

African swine fever virus dual nucleic acid detection Kit

(MIRA-exo) Guide Manual

(Part number: WLP9250KIT)

Product description

The kit designs specific primers and probes in the highly conserved regions of the P72 gene and MGF360-10L gene of African swine fever virus(ASFV), respectively. It can detect the African swine fever virus(ASFV) P72 gene and MGF360-10L gene in swine blood, serum, or tissues in 20 minutes by MIRA technology.

Overview of the MIRA technology

Multienzyme isothermal rapid amplification(MIRA) is based on a combination of polymerases and DNA recombination/repair proteins: Recombinases bind to single-stranded nucleic acid and stimulate the resulting protein-DNA complex to search for homologous sequences in duplex DNA with Single Strand Binding protein. Once homology is located, a strand-switching reaction is performed and the oligonucleotide is paired to its complement permitting a polymerase to begin synthesis from the 3' end. The amplification process is very rapid when optimised and can reach detectable levels of product in a few minutes.

Materials provided

Materials provided	Content
E Buffer	2 tube × 1 mL
B Buffer	1 tube × 150 μL
Positive control DNA template	1 tube × 100 μL
Steel bead reagent	48
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Storage conditions

- Transport temperature: ≤ 20 °C.
- Store at -20 °C without light, no repeated freezing and thawing.
- Full activity is guaranteed for 14 months.

Set up

1. Thaw reagents at room temperature.
2. Shake the lyophilized powder tube to confirm that each tube has a small steel bead. Thoroughly mix the contents of each tube by vortexing, then briefly centrifuge.
3. Add 37.5 μL E buffer (supplied) to a sample response MIRA exo reaction.
4. Add 10 μL sample DNA.

5. Add 2.5 μ l B buffer (supplied) and mix well and spin briefly to start reaction (For multiple reactions, it is recommended to add B buffer to the inner side of the reaction tube cover, and turn the reaction tube upside down to mix well) .
6. Turn on the Amp Future WL-16-II Constant Temperature Fluorescence Detector, put the reaction tube into the device and record the sample placement sequence, select DNA mode, 16-well synchronization, and both of FAM/VIC channel. The amplification temperature of this program was 39 - 42 $^{\circ}$ C, 30 s/cycle, 20 minutes, and a total of 40 cycles.

Reaction mix

E Buffer	37.5 μ L
DNA sample	10 μ L
B Buffer	2.5 μ L
Total volume	50 μ L

Interpretation of results

1. MGF360-10L gene positive:the amplification curve of FAM channel is S-like, and the instrument is judged as "+".
2. P72 gene positive:the amplification curve of VIC channel is S-like, and the instrument is judged as "+".
3. Negative: the instrument both of FAM/VIC channel is judged as "-".

Note: When affected by the color of the sample or other factors, the machine shows the result as "+" but there is no "S"-shaped amplification curve, the result is judged as negative.

Note

1. Reactions start as soon as B buffer is added.
2. Set up a negative control for each experiment.
3. For research and development use only.

Ordering information and technical support

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