

Cloning, expression and purification of outer membrane protein PorA of *Neisseria meningitidis* serogroup B

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Abstract

Introduction: *Neisseria meningitidis* is a major causative agent of bacterial septicemia and meningitis in humans. Currently, there are no vaccines to prevent disease caused by strains of *N. meningitidis* serogroup B. PorA is a major component of the outer membrane of *N. meningitidis* and functions as a cationic porin. This study aimed to clone and determine the expression of PorA.

Methodology: A 1200 bp fragment of *porA* gene was amplified by PCR from serogroup B *N. meningitidis* and then cloned into prokaryotic expression vector pET-32a. For expression of recombinant protein, pET32a-*porA* plasmid was transformed into competent Origami B (DE3) cells. Recombinant protein was overexpressed with isopropylthio-beta-D-galactoside (IPTG) and affinity purified by Ni-NTA agarose. SDS-PAGE and western blotting were performed for protein determination and verification.

Results: Cloning of *porA* was confirmed by colony-PCR and enzymatic digestion. In comparison with the corresponding sequences of original genes, the nucleotide sequence homology of the cloned *porA* gene was 97%. IPTG with a dosage of 1.0 mmol/L could efficiently induce protein expression. SDS-PAGE analysis showed that our constructed prokaryotic expression system pET32a-PorA-Origami efficiently produces a target recombinant protein with a molecular weight of 65 kDa. The recombinant PorA was overexpressed as inclusion bodies and reacted with the serum from a rabbit previously immunized with native outer membrane vesicle.

Conclusion: This prokaryotic expression system provides an easy method for producing recombinant PorA and may also be useful for the production of other bacterial outer membrane proteins for vaccine studies.

Key words: *Neisseria meningitidis*; PorA; Ni-NTA agarose; pET-32a

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Introduction

Neisseria meningitidis is an encapsulated, Gram-negative diplococcus that causes bacterial meningitis and septicemia; therefore, it is considered a major public health problem [1]. Meningococcal strains are classified into 13 serogroups based on the chemical composition of the polysaccharide capsule [2]. Serogroups A, B, C, Y, and W-135 are associated with disease worldwide [3]. Antibiotic therapy is often ineffective to treat meningococcal infections and vaccination is considered an attractive approach for prophylactic purposes [4].

Polysaccharide conjugate vaccines are available for *N. meningitidis* serogroups A, C, Y and W135 [5]. However, due to the structural similarity of the *N*-acetylneuraminic acid component of the serogroup B capsular polysaccharide with surface glycoproteins of human fetal cells, there is no similar vaccine for serogroup B [6]. Alternative antigens are being evaluated as candidates for vaccine [7]. Non capsular candidate vaccines based on outer membrane vesicles

(OMVs) and outer membrane proteins (OMPs) such as PorA, PorB, NadA, NhhA, and AppA have been designed previously [8]. PorA (class 1 OMP) is a major protein of 44kDa that is encoded by the *porA* gene locus and forms trimeric cationic pores in the outer membrane [9]. This transmembrane protein consists of a 16-strand beta-barrel, eight of which are surface exposed loops. Two of these extracellular loops (loops 1 and 4) are highly immunogenic and evoke bactericidal antibodies [10]. PorA has been shown to induce bactericidal immune responses in human serum following natural disease and immunization [11]. In humans, the protein elicits a more protective immune response than any other meningococcal surface protein [12]. Much of this evidence comes from the clinical evaluation of OMV vaccines in which PorA is the immunodominant antigen [13]. Recombinant OMV-formulations with various PorA antigens have been developed in some countries; therefore, it can be considered a reliable candidate antigen as a part of a multi-component

recombinant protein vaccine [14]. The aim of the present study was to construct a prokaryotic high-level expression system for producing recombinant outer membrane protein PorA which can be used for vaccine development in the future.

Methodology

Bacterial strains and vector

N. meningitidis CSBPI, G-245 was kindly obtained from the Pasteur Institute of Iran and was grown on chocolate agar. *Escherichia coli* strains DH5 α (Invitrogen, California, USA) and Origami B (DE3) (Novagen, Wisconsin, USA) were used for cloning and expression of the recombinant protein, respectively. The *E. coli* cells harboring recombinant plasmids were grown aerobically at 37°C in Luria-Bertani (Merck, Darmstadt, Germany) broth with or without 50 μ g/ml Ampicillin (Sigma, Saint Louis, MO, USA). Plasmid pET-32a (Novagen, Wisconsin, USA) was used as an expression vector.

Amplification of porA

Genomic DNA from serogroup B *N. meningitidis* CSBPI, G-245 was extracted using the phenol-chloroform method and dissolved in TE buffer [15]. The full coding sequence of *porA* (1200 bp) was amplified by polymerase chain reaction (PCR) using specific primers. To ensure correct orientation into the multiple cloning sites of the vector, the forward primer (5'-GACGGATCCATGCGAAAAAACTTACC-3') incorporated a *Bam*HI restriction enzyme site, whereas the reverse primer (5'-ATACTCGAGTTAGAATTTGTGGCGCAAACC-3') incorporated an *Xho*I site. Amplification of the DNA was achieved by using Prime STAR DNA polymerase (Takara, Osaka, Japan); a 25- μ l reaction mixture contained 0.5 pmol of each primer, 5 μ l 5X prime STAR buffer, 0.2 mM concentration of each dNTP, 2.5U of prime STAR DNA polymerase, and 100 ng genomic DNA. Amplification was performed by using 30 cycles of denaturation at 94°C for 10 seconds, annealing at 57.5 °C for 15 seconds, extension at 72°C for 90 seconds and 10 minutes at 72°C for the final extension. The PCR products were recovered from the gel and purified by using the PCR purification kit (Bioneer, Taejon, South Korea).

Cloning and expression of porA

The purified *porA* fragment was digested with the restriction enzymes *Bam*HI and *Xho*I and ligated into the *Bam*HI-*Xho*I sites of pET32a vector, which

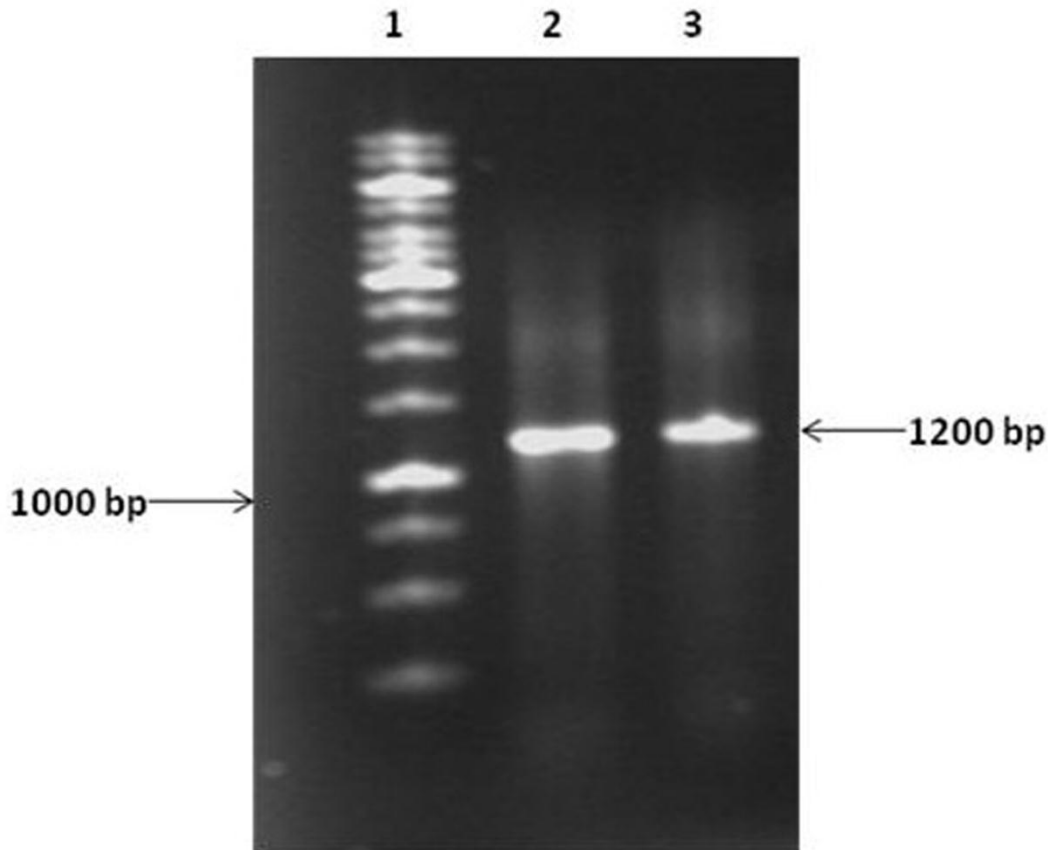
provides six His residues at the N-terminus of the expressed protein. Recombinant vector pET32a-*porA* was transformed into competent *E. coli* DH5 α cells. The integrity of the recovered plasmid was confirmed by restriction endonuclease digestion and sequencing of the *porA* insert by a commercial facility using universal T7-promoter and T7-terminator primers (TAG Copenhagen A/S Symbion, Denmark).

For expression of the recombinant protein, pET32a-*porA* plasmid was transformed into competent Origami B (DE3) cells. Origami cells harboring a pET-32a-*porA* vector were grown in LB medium supplemented with Ampicillin (50 μ g/ml) at 37°C with shaking (250 rpm) to A_{650} of 0.7; then isopropyl- β -D-thiogalactopyranoside (IPTG) was added to a final concentration of 1 mmol/L. The cells were incubated for a further 4 hours before being harvested.

Purification of recombinant PorA

A pellet of Origami cells harboring rPorA was suspended by gentle stirring in lysis buffer (1% Triton X100, 20mM Tris-HCl, 10mM EDTA, pH 7.5). The suspension was subjected to sonication (five cycles, 1 minute each, with intervals of 1 minute on ice) and then centrifuged for 20 minutes at 6000 \times g. After centrifugation, the supernatant and precipitate were examined by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE) to verify the location of the expressed recombinant protein. The crude rPorA was purified by affinity chromatography on a nickel-nitrilotriacetic acid (Ni-NTA) gel matrix (Qiagen, Crawley, United Kingdom) under denaturing conditions. A column containing 5 ml of Ni-NTA resin was equilibrated with 10 volumes of buffer containing 8 M urea, 0.1 M NaH₂PO₄, and 0.01 M Tris (pH 8.0), and the cleared cell lysate was loaded onto the column. The column was washed with 5 volumes of wash buffer (8 M urea, 0.1 M NaH₂PO₄, 0.01 M Tris (pH 8.0) and 20 mM imidazole), followed by 5 volumes of the same buffer, but at pH 6.3. The rPilQ₄₀₆₋₇₇₀ protein was then eluted by increasing the imidazole concentration to 250 mM. Fractions containing the recombinant protein were pooled and dialyzed against PBS, pH 7.4 to remove imidazole. Protein concentrations were determined by Nanodrop analyzer (Bio-Rad, Hercules, CA, USA) and the purity was determined by SDS-PAGE and Coomassie blue staining.

Figure 1. Electrophoresis of outer membrane protein PorA PCR product on agarose gel (1% w/v). Lane 1: 1 kb DNA size marker, Lane 2, 3: Single expected band of *porA* gene (1200 bp)



Anti OMV-PorA antibody production

Female New Zealand white rabbits (Pasteur research institute, Tehran, Iran) were immunized subcutaneously with 100 µg of native meningococcal OMV-PorA obtained by sodium deoxycholate extraction in complete Freund's adjuvant (Sigma, Saint Louis, MO, USA). Booster doses were also given in incomplete Freund's adjuvant at two and 4 weeks. Ten days after the last immunization, the animals were exsanguinated by cardiac puncture under anesthesia and serum samples were collected and stored at -20°C until required for use. All animal experiments were done in accordance with institutional and national ethical guidelines.

Western blot analysis

The separated proteins by SDS-PAGE were blotted onto a 0.45 µm pore size polyvinylidene difluoride (PVDF) membrane (Hi-bond Amersham Biosciences, USA) by using a semi-dry blotter unit (Labconco, Kansas City, Mo). The membrane was blocked by 1% skim milk and then the PVDF

membrane was incubated with native immune serum (diluted to 1:1000 in PBS-Tween 20) for 2 hours at room temperature. The membrane was washed with PBS-Tween 20 and then incubated with sheep anti-rabbit immunoglobulin G (heavy and light chain) horseradish peroxidase (HRP) conjugate antibody (diluted to 1:3000 in PBS-Tween 20) for one hour at room temperature. After washing three times, the membrane was treated using DAB solution (Sigma, Saint Louis, MO, USA) and placed in darkness to develop the protein band.

Results

Amplification of porA and construction of pET32a-porA

Specific primers were designed to amplify *porA* from the *N. meningitidis* CSBPI, G245. The expected size of the *porA* PCR product, approximately 1200 bp, is shown in Figure 1. The integrity of the recombinant vector pET32a-*porA* was confirmed by double digestion using *Bam*HI and *Xho*I restriction

Figure 2. Electrophoresis of recombinant pET32a-porA on agarose gel (1% w/v). Lane 1: 1 kb DNA size marker, lane2: Double digestion of recombinant pET-32a-porA with *Bam*HI and *Xho*I restriction enzymes (pET32a: 5900bp and *porA*: 1200bp)

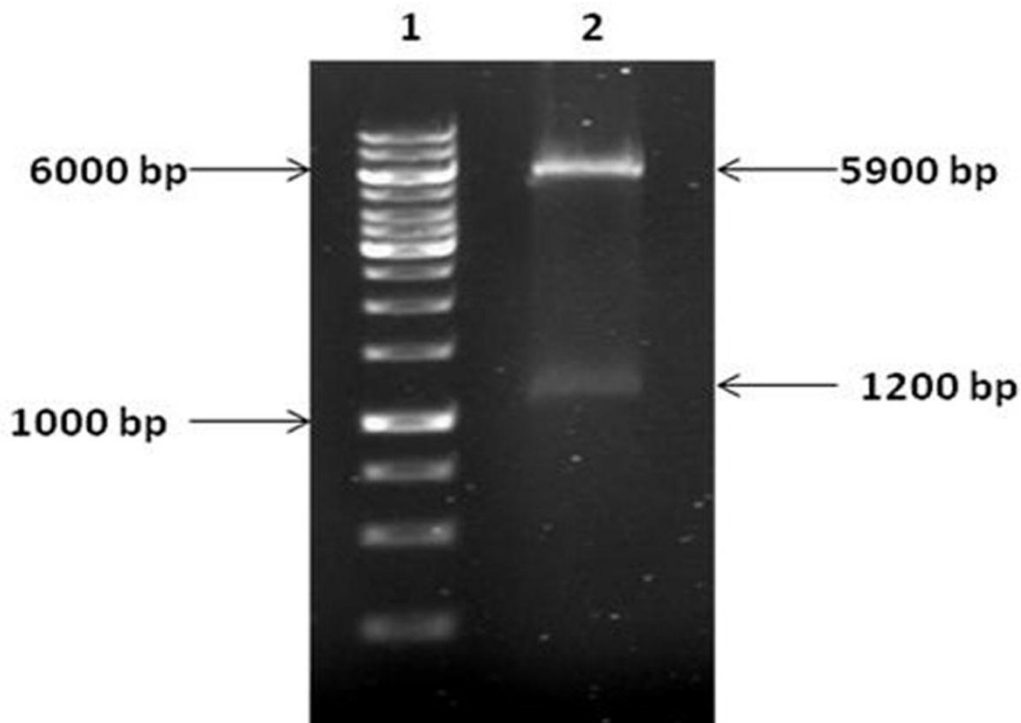


Figure 3. SDS-PAGE analysis of recombinant PorA with Coomassie-stained. Expression of PorA in Origami cells induced with 1.0 mmol/L IPTG. Lane 1: Protein marker, Lane 2, 3: Non-induced with IPTG, lane 4, 5: Induced with IPTG (65 kDa)

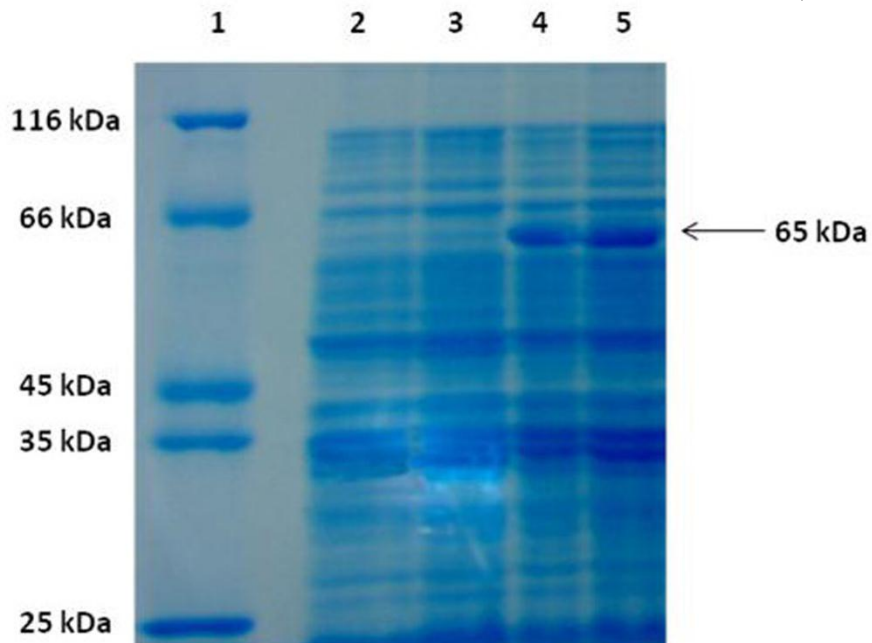
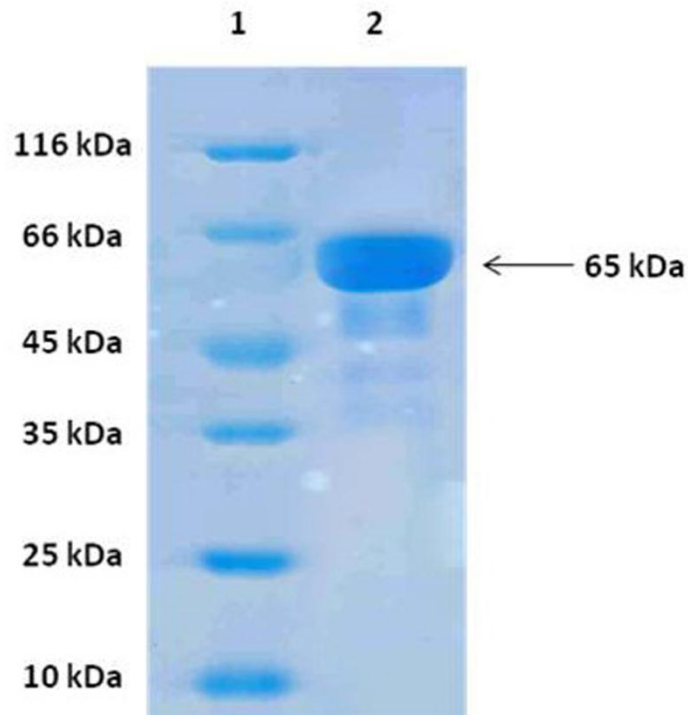


Figure 4. SDS-PAGE analysis of purified recombinant PorA with Coomassie-stained. Lane 1: protein marker, lane 2: recombinant PorA with molecular weight of 65 kDa that purified with Ni-NTA affinity chromatography



enzymes (Figure 2) and colony-PCR with specific primers. Identity and orientation of *porA* in the construct was confirmed by sequencing the recombinant vector. Cloned *porA* gene sequence showed 97% homology with the reference sequences.

Expression and purification of PorA

Origami cells harboring pET32a-*porA* plasmid were cultured at 37°C in the presence and absence of an inducer IPTG. The whole-cell lysates were analyzed by 12% SDS-PAGE. One major band appeared approximately at the 65 kDa position after IPTG induction, which was the expected position of PorA (Figure 3). Induction of the cells with IPTG (1.0 mmol/L) at 37°C for 4 hours was found to be optimal to achieve high-level expression of PorA. Both supernatant and the pellet of cell lysates were tested for the presence of recombinant proteins and the majority of the expressed protein was detected in inclusion bodies. Recombinant protein was carefully purified with Ni-NTA affinity chromatography under denaturing conditions (Figure 4). The output of rPorA was approximately 50% of the total bacterial proteins and the highest detectable level of purified rPorA was up to 0.4 mg/ml.

Western blot analysis

Western blot analysis was performed to detect the antigenicity of the expressed protein. The 65 kDa protein band, observed in SDS-PAGE, was confirmed as PorA protein by western blot analysis using with rabbit anti-OMV antibodies.

Discussion

The outer membrane proteins in Gram-negative bacteria have particular significance as a potential target for protective immunity [16]. Class 1 outer membrane protein (PorA) is a major component of the outer membrane of *Neisseria meningitidis* and functions as a cationic porin [17]. Previous studies have shown that PorA is immunogenic during natural infection and is also a target for bactericidal antibodies, following immunization with experimental OMVs [18,19]. In clinical vaccination trials with OMVs, it has been shown that PorA is critical for the induction of bactericidal antibodies in humans [10]. Arigita *et al.* showed that purified recombinant PorA was immunogenic in the mouse model [20].

The main difficulty with PorA antigens is that they are antigenically diverse and display considerable temporal and geographical variability [21]. A single PorA vaccine component would achieve limited coverage among more heterogeneous populations. To provide protection against heterologous strains, recombinant multivalent OMV vaccines consisting of various PorA antigens (hexavalent and nanovalent) have been developed in the Netherlands [22]. Recent reports on serogroup B meningococcal vaccine candidates have focused on complex mixtures of conserved proteins such as fHbp, NadA, and NHBA [23,24,12].

Traditionally, preparation of native PorA for immunological and *in vitro* studies is time-consuming and needs large volumes of bacterial culture, sophisticated technical equipment and the performance of several purification steps that usually lead to a low yield [25]. Therefore, developing a method to overcome to conventional approaches is an asset. In the present study, recombinant DNA technology was applied to obtain PorA. Among the available protein expression systems, *E. coli* is most commonly used because of its well-characterized genetics, rapid growth characteristics, ability to utilize inexpensive substrates, and the availability of an increasingly large number of cloning vectors and mutant host strains [26,27].

PorA has previously been expressed as a fusion protein in *Bacillus subtilis* and *E. coli* strain E2566 [19]. In this study, we report the cloning of *porA* in a pET-32a vector and its expression in Origami. We believe that this is the first report of overexpression of recombinant PorA with a His-tag in Origami cells. pET-32a vector carries six histidine residues in the N-terminal of the fusion protein. The His tag facilitates purification of the recombinant protein by Ni²⁺-Sepharose resin. Highly purified recombinant protein was obtained after purification. The output of rPorA was approximately 50% of the total bacterial proteins that might be beneficial to industrial production. Our efforts at refolding the solubilized proteins using dialysis led to the precipitation and aggregation of many proteins. In contrast, the on-column purification and refolding approach effectively refolded the desired recombinant protein without protein aggregation or precipitation. In Western blotting, the reaction of rabbit anti native serum with recombinant PorA demonstrates the presence of common epitopes between native and recombinant proteins. One of the limitations of our study was that

we did not have recombinant commercial protein as a control or monoclonal antisera to verify our findings.

In summary, the pET32a-porA-Origami system has been used to express a recombinant outer-membrane protein from *N. meningitidis* to study its potential as a vaccine antigen and may provide an easy method for producing other bacterial outer membrane proteins for the same purpose.

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