

- Number of samples: 36, 72, 100
- PCR assay per sample:
 - 1 (singleplex) \rightarrow most test
 - 6 (multiplex) \rightarrow few test

Challenges with the low-throughput real-time PCR platform

- Limitation of the platform capacity
- Expensive analyses
 - High analysis cost per sample
 - Many farmers keep the sample submissions to a minimum
- The problem with few sample submissions:
 - Fewer animals get diagnosed
 - Lack of laboratory analysis
 - The disease causing pathogen is unknown
 - Medication and vaccination is based on clinical observation
 - Risk of overuse of antibiotics
 - Risk of suboptimal vaccine programs



06-03-2023 6

The change of real-time PCR platform

- Development of high-throughput real-time PCR systems used for detection and typing of pathogens (viruses and bacteria)
 - Pathogens which have importance to the health and welfare of Danish pigs
- Purpose
 - Reduce the analysis cost per sample and working hours

Low-throughput real-time PCR platform
High-throughput real-time PCR platform

Rotor-Gene Q

BioMark

The high-throughput real-time PCR platform

- High-throughput real-time PCR platform BioMark HD (Fluidigm)
 - Requires less sample and reagents (nL vs. μL)
 - Providing up to 9,216 reactions in one chip (96x96)



48 samples x 48 assays



96 samples x 96 assays

TEFFF	10000	111111
FFFFFF	0	111111
FFFFFF		111111
FFFFFF		111111
FFFFFF		333333
TREFFE	A DESCRIPTION OF TAXABLE PARTY.	333333
TEFFFE		333333
SECCEC.		233333
EEEEEE	and the second se	333333
ELECEE		333333
LEEEEE		SECCE
LEELEE		
ELLELL		333333
LALLE	and a second second second	333344
LALLE	0	333344
A A A A A A		

192 samples x 24 assays



24 samples x 192 assays

a,			17
10	erererer O	111111	111 1
e.	erererer .	1 999999	111 1
e	******	333	111 1
e	ererer	333	1 7 7 1
e	ererer .	3.3.3	111 1
e.	e e e e e e e	333	333 3
¢	CCCCCC .	3.3.7	
¢	CCCCCC	222	
¢.	CCCCCC .	222	
8	CECEEE	333	3333
¢	EEEEEE	333	3333
e.	CLLLLL	333	3333
e.	LLLLL	333	1333
e.	LLLLLL		
ē.	LLLLLUULU m	eeccc	
ē.	LLLLLLLL O		
2	I LLC CONTRACTOR		L.F.



8





~ ~ ~

11111 111111 111111

Assay 3

0

e cererere

\$ \$\$\$\$\$¢¢¢



-

A DESCRIPTION OF

11 A 1

The Workflow



The first porcine high-throughput real-time PCR system

Swine influenza A virus (swIAV)

• Establishment of a high-throughput system for detection and subtyping of swIAVs





Subtyping of swine influenza A viruses

- Currently 18 hemagglutinin (HA) and 11 neuraminidase (NA) genes identified
- Different variants of the HA and NA genes exist
 - H1 lineage
 - H1-human like
 - H1-avian like
 - H1-pandemic 2009
- Subtypes **H1N1**, **H1N2** and **H3N2** are commonly found in pigs



06/03/2023 13

Subtyping of swine influenza A viruses

- A passive surveillance program for swIAVs has been conducted in Denmark since 2011
 - The aim is to identify which subtypes circulate in Danish pigs
 - Be able to act fast on new virus variants

- Subtyping of swIAVs
 - Subtyping is conducted in a multiplex setup
 - Several PCR assays are needed to cover the wide range of circulating subtypes
 - High analysis cost \rightarrow A limited number of samples can be subtyped

Subtyping of swine influenza A viruses

• Change of real-time PCR platform



A more detailed characterization

Internal genes:

- PB2
- PB1
- PA
- NP
- M
- NS



The second porcine high-throughput real-time PCR system

Development of a porcine high-throughput system

- Used for detecting of 18 different porcine viruses and bacteria
- All the relevant pathogens causing respiratory and intestinal diseases in pigs are included
 - Reduce the risk of not detecting the relevant pathogen(s)





The use of the porcine high-throughput real-time PCR system

- Test the performance of the high-throughput diagnostic system on field samples
 - Samples collected from ten Danish pig production units (6 herds)
- Investigate the connection between prevalence of different pathogens and clinical signs observed
- Test if monthly monitoring of pathogens on herd level could be a supportive tool for the veterinarians
 - Create a more objective basis for intervention

(2020) 6:23 F	orcine Health Management
	Open Access
ogen monitoring in Is by monthly labor ng	nursery atory
D I I	gen monitoring in s by monthly labor



SOS "Swine, Objective Surveillance"

The concept behind SOS:

- Herds are sampled at regular intervals throughout the year
- Common guidelines for all herds
- The farmer is responsible for the sampling









The pathogens in **S**

Oral fluid samples:

- Influenza A virus
 - Pandemic H1 (H1pdm)
- Porcine Circovirus Type 2 (PCV2)
- Porcine Circovirus Type 3 (PCV3)
- Porcine Cytomegalovirus (PCMV)
- Streptococcus suis type 2
- Haemophilus parasuis
- Pastuerella multocida
- Actinobacillus pleuropneumoniae
- Mycoplasma hyopneumonia
- Mycoplasma hyorhinis
- Bordetella bronchiseptica

Sock samples:

- Escherichia coli F4
- Escherichia coli F18
- Lawsonia intracellularis
- Brachyspira pilosicoli
- Rotavirus A
- Porcine Circovirus Type 2 (PCV2)
- Porcine Circovirus Type 3 (PCV3)

PCV2 – Result example for four herds









- Overview of the pathogen dynamic in the herd
- Monthly monitoring of pathogens provides information about:
 - The distribution of pathogens in a healthy status
 - An outbreak situation \rightarrow benchmarking the findings of pathogens
 - \rightarrow More specific treatment
- Enables follow-up on interventions and change in management

The bovine high-throughput real-time PCR system

Development of a bovine high-throughput system

- Used for detecting of 11 different bovine viruses and bacteria
- All the relevant pathogens causing respiratory and intestinal diseases in calves are included
 - Reduce the risk of not detecting the relevant pathogen(s)



Viruses

Bovine coronavirus (BCoV)

Study design

- Sampling
 - Nasal swab, serum and fecal samples were collected from 100 Danish herds (83 dairy and 17 slaughter herds)
 - Dairy herds: three age groups (0-10 days, 3 weeks and 3 months of age)
 - Slaughter herds: two age groups (2 weeks after arrival and 3 months of age)
 - Sample periods: September-April 2018-2019 and 2019-2020
- Clinical observations
 - Rectal temperature, coughing, nasal and eye discharge, diarrhea, hair coat, joints, body condition

Occurrence of respiratory and enteric pathogens in pools



Occurrence of respiratory and enteric pathogens in pools



Influenza D virus

- The family Orthomyxoviridae
 - Influenza A virus 8 gene segments
 - Influenza B virus 8 gene segments
 - Influenza C virus 7 gene segments
 - Influenza D virus 7 gene segments encodes only one glycoprotein (HEF)
- Isolated for the first time in 2011 in the US (Oklahoma)
- Detected in Asia, Europe and USA
- Detected in bovine, pigs, horses, sheep and goats



D/bovine/Italy/16RS2359/2016
 D/bovine/Italy/4128547/2018
 D/bovine/Italy/18RS182-14/2017
 D/bovine/Italy/59542/2018
 D/bovine/Italy/50543-3/2018

D/swine/ltaly/254578/2015

D/bovine/Italy/31525/2018
 D/bovine/Italy/18RS182-13/2017
 D/bovine/Italy/18RS182-8/2017
 D/bovine/Italy/16RS2581-9/2016

D/bovine/France/5920/2014

D/swine/Italy/218655/2017
 D/bovine/Italy/37760/2019
 D/swine/Italy/173287-4/2016

Influenza D virus – HEF gene

Red: Samples collected in 2019 (Robust calves) Blue: Samples collected in 2020 (Robust calves) Green: Sample collected in 2015 (another project)



An AMR high-throughput real-time PCR system

• AMR: Antimicrobial resistance genes



Antimicrobial classes (55 genes)

Aminoglycoside (5) Macrolide (4) Phenicol (3) Sulphonamide (2) Tetracycline (5) Trimethoprim (3) Vancomycin (2) Beta-Lactamase (4) Low affinity PBPs (2) ESBL (7) Quinolone (6) Ampc Beta-Lactamase (3) Carbapenemase (6) Colistin (2) Fosfomycin (1)

06/03/2023 28

SEGES

Acknowledgment

Lars Erik Larsen Charlotte Hjulsager Kerstin Skovgaard Jesper Schak Krog **Rikke Søgaard** Maja Kobberø Thomas Kusk Yuan Liang Lise Kirstine Kvisgaard Marlene Rask Andersen Pia Ryt-Hansen Simon Welner Karin Tarp Tine Skotte Hammer Hue Thi Thanh Tran Nina Dam Grønnegaard Sari Mia Dose Jonas Høgberg Ivan Larsen Sofie Hagedorn Nielsen Katrine Fog Thomsen Sven Erik Lind Jorsal

Charlotte S. Kristensen Ken Steen Pedersen Solvej Ø. Breum Jonathan Rogersen Bodil H. Nielsen Mette B. Petersen Liza R. Nielsen Nina D. Otten Anne Marie Michelsen Maria Brydensholt Alicia F. Klompmaker Mogens A. Krogh Mogens Vestergaard Henrik L. Martin Martin Bjerring Henrik H. Møller Dorte B. Lastein Nynne Capion Franziska H. S. Pedersen Stine Lindgren Masja F. R. Søndergaard Thomas D. Poulsen

Jesper K. Davidsen Lucie J. M. Dupont Anaëlle Bouqueau Sascha Coes Jensine Wilm Mette Gerdes Wilson Emma Madsen Helge Kromann Sofie Jeppesen Maëva Durand Helene Larsen





UNIVERSITY OF COPENHAGEN

DTU



Thank you for your attention



Email: nbgo@sund.ku.dk