Production of monoclonal antibodies specific to *Macrobrachium rosenbergii* nodavirus using recombinant capsid protein

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ABSTRACT: The gene encoding the capsid protein of *Macrobrachium rosenbergii* nodavirus (MrNV) was cloned into pGEX-6P-1 expression vector and then transformed into the *Escherichia coli* strain BL21. After induction, capsid protein-glutathione-S-transferase (GST-MrNV; 64 kDa) was produced. The recombinant protein was separated using SDS-PAGE, excised from the gel, electro-eluted and then used for immunization for monoclonal antibody (MAb) production. Four MAbs specific to the capsid protein were selected and could be used to detect natural MrNV infections in *M. rosenbergii* by dot blotting, Western blotting and immunohistochemistry without cross-reaction with uninfected shrimp tissues or other common shrimp viruses. The detection sensitivity of the MAbs was 10 fmol µl⁻¹ of the GST-MrNV, as determined using dot blotting. However, the sensitivity of the MAb on dot blotting with homogenate from naturally infected *M. rosenbergii* was approximately 200-fold lower than that of 1-step RT-PCR. Immunohistochemical analysis using these MAbs with infected shrimp tissues demonstrated staining in the muscles, nerve cord, gill, heart, loose connective tissue and inter-tubular tissue of the hepatopancreas. Although the positive reactions occurred in small focal areas, the immunoreactivity was clearly demonstrated. The MAbs targeted different epitopes of the capsid protein and will be used to develop a simple immunoassay strip test for rapid detection of MrNV.

KEY WORDS: Capsid protein · Extra small virus · XSV · Immunohistochemistry · *Macrobrachium rosenbergii* nodavirus · MrNV · Monoclonal antibody · Western blot

INTRODUCTION

Outbreaks of disease due to *Macrobrachium rosenbergii* nodavirus (MrNV) were first reported from hatchery-reared *M. rosenbergii* postlarvae (PL) in the French West Indies (Arcier et al. 1999, Bonami & Sri Widada 2011) and later in Taiwan (Wang et al. 2008), China (Qian et al. 2003), India (Sahul Hameed et al. 2004), Thailand (Yoganandhan et al. 2006) and Australia (Owens et al. 2009). MrNV infection may cause nearly 100% mortality of PL in hatcheries and nursery ponds within 5 d after appearance of the first gross signs of the disease. These signs include whitish appearance of muscle tissue, particularly in the tail portion, leading to the name white tail disease (WTD).
Genome-based detection methods with high specificity and high sensitivity for detection of MrNV include dot blot hybridization, in situ hybridization (Sri Widada et al. 2003) and 1-step RT-PCR in the form of single tests for MrNV or XSV (Sri Widada et al. 2003, Sahul Hameed et al. 2004) or multiplex tests for both viruses (Yoganandhan et al. 2005, Tripathy et al. 2006, Senapin et al. 2010). A 2-step real-time RT-PCR method (Zhang et al. 2006) and a reverse transcription loop-mediated isothermal amplification (RT-LAMP) method (Pillai et al. 2006, Puthawibool et al. 2010) have also been described. Although a thermal cycler is not required for RT-LAMP, the technique is still expensive and fairly complicated, and a laboratory is required. Therefore, these molecular-based techniques are not useful for pond-side detection by shrimp farmers, who require simpler and quicker disease monitoring and disease-outbreak confirmation methods that are easy to perform with high specificity and optimal sensitivity. These objectives may be achieved using immunoassay methods. Use of a polyclonal antibody (PAb) (Romestand & Bonami 2003) and monoclonal antibodies (MAbs) (Qian et al. 2006) to detect purified MrNV using sandwich ELISA and Western blotting (Sahul Hameed et al. 2011) have been described. Although the cost per sample of these antibody-based detection methods is much lower than that of genome-based detection methods, these methods still require a well-equipped laboratory.

In the present study, MAbs specific to MrNV were developed by mouse immunization with recombinant capsid protein of MrNV expressed heterologously in *Escherichia coli*. These MAbs were selected and characterized using various immunological methods including dot blotting, Western blotting and immunohistochemistry. The objective was to obtain MAbs that targeted a variety of MrNV epitopes for use in the development of a simple and rapid strip test that required at least 2 MAbs, one as a captured antibody and another as a labeled antibody, that recognized different epitopes of the antigen in sandwich format (Sithigorngul et al. 2006, 2007, 2011).

**MATERIALS AND METHODS**

**Sample preparation**

*Macrobrachium rosenbergii* PL samples infected naturally with MrNV preserved in 70% ethanol at 4°C for 3 d were obtained from a shrimp hatchery in Chachoengsao Province, Thailand. The preservative solution was changed to 95% ethanol before storage at −70°C prior to use. Part of each sample batch was homogenized in PBS (0.15 M phosphate-buffered saline, pH 7.2) at approximately 1:10 (V:V) and centrifuged at 3000 × g for 30 min. The supernatant was aliquoted into small vials and stored at −70°C. Another part of each sample batch was fixed in Davidson’s fixative and processed for immunohistochemical analysis. Uninfected *M. rosenbergii* PL were also collected and used as negative controls in various assays. Adult *M. rosenbergii* naturally infected with MrNV-were obtained from the local market in Bangkok and also fixed in Davidson’s fixative for use in immunohistochemical assays. Nested RT-PCR for MrNV and XSV (Senapin et al. 2010) was used to verify the presence of both viruses in the samples.

**MrNV RNA preparation**

MrNV-infected *Macrobrachium rosenbergii* PL were homogenized in lysis buffer (50 mM Tris–HCl pH 9, 100 mM EDTA, 50 mM NaCl, 2% SDS), and RNA was extracted from 200 µl of the homogenate using a High Pure viral nucleic acid kit (Roche Molecular Biochemicals) as described in the product manual. The extracted nucleic acid was stored at −70°C.

**Cloning and expression of MrNV capsid protein gene**

RNA from MrNV-infected shrimp was reverse transcribed into cDNA using Superscript III reverse transcriptase (Invitrogen) following the manufacturer’s directions. Polymerase chain reaction (PCR) was performed to amplify the MrNV capsid protein gene without the corresponding transmembrane region at the N-terminus (nucleotides 91 to 1116 of the open reading frame [ORF] of MrNV capsid protein gene; GenBank accession number NC005095.1), using this cDNA as a template, *Pfx* polymerase (Invitrogen) and the primers MrNVF91 (5'-CG GGA TCC CCG CAG ACG GTT CCC AAC-3') and MrNVR (5'-G GAA TTC CTA ATT ATT GCC GAC GAT AG-3') containing added restriction sites (underlined). The PCR conditions were 94°C for 2 min and 35 cycles of 94°C for 15 s, 57°C for 30 s and 68°C for 1 min, followed by an extension at 68°C for 15 min. The 1041 bp PCR product was digested with restriction enzymes *Bam*HI and *Eco*RI and then cloned into the corresponding restriction sites of the pGEX-6P-1 expres-
sion vector and transformed into *Escherichia coli* strain BL21. The integrity of the ORF of the recombinant plasmid was verified using DNA sequencing with 5′pGEX sequencing primer (5′-GGG CTG GCA AGC CAC GTT TGG TG-3′). For comparison, a pGEX-6P-1 expression vector without the inserted gene was also concurrently transform the *E. coli* strain BL21 with the MrNV-pGEX-6P-1 plasmid.

**Preparation of recombinant MrNV capsid protein**

*Escherichia coli* transformed with MrNV-pGEX-6P-1 was cultured in Luria-Bertani (LB) broth to the exponential phase. Expression of recombinant protein was induced with 1 mM isopropyl-β-D-thiogalacto-pyranoside for 4 h. After centrifugation at 3000 × g for 20 min at room temperature, the bacterial pellet was resuspended in a buffer containing 100 mM NaH2PO4, 10 mM Tris-HCl, 8 M urea (pH 8) and 1 mM phenylmethylsulfonyl fluoride and sonicated until a clear lysate was obtained. The lysate was separated using 12% sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE). After treatment with 0.3 M KCl, the band of recombinant fusion protein called glutathione-S-transferase (GST)-MrNV was excised and collected in dialysis bags. The recombinant protein solutions were eluted with a Transblot apparatus (BioRad) at 70 V for 6 h and dialyzed in PBS. The protein concentration was determined using the Bradford assay (Bradford 1976). The recombinant protein solutions were adjusted to 0.4 mg ml−1, divided into 0.5 ml aliquots and stored at −70°C.

**Production of monoclonal antibodies**

A cell fusion protocol was adapted from the method developed by Köhler & Milstein (1976) with modifications described by Mosmann et al. (1979). A P3X myeloma cell line was used as the fusion partner. Fusion products from 1 mouse were plated onto 20 microculture plates (96 wells per plate). The identification of positive cultures was performed by dot blotting against GST and GST-MrNV, by dot blotting and Western blotting against homogenates from MrNV-infected and uninfected PL and by immunohistochemistry, as described below. The selected hybridoma clones were recloned twice using the limiting dilution method to establish cell lines that were stored in liquid nitrogen.

The class and subclass of the mouse immunoglobulins produced by hybridomas were determined using sandwich ELISA with a Mouse MonoAb ID Kit: HRP (Zymed).

**Specificity testing**

**Dot blotting**

Lysates of *Escherichia coli* BL21 containing either the GST, GST-MrNV proteins or homogenate samples from uninfected and MrNV-infected PL were applied to nitrocellulose membranes (1 µl per spot) that were baked at 60°C for 10 min and then incubated in MAb in hybridoma-conditioned culture medium from each culture well at a dilution of 1:20 in 1% blocking solution (1% nonfat dry milk, 0.1% Triton X-100 in PBS) for 5 h. After extensive washing in 0.5% blocking solution, the membrane was incubated in horseradish-peroxidase-labeled goat anti-mouse gamma immunoglobulin heavy and light-chain-specific antibody (GAM-HRP; Bio-Rad) at 1:1500 dilution in 1% blocking solution for 3 h. The membrane was then washed for 5 min in 0.5% blocking solution and incubated in substrate mixture containing 0.03% diaminobenzidine (DAB), 0.006% hydrogen peroxide and 0.05% cobalt chloride in PBS (Sithigorngul et al. 2002).

**Western blotting**

Lysates of *Escherichia coli* BL21 containing either the GST or GST-MrNV proteins or homogenate samples from uninfected and MrNV-infected PL were separated using 12% gel SDS-PAGE according to the method described by Laemmli (1970). The samples
were electrophoresed for 3 h at 60 V, and one part of the gel was stained using Coomassie Brilliant Blue R-250. For Western blot analysis, the samples resolved using SDS-PAGE were transferred onto nitrocellulose membranes using a Transblot apparatus (BioRad). The nitrocellulose membranes were incubated in 5 % blocking solution for 10 min and treated with MAbs in culture medium (1:200 dilution) or mouse anti-GST-MrNV capsid protein antisera (1:5000 dilution and preabsorbed with E. coli lysate containing GST) for 5 h. After extensive washing in 0.5 % blocking solution, the membranes were incubated with GAM-HRP at 1:1500 dilutions for 3 h. They were then washed extensively as before and incubated in a substrate mixture containing 0.006 % hydrogen peroxide, 0.03 % DAB and 0.05 % cobalt chloride in PBS.

**Immunohistochemistry**

Alcohol-preserved PL specimens infected naturally with MrNV were further fixed in Davidson’s fixative solution for 24 h before processing for paraffin sectioning. Serial sections of tissues (8 µm thickness) were prepared and processed for indirect immunoperoxidase staining using MAbs obtained from the present study at a 1:100 dilution and then GAM-HRP at a 1:1000 dilution in 10 % calf serum in PBS for 5 h at 37°C each step. After extensive washing with PBS, peroxidase activity was visualized by incubation with 0.03 % DAB and 0.006 % hydrogen peroxide in PBS for 5 min. The preparations were counterstained with hematoxylin and eosin Y (H&E Stains), dehydrated in a graded ethanol series, cleared in xylene and mounted in Permount (Sithigorngul et al. 2000). Positive reactions were visualized as brown coloration against pink cytoplasm and purple nuclei. Pieces of tissues dissected from adult Macrobrachium rosenbergii infected naturally with MrNV were also directly fixed in Davidson’s fixative solution for 24 h and processed using the same protocol as the alcohol-preserved PL.

Cross reactivity of the MrNV-specific MAbs to various shrimp viruses was determined using immunohistochemistry. Shrimp samples infected with Penaeus monodon densovirus (PmDNV; also called hepatopancreatic parvovirus), Penaeus monodon nucleopolyhedrovirus (PemoNPV; also called monodon baculovirus), Taura syndrome virus (TSV), white spot syndrome virus (WSSV), yellow head virus (YHV) and infectious myonecrosis virus (IMNV) were processed for paraffin sectioning and immunohistochemistry using MAbs specific to MrNV. The results were compared to those from MAbs specific to PmDNV (Rukpratanporn et al. 2005), TSV (Longyant et al. 2008), WSSV (Chaivisuthangkura et al. 2004), YHV (Sithigorngul et al. 2002), PemoNPV (Boonsanongchokying et al. 2006), Penaeus stylirostris densovirus (PstDNV; Sithigorngul et al. 2009) and IMNV (Kunanopparat et al. 2011).

**Sensitivity testing with recombinant MrNV capsid protein**

Purified GST-MrNV protein was serially diluted with PBS, and 1 µl of each dilution was spotted onto nitrocellulose membranes and processed for dot blotting using the MrNV-specific MAbs generated in the present study. The last dilution yielding a clear positive result was determined.

**Sensitivity comparison between MAbs and 1-step RT-PCR**

The sensitivity of MrNV detection in a shrimp infected naturally was determined using MAbs specific to MrNV. The MrNV-infected shrimp PL homogenate was diluted serially with uninfected shrimp homogenate and processed for dot blotting as described above. The last dilution of shrimp homogenate yielding a clear positive result was determined. RNA from the same PL homogenate was also extracted with a High Pure viral nucleic acid kit (Roche Molecular Biochemicals), serially diluted with uninfected shrimp nucleic acid and tested for MrNV using single step RT-PCR with primers MrNVF91 and MrNVR as described above. The RT-PCR conditions were 50°C for 30 min, 95°C for 5 min, 4°C for 5 min and 35 cycles of 95°C for 1 min, 55°C for 45 s and 72°C for 1 min, followed by an extension at 72°C for 15 min.

**RESULTS**

**Capsid protein gene cloning and expression**

The expected PCR amplicon of 1041 bp was obtained for the MrNV capsid protein gene (Fig. 1). Expression of the GST-tagged-MrNV fusion protein (GST-MrNV) was visualized by Coomassie Brilliant Blue staining as a band with the expected molecular mass of 64 kDa (Fig. 2A, Lane 3). After the recombinant band was cut from the gel and eluted, a highly purified fusion protein was obtained (Fig. 2A, Lane 4) and used at 0.4 mg ml⁻¹ protein for immunization.
Monoclonal antibody production

After the fourth immunization, the mouse that produced the strongest immunoreactivity was used for hybridoma production (Fig. 2B). Approximately 1500 hybridoma-containing wells were obtained, and ~20 wells gave positive binding results when initially screened with GST-MrNV. Hybridomas producing 4 MAbs specific to MrNV with high avidity to GST-MrNV but with different characteristics, classes, avidities and abilities for use in immunohistochemistry were selected and cloned to establish cell lines. All MAbs belonged to the IgG class (Table 1). Because MAbs MrNV-2 and MrNV-12 gave the best results in the various immunoassays described above, MAb MrNV-12 was used as a representative MAb for further experiments.

Specificity of the monoclonal antibody

In dot blotting, MAb MrNV-12 bound and gave an intense spot with *Escherichia coli* lysates containing GST-MrNV and with homogenates from MrNV-infected PL but did not bind to *E. coli* lysate containing GST or homogenate from uninfected PL (Fig. 3). Similar results were obtained using Western blotting.
with MAb MrNV-12. The MAb bound to GST-
MrNV (a single band at 43 kDa) in homogenate
from MrNV-infected PL but did not bind to
E. coli
lysate containing GST or homogenate from unin-
fected PL (Fig. 2C). In the immunohistochemi-

cal analysis, 2 MAbs, MrNV-12 and MrNV-2, gave sim-
ilar results, with strong immunoreactivities in
infected muscles, gills, the heart, loose connective

tissue, the nerve cord and the intertubular cells
of the hepatopancreas of MrNV-infected shrimp
(Fig. 4). None of the MAbs exhibited cross-reactivity
to tissues from uninfected shrimp or shrimp infected
with IMNV, WSSV, YHV, TSV, PmDNV, PstDNV or
PemoNPV (data not shown). Of the other 2 MAbs,
MrNV-15 gave a poor staining, and MrNV-24 did
not give any staining for immunohistochemical
localization of the viral infection (Table 1). Despite
their lower avidity, the poor staining may have been
the result of target epitope modification or epitope
inaccessibility that can occur during the processes
of preparing paraffin sections.

### Sensitivity testing with recombinant MrNV
capsid protein

To determine the sensitivity of the MrNV-specific
MAbs, dot blotting against purified recombinant
GST-MrNV protein was performed. The detection
limit of the first 2 MAbs, MrNV-2 and MrNV-12, was
approximately 600 ng ml⁻¹, which is equivalent to
600 pg µl⁻¹ or 10 fmol µl⁻¹. The other MAbs, MrNV-
15 and MrNV-24, gave 2-fold less detection sensitiv-
ty (Table 1).

### Sensitivity comparison between MAbs and
1-step RT-PCR

To compare the sensitivity of MrNV detection
between dot blotting and 1-step RT-PCR, parallel,
diluted homogenates from naturally infected PL
were used for both assays. In the dot blot assay, the
lowest limit of detection of each MAb, MrNV-2 and
MrNV-12, was at a 1:320 dilution. However, when
both MAbs were combined, the detection sensitivity
limit increased to a 1:640 dilution (Fig. 5A–C). The
other 2 MAbs, MrNV-15 and MrNV-24, gave similar
results but with one dilution-step lower sensitivity
(Fig. 5D–F). The improvement in sensitivity with
combined pairs of MAbs demonstrated that they
bound to non-overlapping epitopes of the antigen.
However, a combination of all 4 MAbs did not yield
any further significant increase in detection sensitiv-
ty (not shown).

In 1-step RT-PCR detection using the same PL
homogenate, an amplicon band (1041 bp) was still
observable at a dilution of 10⁻⁵ (Fig. 6). Therefore, in
comparison with 1-step RT-PCR, the dot blot assay
was approximately 200-fold less sensitive.

### DISCUSSION

Four MAbs specific to MrNV capsid protein were
produced from a mouse immunized with a fusion
GST-MrNV capsid protein. These MAbs could be
used for detection of MrNV infection in PL using dot
blotting, Western blotting and immunohistochem-
istry. MrNV infections have been localized using in
situ hybridization only in striated muscle (Sri Widada

<table>
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<th>MAb (subclass)</th>
<th>Sensitivity (fmol µl⁻¹)</th>
<th>Western blot</th>
<th>IHC reactivity</th>
<th>Cross-reactivity</th>
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<tr>
<td>MrNV-2 (IgG2a)</td>
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<td>+++</td>
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<tr>
<td>MrNV-12 (IgG2b)</td>
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<td>+++</td>
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<tr>
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<td>+++</td>
<td>+</td>
<td>None</td>
</tr>
<tr>
<td>MrNV-24 (IgG1)</td>
<td>20</td>
<td>+++</td>
<td>–</td>
<td>None</td>
</tr>
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Table 1. Monoclonal antibodies (MAbs) obtained from a
mouse immunized with GST-MrNV. IHC: immunohisto-
chemistry; –: no immunoreactivity; +: light immunoreactiv-
ty; +++: strong immunoreactivity.
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et al. 2003, Bonami & Sri Widada 2011). Even though immunohistochemistry is simpler and easier for handling than in situ hybridization, and both MAbs and PAbs against MrNV capsid protein were produced for various types of ELISA (Romestand & Bonami 2003, Qian et al. 2006, Sahul Hameed et al. 2011), immunolocalization of the MrNV in other infected tissues has not been reported previously using these 2 methods. In the present study, we used the MAbs to localize MrNV infection in additional tissues, including the nerve cord, loose connective tissue, intertubular tissue of the hepatopancreas, gills and the heart, resulting in an overall tissue distribution that was similar to those for infections of PvNV (Tang et

Fig. 4. Examples of immunohistochemical analysis using MAb MrNV-12. MrNV immunoreactivity was observed in adult *Macrobrachium rosenbergii* (A) muscle and (B) connective tissue and *M. rosenbergii* postlarvae (C) heart muscle, (D) nerve cord, (E) gills and (F) intertubular tissue of the hepatopancreas. Positive immunohistochemical reactions appeared as brown staining. Preparations were counterstained with hematoxylin and eosin (A,B,D) or only eosin (C,E,F). Arrowhead: light infection; m: striated muscle; n: nerve cord; t: thoracic ganglion
al. 2007) and IMNV (Tang et al. 2005, Kunanopparat et al. 2011) in *Penaeus vannamei*, except that the MrNV infection in heart muscle was very rare (2 out of 15 samples) and the immunoreactivity occurred in the heart muscle cells rather than in the fixed hemocytes. The MrNV immunoreactivity in the heart muscle was specific because several nearby uninfected heart muscle fibers did not show MrNV immunoreactivity (Fig. 4C). In addition, the immunoreactivity in infected heart muscle fibers co-localized with the MAb specific to XSV (data not shown). Immunohistochemical detection could be observed even though the infection was located in very small focal areas or few cells (Fig. 4D,F). In some places, the immunoreactivity was observed around the cells or in the hemolymph. This phenomenon may have been a preparation artifact because the sample was collected in 70% ethanol before being fixed in Davidson’s fixative. This may have caused the release of the virions from the infected cells into the surrounding tissues and hemolymph.

Purification of the recombinant GST-MrNV capsid protein using SDS-PAGE and electro-elution is a convenient method. This method could provide a sufficient amount of any recombinant protein, including intein-tagged or histidine-tagged recombinant proteins (Chaivisuthangkura et al. 2004, Kunanopparat et al. 2011) or natural capsid proteins of MrNV and XSV (Sahul Hameed et al. 2011). The
The avidity of 2 of our MAbs (MrNV-2 and MrNV-12) were 2-fold higher than the other 2 (MrNV-15 and MrNV-24). The detection sensitivity of the first 2 was 10 fmol µl⁻¹, as determined using dot blotting. In a sandwich ELISA using a PAb for MrNV, the lowest detection limit was 10 µg ml⁻¹ (10 ng ml⁻¹) of viral preparation (Romestand & Bonami 2003), and in a triple MAb sandwich-ELISA, the detection limit was 0.98 ng for purified MrNV and 1.2 µg total protein ml⁻¹ for infected PL homogenate (Qian et al. 2006). The sensitivity among these 3 MrNV assays could not be precisely compared due to differences in the purity and volume of antigen preparations used. However, the Western blot technique and indirect ELISA using PAb against MrNV or XSV could be used to detect both viruses in experimentally infected PL at 24 h post infection; these results were confirmed by results from 1-step RT-PCR (Sahul Hameed et al. 2011).

Although the sensitivity of the antibody-based assay was poorer than that of PCR-based methods, the immunobased assays, including dot blotting, offer some advantages for farm management of viral infection risks, such as WSSV. Patil et al. (2008) compared the use of PCR tests and an MAb-based immunodot test for detecting WSSV in Penaeus monodon, and both were found to have value. For example, the lower detection of the immunodot test proved to be instructive regarding the likelihood of disease occurrences: all 6 farms at which WSSV was detected at various times post stocking had failed harvests. Conversely, the very high sensitivity of PCR can be instructive regarding the unlikelihood of disease occurrences: at 4 farms at which WSSV was only detected at various times following the use of a more sensitive 2-step PCR, shrimps were harvested successfully at 105 d post stocking (Patil et al. 2008). Similar PCR evidence of the long-term persistence of WSSV infections in the absence of disease, unless environmental stressors trigger elevated virus replication leading to the disease, has been obtained in other studies (Peng et al. 1998, Tsai et al. 1999). Therefore, the lower sensitivity of various immunoassays would provide shrimp farmers with a lower cost per test, while still being an effective diagnostic tool for monitoring viral infection, predicting impending disease and hence managing against production losses.

The sensitivity of the MAbs obtained in the present study (600 pg µl⁻¹ or 10 fmol µl⁻¹) was in the range of sensitivity limits reported for various other shrimp viruses, such as WSSV at 1.2 to 5 fmol per spot for VP19 (Chaivisuthangkura et al. 2010a), 5 fmol per spot for VP28 (Chaivisuthangkura et al. 2004), 500 pg for VP28 (Anil et al. 2002), 400 pg for VP28 (Liu et al. 2002) and 625 pg per spot for VP28 (Makesh et al. 2006). For TSV, the detection limit per spot for MAbs against VP3 was 400 to 800 pg (Longyant et al. 2008) and against VP2 was 400 to 800 pg of VP2 (Chaivisuthangkura et al. 2010b), while that for the capsid protein of PstDNV was 300 pg (Sithigorngul et al. 2009). Because the MAbs targeted different epitopes on the MrNV capsid protein, they are expected to be useful for the development of an immunochromatographic strip test for simple and rapid detection of MrNV infection, as has been reported previously for WSSV and YHV (Sithigorngul et al. 2006, 2007, 2011).

In conclusion, MAbs specific to MrNV were generated from a mouse immunized with recombinant GST-MrNV capsid protein. The MAbs could be used to detect MrNV infection by Western blotting in a manner similar to that used for detection using 1-step RT-PCR in time-course infectivity studies. The MAbs were also used to expand the list of target tissues for MrNV infection. Because these MAbs target different capsid protein epitopes, they are promising for use in the development of an immunochromatographic strip test for simple and rapid detection of MrNV.

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