

Immunodiagnosis in Jembrana Disease: A Review

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Article history

Received: 30-10-2014

Revised: 16-08-2015

Accepted: 17-08-2015

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Abstract: Jembrana disease is a bovine disease that affects Bali cattle (*Bos javanicus*). The causal agent is named Jembrana Disease Virus (JDV), a lentivirus member. The disease development of Jembrana disease in Bali cattle is unique for a lentivirus infection as it is related with severe clinical syndrome in an acute period. In experimentally JDV-infected Bali cattle, the death rate was about 21% and occurred within only 1 to 2 weeks post-infection. Indeed, the mortality of more than 60 000 cattles in a year was observed during the first outbreak and the disease is now endemic throughout parts of Indonesia. Early diagnosis constitutes a preventive method of further disease outbreaks. The Jembrana disease can be diagnosed by the clinical symptoms but more reliable diagnostic tools are available, based on either antigen (immunodiagnosis) or viral genome (molecular diagnosis). In this review, we summarize about immunodiagnostic tools of Jembrana disease which has been developed so far.

Keywords: Immunodiagnosis, Bovine Disease, Humoral Immune Response, Immunologic Test, Immunoassay

Introduction

Jembrana disease is a bovine disease that affects Bali cattle (*Bos javanicus*). It was first identified in the Jembrana district of Bali island (Indonesia). Thus the causal agent is named Jembrana Disease Virus (JDV), a lentivirus member of the family Retroviridae (Kusumawati *et al.*, 2014a). The disease is now endemic through-out parts of Indonesia, particularly in Java, Sumatra, Kalimantan (Hartaningsih *et al.*, 1993) and also in Australia (Chadwick *et al.*, 1998). The disease development of Jembrana disease in Bali cattle is unique for a lentivirus infection as it is related with severe symptoms in an acute period (Dharma *et al.*, 1991). The mortality of more than 60 000 cattles in a year was observed during the first outbreak (Hartaningsih *et al.*, 1993). In experimentally JDV-infected Bali cattle, the death rate was about 21% (Soesanto *et al.*, 1990; Soeharsono *et al.*, 1990). This finding was supported with

evidence acquired in several epidemiological studies (Soeharsono *et al.*, 1995a; 1995b; Chadwick *et al.*, 1998). In lethal infection, mortality was occurred within only 1 to 2 weeks after infection and was correlated to multi-organ failure (Wilcox *et al.*, 1995; Wilcox, 1997). The economical aspect of Jembrana disease in Bali cattle in Indonesia is also important to be considered since these cattle have been widely spread throughout Indonesia. Bali cattle comprise about 27% of the total cattle population of Indonesia-the highest contribution to beef production in Indonesia (Desport and Lewis, 2010).

Virus-borne diseases are among the most difficult to overcome as drugs are usually expensive or not available in most cases. Early diagnosis constitutes a preventive method of further disease outbreaks. Bioassay provides a method for titration of infectious virus but it is time-consuming and expensive. Techniques that do not require the use of animals for assay are needed for easy and efficient detections and routine health controls of

JDV infections in cattle industry. Techniques that enable easy virus quantification are also needed for further studies on the kinetics of virus replication during the acute phase of the disease process and for understanding the persistence of virus in recovered animals. The Jembrana disease can be diagnosed by the clinical symptoms but more reliable diagnostic tools are available, based on either antigen (immunodiagnosis) or viral genome (molecular diagnosis). In this review, we summarize about immunodiagnostic tools of Jembrana disease which has been developed so far.

Clinical Symptoms

JDV infection is characterized by acute, severe disease syndrome in Bali cattle (Chadwick *et al.*, 1998). Briefly, the main clinical symptoms in Bali cattle include elevated body rectal temperature, lethargy, anorexia, diarrhoea with blood in feces, pallor of the mucus membrane, swollen of the superficial lymph nodes during the febrile phase, high titer of infectious virus in blood and secreted fluids (milk, saliva) (Soeharsono *et al.*, 1990; Soesanto *et al.*, 1990). In experimental infections, study showed that JDV had the capability to infect other type of cattles, including the most prevalent farmed cattle, such as Ongole cattle (*Bos indicus*) and Friesian cattle (*Bos taurus*), buffaloes (*Bubalus bubalis*), sheep, goats and pigs (Soeharsono *et al.*, 1990; Wilcox *et al.*, 1995). However, it is hard to establish the clinical diagnosis especially in other cattle species and pigs infected with JDV as it is only develop a mild febrile response (Wilcox *et al.*, 1995; Soeharsono *et al.*, 1990). The clinical symptoms are not always precise and may be difficult to establish at the initial time course of infection.

Antibody Response to JDV Infection

Lethal infection often occurs in JDV-infected cattle due to secondary infections (Dharma *et al.*, 1991), resulting from temporary immunosuppression characteristic to Jembrana disease (Wareing *et al.*, 1999). The temporary immunosuppression occurring during the acute phase was demonstrated by a decline of Immunoglobulin-G (IgG) containing cells in the lymphoid organs proven by immunohistochemistry method (Dharma *et al.*, 1994; Desport *et al.*, 2009a). This finding was supported with studies by Enzyme-Linked Immunosorbent Assay (ELISA) and Agar Gel Immunodiffusion (AGID) which showed a slowed antibody response (Desport and Lewis, 2010). JDV-specific antibodies were remained undetected until 11 weeks post-infection in most infected cattle. This antibody response was maximal at 23-33 weeks post-infection and was still detectable at 59 weeks post-infection (Hartaningsih *et al.*, 1994).

Immunodiagnostic Tool: Viral Detection Based on Antibody Response

Use of Whole Viral Particles

The first developed serodiagnosis used plasma-derived virus, purified by sucrose gradient, to detect antibody response in infected Bali cattle (Hartaningsih *et al.*, 1993; 1994). It is based on Enzyme-Linked Immunosorbent Assay (ELISA) and Agar Gel Immunodiffusion (AGID). This diagnostic method allowed to distinguish clinically-positive cattle for Jembrana disease, which gave a positive response, from cattle originated from Jembrana disease-free areas in Bali which were negatively responding (Hartaningsih *et al.*, 1994). According to this study, AGID was less sensitive than ELISA. The method has also been used to detect JDV infections in other parts of Indonesia, including Java (East provinces) and Sumatra (Lampung and West Sumatra) (Hartaningsih *et al.*, 1993). The drawback of the method is its usefulness for detecting JDV infection during the initial stage of disease development and during the acute phase because JDV infections induce a temporary immunosuppression in cattle, resulting in delayed antibody response (Dharma *et al.*, 1994; Hartaningsih *et al.*, 1994; Wareing *et al.*, 1999).

Optimization by Recombinant Antigens

Optimizing the serodiagnostic tool was further carried out by using recombinant JDV antigens, i.e., the Capsid (CA) and Transmembrane (TM) subunits of the respective *gag* and *env* ORF (Burkala *et al.*, 1998). Recombinant proteins were produced by a prokaryotic expression in *E. coli* as soluble proteins fused to Glutathione-S-Transferase (GST) (Burkala *et al.*, 1998). Using antisera from JDV-naturally infected Bali cattle (*Bos javanicus*) or Bovine Immunodeficiency Virus (BIV)-infected taurine cattle (*Bos taurus*) in Western blot analysis (Burkala *et al.*, 1998) and ELISA (Barboni *et al.*, 2001), it appeared that the recombinant JDV CA and TM reacted well with antisera from animals infected by JDV or BIV. This finding is indeed not surprising as the two bovine lentiviruses are genomically and antigenically very closely related (Chadwick *et al.*, 1995). Antibodies produced by infected cattle recognize antigens originating from both lentiviruses. Sequence comparison of the expressed regions showed 63% identity for TM and 66% for CA proteins of JDV and BIV. This identity is indicative of shared common immunogenic epitopes (Burkala *et al.*, 1998; Kertayadnya *et al.*, 1993). The lentivirus CA proteins contain a conserved epitope (Grund *et al.*, 1994) and the antigenic cross-reactivity can, to a certain extent, be attributed to this epitope. The TM cross-reactive epitope was not identified but is possibly the principal TM immunodominant domain.

Immunodiagnostic Tool: Detection of Viral Antigens

Improvement of the Diagnosis Efficacy by Antigen Capture ELISA

Immunodiagnostic tools are convenient and relatively easy to perform. However their sensitivity still needs to be increased, firstly for detecting low amounts of viral particles and also for their possible quantification. Improvement was achieved by antigen capture ELISA. The method is based on the recognition of JDV-p26 (capsid protein), prepared from plasma, by a specific monoclonal antibody coated to the well of ELISA tray. The captured antigen was then identified by rabbit anti-JDV-p26 antiserum (Stewart *et al.*, 2005). Though not allowing to distinguish JDV- from BIV-infections, the antigen capture ELISA is reliable for quantification of JDV-p26 over a linear range of 10 to 200 ng mL⁻¹ of plasma, allowing to correlate the antigen concentration with the number of viral RNA genome copies. Using this method, it has been shown that a peak of p26 concentration is attained during the acute phase and indeed at this stage the titer of infectious units is the highest as attested by analysis by real-time RT-PCR (Stewart *et al.*, 2005). During the acute phase, in plasma, the p26 concentration, the RNA copy number (Stewart *et al.*, 2005) and the infectious units (Soeharsono *et al.*, 1990) are respectively 3.5 µg mL⁻¹, 10¹² copies/ml and 0.5×10⁹ ID₅₀/mL. It proved so that antigen capture ELISA is not only highly sensitive but importantly it also enables to quantify the viral load in plasma and consequently to determine the stages of the disease development.

Differentiating Antigen

A study suggested a possibility distinguishing antibody to BIV and JDV (Barboni *et al.*, 2001). However attempts to locate JDV-specific antigenic determinants confirming this finding were unsuccessful. Various recombinant proteins have been tested but they were unable to differentiate antisera of JDV-infected from BIV-induced antisera (Desport *et al.*, 2005). Attempts to identify distinguishing epitopes are only made feasible by using monoclonal antibody approach. Indeed, a monoclonal antibody raised against BIV *gag* protein - a capsid protein, was found to only recognize BIV protein and not the JDV counterpart in Western blotting (Zheng *et al.*, 2001). The differentiating BIV *gag* epitope appeared to be the unique epitope that is not shared by JDV (Lu *et al.*, 2002). Further identification by chemical cleavage analysis, recombinant overlapping peptides and synthetic peptide showed that the specific epitope is 26 amino acids in length and composed of 6 C-terminal amino acids of the matrix

protein (p16^{MA}) followed by 20 N-terminal amino acid residues of the p2L peptide (Lu *et al.*, 2002). This epitope is absent in JDV because the intragenic p2L is missing in JDV (Lu *et al.*, 2002). The BIV-specific epitope is of great interest in confirming or disproving JDV infection by defect. Indeed, JDV infections will be characterized by a positive response in e.g., capsid-based detection but by a negative response to BIV-specific epitope monoclonal antibody (Zheng *et al.*, 2001; Lu *et al.*, 2002).

Future Direction

JDV antigen detection using ELISA was found to be more sensitive than Western blot (Barboni *et al.*, 2001; Lewis *et al.*, 2009). In addition, latest achievement method in immunodiagnosis is gain by JDVp26 capture ELISA that facilitated monitoring and detection of circulating viral antigen during the acute phase of the disease. An understanding of the use of a tyramide-based signal amplification substrate which allow the detection of little titer of HIV-1 p24 antigen may provide important clue to increase the sensitivity of the current method JDVp26 capture ELISA. Substantial progress can also be made by continuing research to improve JDVp26 antigen ELISA which may be affected by JDVp26 monoclonal antibodies (Stewart *et al.*, 2005). Furthermore, antibody-based diagnostic methods do not enable to distinguish JDV- from BIV-infection as the two bovine lentiviruses are antigenically very closely related (Fultz, 1991; Soeharsono *et al.*, 1990). Distinguishing BIV-infection was only made feasible by using a BIV-specific monoclonal antibody that only recognizes the unique BIV *gag* epitope, which is not shared by JDV (Lu *et al.*, 2002).

JDV infection induces a delayed humoral response (Dharma *et al.*, 1994; Hartaningsih *et al.*, 1994). The majority infected animals cultivate possible amount of antibodies to be detected only 6 weeks or more after recovery from the acute phase of the disease (Desport *et al.*, 2009b; Desport and Lewis, 2010; Tenaya *et al.*, 2012). This ensue severe disease as the limitation of immunodiagnosis that cannot be used in acute stage of the disease. Moreover, immunodiagnosis are comprised by cross-reactive epitopes in the CA protein of JDV and BIV (Burkala *et al.*, 1998; Kertayadnya *et al.*, 1993). During the acute phase, high titer of infectious JDV viral particles is found in plasma (Kusumawati *et al.*, 2014b). Viruses are also abundantly present in secreted fluids, namely milk and saliva (Soeharsono *et al.*, 1995b). This make viral antigen identification by molecular method is ideal detection tool in order to detect viral infection as early as possible during the course of the disease. However, molecular detection method is not always applicable as it requires a high technical skill and

expensive equipments. Several viral genome amplification tools have been developed for the identification of JDV, such as *in situ* dot-blot hybridization (Chadwick *et al.*, 1998), polymerase chain reaction (Stewart *et al.*, 2005) and loop-mediated isothermal amplification which was the more recently developed and was the more sensitive (Kusumawati *et al.*, 2015). Nevertheless, finding from a comparative study (immunological *vs* molecular detection methods) recommended that a combination of molecular detection and immunodiagnosis method is used for routine control of Jembrana disease in Bali cattle (Lewis *et al.*, 2009).

Conclusion

Immunodiagnosis is more convenient for use in routine health controls than bioassays or anapathological analyses even if these latter are suitable techniques for post mortem diagnosis. The methodologies rely on either the humoral immune response of JDV-infected cattle or the presence of JDV antigens. However, given that JDV-specific antibodies are not produced in most infected cattle until 11 weeks post infection (Dharma *et al.*, 1994; Hartaningsih *et al.*, 1994; Wareing *et al.*, 1999), immunodiagnosis based on host humoral response can not be used in early stages of the disease. Maximal production of antibodies only occurs at 23-33 weeks post-infection even if antibodies are still detectable at 59 weeks post-infection (Hartaningsih *et al.*, 1994). Detection of JDV infection is so preferably based on the presence of viral antigens in order to detect viral infection as early as possible during the course of the disease to prevent further disease outbreaks. Optimizing the serodiagnosis is achieved by antigen capture ELISA which also allows the quantification of JDV viral particles although the method is unable to distinguish JDV- from BIV-infection (Stewart *et al.*, 2005). Differentiating BIV infections is feasibly attained by the use of a specific monoclonal antibody, only recognizing BIV capsid protein epitope that is absent in JDV (Zheng *et al.*, 2001; Lu *et al.*, 2002). Though relatively insensitive, ELISA provides an economical and feasible method for monitoring the virus in the absence of more sensitive methods (Barboni *et al.*, 2001; Lewis *et al.*, 2009).

Acknowledgment

We acknowledge to Professor Sri Hartati, DVM, MSc, PhD for provided general support and aid.

Funding Information

This work partly funded by a grant from Directorate General Higher Education (DIKTI), Ministry of Education and Culture of Indonesia.

Author's Contribution

Asmarani Kusumawati: Participated in all acquisition of literatures, reviewed the literatures, coordinated and contributed to the writing of the manuscript.

Tenri Ashari Wanahari: Participated in all acquisition of literatures, reviewed the literatures and contributed to the writing of the manuscript.

Widya Asmara: Reviewed the literatures and provided the technical guidance in literature review study.

Surya Agus Prihatno: Reviewed the literatures and provided the technical guidance in literature review study.

Basofi Ashari Mappakaya: Contributed to the writing of the manuscript.

Bambang Hariono: Provided the technical guidance in literature review study.

Ethics

This study was approved by Research Ethics Committee, Faculty of Veterinary Medicine, Gadjah Mada University, Indonesia.

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