

PROCEEDING

INTERNATIONAL SEMINAR

ADVANCED

TECHNOLOGY ON

VETERINARY AND LIFE

SCIENCE

NO. ISBN: 978-979-96104-4-7

YOGYAKARTA
MARCH 12nd, 2011



FACULTY OF VETERINARY MEDICINE
UNIVERSITAS GADJAH MADA



ISBN: 978-979-96104-4-7

Title **Proceedings on International Seminar Advanced Technology on
Veterinary and Life Science**

Published By **Faculty of Veterinary Medicine Universitas Gadjah Mada**
Address

Jalan Fauna 2 Karang Malang Yogyakarta 55281

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March, 2011

each authors

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DETECTION OF JEMBRANA VIRUS USING RT-PCR

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Introduction

Jembrana virus (JV) is a bovine lentivirus genetically similar to bovine immunodeficiency virus (Chadwick et al., 1995) that causes an acute, severe disease syndrome in infected Bali cattle (*Bos javanicus*) in Indonesia with a mortality of about 17% (Soesanto et al., 1990). The acute disease associated with JV infection has a short incubation period of 5-12 days and a duration of about 7 days (Soeharsono et al., 1990). The infection is characterized by fever, lethargy, anorexia and enlargement of superficial lymph node (Soesanto et al., 1990) which in recovered animal the virus is still detectable in blood for 2 years after infection (Dharma et al., 1991). Although JV infect Ongole cattle (*Bos indicus*), Friesian cattle (*Bos taurus*) and buffaloes (*Bubalus bubalus*), the clinical signs are milder and the infectious agent only persists for maximum of 9 months in buffaloes or shorter period in other species (Soeharsono et al., 1995). Diagnosis of Jembrana disease is an important constituent to control disease and avoid serious economic losses in farming. Various diagnostic methods based on clinical signs and antigen-antibody reaction have been as routine procedure but has disadvantage because the clinical sign could be similar to other illness and antibody to JV is not detected in cattle until

after recovery from the acute disease, and is not seen in a majority of cattle until 11 weeks after infection (Hartaningsih et al., 1994). Molecular techniques based on genomic sequence such as revers transcription PCR are sensitive, accurate and rapid. Recent this technique has been a usual standard in laboratory practice especially due to portable devices designed for small laboratory.

Material and Methods

2.1. Isolation of total RNAs

Total RNAs were isolated tissues of infected animal using TRIzol reagent (Life Technologies), according to the manufacturer recommandation. Briefly, approximately 70 mg of tissue were homogenized in 1 mL of TriZol then let for 10 min. at room temperature. After centrifugation at 12,000 rpm, for 10 min. in a bench minicentrifuge, the supernatant was treated with 0.2 mL chloroforme then precipitated with 0.5 mL isopropanol. The pellet was washed with 70% ethanol and dissolved in 100 μ L of steril DEPC-treated water. The purity and concentration of the RNA preparation were spectrophotometrically established at 260 and 280 nm.

2.2. Recombinat plasmid preparation

For the experiment , recombinant plasmid, pGEX-CA, contain *gag-ca* gen along 670 bp has constructed from research before was used as template. This plasmid is isolated and purified from E. Coli DH5 α using GeneJET™ Plasmid Miniprep Kits (Fermentas), followed by spectrophotometric analysis at 260 and 280 nm to determine purity and concentration.

2.3. PCR reaction

A set of two primer, forward (CCAAGAATGCAGAGACT) and backward (GGCAGTCCTCATTTGCATG) was designed to amplify 232 bp in *gag-ca* gen region. PCR reaction using Dream Green PCR Mix reagent (Fermentas) and pGEX-CA plasmid (10 ng/ μ l) was conducted to obtain annealing temperature and optimized cycle number. Negatif control were used water and pGEX-2T plasmid as template for each reaction to monitor cross-contamination. In order to test sensivity, serial dilution of pGEX-CA plasmid (10^{-1} to 10^{-9}) were used as template (original concentration sampel was 10 ng/ μ l).

2.4. RT-PCR reaction

The obtained of optimized PCR conditions were used to RT-PCR reaction. As RT-PCR reaction, we used previously reverse transcripts reaction and followed amplification of cDNA using various infected tissues as template. This reaction was done in one tube reaction using One Step RT-PCR kit (Roche).

Result and Discussion

3.1. PCR reaction

The optimized PCR reaction was carried out in a final volume of 25 μ L, using Dream Green PCR Mix reagent (Fermentas) with the following conditions : firts denaturation at 94°C for 4 min, denaturation at 94°C for 45 sec, annealing at 56°C for 30 sec, elongation at 72°C for 45 sec and additional elongation at 72°C for 10 min. The a set of primer was designed to amplify *gag-ca* gen , a conserve (Desport *et.al*, 2007) structural gen of JV that encode capsid protein virus (Chadwick *et.al*,1995).The sensivity PCR was performed under the same conditions

which it was able to detect template at 10^{-5} dilution DNA plasmid sample. This result showed the efficiency of PCR slightly high, which based on jurnal report the spesifity of PCR averagely fall at 10^{-2} to 10^{-6} . The PCR efficiency was achieved by beside accuracy of optimized reaction conditions also by primer designing.

3.2. RT- PCR reaction

The following RT-PCR conditions (final volume was 25 μ L), first cDNA synthesis at 45°C for 30 min, AMV inactivation 94°C for 2 min, cDNA amplification 40 cycles using PCR conditions discribed previously. RNA virus that extracted from infected tissues such as lymph node, whole blood, meat, and brain was used as template for RT-PCR reaction. The PCR method could detect existency of virus that showed a clear band in 1,8 % agarose with etidium bromide staining after electrophoresis. Different from DNA plasmid as template ,using RNA total usually has lower sensitivity for RT-PCR to detect the viral particles in organs or tissues that may arise from the intricate population of genes and sequences within the total RNA extracted from infected organs or tissues which comprises virus as well as host-cell RNA. But the RT-PCR method was still well detecting the virus.

Conlusions

The RT-PCR method could detect RNA virus from infected tissues and should be value in high sensitivity, rapid, and accurate than conventional detection by clinical sign and antigen-antibody reaction to constitute diagnosis in preventing spread this severe virus. It appropriate for field application in Veterinary Medicine due to portable devices designed for small laboratory.

Acknowledgment

This research was partly financed by the Fundamental Research Grant from Gadjah Mada University, and supported French Embassy in Indonesia .

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