



Western Blot Analysis to Detect Cross-reaction in *Toxocara vitulorum* Protein with Anti-*Mecistocirrus digitatus* Serum

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ABSTRACT

Worm infections are found in livestock and can be transmitted to humans. *Toxocara vitulorum* is a worm species which commonly infected people. Cross-reaction among worms can generate false positive to establish helminthiasis diagnosis through antibody inspection. This study aimed to determine specific proteins that caused cross-reaction between *Toxocara vitulorum* antigen and anti-*M. digitatus* serum by using the western blot technique. In the present study, the whole worms extracted of *T. vitulorum* and *M. digitatus* have been used to make polyclonal antibodies from *M. digitatus* with Wistar rats as hosts. The cross-reaction between whole worm extract of *T. vitulorum* protein and anti-*M. digitatus* serum obtained 12 protein bands that each relative molecular mass (Mr) valued of 176, 124, 92, 68, 59, 47, 31, 29, 26, 16, 12, and 10 kDa. Cross-reaction occurred between *T. vitulorum* protein and anti-*M. digitatus*.

Key words: Cross-reaction, *Mecistocirrus digitatus*, Specific protein, *Toxocara vitulorum*, Western blot

INTRODUCTION

Worm infections can decrease productivity and cause inefficiency on livestock (Rast et al., 2014). The disease that causes low meat production is digestive tract infection by parasite worms, including *Toxocara vitulorum* worm, which can infect cows and buffalos (Mufidah et al., 2013).

Toxocarosis is a zoonosis disease because the larvae can cause Visceral larvae migrans and Ocular larva migrans in people. It can cause permanent eye damage in humans (De Souza et al., 2004). The larvae of *T. vitulorum* can be transmitted through colostrum and milk. The second larva becomes dormant in the non-definitive host. As the larvae migrate to the host system, it would be difficult to be diagnosed and controlled (Wickramasinghe et al., 2009).

Mecistocirrus digitatus is a blood-sucking cattle nematode located in the abomasum of infected ruminants and causing severe micro- and macroscopic lesions such as mucosal inflammation, hemorrhage, ulcers, and necrosis. The high pathogenicity of this nematode generates important losses in the cattle industry. *Mecistocirrus digitatus* has a wide distribution in Asia, Central, and northern South America, occurring mainly in young cattle (Fernex et al., 2014).

Each relative molecular mass (Mr) of *Mecistocirrus digitatus* protein profile were 107.74, 72.88, 64.68, 51.39, 47.1, 43.52, 38.55, 36.27, 32.97, 28.95, 25.8, 23.25, 20.95, 15.65, 12.85, and 9.93 kDa. On the other hand, the *T. vitulorum* protein profile were 224, 227, 198, 155, 104, 87, 73, 67, 58, 55, 51, 48, 43, 37, 26, 16, 13, and 11 kDa (Nugroho, 2012). *Mecistocirrus digitatus* and *T. vitulorum* proteins with the same Mr were 73, 51, 43, 26, 16, and 13 kDa. The protein diversity might cause cross-reaction; it is about those antibodies that could react with more than one antigenic determinant. The epitope-specific antibody could bind other epitopes that are not related but have the same structure. The concept of antigen-antibody bonds is similar to keys and padlocks that are mutually bound (Mayer, 2010).

Worm antigens injected into animals could affect the immune response and stimulate animal antibodies (Darmawi et al., 2013). Western blot has 3 methods such as nitrocellulose membrane used to bind nonspecific antibody, primary antibody for incubating process and secondary antibody used to antigen-antibody reaction (Irnidayanti et al., 2018; Lastuti et al., 2018). This study was conducted to determine the specific proteins that caused cross-reaction between *T. vitulorum* antigens and anti-*M. digitatus* serum by using the western blot technique.

MATERIALS AND METHODS

Ethical approval

All experimental protocols and procedures were approved by the Institutional Animal Care of Indonesia.

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The present study was conducted from December 2016 until April 2017 in Faculty of Veterinary Medicine Laboratory of Universitas Airlangga. The type of study was explorative research.

Tools

The tools utilized for this study included binocular microscopes (Olympus), Petri cup, trays, plastics, surgical instruments, tweezers, scissors, mortars, carmine staining sets, disposable syringe, tissue, plastic, object glass, glass cover, 2 mL microtube, microplate, nitrocellulose membrane, SDS-PAGE chamber running (Biometra), electrophoresis device, trans blotter (Biometra), Whatman paper, shaker, water bath, yellow tip, blue tip, 5 mL scale pipette, Eppendorf pipette. The main materials used in this study were Whole Worm Extract (WWE), which was from both male and female *Toxicara vitulorum* and *Mecistocirrus digitatus* worms, and six Wistar rats. On the other hand, the chemicals required included phosphate buffer saline (PBS, Merck), distillate water (Merck) in 70%, 85%, 95%, glycerin alcohol (Merck), Hung's I, Hung's II (Merck), Carmine seed solution (Merck), acid alcohol, alkaline alcohol, HCl (Gibco cat no. 1640-430 -1800).

The materials for SDS-PAGE with Biometra devices included Tris aminomethane electrophoresis buffer solution (Sigma cat no. 172-2051), glycine, and SDS (Biorad), acrylamide/bis-acrylamide (Sigma cat no. A-2917), Tris-HCl (Promega 608-274-4330), SDS (Sigma cat no. G-7403), TEMED (Sigma cat no. 7024), Ammonium persulfate (APS) (BioRad cat no. 161-0700, aquadest (Merck), glycerin (Merck), bromfenolblue (Merck), mercaptoethanol (Sigma cat no.M-7154), NaOH (Merck), NH₃ (Merck), citric acid (Merck), formaldehyde (Merck), and acetic acid (Merck). The materials which used for ELISA included 2µg/mL of *T. vitulorum* antigen, carbonate buffer, 4% of creamer, washing buffer (0.15 M NaCl, 0.05% Triton x-100, 0.02% NaN₃), IgG anti-mouse labeled enzyme alkaline phosphatase, substrate (2.7 mmol/l 4-nitrophenyl phosphate in 1 M diethanolamine; 0.5 M MgCl₂; 0.02% NaN₃; pH 9.8). The materials needed for the Western Blot test were the SDS-PAGE gel which has been running containing the analyzed protein, transfer buffer (Tris aminomethane, glycine, methanol, and aquades, pH 8.3, ethanol, 5% blotto, western blue, anti-*M. digitatus* serum, conjugate (IgG anti-mouse), and Alkaline Phosphatase (AP) substrate (Sigma cat no.AB0300).

Work procedures

Collection and identification of Toxicara vitulorum and Mecistocirrus digitatus

Toxicara vitulorum adult worms were obtained from cows' small intestine suffering from toxocariasis. *M. digitatus* worms were also extracted from cow abomasum obtained from an abattoir in Surabaya. *T. vitulorum* and *M. digitatus* worms were cleaned by putting them in Petri cup, which contained 10% PBS media with a pH level of 7.2 by using anatomical tweezers. The obtained *T. vitulorum* and *M. digitatus* worms were then carefully identified based on morphology to ensure each species (Kusnoto, 2008).

Homogenate preparation and worm protein level measurement of Whole Worm Extract

Each homogenate was made of four *T. vitulorum* worms and 100 *M. digitatus* worms, both male and female, manually ground using mortar. Then, the ground worms were put into a tube (15 mL) and suspended with 5 mL of PBS. The suspension was centrifuged at 4,000 rpm for 15 minutes. The pellets and supernatants were separated and the supernatant was stored at 2-8 °C. Then, the homogenate protein was concentrated, and the results of protein concentration were calculated based on the standard protein concentration curve (Maehre et al., 2018). The process was made based on the standard concentration absorbance by using a spectrophotometer with 590 nm wavelength (Bradford, 1976).

Preparation of polyclonal antibodies by Mecistocirrus digitatus

The polyclonal antibody was prepared by injecting *M. digitatus* homogenate into rats at the dose of 200 µg/rat. The injections were subcutaneous with the addition of Complete Freund's Adjuvant (CFA) with the ratio of 1:1 on the first injection, and Incomplete Freund's Adjuvant (IFA) in the booster injections. The re-injection was done three times with two weeks interval times. After the last booster was injected in the sixth weeks, 0.5 mL of each rat's blood (three blood collection with three days interval) was extracted from the tail to obtain the serum (Kusnoto et al., 2011).

Visualization of Toxicara vitulorum and Mrci.stocirrus digitatus protein homogenate

The visualization was carried out by using the SDS-PAGE technique with the composition of 12% separating gel and 5% stacking gel. Samples of *T. vitulorum* and *M. digitatus* homogenate were 15 µL. As a marker, a protein with a relative molecular period (Mr) ranging from 10 to 245 kDa was utilized to produce the Vivantis. The electrophoresis was set on 100 V and 40 mA current for an hour.

Indirect-ELISA

The 2 µg/mL of *T. vitulorum* antigen was diluted with carbonate buffer (50 mmol/L carbonate, pH 9.6). Then, the mixture was adsorbed on ELISA microplate at 100 µL/well and incubated at 4°C overnight. The blocking used 4% of creamer and was incubated at 37°C for an hour. The microplate was washed with washing buffer (0.15 M NaCl, 0.05% Triton x-100, 0.02% NaN₃) for 3 times. Then, the antibodies were inserted in each well, amounting to 100 µL and incubated at 37°C for an hour after being washed with a buffer for three times. After that, the microplate was washed

with washing buffer. Next, substrates were added (2.7 mmol/L 4-nitrophenyl phosphate in 1 M di-ethanolamine; 0.5 M MgCl₂; 0.02% NaN₃; pH 9.8) amounting to 100 µL/well and was incubated for 10-30 minutes in the dark room. The absorbance was read by using ELISA-reader at 405 nm wavelength.

Cross-reaction with western blot

A running process of the *T. vitulorum* worm homogenate was conducted with SDS-PAGE. Then, the gel containing the protein fragment was removed from the glass plate. After that, it was soaked for 40 minutes in Trans blot buffer and become ready to be transferred to the nitrocellulose membrane. Then, the gel was covered by three sheets of Whatman paper which previously soaked with a Trans blot buffer for 40 minutes and was placed perfectly flat. The transfer process was carried out with 100 V constant voltage and 40 mA current for 90 minutes. The nitrocellulose membrane was soaked with 5% blotto for 60 minutes. The membrane was then washed with TBS, and the nitrocellulose membrane blocking was done by adding 1% Creamer and 0.05% TBS-Tween. Next, it was incubated at room temperature overnight. Then, the nitrocellulose membrane was washed with 0.05% TBS-Tween five times. The nitrocellulose membrane was inserted in a Petri dish, containing anti-*M. digitatus* serum which has been diluted (50 times) with TBS-Tween and was incubated for an hour. It was washed again with 0.05% TBS-Tween and added with enzyme-labeled antibodies (conjugates) that had been diluted 1,000 times with TBS-Tween. Then, it was incubated at room temperature for an hour. After that, it was washed with 0.05% TBS-Tween. The membrane was placed on a Petri dish which had been filled with substrate and then incubated at room temperature in the dark room. The Petri dish was then shaken until the color was visible by applying western blue. To stop the process, after enough color was visible, the membrane was placed on a Petri cup containing non-deionized aquadest (Susana et al., 2019).

Data analysis

To calculate the Mr protein, a regression formula was applied to determine the Mr protein from running SDS-PAGE and Western Blot with the help of standard protein (marker, Vivantis) according to Mishra et al. (2017). The Mr protein calculation was determined through a regression equation between the Rf (retardation factor) value and the log Mr Data in the marker. The obtained formula could be linear, quadratic, or cubic and could be used to calculate the Mr in the sample by determining the sample Rf value (X) and relative molecular period (Y).

RESULTS AND DISCUSSION

The anti-*M. digitatus* serum was administered to the rats. The blood was taken two weeks after the last booster of anti-*M. digitatus* serum and then the serum was tested with indirect ELISA technique. The results of ELISA-reader at 450 nm wavelength indicated the results of anti-*M. digitatus* serum with *T. vitulorum* antigen as seen in table 1.

Based on table 1, it has been indicated that the OD serum value of anti-*M. Digitatus* with *T. vitulorum* antigen reached the lowest level by 0.376, the highest level was 0.530 and the average OD value in all tested samples tested were 0.450. The serum results were visualized with SDS-PAGE while the cross-reaction was demonstrated with Western blot techniques produced bands. After that, relative molecular mass (Mr) calculations were carried out with regression analysis.

The regression calculation analysis on Western blot techniques was conducted through a curve fit analysis to determine the shape of the correlation between Rf and log Mr at the marker. Based on the results of the analysis known that the relationship between the two variables, namely Rf and log Mr was followed by a regression equation. The results of cross-reaction in WWE *T. vitulorum* worms with anti-*M. digitatus* serum based on Western blot technique revealed in 12 protein bands as has been demonstrated in figure 1. Based on figure 1, there was a protein band with Mr 10 kDa under marker 11 kDa and above 12 kDa protein. Under marker 17 kDa, there was 16 kDa protein. Between marker 17 and 25 kDa, there were no cross-reacting proteins. Between marker 25 to 32 kDa, there were protein bands of 26, 29, and 31 kDa which were thinly expressed. Above the 46 kDa marker, there was a protein which has cross-reaction in 47 kDa which expressed thickest compared to other bands. Between marker 46 kDa and 135 kDa, there were cross-reaction proteins in Mr 47, 59, 68, 89, and 125 kDa. Between marker 135 kDa to 245 kDa, there was 176 kDa protein cross-reaction. *T. vitulorum* cross-reaction protein with anti-*M. digitatus* serum obtained 12 protein bands with Mr 176, 124, 92, 68, 59, 47, 31, 29, 26, 16, 12 and 10 kDa.

Table 1. The result of optical density *Mecistorricus digitatus* serum with *Toxocara vitulorum* antigen with indirect-ELISA on Wistar rats.

First blood collection (63 rd day)	Second blood collection (66 th day)	Third blood collection (69 th day)
0.396	0.434	0.507
0.451	0.432	0.479
0.427	0.507	0.518
0.411	0.449	0.483
0.386	0.530	0.451
0.376	0.448	0.417

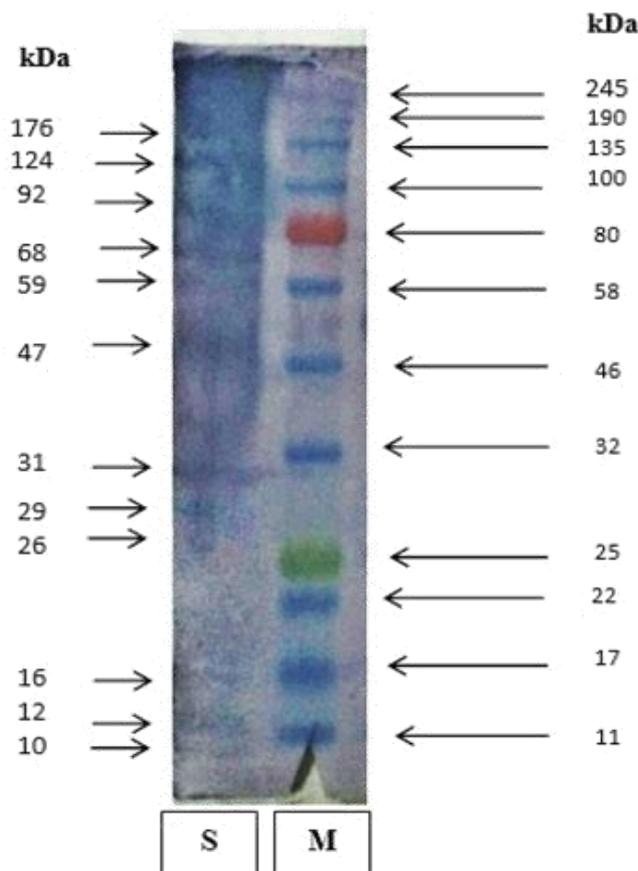


Figure 1. The results of protein cross-reaction from Whole Worm Extract (WWE) *Toxocara vitulorum* worms with anti-*Mecistocirrus digitatus* serum by using western blot techniques. M: Marker, S: *Toxocara vitulorum* and *Mecistocirrus digitatus* cross-reaction.

DISCUSSION

Cross-reaction Protein from WWE *T. vitulorum* worms with anti-*M. digitatus* serum by using the Western blot technique obtained 12 protein bands. The *T. vitulorum* worm protein profile by using Western blot technique indicated 10 similar protein bands between uterine antigen and body wall antigen, i.e., 20.3; 43.6; 46; 53.1; 57.4; 60.3; 66.9; 80.1; 111.3; and 126.3 kDa. The Mr values were recorded at 20.3; 24.1; 25.6; 33.7; 37.2; 43.6; 53.1; 57.4; 66.9; 105.5; 111.3; 126.3 and 133.9 kDa, which were the same protein bands between perienteric fluid antigen and *T. vitulorum* body wall antigen (Jyoti et al., 2011).

The calculation of Mr Protein by using the regression correlation formula might cause relative differences in determination of protein band distance, the length, and the beginning of gel measurement. Therefore, there was a risk of having different difference in actual weight (Kusnoto et al., 2011).

There was cross-reaction between *T. vitulorum* antigen and *Fasciola gigantica* antibodies on Mr 133 kDa protein. On the other hand, 143 kDa protein was recognized by *Moniezia expansa* antibody (Abdel-Rahman and Abdel Megeed, 2000). *T. vitulorum* WWE antigen with Mr 57.4 kDa was detected as an immune-dominant antigen with anti-*T. Vitulorum* serum by using Western blot techniques. The antigen 66.9 kDa was found in somatic *T. vitulorum* antigen. It might be a specific immune-dominant antigen because it was reported not to have a cross-reaction with *Haemonchus* spp. and *paramphistomum epiclitum* (Jyoti et al., 2011). Cross-reaction could occur if two antigens have the same or identical epitopes. Moreover, they may have specific antibodies for one epitope that were initially used to bind other unrelated epitopes. Nevertheless, having the same chemical properties could produce cross-reaction (Huebner, 2004). Antibodies which were formed in response to antigens from one parasitic species, could allow reaction with different parasitic species antigens that caused immunological cross-reaction (Noble and Noble, 1982).

Antibodies would be produced optimally when antigens enter the body. Chronic worm infection would be the cause of persistent antigens stimulation that have increased levels of immunoglobulins in immune complexes, circulation and formation. The defense against most of worm infections was performed by activation of Th2 cell (Abbas et al., 2019). Parasites have a larger size than bacteria and viruses, so they contain more antigens in both in terms of numbers and species (Delves and Roitt, 1998).

The existence of cross-reaction in worms was the basis of the polyvalent vaccines development. They were highly crucial components in cross-reaction that were expected to be useful as reagents (Hillyer, 1995). Immunodiagnostic, as an early diagnosis of worm infection had the potential to be deteriorated by non-specific and cross-reaction, due to the

distribution of antigen epitopes in several worms (Ghosh et al., 2005). Further research is required for purification with affinity chromatographic techniques. In addition, determining the sensitivity and specificity of antigens as candidates for latent toxocariasis immunodiagnosis cases in an adult cow is necessary. This research could lead to chemotherapy with some effective anthelmintic and control of disease in adult cows and calves (Jyoti et al., 2011).

The observations from Western blot indicated high coloring background caused by the unbound antibodies due to lack of washing. Double band visualization was caused by high primary and secondary antibody concentrations bound to specific proteins. Black spots on the blot were caused by antibodies bound to the blocking agent. Meanwhile, the white band on the black blots was possibly caused by too much secondary antibody (Kalanjati, 2011).

The proteins that revealed cross-reaction were not recommended to be used as diagnostic kit material, but they could be developed as vaccine candidates (Abdel-Rahman and Abdel Megeed, 2000). *T. vitulorum* specific proteins were with Mr 244, 227, 198, 155, 104, 87, 73, 51, 43, 37, 28, 13 and 11 kDa and because they were not recognized by *M. digitatus* antibodies, *Fasciola gigantica*, *Moniezia expansa*, *Haemonchus spp.* and *P. eplicitum*, known as specific proteins. The proteins that do not show cross-reaction are specific proteins (Susana et al., 2019). So, it is necessary to conduct an experiment with cross-reaction with other worm species as well as further research that could test the protein specificity. Purification methods will be required to obtain these specific proteins. These specific proteins could be used as a toxocariasis diagnostic kit material.

Purified proteins were better antigens for serological diagnosis than somatic antigens. The existence of cross-reaction that occurred in helminthiasis infection could overcome by the use of purified proteins. The use of serological diagnosis techniques on a large scale could be done by using serum obtained from farms, abattoir and experimental animals to determine the level of sensitivity and specificity of the test (Arora et al., 2010).

CONCLUSION

The cross-reaction between Whole Worm Extract (WWE) of *Toxocara vitulorum* protein and anti-*Mecistorricus digitatus* serum resulted in 12 protein bands with each Mr of 176, 124, 92, 68, 59, 47, 31, 29, 26, 16, 12 and 10 kDa. Suggestions for future research include examination of the cross-reactions from other nematodes, cestodes, and trematodes. Therefore, the test specificity will be high. It is necessary to purify *T. vitulorum* proteins to obtain specific band results to be used as diagnostic materials. Further research could determine the level of immunogenicity, antigenicity and sensitivity of proteins. The specificity of each protein could be used as a basis for selecting diagnostic materials.

DECLARATIONS

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Authors' contribution

Ir.H Kusnoto was the consulting advisor. Desy Meta Anggraini and Suryanie Sarudji appreciated the cooperation for taking care of fishes during the experiment period and helping in the process of experimental work. Ethical issues (Including plagiarism, consent to publish, misconduct, data fabrication and/or falsification, double publication and/or submission, redundancy, etc.) have been checked by the authors.

Conflict of interests

The authors declared there was no conflict of interests.

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