



Detection of avian Influenza A virus by RT-LAMP on VETPOD system

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PLOUFRAGAN - FRANCE

VIVALDI project

Veterinary Validation of Point of Care Detection Instrument

- Partners



anses



- Validate new equipment (the VETPOD platform) for rapid on-site detection of zoonotic pathogens (**Avian influenza**, Salmonella and Campylobacter) in industrial food and animal production chains

VETPOD



- VETPOD containing both
 - **Multiplication** of nucleic acids in sample by RT-LAMP (RT-Loop Mediated Isothermal Amplification)
 - **Detection** of turbidity formed by precipitates during the RT-LAMP reaction

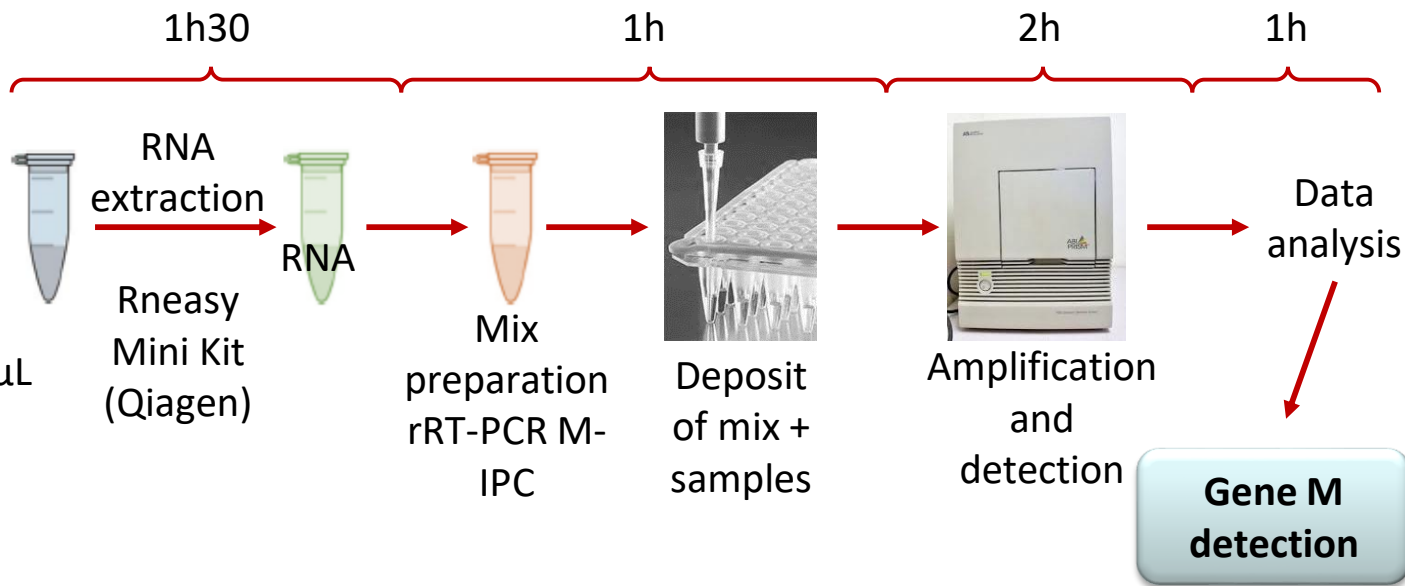
VETPOD



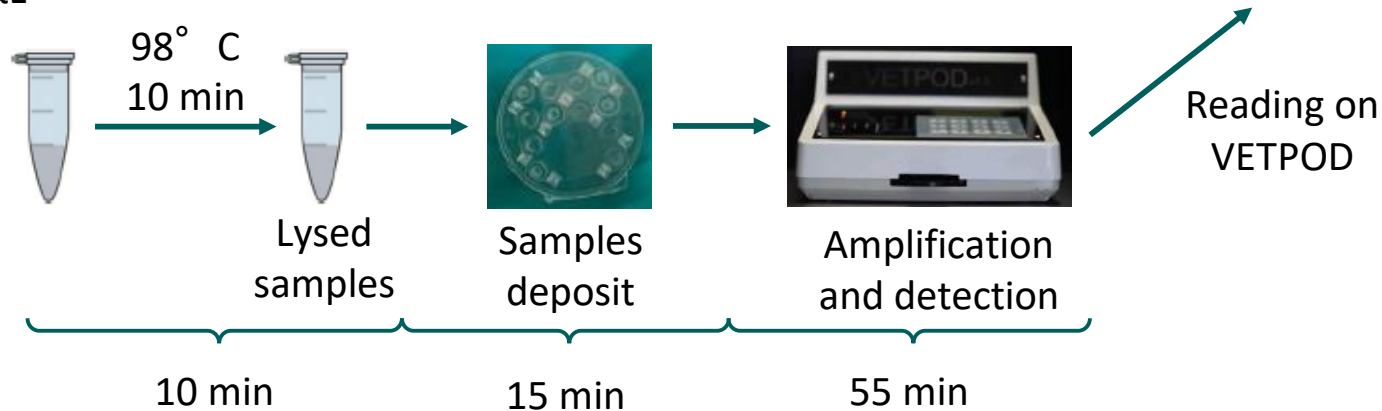
- Disposable polymeric **cartridges** (a Lab-on-Chip) performing the biological reactions of pathogen detection.
- The VETPOD system is **generic**, i.e. the cartridges may be loaded with reagents targeting almost any pathogen of choice.

Avian influenza RT-LAMP VETPOD validation method

Reference method
rRT-PCR M-IPC
(adapted from Spackman)

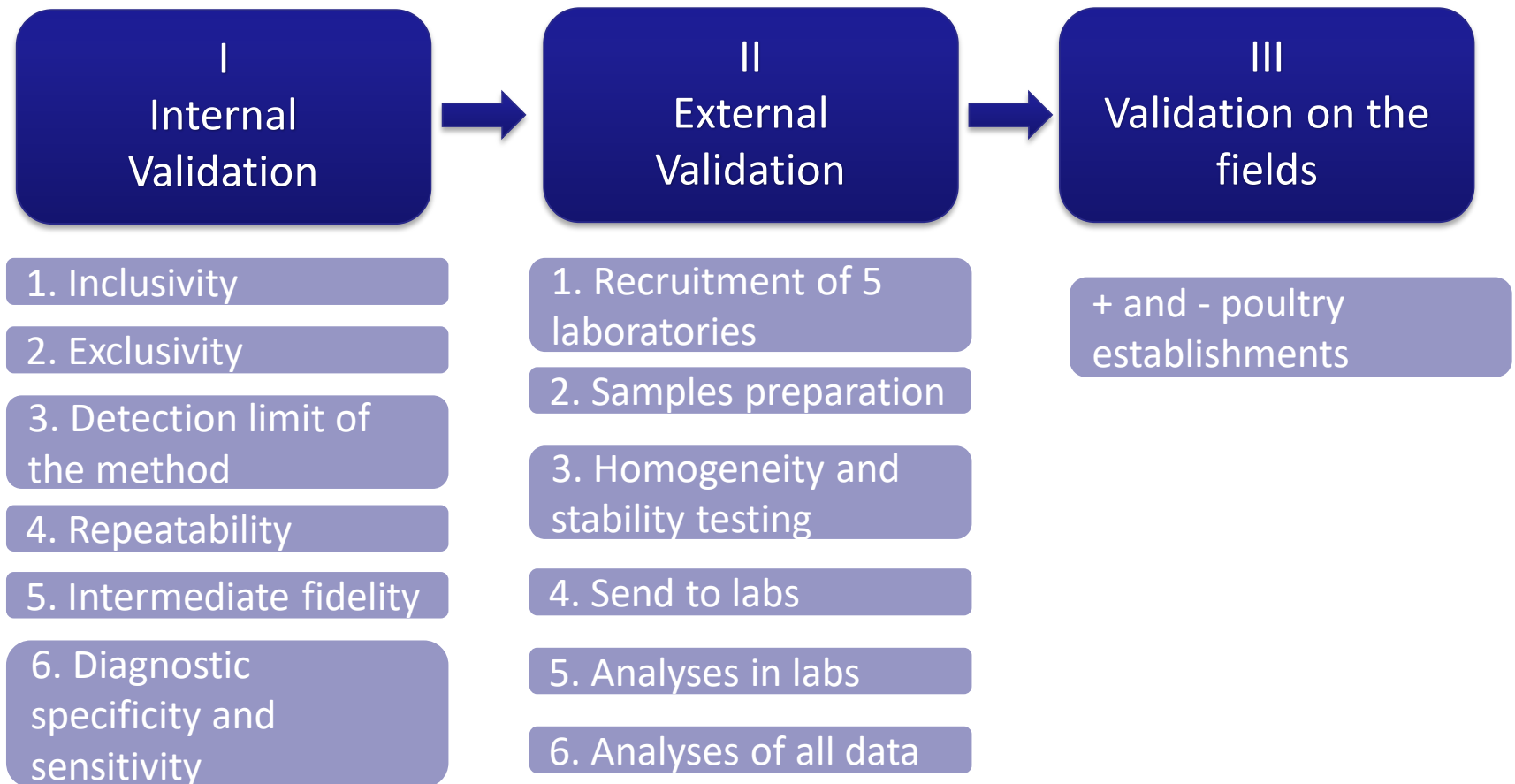


Alternative method
RT-LAMP with
VETPOD



Avian influenza RT-LAMP VETPOD validation plan

- Validation following
 - OIE Terrestrial Manual
 - French standard NF U47-600



I – Internal Validation: Results

1. Inclusivity

28 AIV strains detected
2 strains not detected

H1N1_130045 : Pandemic strain

H6N2_0139/01 : American lineage

Hypothesis : virus quantity or lack of specificity or lysis efficiency

2. Exclusivity

25 non-AIV strains

All strains not detected as expected

3. Detection limit of the method

H5N3_100205b $10^{4.53}$ EID₅₀/mL

H4N2_100272p $10^{4.53}$ EID₅₀/mL

H7N9_090034d $10^{2.46}$ EID₅₀/mL

Same limit of detection as the reference method

4. Repeatability

92.3 % of positive samples were detected with the VETPOD system

100 % of negative samples were not detected

5. Intermediate reproducibility

1 sample (/60) excepted positive was detected negative

98.3% of samples were detected

6. Diagnostic specificity and sensitivity

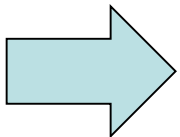
DSp : 100 %

DSe : 73.68 %

I – Internal Validation: Results

6. Diagnostic sensitivity

- Why were some samples not detected for M gene?
- Why was estimated DSe lower than expected?
 - Inhibition of samples ?
 - Mismatches in primers sequence ?
 - Lysis not efficient ?
 - Comparison of samples: direct thermal lysis vs RNA extraction
 - Test with added sonication step



Test of the 3 hypothesis

I – Internal Validation: Lack of detection hypothesis

6. Diagnostic sensitivity

Hypothesis 1
Inhibition of RT-LAMP

Hypothesis 2
Mismatches

Hypothesis 3
Lysis not efficient

No inhibitor effect

No major shift
between primers
and M gene
sequence

Lysis not fully
effective for
diagnostic samples

Ultrasound seems
to improve lysis

II. External Validation : Plan

- 25 samples prepared
 - H5 / H7 / AIV / no-AIV / negative
 - Dilutions (Low/Medium/High)
 - Duplicate samples
- Homogeneity / Stability
- Samples received in the 5 laboratories :
 - 2 Laboratories in France (Labocéa/Labofarm)
 - IZSVe and FLI
 - ANSES (analyses done)

Conclusion

- Fast / effective / easy to implement / inexpensive method
- Adaptation required but high potential to be usable directly on farm



VIVALDI 

Validation plan of VETPOD system for avian Influenza A viruses

THANK YOU FOR YOUR ATTENTION



Project managed by Marion Flodrops
(Post-doctoral position)