Cloning, Expression and Purification of Recombinant HMW1, HMW2 and Hia as a Fusion protein for Vaccine Candidate of Nontypeable Haemophilus influenza

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Nontypeable H.influenzae (NTHi) is an important pathogen in children causing otitis media. However, there is no vaccine against NTHi-induced diseases. HMW1, HMW2 and Hia are adhesin proteins of NTHi-strains and have potentiality to provide protection against NTHiinfections. The aim of present study was to construct recombinant HMW1-HMW2-Hia as a fusion protein and evaluate immune responses against recombinant fusion protein as vaccine candidates antigens. Binding domain of hmw1 gene (1080 bp) was amplified by PCR from genomic DNA and cloned into PET-28a vector, this vector was already modified by the insertion of a fragment named hmw2-hia into multiple cloning sites, and then was confirmed by colony-PCR, enzymatic digestion and gene sequencing. To express recombinant fusion protein, PET-28a-hmw1-hmw2-hia vector was transformed into competent BL12(DE3). Expressed protein was purified by affinity chromatography. BALB/c mice were subcutaneously immunized by purified protein in combination with Freund's adjuvant. Serum antibody responses and functionality of antibodies were determined by ELISA and SBA, respectively. In immunized group of Balb/c mice, the serum IgG responses was significantly increased against recombinant fusion protein in comparison with control, and also the antisera have shown strongly bactericidal activity against NTHi strain. The results have shown that recombinant fusion protein could induce the immune system to produce antibodies which was funtional to kill NTHi-strains. Based on the observation, the antisera have the possibility to provide protection against infections caused by NTHi.

Key words: NTHi; fusion Protein; Expression Vector; affinity chromatography.

Haemophilus influenzae is a Gramnegative bacterium that divided to nonencapsulated (nontypeable) and encapsulated (typeable) strains. Nontypeable H.influenzae (NTHi) is an important pathogen in children, causing otitis media, sinusitis, conjunctivitis, pneumonia, and typeable H.influenzae strains are responsible for invasive diseases, such as sepsis and meningitis¹. Bacterial adherence to the epithelium of the upper respiratory tract is the first step of colonization and is mediated by both pilus

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and nonpilus adhesive factors, followed by invasion to the bloodstream or contiguous spread within the respiratory tract^{2, 3}, and these adhesin proteins have been proposed as vaccine candidates antigens for nontypeable H.influenzae diseases⁴.

In nontypeable H.influenzae strains, the HMW1, HMW2 and Hia proteins are major nonpilus adhesins which play an important role in adherence to epithelial cells in a direct way^{5, 6}. The children who recovered from acute otitis media caused by NTHi strains demonstrated development of the bactericidal antibodies in the sera against highly immunogenic high-molecularweight (HMW1/HMW2) proteins7, Subsequently, these studies led to characterization and identification of the HMW1 and HMW2 proteins. Further studies have shown that the HMW1/ HMW2 proteins are the main adhesin proteins of nontypeable H.influenza as well as targets of bactericidal and protective antibodies^{8, 9}. The HMW1/HMW2 proteins are expressed by nearly 75% of NTHi strains and those NTHi strains (25%) not expressing HMW1/HMW2 proteins have been shown to express a second different class of adhesins known as Hia proteins⁹. The Hia proteins have been shown to serve as a target for opsonophagocytic and bactericidalitic antibodies. All of the NTHi strains that lack HMW1/HMW2 proteins contain a hia gene expressing a Hia protein, and vice versa¹⁰.

Binding domain of these proteins are very important for adhesion to the epithelial cells of the respiratory tract for the onset of diseases. The binding domain of HMW1 and HMW2 consist of amino acids residues 555-914 and 553-916 of fulllength protein, respectively, and these residues corresponding to amino acids 114-473 and 112 to 475 of mature HMW1/HMW2 proteins in order^{(11,} ¹². Hia contain tw homologous binding domains, named HiaBD1 and HiaBD2 interacting with the same host cell receptor. The HiaBD1/HiaBD1 is in 585-705 and 51-166 amino acids of full length protein. Because of the greater affinity of HiaBD1, this binding domain, HiaBD1, was selected as a partner of recombinant fusion protein¹³. The purpose of the current study was to produce a recombinant fusion protein which composed of binding domain of HMW1, HMW2, and Hia of nontypeable H.influenzae and to evaluate the antisera production against fusion protein in Balb/ c mice. Finally we aimed, to analyze whether the production of antibodies against recombinant protein is functional or not.

MATERIALSAND METHODS

Bacterial strains and vectors

Nontypeable H.influenzae strain (ATCC 49766) was kindly obtained from the Department of microbiology, Pasteur Institute of Iran. The strain was grown at 37°C on chocolate agar. E. coli Top10 (Invitrogen, USA) and BL-21(DE3) (Novagen, USA) were used as a cloning and expression host, respectively. pTG19-T (Malaysia, Vivantis) and pET-28a (Novagen, USA) were used as a cloning and expression vectors, respectively.

Amplification of binding domain of HMW1 555-914

Genomic DNA of the nontypeable H. influenzae strain (ATCC 49766) was extracted by high pure PCR template preparation kit (DNA Technology, Russia). Binding domain of hmw1 gene by the length of 1080 bp was amplified by polymerase chain reaction (PCR) from genomic DNA of nontypeable H.influenzae.

To ensure the correct orientation of cloning the the forward primer (5'-ACTGCCCATGG TTGATGTTCATAAAAAT-3') contained a NcoI restriction enzyme site, whereas the reverse primer (5'-ACTGCGCTAGCTA CATTAAAAGT GAAATT-3') harbored a NheI site. Amplification of the mentioned gene was achieved by using Pfu DNA polymerase (Thermo scientific, USA), with 1 μ l of DNA template (100 ng genomic DNA), 2 μ L of 10x reaction buffer, 0.4 µL of dNTPs (10 mM), 0.6 µL of MgCl2 (50 mM), 1 µL of each primer (10 pmol), and 1 U of pfu DNA polymerase in final volume of 20 µL. Hot Start Amplification was performed in 94°C for 5 min, followed by 30 cycles of denaturation at 94°C for 30 seconds, annealing at 55°C for 60 seconds, extension at 72°C for 120 seconds, and 10 minutes at 72°C for final extension. The PCR product was recovered from the gel and purified with a high pure PCR product purification kit (Vivantis, Malaysia) based on the manufacturer's instructions.

Cloning binding domain of hmw1 into PTG19-T vector

The polymerized DNA corresponding to

the binding domain of hmw1(HMW1 555-914) gene was treated with dATP, and then ligated into pTG19-T vector, The ligation was performed by T4 DNA ligase. Recombinant construct as a cloning vector was transformed into competent E. coli TOP10 cells by the CaCl2 method. TOP10 cells harboring recombinant vector were grown in LB agar medium supplemented with ampicillin ($50 \mu g/mL$). After colony PCR, the recombinant vectors in positive colonies were amplified, and then purified. In order to confirmation the insertion of hmw1 gene into pTG19-T vector, double digestion by Nc°CoI and NheI restriction enzymes and sequencing were performed.

Subcloning and Expression of Recombinant Fusion Protein

The recombinant cloning vector, rpTG19-HMW1 555-914, was digested with NcoI and NheI and the gene corresponding to binding domain of hmw1 ligated into the NcoI- NheI sites of pET28ahmw2-hia construct, which was already synthesized by as Biomatik, (Canada). Recombinant vectors were transformed into competent E.coli TOP10 cells. After colony PCR, the positive colonies were amplified, and then purified, and finally confirmed by sequencing analysis and double digestion NcoI and NheI. In order to express the recombinant fusion protein HMW1-HMW2-Hia, rpET-28a-hmw1-hmw2-hia, was transformed into competent BL-21(DE3) bacteria. BL21(DE3) cells harboring recombinant vector were grown in LB medium supplemented with kanamycin (50µg/ml) at 37 æ%C with shaking (150 rpm) to reach OD of 0.6 in A600. Thereafter, isopropyl â-d-thiogalactopyranoside (IPTG) was added to the final concentration of 1 mM. The BL21(DE3) cells were incubated for 4 hours and then being harvested.

SDS-PAGE and western blotting

The expression of the recombinant fusion protein, HMW1-HMW2-Hia, was analyzed by SDS-PAGE. The bacterial pellets were suspended in a loading buffer sample, heated for 10 minutes at 95°C, and 20 μ L of each sample was subjected to 12% SDS-PAGE gel following by western blotting. The samples were separated by SDS-PAGE method and then proteins on SDS-PAGE gel transferred by wet system to polyvinylidene difluoride (PVDF) membranes by using a liquid transfer system (Bio-Rad, USA). The membranes were blocked with BSA(2% w/v) in phosphate buffered saline (PBS), and then washed several times with PBS-T. The membranes were incubated with his-tag antibody-HRP conjugate (1:3000 diluted in PBS) for 1 hours at room temperature and developed by diaminobenzidine (DAB) solution (Roche, Germany).

Purification of Recombinant Fusion Protein

A pellet of BL21(DE3) cells harboring recombinant fusion protein, HMW1-HMW2-Hia, were suspended by gentle stirring in lysis buffer (pH 8, containing 300 mM NaCl, 50 mM NaH2Po4, 10 mM Immidazol) followed by freezing, thawing, and sonication on ice in the presence of PMSF (1 mM) as a protease inhibitor. The suspension was subjected to sonication (Six cycles, 30 seconds each, with the intervals of 30 seconds on ice) and then centrifuged for 30 minutes at 10,000 g. After centrifugation to verify the location of the expressed recombinant fusion protein, the precipitate and supernatant were examined by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE). The recombinant fusion protein was purified by affinity chromatography on a nickel-nitrilotriacetic acid (Ni-NTA) gel matrix under denaturing conditions. A pellet of BL-21 cells harboring recombinant fusion protein were suspended by denaturing binding buffer (pH 8, 8 M urea, 100 mM NaH2PO4, 10 mM Tris-HCI), and then centrifuged for 30 min at 10,000 g. A column containing 6 ml of Ni-NTA resin was equilibrated with 9 mL of denaturing binding buffer (pH 8, 8 M urea, 100 mM NaH2PO4, and 10 mM Tris-Hcl), and then the cleared cell lysate was loaded onto the column. Following adding the samples to the column, it was shook for 60 minutes. The column was washed by washing buffer (pH 6.3, 8 M urea 100mM NaH2PO4, 10mM Tris-Hcl) Finally, The recombinant fusion protein of interest was eluted by elution buffer (pH 5.7, 8 M urea, 100 mM NaH2PO4, 10mM Tris-Hcl). The purified recombinant fusion proteins were pooled and were subjected to dialysis following a linear gradient of urea from 8 M to 0 M in the refolding buffer (pH 7.2, 20 mM sodium phosphate, and 300 mM NaCl). Finally the protein concentration was determined by Nanodrop spectrophotometry (Thermo scientific, USA). The concentrations of recombinant proteins were determined by Bradford assay with bovine serum albumin (BSA) as the

standard protein in NanoDrop spectrophotometry (Bio-RAD, USA)

Mouse immunization

Groups of six-week-old, female BALB/c mice were immunized subcutaneously with 50µg of purified recombinant fusion protein, HMW1-HMW2-Hia, mixed with an equal volume of complete Freund's adjuvant for the first injection and incomplete Freund's adjuvant for the second and third injections as a booster 14 and 21 days following after the first injection. Control groups of BALB/c mice received Freund's adjuvant and PBS alone. Antisera were collected 14 and 28 days after injection to evaluate the antibody responses and functionality of those antibodies. Aliquots of antisera were stored at "70 æ%C for assessment. **Determination of serum IgG levels against purified Recombinant Fusion Protein**

The IgG responses against purified recombinant fusion proteins, HMW1-HMW2-Hia, were determined by ELISA. Serum of female BALB/ c mice were collected 14 and 28 days after the first injection, To assess the individual serum samples of immunized and control groups, 96-well flatbottomed plates were coated overnight at 37 æ%C with 1µg/100 µl of purified recombinant fusion protein, HMW1-HMW2-Hia, in PBS. Following the coating process, the plates were blocked with 1% (w/v) bovine serum albumin (BSA) in PBS for 2 hours at room temperature. After washing, the plates were incubated with antisera (1/256 diluted in PBS) for 2 hours at room temperature. The wells were washed three times with PBS-T thereafter plates were incubated by goat anti-mouse IgG conjugated to Horseradish Peroxidase (Bio-Rad, Hercules, Calif.) in 1:3000 dilution in PBS. Plates were incubated for 1 h at 37 æ%C and washed three times in PBS-T, and then tetramethylbenzidine (TMB) substrate was added to each well. Finally, reaction was stopped after 15 min by adding of H2SO4 2M and the absorbance was measured at 450 nm. All assays were performed in duplicate. Serum bactericidal assay

Serum bactericid

Serum bactericidal assay (SBA) is a functional measurement of the ability of antibodies, in combination with complement, to kill bacteria. Based on the previous protocols, the test was performed(14). Following overnight incubation of nontypeable H.influenzae strain on chocolate agar, single colonies of nontypeable H.influenzae inoculated into LB broth. The NTHi were grown for 5 hours to get to the log phase, it was pelleted by centrifugation and washed and resuspended in PBS containing 9 mM of CaCl2, 4.9 mM of MgCl2 (Dulbecco's buffer) and 1% (w/v) bovine serum albumin (BSA). In the present study, the SBA was done on pooled serum specimens collected at weeks 0, 2 and 4. All sera were inactivated, by heating for 30 minutes in 56 æ%C all. As a source of complement, 5-week-old serum of baby rabbit was used 10% (v/v). In 96-well, sterile plates SBA assay was done. The mixture in each well contained 25 μ l of complement (20%, v/v), 50 μ l of serially diluted sera in buffer, and 25 µl of buffer containing 500 CFU of bacteria, in final volume of 100 µl. After adding components to the wells, microplates were incubated for 60 minutes at 37 æ%C in a 5% CO2, and then 5 µl aliquot of each well was plated onto a chocolate agar and incubated 16 hours at 37 æ%C. The surviving percentage of bacteria assessed by comparing the respective CFU at 60 minutes compared to those at zero time in the control group. Bactericidal titers defined as the reciprocal of the highest dilution of serum that resulted in e"50% killing effect in comparison to the control group. Specimens that showed d"50% bacterial killing effect at the lowest dilution of the tested serum (the lowest dilution which wa tested for serum samples was 1:8) were reported as having a SBA titer of d"8.All assays were performed in duplicate. **Statistical analysis**

All the statistical analysis was done by using SPSS statistical software package (version 17.0). Following normalization of the data, one-way ANOVA and t-tests were used to check for differences between data sets. P values d'' 0.05 were considered statistically to be significant.

RESULTS

Amplification of hmw1 and the construction of rpET28a-hmw1-hmw2-hia

A 1080 bp fragment as a binding domain of HMW1 555-914 was amplified by PCR using specific primers. Electrophoresis of PCR products showed that the length of HMW1 binding domain was approximately 1080 bp as expected (Figure 1), and then this fragment was cloned into pTG19T as a cloning vector. The positive colonies were selected on LB/Amp and existence of HMW1 555-



Fig. 1. Electrophoresis of the HMW1 PCR products on agarose gel (1% w/v) lanes 1-3: binding domain of hmw1gene (PCR product 1080bp), Lane 4: 1 kb DNA size marker



Fig. 2. Electrophoresis of the HMW1 Colony PCR products on agarose gel (1% w/v) lanes 1-6: binding domain of hmw1gene (PCR product 1080bp), Lane 7: 1 kb DNA size marker



Fig. 3. Double digestion of the recombinant PTG19hmw1 vector with *Nco*I and *Nhe*I restriction enzymes., lane 1:: Non digest recombinant PTG19-hmw1 vector , lane 2: Double digestion of the recombinant PTG19_hmw1 (PTG19T: 2880 bp and binding domain of hmw1: 1080 bp), lane 3: Single digestion of the recombinant PTG19-hmw1 (rPTG19-hmw1: 3960 bp) with *Nco*I enzyme, Lane 4: 1 kb DNA size marker

914 insert in recombinant vector, pTG19T-HMW1 555-914, was confirmed by colony-PCR, double digestion using *NcoI* and *NheI* restriction enzymes (Figure 2, 3). The orientation of HMW1 555-914 in the construct was confirmed by sequencing analysis. Then binding domain of HMW1 555-914, was subcloned into pET28a-hmw2-hia. The positive colonies were selected on LB/Kan and the existence and the orientation of insert in rPET-28a-hmw1-hmw2-hia, was confirmed colony PCR,





Fig. 4. Electrophoresis of the HMW1-HMW2-Hia Colony PCR products on agarose gel (1% w/v) lanes 1-6: recombinant fusion of HMW1-HMW2-Hia gene (PCR product 2673bp), Lane 7: 1 kb DNA size marker



Fig. 5. Double digestion of the recombinant pET28a(+)HMW1-HMW2-Hia vector with *NcoI* and *XhoI* restriction enzymes., Lane 1: 1 kb DNA size marker, lane2: Double digestion of the recombinant pET28a(+)HMW1-HMW2-Hia vector (pET28a(+) :

5229 bp and recombinant fusion of hmw1-hmw2-hia:
2673bp), lane 3: single digestion of the recombinant
pET28a(+)HMW1-HMW2-Hia vector:7900 bp) with
*Xho*I Enzyme , lane 4: undigested the recombinant
pET28a(+)HMW1-HMW2-Hia vector

digestion analysis and sequencing (Figure 4, 5). Expression and purification of recombinant fusion protein HMW1-HMW2-Hia

BL21 (DE3) cells that harbor pET28a-HMW1-HMW2-Hia were cultured at 37 æ%C in



Fig. 6. SDS-PAGE Analysis of the recombinant fusion protein HMW1-HMW2-Hia protein with Coomassie Staining. Expression and purification of the recombinant fusion proteins in E. coli BL21(DE3). Lane 1,5: protein marker, Lane 2-4: purified recombinant fusion protein marker, Lane 6-8: recombinant fusion protein HMW1-HMW2-Hia induced with IPTG 1mM



Fig. 7. Western Blot Analysis of recombinant fusion purified proteins of HMW1-HMW2-Hia. Lane 1,2: recombinant fusion protein of HMW1-HMW2-Hia with molecular weight of 95kDa, Lane 3: protein marker

the absence and presence of an IPTG with the concentration of 1 mM. One major band appeared at the 95 kDa position after IPTG induction, as expected at the position of recombinant fusion protein HMW1-HMW2-Hia. Induction of the cells with 1mM IPTG for 4 hours at 37 æ%C, was found to be optimal to achieve high level expression of recombinant fusion protein HMW1-HMW2-Hia. Recombinant fusion protein HMW1-HMW2-Hia was mainly presented in the form of dissoluble inclusion body. Thus, inclusion bodies were carefully purified with Ni-NTA affinity chromatography under denaturing condition The output of recombinant fusion protein HMW1-HMW2-Hia was approximately 30% of the total bacterial proteins. The highest detectable level of Recombinant fusion protein HMW1-HMW2-Hia was up to 3 mg/L in culture medume (Figure 6). Western blot analysis

The bands of recombinant fusion protein, HMW1-HMW2-Hia, was detected respectively to be 95 kDa, as expected (Figure 7). The identification of the expressed fusion proteins further confirmed with western blot analysis by using anti-histidine tag monoclonal antibody, where bands corresponding to the molecular mass of the recombinant fusion protein HMW1-HMW2-Hia was respectively detected.

Serum IgG responses after immunization with recombinant fusion protein

Those BALB/c mice which immunized subcutaneously with recombinant fusion protein, HMW1-HMW2-Hia, mixed with Freund's adjuvant exhibited high levels of specific antibody responses. Serum IgG responses were significantly increased in immunized groups with recombinant fusion protein, HMW1-HMW2-Hia in comparison with control groups ($P \le 0.05$). The booster



Fig. 8. The IgG antibody responses against recombinant fusion protein were determined by ELISA of individual serum samples collected at days 0, 14 and 28

injections were effective to significantly increase the responses of anti- recombinant fusion protein HMW1-HMW2-HiaIgG (Figure 8).

Serum bactericidal assay

The antisera were tested for their ability to promote complement-mediated killing of nontypeable H.influenzae (NTHi) strains in vitro. The antisera from immunized animals (Balb/c mice) with recombinant fusion protein,HMW1-HMW2-Hia were relatively bactericidal against nontypeable H.influenzae (NTHi) strains in comparison with the control group. Antisera raised against recombinant fusion protein, HMW1-HMW2-Hia, demonstrated bactericidal activity with titers of 1/128.

DISCUSSION

Nontypeable H. influenzae (NTHi) strains are responsible for localized respiratory tract diseases such as otitis media, pneumonia and sinusitis. The colonization of nontypeable H. influenza in the upper respiratory is being mediated by adhesin proteins which play an important role in the pathogenesis of diseases^{2, 15, 16}. Therefore, these adhesin proteins have the potentiality to be utilized as vaccine candidates. Among these proteins, there are a class of proteins named on the basis of their high molecular weight, these adhesin protein are HMW1 and HMW2 and Hia which are highly immunogenic, surface adhesins and expressed abundant among NTHi strains. The HMW1 and HMW2 proteins are expressed by approximately 75% of nontypeable H.influenzae strains^{17, 18}. Those strains of Nontypeable H.influenzae strains that do not express HMW1 and HMW2 proteins, express immunogenic adhesins known as Hia proteins, explaining why fusion proteins should be used for designing a vaccine aiming at protection against widespread spectrum of diseases caused by NTHi strains7, 19.

In the present study, we constructed a recombinant fusion protein HMW1-HMW2-Hia from amino acid residues of binding domain of HMW1 555–914, HMW2 553–916 and Hia 585–705, respectively. Evaluation of IgG antibodies have shown the efficacy of vaccine candidate for the induction of immune responses against recombinant fusion protein HMW1-HMW2-Hia. Also serum bactericidal assay (SBA) confirmed

that antibodies responses corresponding to immunization with the recombinant fusion protein HMW1-HMW2-Hia was capable of inducing bactericidal activity against NTHi strains. These data suggest that recombinant fusion protein HMW1, HMW2, and Hia purified as vaccine candidates have a high potential to provide protection against infections caused by NTHi strains.

At this time, there is no vaccine providing protection against NTHi diseases. Our current study has focused on the capability of antisera directed against HMW1-HMW2-Hia fusion protein which also evaluate complement-mediated killing of nontypeable H.influenzae (NTHi) strains in vitro in order to confirm that whether mentioned proteins are good enough to make protection. However, there is a long way to consider these adhesin proteins as good candidates against diseases caused by NTHi. Our present study was adapted from previous study which demonstrated that a group of common epitopes existing on the HMW1 and HMW2 and Hia proteins of NTHI could induce active antibodies¹ and also the results of our study coincide with the fact that recombinant adenoviruses expressing recombinant HMW1, HMW2, or Hia proteins are particularly attractive vaccine candidates providing protection against upper respiratory tract diseases²⁰. Antibodies against native HMW1, HMW2 proteins and recombinant Hia are suitable of mediating broadbased opsonophagocytic killing of homologous and heterologous NTHi strains^{21, 22}. A vaccine constructed of HMW1 and HMW2 and Hia proteins might provide protection against diseases caused by many Nontypeable H.influenzae strains. In spite of the fact that binding domain has diversity, it has conserved amino acid motifs which can help us to design a fusion protein as potential vaccine candidates against NTHi.

The results of ELISA and SBA have shown that recombinant fusion protein could induce the immune system to produce antibodies killing nontypeable H.influenzae strain by complement system in vitro. Based on the observation, the antisera against recombinant fusion protein have the possibility to provide a good protection against infections caused by NTHi. Therefore, the results argue for the further investigation designing effective vaccines for NTHi-induced diseases based on these proteins or combination of them.

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Conflicts of interest

There are no conflicts of interest in this research project.

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