Preferential suppression of yellow head virus (YHV) envelope protein gp116 in shrimp that survive challenge with YHV

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ABSTRACT: The DNA sequence that encodes the first 406 amino acid residues at the N-terminus of yellow head virus (YHV) protein gp116, namely N/2 gp116ΔTM, and the DNA sequence that encodes the next 392 amino acid residues at the C-terminus of gp116 (without the transmembrane region), namely C/2 gp116ΔTM, were cloned into pGEX-6P-1 plasmid and expressed in E. coli. Both recombinant proteins were expressed, purified by SDS-PAGE and used to immunize mice. The mouse anti-recombinant N/2 gp116 and C/2 gp116 antisera bound specifically to both the recombinant proteins and to natural gp116 protein in YHV-infected haemolymph as shown by Western blotting and in tissues as shown by immunohistochemistry. Immunohistochemical localization of YHV using anti-gp116 antiserum or monoclonal antibodies specific to gp116 (V3-2B), gp64 (Y18) and p20 (Y19) revealed similar immunoreactivity patterns for all these reagents in muscle and mandibular tissue in shrimp showing gross signs of yellow head disease. However, in gill, hepatopancreas, lymphoid organ and thoracic ganglion tissues from experimental YHV-infected shrimp (Penaeus vannamei and Palaemon serrifer) that did not show signs of disease, immunoreactivity to gp116 was reduced or absent while that for gp64 and p20 remained intense. Thus, some shrimp species were able to selectively inhibit the synthesis of gp116 in a manner that was associated with absence of gross signs of disease.

KEY WORDS: Penaeus vannamei · Palaemon serrifer · Penaeus monodon · Polyclonal antibody · Yellow head virus (YHV)

INTRODUCTION

Yellow head virus (YHV) is a highly pathogenic virus that causes mass mortality in Penaeus monodon and a variety of other penaeid shrimp (Boonyaratpalin et al. 1993). Infection in P. monodon has been reported in southeast Asian and Indo-Pacific countries such as Thailand (Flegel et al. 1997), India (Mohan et al. 1998) and Taiwan (Wang & Chang 2000). Along with the closely related virus, gill associated virus (GAV) from Australia (Cowley et al. 1999, 2000), YHV is classified in the new Family Roniviridae, genus Okavirus, within the Order Nidovirales (Cowley et al. 2000, Sittidi-lokratna et al. 2002). The YHV virions contain 3 major structural proteins with estimated molecular masses of 116 kDa (gp116), 64 kDa (gp64) and 20 kDa (p20) (Jitrapakdee et al. 2003), formerly identified as 135, 67 and 22 kDa proteins, respectively (Nadala et al. 1997). The glycoproteins gp116 and gp64 are proteolytic products of a polypeptide encoded by a gene in ORF3 of the YHV genome (Jitrapakdee et al. 2003). Recently, monoclonal antibodies (MAbs) specific to gp116 (V3-2B), gp64 (Y18) and p20 (Y19) were produced (Sithigorngul et al. 2000, 2002), and used for detection of
YHV infection. By means of immunohistochemistry, MAbs Y18 and Y19 were shown to react strongly with P. monodon tissues infected with YHV or GAV, while MAb V3-2B reacted with YHV only (Soowannayan et al. 2003). A similar pattern of reaction was also observed in palaeonimid shrimp experimentally infected with YHV that showed different degrees of tolerance to the YHV infection (Longyant et al. 2005). Phage display analysis revealed that MAb V3-2B recognized an epitope in the first 25 amino acid residues from the N-terminus of gp116 that was either missing from or different in GAV (Jitrapakdee et al. 2003). From our earlier studies in experimentally YHV-infected Palaeomon serrifer, Palaeamon styliferus, Macrobrachium lanchesteri, Metapenaeus affinis and Metapenaeus brevicornis, immunoreactivities against gp64 (using MAb Y18) and p20 (using MAb Y19) were easily observed, whereas immunoreactivity against gp116 (using MAb V3-2B) was very weak and inconsistent (Longyant et al. 2005, 2006). However, it was difficult to conclude with confidence whether lack of MAb V3-2B immunoreactivity was due to lack of gp116 in infected shrimp tissues or due to altered gp116 epitopes. To answer this question, we prepared polyclonal antibodies to different epitopes of gp116 and used these to study gp116 expression by immunohistochemistry. Specifically DNA sequences encoding the first 406 amino acid residues at the N-terminus (N/2 gp116ΔTM) and the next 392 amino acid residues at the C-terminus (C/2 gp116ΔTM) of gp116 were cloned and expressed in E. coli. The polyclonal antiserum raised against the expressed recombinant proteins were used to locate gp116 in YHV-infected shrimp.

**MATERIALS AND METHODS**

**Monoclonal antibodies.** MAb V3-2B specific to the gp116 envelope protein (Sithigorngul et al. 2000), MAb Y18 specific to the gp64 envelope protein and MAb Y19 specific to the p20 structural protein of YHV (Sithigorngul et al. 2002) were used for comparison with the polyclonal antibodies.

**Preparation of YHV infected haemolymph and tissues from Penaeus monodon.** Penaeus monodon (15 to 25 g), purchased from farms near Bangkok, were maintained in 10 ppt diluted seawater. At 2 to 3 d after inoculation with YHV, shrimp that showed slow movement were selected for antibody analysis. MAb Y19 was injected into P. monodon at approximately 2 µl g–1 body weight. At 2 to 3 d after inoculation with YHV, shrimp that showed slow movement were selected for collection of haemolymph. Collected haemolymph was mixed with anticoagulant (Rodriguez et al. 1995) at the ratio of 1:1. The haemolymph from each shrimp was tested for YHV by dot blottting using MAb Y19 as described previously by Sithigorngul et al. (2002). The strongly YHV-positive haemolymph samples were pooled and stored at –70°C until used.

**Viral preparation.** The virus in the pooled haemolymph was concentrated using a 33% saturated solution of ammonium sulfate. The resulting precipitate was harvested by centrifugation at 3000 × g for 15 min and the pellet was re-suspended in 2× PBS and then centrifuged at 100,000 × g for 2 h.

**RNA isolation and cDNA synthesis.** RNA from the partially purified virus was isolated using an RNaseasy kit (Qiagen) as described in the product manual. cDNA was synthesized using Superscript II reverse transcriptase (Invitrogen), oligo d(T) primer and E. coli RNase H (Sambrook & Russell 2001).

**PCR amplification of N/2 gp116ΔTM and C/2 gp116ΔATM.** The N/2 gp116ΔTM DNA sequence, corresponding to nucleotide positions 685 to 1218 of YHV ORF3 (Genbank accession no. AF540644), was amplified by 2 primers (135F685: CGC GGA TCC ACG ATT CTA AGT GGA ATT CCT G; and 116R4Eco: GGA A CAC TGA TAG) with added restriction sites (underlined) and Pfx polymerase (Invitrogen) by using cDNA as the template. The C/2 gp116ΔTM DNA sequence, corresponding to nucleotide positions 1219 to 2394 of YHV ORF3, was amplified by 2 primers (116FC: CGC GAT CTT ATC ATG ACG GAG ATT C and 135R4Eco: GGA ATT CAA ACA GCA TAG CTA CTG CCT G) with added restriction sites (underlined) and Pfx polymerase (Invitrogen).

The PCR protocol consisted of initial denaturation at 94°C for 2 min followed by 35 cycles of 94°C for 15 s, 55°C for 30 s and 68°C for 1 min 15 s, and a final extension at 68°C for 20 min.

**Expression and purification of N/2 gp116ΔATM and C/2 gp116ΔATM.** The amplified fragment of N/2 gp116ΔATM was cloned into pGEX-6P-1 (Amersham Biosciences) at the BamHI and XhoI sites while the amplified fragment of C/2 gp116ΔATM was cloned into pGEX-6P-1 at the BamHI and EcoRI sites. Each recom-
binant plasmid was transformed into *E. coli* strain BL21. The integrity of the open reading frame in the recombinant plasmids was verified by DNA sequencing.

*E. coli* with each recombinant plasmid was cultured in Luria-Bertani (LB) broth to exponential phase and expression of the recombinant proteins was induced with 1 mM isopropyl-β-D-thiogalacto-pyranoside (IPTG) for 4 h. After centrifugation at 4000 × g for 20 min, the bacterial pellet was resuspended in 100 mM NaH₂PO₄, 10 mM Tris-HCl, and 8 M urea (pH 8) containing 1 mM phenylmethylsulfonyl fluoride (PMSF), and sonicated until a clear lysate was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie brilliant blue, the recombinant protein band was obtained. The lysate was separated by SDS-PAGE with 10% polyacrylamide gel. After staining with Coomassie bright
nant was separated by SDS-PAGE and tested for the presence of YHV by Western blotting in comparison with the viral preparation as described previously. The walking legs from the same shrimp were also subjected to tests for YHV infection by RT-PCR using an IQ2000 YHV/GAV test kit (Farming IntelliGene Technology Co-operation). Using this kit, YHV-positive samples exhibit a band of 277 bp while GAV-positive samples exhibit a band of 406 bp.

RESULTS AND DISCUSSION

Cloning and expression of gp116

Several attempts were made to express the whole molecule of gp116 without the transmembrane region (gp116ΔTM) in various E. coli systems, but they were not successful. However, in this study the gp116ΔTM was expressed in 2 parts, each consisting of approximately half of the gp116ΔTM linked with glutathione-S-transferase (GST) at the N-terminus. The N/2 gp116ΔTM and the C/2 gp116ΔTM DNA sequences could be amplified as 1234 bp and 1191 bp PCR products, respectively (Fig. 1). These were cloned and expressed in E. coli and recombinant proteins were visualized by Coomassie blue staining. The rN/2 gp116 and rC/2 gp116 proteins were expressed in the form of GST tag fusion proteins with molecular masses of 72 and 70 kDa, respectively (Fig. 2). These recombinant fusion proteins were combined and used to generate polyclonal antibodies.
Production of mouse anti-rN/2 + rC/2 gp116 antiserum

After immunization of mice with rN/2 gp116 mixed with rC/2 gp116, antisera from 4 mice (mouse anti rN/2 + rC/2 gp116 antiserum) demonstrated specific binding to rN/2 gp116, rC/2 gp116 in bacterial lysates and the natural gp116 from partially purified YHV (Fig. 2B). The MAb V3-2B bound only to rN/2 gp116 (Fig. 2C, lane 2) and the natural gp116 (Fig. 2C, lane 4). The MAbs Y18 and Y19 bound to gp64 and p20 structural proteins of the virus, respectively (Fig. 2B,C, lane 4). This evidence confirmed that the N and C termini of gp116 protein were correctly expressed and that the expressed proteins could be used as antigens for production of specific antibodies.

Sensitivity testing of mouse anti-rN/2 + rC/2 gp116 antiserum

The detection sensitivity of YHV in viral preparations using mouse anti-rN/2 + rC/2 gp116 antiserum was similar to that of MAbs V3-2B, Y18 and Y19 (detection limit, ~1:2000 dilution). However, MAbs V3-2B and Y19 showed slightly darker spots than those for MAb Y18 and mouse anti-rN/2 + rC/2 gp116 antiserum (Fig. 3). The result corresponded to Western blot results for viral preparations (see Figs. 2 & 5) in that the intensity of all 3 viral structural proteins in viral preparations was similar.

Reduction or absence of gp116 immunoreactivity in immunohistochemical studies

Immunohistochemical localization of YHV infection in gill tissues of most of the challenged shrimp revealed strong immunoreactivity with MAbs Y18 and Y19 and weak or absent immunoreactivity with MAb V3-2B and mouse anti rN/2 + rC/2 gp116 antiserum (Fig. 4, column B). Only in some tissues such as those in the mandible region (Fig. 4, column A) nervous tissue, (Fig. 4, column C) and muscle did all these reagents (i.e. MAbs V3-2B, Y18, Y19 and mouse anti-rN/2 + rC/2 gp116 antiserum) exhibit strong immunoreactivity. Since the mouse anti rN/2 + rC/2 gp116 antiserum contained antibodies directed to several epitopes on N and C termini of gp116, lack of immunoreactivity was probably due to a reduced quantity or lack of gp116, and not due to either absence of specific epitopes or lower affinity of MAb V3-2B and the antisera. Interestingly, nervous tissue from the same shrimp specimens gave overall immunoreactivities that were comparable for the 4 antibodies (Fig. 4, column C). However, some small neuronal processes in the gills exhibited weak or absent immunoreactivity with MAb V3-2B and mouse anti rN/2 + rC/2 gp116 antiserum (Fig. 4, column C, rows 1 & 3, arrowheads) when compared with the strong overall immunoreactivity in the gills with MAbs Y18 and Y19 (Fig. 4, column C, rows 2 & 4, arrowheads). The suppression of gp116 production varied from tissue to tissue, from shrimp to shrimp and between shrimp species. The reasons for these differences are currently unknown.

Reduction or absence of gp116 immunoreactivity by Western blotting

Western blot analysis revealed clearly that immunoreactivity to gp116 by mouse anti rN/2 + rC/2 gp116 antiserum (Fig. 5B) and MAb V3-2B (Fig. 5C) in gill extracts from YHV infected Penaeus vannamei was absent or reduced, while immunoreactivities of gp64 and p20 were strong (Fig. 5B,C). Partially purified YHV served as a positive control for Western blot analysis and showed strong immunoreactivity with all 4 antibodies (Fig. 5B,C, lane 2). RT-PCR tests for YHV using the IQ2000 test kit confirmed YHV infections in these shrimp was caused by the virulent Thai YHV strain used as inoculum (band at 277 bp) and not by GAV or other nonvirulent strains of YHV (Fig. 6). These results supported the results obtained by immunohistochemical localization tests by showing that the quantity of gp116 structural protein was indeed reduced or absent in the gill tissue.
Fig. 4. Immunohistochemistry of consecutive sections of (column A) mandibular tissue from YHV-infected *Penaeus vannamei*, (column B) gill from the same individual of YHV-infected *P. vannamei*, and (column C) thoracic ganglion (T) and hepatopancreas (H) of YHV-infected *Palaemon serrifer*. Each section was treated with (row 1) MAb V3-2B, (row 2) MAb Y19, (row 3) mouse anti-rN/2 + rC/2 gp116 antiserum and (row 4) MAb Y18. Arrows indicate light immunoreactivity of anti-rN/2 + rC/2 gp116 antiserum and MAb V3-2B in gill tissues and the small neuronal process. Arrowheads indicate areas where strong immunoreactivity with MAbs Y18 and Y19 in small neurons did not occur with anti-rN/2 + rC/2 gp116 antiserum and MAb V3-2B. Asterisks (*) indicate the strong immunoreactivity in the same neuron.
Evidence for preferential reduction in YHV envelope protein gp116 agreed with the previous reports showing that several palaemonid shrimp species and some penaeid shrimp that survived YHV challenge also showed a preferential absence of or reduction in gp116 structural protein (Longyant et al. 2005, 2006). Polyclonal antibody raised against gp116 could neutralize YHV infection in primary cultures of lymphoid organ cells, suggesting that anti-gp116 antiserum blocked the binding of YHV virions to cellular receptors that facilitate YHV entry into lymphoid organ cells (Assavalapsakul et al. 2005). Therefore, the ability to suppress gp116 in infected cells may be involved in the ability of shrimp to survive YHV challenge. Curiously gp116 is encoded upstream of gp64 in ORF3 of YHV as a polyprotein transcript (Jitrapakdee et al. 2003) and the 2 proteins are released post-translation by proteolytic cleavage. Given this background, the simplest explanation for the absence of gp116 would be that it can be selectively degraded after its formation in some species or tissues, but not others. Further tests would be required to distinguish this possibility from other more complex possibilities that might be proposed to explain this very interesting phenomenon.

Since YHV is a significant pathogen in the southeast Asian and Indo-Pacific regions, the development of methods to prevent YHV infection would be beneficial. Since polyclonal antibodies raised against gp116 were successful in preventing YHV infection in cell culture (Assavalapskul et al. 2005), the recombinant gp116 plasmids and proteins developed in this study may prove useful in development of reagents for prevention of YHV infection in shrimp, as has been reported for white spot syndrome virus (Witteveldt et al. 2004a,b).

Acknowledgements. This work was supported by the Srinakharinwirot University Research Fund.

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Editorial responsibility: Timothy Flegel, Bangkok, Thailand


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Editorial responsibility: Timothy Flegel, Bangkok, Thailand

Submitted: June 30, 2005; Accepted: November 7, 2007
Proofs received from author(s): January 22, 2008