

Detection of novel ASF variant via triplex Real-time PCR assay

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Triplex real-time PCR assay

This assay can be used to detect and identify viral nucleic acid (DNA) of ASF virus variants containing genome deletions in the *MGF360/505* and *EP402R (CD2v)* genes.

- The assay cannot be used to determine genotype. Additional conventional PCR testing and sequencing for p72 genotyping is required using specific primers.
- No cross-reactions with other porcine viruses have been detected.

PCR Protocol

Samples

- Tissue (spleen, lymph nodes, liver, tonsils and lung *et al*)
- Whole Blood in anti-coagulant
- Blood swabs or dried blood spots on filter paper
- Serum
- Oral-nasal swabs, oral fluids from rope chew collections, deep throat swabs

PCR Protocol

- **DNA Extraction**

- There are several commercially available nucleic extraction kits that are suitable for the extraction and purification of ASF viral DNA.
- Manually
- automatically

Primer and Probe Mix for detection of ASFV variant

| Primer | Sequence (5' to 3') | Volume per reaction (μl) | Final Conc. (nM) |
|-----------------------|--|--------------------------|------------------|
| CD2v_F (20 μM) | AGAAGAACAATGTCAGCATGATGAC | 0.5 | 400 |
| CD2v_R (20 μM) | CGACTGTAAGGCTTAGGAAGTAATGG | 0.5 | 400 |
| CD2v_P_VIC (10 μM) | VIC-CCACTTCCATACATGAACCATCTCCCAGA-BHQ1 | 1 | 400 |
| MGF-14L_F (20 μM) | AGAAGACGGGGTTCGGATACAG | 0.5 | 400 |
| MGF-14L_R (20 μM) | GCAAATCCTGAATATGGGCTTATACG | 0.5 | 400 |
| MGF-14L_P_Cy5 (10 μM) | CY5-CTCCCAGTTCCGCACACAGCCGC-BHQ2 | 0.8 | 320 |
| p72_F (20 μM) | AAGTTTCGGTACGCATTCTTTGT | 0.4 | 320 |
| p72_R (20 μM) | TATGACTGGGACAACCAAACACC | 0.4 | 320 |
| p72_P_FAM (10 μM) | FAM-ACAAGCGTGTAACGGCGCCCTC-BHQ1 | 0.4 | 160 |

Designed and verified by Li Lin

Preparation of mastermix

| Reaction Component | Volume (μ l) |
|----------------------------------|----------------------------|
| Nuclease-Free Water | 2.5 |
| 2X RT-PCR Buffer | 12.5 |
| ASF CD2v Primer Probe Mix | 2 |
| ASF MGF360-505R Primer Probe Mix | 1.8 |
| ASF p72 Primer Probe Mix | 1.2 |
| Total volume mastermix | 20.0/reaction tube or well |
| Addition of template | Volume (μ l) |
| Sample DNA | 5.0 μ l |
| NC | 5.0 μ l |
| PC | 5.0 μ l |

Total volume of reaction mixture 25 μ l

Program of the Real-time PCR

| Setting | Temperature (°C) | Time | Cycle |
|--------------------------|------------------|--------|----------|
| Pre-incubation | 50 | 2 min | 1 cycle |
| Pre-denaturation | 95 | 5 min | |
| Denaturation | 95 | 15 sec | 45 cycle |
| Annealing and elongation | 60 | 60 sec | |

Fluorescence is recorded in the FAM, VIC and Cy5 channels during annealing.

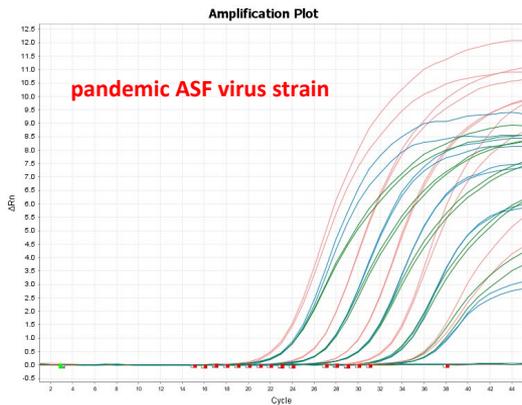
Result Analysis

- **Positive result:** Cts < 35 for CD2v/MGF-14L/p72 **AND** the appearance of specific amplification curves for each assay
- **Negative result:** Ct ≥ 40 or no value
- **Indeterminate result:** Ct between >35 and 40, if indeterminate result is obtained, repeat the test to confirm.
- Where the p72 assay is positive AND one or both the MGF and CD2v assays are negative, this indicates that a **variant ASF virus** has been detected.
- Where all assays are positive, this indicates that the **parental or pandemic ASF virus strain** has been detected.
- Conventional PCR and sequencing for the variant strains.

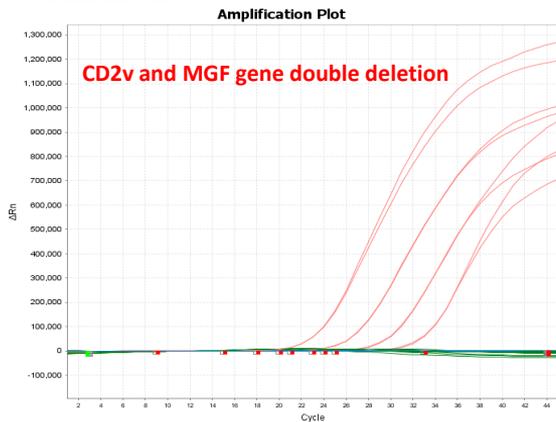
Interpretation of ASF virus triplex PCR results

| Interpretation | Test Results | | |
|--|--------------|----------|---------|
| | P72-FAM | CD2v-VIC | MGF-Cy5 |
| ASFV pandemic strain positive | + | + | + |
| ASFV CD2v gene deletion strain positive | + | - | + |
| ASFV MGF gene deletion strain positive | + | + | - |
| ASFV CD2v and MGF gene double deletion strain positive | + | - | - |
| ASFV negative | - | - | - |

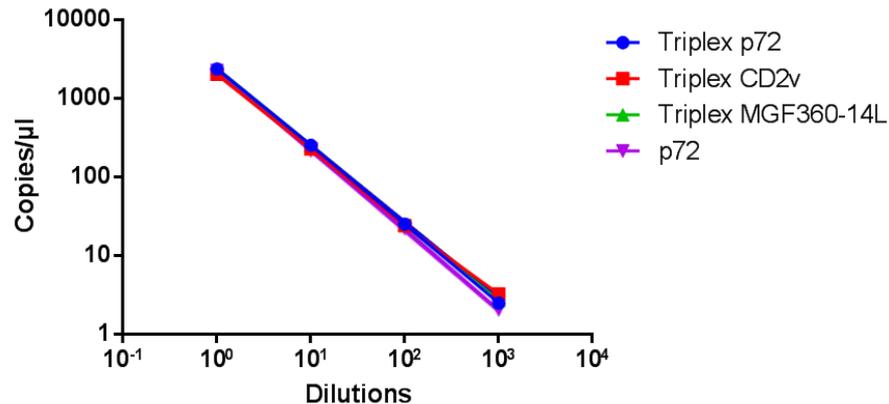
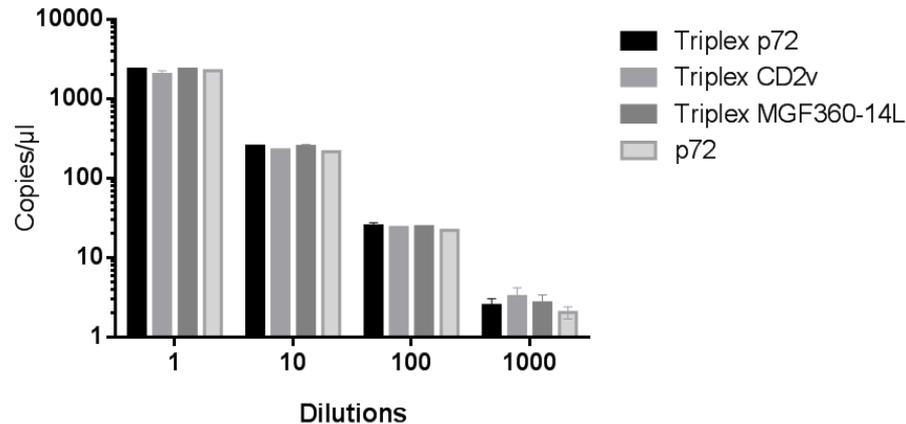
Method Validation



■ CD2v ■ MGF360-14L ■ P72



■ ASFV-MGF360-14L ■ ASFV-p72 ■ ASFV-CD2v



Thank you!

