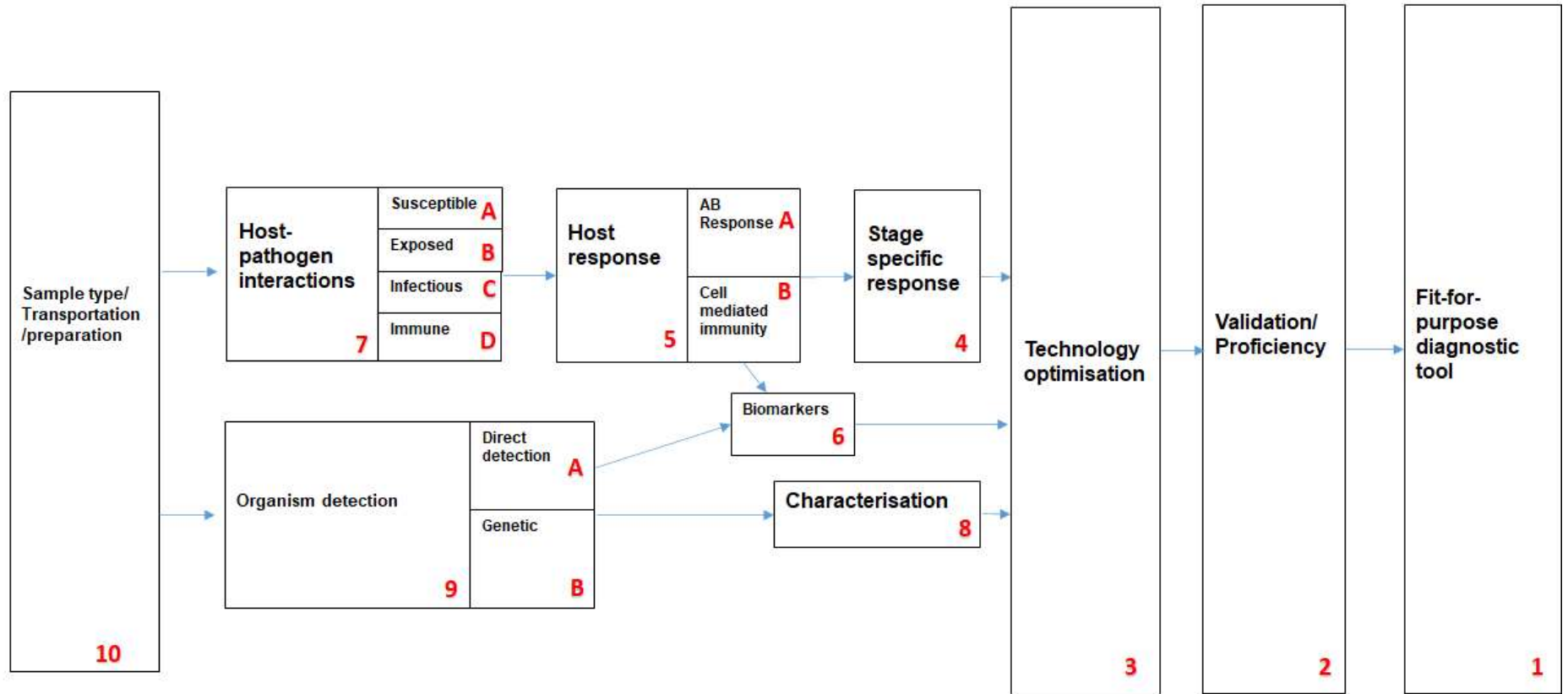




## Roadmap Lead Summaries

<b>Disease/pathogen</b>	Influenza				
<b>Roadmap type</b>	Development of Diagnostic tests				
<b>Version: Date</b>	V1	17/03/2023			

## Diagnostic Test Development Roadmap



## Lead Summary 1 - Diagnostic

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Development of rapid, reliable diagnostic test for early detection of influenza virus including viruses with zoonotic and/or pandemic potential and including pen-side tests

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Subtype-specific testing capability, including antibody tests for sera  
Automatic statutory reporting requirements for High Path viruses  
Commercial market of diagnostic tests  
Multiple antibodies to different IAV  
Utility/sensitivity of in-field diagnostic solutions

### Solution Routes

*What approaches could/should be taken to address the research question?*

Development of tests for use in the field  
Development of DIVA tests  
Fast typing of AI virus using molecular tools without virus isolation, applicable for

Pets diagnostic tests may need to be developed, but lack of commercial market is disincentive for development  
Tests for the identification of seroconversion resulting from IAV infection include both non-specific and multiplex assays  
Assays to identify HA binding specificity for sialic acid-terminated glycans  
Improve Real-Time PCR tests for subtype-specific and internal controls  
In-field direct to sequencing, improved laboratory sequencing  
Testing for low resource settings that don't require infrastructure

### Dependencies

*What else needs to be done before we can solve this need?*

DIVA surveillance protocols acceptable to trade partner  
Sample transportation and quality  
Local test availability  
Way to capture and share field observations and knowledge obtained during an outbreak to help prevent and control disease in subsequent events  
Adequate/standardized protocols for sampling and extraction  
The lower accuracy of the rapid test ,a kind of limitation, needs to be understood and accepted by the users and governments

### State of the Art

*Existing knowledge including successes and failures*

Serologic DIVA strategies are available, but little information is available to understand how this can be used in the field ( How many samples need to be taken to meet surveillance needs )

**Projects**

*What activities are planned or underway?*

## Lead Summary 2 – Validation

**Research Question**

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Validation of diagnostics tools

**Challenge(s)**

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Validation of existing diagnostics against novel isolates  
Lack of harmonized validation protocols  
Development of internal controls that are standardized and readily shared  
Validation of portable PCR  
Validation of serologic DIVA tests to understand levels of sampling to achieve goal to show freedom from disease  
Understand how NGS can be used to supplement/augment PCR testing  
Validation of use for minor species (9000 avian species!!) and age of animal (e.g. piglets and sows)  
The chances of independent and external validation are rare and its cost may be expensive  
Lack of reference material

**Solution Routes**

*What approaches could/should be taken to address the research question?*

Standardization of methods across labs – ring trials

Co-ordinated lines of reference material production  
Orchestrated collections of field material  
Improve broad detection, sensitivity and specificity of diagnostic tests  
Validation of portable PCR  
Validation of serologic DIVA tests to understand levels of sampling to achieve goal to show freedom from disease  
Understand how NGS can be used to supplement/augment PCR testing  
Public/private partnerships – letting private companies have access to samples and share technologies with companies despite restrictions in sampling handling of material from HPAI/regulated diseases

**Dependencies**

*What else needs to be done before we can solve this need?*

Reference materials  
Organizational structures of concerted validation lines.  
Availability of economic resources

**State of the Art**

*Existing knowledge including successes and failures*

**Projects**

*What activities are planned or underway?*

## Lead Summary 3 - Technology optimisation

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Continued improvement of technologies

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Decrease time and costs

Improve NextGen Sequencing for a cost-effective use in the field

Continue to optimize rRT-PCR to support routine and targeted surveillance

Pen-side application: RT-rtPCR and sequencing.

What can go wrong will go wrong in the field

### Solution Routes

*What approaches could/should be taken to address the research question?*

Benchmark gold standard versus new technology (PCR vs Metagenomics)

Overcoming patent issues? Capitalising on human medicine/dx to bankroll animal medicine/dx

Validation testing conducted in parallel to reduce time to release on market

Improvement of NextGen Sequencing: a ) decrease the costs b )

Continue to develop Minion and other rapid NGS platforms for more rapid targeted and random sequencing c ) Improve sensitivity of NGS platforms d ) Improve bioinformatic analysis pipelines and provide

technology to be usable at state diagnostic laboratories e ) Identify faster and more efficient sample preparation and testing  
Optimisation of rRT-PCR: a)Continue to review and update as needed for type A and subtyping tests ( use of rapid sequencing approaches ) b) Continue to explore and validate environmental testing c ) Integrate internal positive control to identify false negative results ( possible host control ) d) assemble Rt-qPCRs to match the circulating HPAIV sub- a Local and centralized tools/staff support for data evaluation and sequence interpretation and visualization/sharing  
Improvement of communication and data sharing: selected tests or diagnostics need to be used and shared by many players

### Dependencies

*What else needs to be done before we can solve this need?*

Continue to harmonize protocols and adapt to new technologies

Sample collection and transportation independent of cold chain

Provide copy-based, non-infectious reference material and controls

Public/private sharing

Develop strategies to use tests (screening to characterization)

Reduce cost of rRT-PCR to increase use

Development of alternative reagents to ~~de~~ not be dependent on single manufacturer

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

(1)A research project exploiting Nanopore technology to identify and characterise the field clinical samples quickly and accurately ('23-24)

## Lead Summary 4 - Disease stage specific response

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

### Solution Routes

*What approaches could/should be taken to address the research question?*

### Dependencies

*What else needs to be done before we can solve this need?*

Clear definition of disease stages across species

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 5 - Host response

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Identify a measurable host response

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Complexity of interwoven factors

Inherent variability of maternal antibodies

Varying types of antibodies, deltaFcgY and their roles, mucosal immunity

Need for animal models in the laboratory

Many different wild bird species as a host to understand

### Solution Routes

*What approaches could/should be taken to address the research question?*

Antibody detection vs. Protection, methods for hi throughput HIs?

Other?

Utility of in vitro systems

### Dependencies

*What else needs to be done before we can solve this need?*

Avian immunology for various species

Understanding of the immune system of migrating birds

Research facilities that can house dams, mares, hens and not just their offspring

Understanding the nature and effectors of resilient AIV immunity

Factors that contribute to failure to mount such immune responses in a species-dependent manner

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*



## Lead Summary 5A - Antibody response

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Rapid and easy antibody response detection

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Rapid and easy detection of neutralization antibody, cross-reactivity antibody and antigenic characterization.

Sensitivity of rapid assays

Inability to interpret antibodies in young animals – are they neutralizing? Are they maternally derived? Does their presence impact disease expression (VAERD or protection)?

Lack of true correlates of protection versus seroconversion

Understanding of the generation and decline of the humoral immunity of the infected and vaccinated birds

### Solution Routes

*What approaches could/should be taken to address the research question?*

ELISA and neutralizing antibody test development.

In-field rapid test development and improvement

Antibody test platforms to evaluate cell mediated immune responses.

Antibody test platforms to evaluate neutralizing antibodies.

Orchestrated development of monoclonal antibodies

Roadmap for the varying uses of antibody diagnostic approaches for veterinarians/responders based on purpose (DIVA vs. Early detection, etc)

### Dependencies

*What else needs to be done before we can solve this need?*

Monoclonal antibodies availability

Reference material collection

Field sample collection

### State of the Art

*Existing knowledge including successes and failures*

(2)Competitive ELISA based on NP protein is developed, and this can be used for the serum samples for many different avian species to see the seroconversion due to vaccination or natural infection.

### Projects

*What activities are planned or underway?*

## Lead Summary 5B - Cell-mediated immunity

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Diagnose the presence and valence of cell-mediated immunity against AIV

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

High laboratory specialization and technical skills of staff required  
Limited understanding of variation of mechanisms between species and viral strains  
Need for specific equipment to measure immune cells (I.e.: FACS, etc)  
Variable genetic backgrounds of host (I.e: commercial breeds versus backyard stocks)

### Solution Routes

*What approaches could/should be taken to address the research question?*

Develop technically undemanding and cost-effective tools to define (parts) of CMI  
Role of cell-mediated immunity (CMI) in individual animal recovery and outbreak outcomes  
Tools to define immune cell populations in avian species  
Test to predict cell mediated levels and correlate with vaccine protection

### Dependencies

*What else needs to be done before we can solve this need?*

Basic data on the development of cellular immune effectors in various avian species  
Research facilities that can hold animals for a long period of time (months) to determine if CMI is protective against heterologous challenge

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 6 – Biomarkers

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Biomarkers for influenza that can capture different infection phases

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Biomarkers not known

### Solution Routes

*What approaches could/should be taken to address the research question?*

Determine biomarkers that can better capture infection phases, particularly for low pathogenic influenza viruses, i.e. acute phase proteins.

### Dependencies

*What else needs to be done before we can solve this need?*

Validated surrogate markers

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 7 - Host-pathogen interaction

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Identify the most appropriate target for diagnosis based on the Host-pathogen interaction

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Lack of knowledge on the impact of the host's infection status on the (immune) response

Virus-host interactions control species-specific pathology.

Virus-specific host susceptibility, including carrier state.

### Solution Routes

*What approaches could/should be taken to address the research question?*

Need to better understand the relevant mechanisms for an effective immune response, role of cross-reactivity and cell-mediated immunity.

Need to understand resistance or susceptibility factors (e.g. intracellular enzymes in different organs, endosomal pH, adhesion molecule 1 (VCAM-1), intercellular adhesion molecule 1 (ICAM-1), tumor necrosis factor (TNF) etc.) towards HPAI viruses in different wild (main reservoirs) and domestic bird and mammal species

Role of microRNAs in modulating pathogenesis

Host responses for diagnosis of infections, for clues about viral pathogenesis

### Dependencies

*What else needs to be done before we can solve this need?*

Challenge studies that provide foundational information

Longitudinal studies

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 7A- Susceptible

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

### Solution Routes

*What approaches could/should be taken to address the research question?*

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### Dependencies

*What else needs to be done before we can solve this need?*

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### State of the Art

*Existing knowledge including successes and failures*

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### Projects

*What activities are planned or underway?*

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## Lead Summary 7B – Exposed

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

### Solution Routes

*What approaches could/should be taken to address the research question?*

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### Dependencies

*What else needs to be done before we can solve this need?*

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### State of the Art

*Existing knowledge including successes and failures*

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### Projects

*What activities are planned or underway?*

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## Lead Summary 7C – Infectious

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

### Solution Routes

*What approaches could/should be taken to address the research question?*

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### Dependencies

*What else needs to be done before we can solve this need?*

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### State of the Art

*Existing knowledge including successes and failures*

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### Projects

*What activities are planned or underway?*

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## Lead Summary 7D – Immune

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Define duration and valence (“how protective is it?”) of immunity following

- Natural infection
- Vaccination
- After multiple exposures

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Define assays that help to evaluate the valence (protectivity) of immunity after multi exposures or multi immunizations

### Solution Routes

*What approaches could/should be taken to address the research question?*

Conduct longitudinal studies in naturally infected and vaccinated avian species

Determine the efficacy of natural and vaccinal immunity by virus exposure (challenge studies)  
Maternal antibody interpretation, efficacy

### Dependencies

*What else needs to be done before we can solve this need?*

Challenge studies  
Availability of field viruses  
Facilities to challenge various ages of animals  
Establish standardized challenge protocols that take into account harmonized calculation of  $R_0$ .

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*



## Lead Summary 8 – Characterisation

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Characterisation of influenza virus (biological, sequences, viruses, phenotype)

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Characterise determinants of high- vs. low-pathogenicity avian influenza also outside the HA cleavage site  
Inability/ lack of ease in detecting true reassortment vs. Contamination of sample  
Characterize zoonotic propensity  
Limitations for in vivo testing due to animal welfare laws  
Characterize antigenic properties with respect to vaccine susceptibility  
Failure to maintain sample authenticity thus inability to study observed outcomes of infection  
Transparency about human impacts  
Lack of understanding how genomic reassortment is made exactly

### Solution Routes

*What approaches could/should be taken to address the research question?*

Loss of function experiments both in vitro and in vivo  
Study virus transmission across interfaces  
Identification of antibody epitopes important for antigenic drift in swine and poultry species

Development of models to predict epitopes based on HA sequence evolution

Promotion and adoption of a standard classification scheme (e.g., the global swine clade classification scheme and the H5 clade classification scheme) - and have this classification updated in real time (i.e.: standardized nextstrain, etc)

### Dependencies

*What else needs to be done before we can solve this need?*

Virus isolates  
Pathogenicity models  
Interface transmission models  
Genomic surveillance in multiple hosts and even different sites within host

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 9 - Organism detection

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Detection of influenza viruses in various species and matrices including environmental sources

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Sensitivity of detection versus specificity versus contamination risks  
Cost effectiveness, commercialization  
How many/much samples are needed to determine prevalence/risk?  
I.e.: pools, environmental samples, versus direct from animal

### Solution Routes

*What approaches could/should be taken to address the research question?*

Sampling methods and transport for low resource settings  
Assessment of local need to reach acceptable confirmatory diagnosis  
Implementation of routine targeted testing for endemic IAVs that can be scaled up during an outbreak and/or serve as background data when a new IAV strain emerges or is a concern  
Validate and verify novel solutions (field PCR, LAMP, etc) for in-field and laboratory methods

Standardize diagnostics from environmental samples

Surveillance sensitivity should be decided

Protocols to work through co-infections in order to capture all pathogens

Rapid sensitive and specific methods for field detection (subtype specific if possible?)

### Dependencies

*What else needs to be done before we can solve this need?*

Clear country requirements for diagnostic (matrices and species)

Defining diagnostic needs according to region, poultry vs wild birds vs livestock vs humans etc

Understanding IAV ecology for a location/system in order to target surveillance

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 9A - Direct detection

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Rapid direct detection of influenza viruses and its variants

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Cost-effective and rapid direct detection of replication-competent influenza virus

Matrix-specific effects

Overcoming embryonated eggs as the main propagation vehicle-retaining sample authenticity (re sample passage in cells/eggs)

Host antibodies that neutralize virus

Field based diagnostic for rapid early assessment with perhaps lab based confirmation/subtyping

Genetic changes associated with different systems

Low viral load in the field samples

Mixed subtypes in field samples (coinfections)

### Solution Routes

*What approaches could/should be taken to address the research question?*

Improvement of virus isolation and propagation by new in vitro cell culture techniques

Improvement of systems for viral replication/growth

Build cell culture collections

Modify existing cell lines to be more sensitive by expressing or repressing genes

Pen-side tests?

### Dependencies

*What else needs to be done before we can solve this need?*

Availability of SPF eggs in low resource settings

Cell culture collections

Crispr/Cas technology with avian cell culture systems

Germ line manipulation tools in avian species

Sharing of resources

Early assessment linked with automatic alerts to advisors/central labs for additional tests

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

## Lead Summary 9B – Genetic

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Rapid, cost-effective tools for

- genetic detection
- Sequence analysis and characterization
- Sequence reporting/databases that communicate with each other (I.e., sequences from one database can be downloaded and analyzed in another without loss of integrity)

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Systematic approaches for analysing and reporting viral genome sequences

Issues around samples most relevant for infection in different species with swabs alone not necessarily enabling detection

Approaches that are specific ie need to know what you have before you can test

### Solution Routes

*What approaches could/should be taken to address the research question?*

Effective deployment of 3rd -gen sequencing technologies for both wet- and dry-lab protocols, allowing point-of-incidence analysis

Rapid characterization of variants circulating in the field via high-throughput point-of-care technology

Improve sensitivity and purification methods for whole-genome sequencing

Evaluation of whole genome sequencing platforms

Develop highly multiplexed PCR assays

Improved and easily employed metagenomic pipelines

Commercialization or Increased availability of novel technology like viability PCR tests

### Dependencies

*What else needs to be done before we can solve this need?*

Training of more bioinformaticists to mine and interpret the data

Understand the relationship between genetic detection and transmission IRL

Improved reference databases/faster assignment of genetic clades as they appear

Turnkey solutions to allow use in more veterinary labs

Reduce complexity of testing procedure to make it more accessible

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

(3)A Research project for development of multiplex real-time RT-PCR for subtypes H5/H7/H9 and multiplex end-point RT-PCR of HA subtyping ('22-23)

## Lead Summary 10 - Sample type/transportation/preparation

### Research Question

*What are we trying to achieve and why? What is the problem we are trying to solve?*

Improvement of viral sampling, transportation and isolation of nucleic acids and viruses

### Challenge(s)

*What are the scientific and technological challenges (knowledge gaps needing to be addressed)?*

Sampling sparing methods  
Improvement of safe transport of virus  
Pools and matrix of choice for environmental sampling, Virus extraction from environmental matrices (e.g. water, mud, aquatic microfauna, etc.)  
Poultry environs that also include varying types of chemicals  
Limitations defined by CITES (species protection and conservation acts)  
Limitations defined by the NAGOYA protocol (WHO exemptions etc.)  
Preserve the quality and viability of biological samples during transportation over long distances and temperature extremes

### Solution Routes

*What approaches could/should be taken to address the research question?*

In-field extraction and diagnostics (PoC/Penside) to minimize transport  
Development of transport media that stabilise the virus without refrigeration  
Non-invasive samples from wild birds  
Non-invasive samples from swine  
Environmental sampling methods including

- Air sampling and analysis protocols
- Waste water
- Poultry house sampling, where are the best samples

### Dependencies

*What else needs to be done before we can solve this need?*

Standardise environmental sampling procedures  
Easy legal frameworks for transport  
Manageable CITES and NAGOYA rules for avian samples  
Data management for sampling/testing coordination  
Human resources

### State of the Art

*Existing knowledge including successes and failures*

### Projects

*What activities are planned or underway?*

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