

Immunogenic Outer membrane Proteins (Omps) of Salmonella: Potential Candidate for sub-unit vaccine

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Abstract

The outer membrane is a continuous structure on the surface of Gram-negative bacteria and has particular significance as one of the potential targets for protective immunity. Recent studies have focused attention on the outer membrane proteins (OMPs) suggesting the existence of protective immunogenic components in Salmonella. Outer membrane protein (OMPs) has been reported to be immunogens for eliciting active/protective immunity against Salmonella and thus, have great potential for use in vaccination. OMPs have been investigated as potential vaccine candidates, virulence factors and these surface exposed proteins play a critical role in pathogenic processes such as motility, adherence and colonisation of the host cells, injection of toxins and cellular proteases, as well as the formation of channels for the removal of antibiotics (antibiotic resistance).

Keywords: Outer membrane protein; Immunogens; Salmonella

Introduction

The outer membrane (OM) of Gram-negative enteric bacteria is a highly asymmetric lipid bilayer. Its outer leaflet is occupied by lipopolysaccharide (LPS), a unique constituent of the OM, whereas the inner leaflet is covered by phospholipids, mainly phosphatidylethanolamine [1-3]. About 50% of the outer membrane mass consists of protein, either in the form of integral membrane proteins or as lipoproteins that are anchored to the membrane by means of N-terminally attached lipids [4]. The OMPs of Gram-negative bacteria are synthesized in the cytoplasm and have to cross the inner membrane before being assembled into a correctly folded state in the outer membrane. Proteins are usually unfolded by chemical and physical treatments, which lead to alterations on their

conformations [5,6]. The outer membrane proteins of bacteria function as the dynamic interface between the bacterium and its surroundings and are involved in the maintenance of cell structure, binding a variety of substances adhesion to other cells, and regulation of transport of both nutrients and bactericidal agents. [7]. The outer membrane protein (OMP) from Gram-negative bacteria (Salmonella Typhi) is a major immunogenic target to synovial fluid lymphocytes of patients with reactive arthritis (ReA)/undifferentiated spondyloarthropathy (uSpA) [8] and immunologically important because of their accessibility to the host defense system.

The outer membrane proteins (OMPs) of *Salmonella* have been considered possible candidates for conferring protection against typhoid. Vaccines for nontyphoidal *Salmonella* are urgently required especially in Africa. Besides, O antigen polysaccharide-based vaccine might be ineffective and increase susceptibility to life-threatening extracellular *Salmonella* growth, and an outer membrane protein-based vaccine could induce protective antibodies [9,10]. Large number of studies with bacterial outer membrane molecules as candidate vaccines has shown considerable promise [11,12]. Over the past years, several *Salmonella* OMPs have been investigated as potential vaccine candidates, virulence factors, and diagnostic antigens and the molecular structure and function of OMPs and their respective genes have been studied. However, only a small number of OMPs have so far been characterized [13-17]. Study of other gram-negative bacteria demonstrated that porins represent the most abundant class of OMPs that are protective and show some degree of antigenic heterogeneity among different strains [18]. The outer membrane (OM) of *Salmonella* and other Gram negative bacteria contains a family of pore-forming proteins called porins [19]. These porins are present in mostly trimeric forms in the outer membrane [20] and are involved in the permeability of low molecular mass substances [21]. Porins generally have multiple surface epitopes which are usually 6 to 25 residues in length [22]. Several workers have successfully demonstrated the induction of high levels of anti-porin antibodies and enhanced cell-mediated immunity along with protection afforded by porins against *Salmonella* [23]. The major porins in *Salmonella* are OmpC, OmpF, OmpD, out of which OmpC was found to be major surface antigen with unique exposed epitopes expressed in more amounts regardless of the growth condition [24]. Study of other gram-negative bacteria demonstrated that porins represent the most abundant class of OMPs that are protective and show some degree of antigenic heterogeneity among different strains [18]. However, these are relatively nonspecific and have interspecies cross-reactivity. The porin proteins may also have potential use in the development of oral vaccines, biosensors, and nano-reactors [25]. Antibody responses are important to achieve protection against *Salmonella* infection [13]. Because of the ability of the porins to elicit antibody response that has shown to be protective in nature, several groups have studied native *S. Typhi* porins prepared by conventional method as vaccine candidates against typhoid fever [26,27].

OmpC

OmpC is a major porin protein of *S. Typhi* and also a major surface antigen, expressed throughout the infection

period in typhoid patients. It is a good candidate to display heterologous epitopes on the cell surface [28,29]. The functional and mature OmpC is a homotrimer. OmpC monomer has 357 aa without signal peptide, with a molecular mass of 39 kDa. It is a barrel protein having pore radius estimated to be 1.1 nm [19], which is significantly larger than *E. coli* [30]. LPSs have been shown to provide structural stability to OmpC [31]. It was reported through hybridization studies with gene segments on OmpC, that OmpC protein is highly conserved in 11 different *Salmonella* serovars [29]. It was reported that heterologous expression of OmpC with signal peptide is toxic to the cells [32]. It has been reported through multiple sequence alignment tools that *S. Typhi* OmpC consists of 8 variable regions on comparison with other porins with well-known crystal structures [33]. These variable regions have been found to be on the outer side of the membrane and therefore they have high probability to be presented for B-cell recognition and elicit immune response. These results depict that OmpC has recognized B-cell epitopes and as it shares maximum similarity with OmpC of *S. Typhimurium* (98%) these variable regions can be strongly predicted to act as possible B-cell epitopes capable of evoking immune response. This porin is expressed under low and high osmolarity conditions [34, 29]. Hence OmpC is expressed not only under free living conditions, but also during infection, since the osmolarity of the human serum is equivalent to high salt conditions maintained in laboratory [1]. OmpC of *S. Typhi* was successfully expressed with signal peptide by Verma et al. in 2009. The immunogenic nature of the recombinant porin protein was evaluated by ELISA by raising hyperimmune sera in Swiss Albino mice [35]. These reasons suggest that OmpC could be a potential candidate for development of r-DNA vaccine against *Salmonella*. And also can act as candidate antigen for diagnostics and vaccination.

OmpF

Outer membrane protein F (OmpF) is a major porin in *S. typhi* responsible for the translocation of antibiotics. The functional unit of OmpF is a homotrimer. Each monomer of molecular weight 37 kDa forms a β -barrel structure having 16 membrane spanning β -strands. The OmpF porin forms three large water-filled channels per trimer, allowing the diffusion of small hydrophilic molecules such as nutrients, antibiotics and waste products across the outer bacterial membrane [36]. OmpF allows the passage of drugs such as quinolones, tetracyclines, and β -lactams [37,38]. The understanding of the structure function relationship of *S. typhi* OmpF is important for the development of new drugs against

typhoid. The mutants of OmpC and OmpF of *S. Typhimurium* showed attenuated virulence [39]. OmpC and OmpF of *S. Typhi* were shown to confer lifelong, specific bactericidal antibody response [40]. Researchers showed that denatured forms of recombinant OmpC and OmpF of *S. enterica* ssp *Typhi* are immunogenic proteins in murine models. They suggest those two OMPs as suitable vaccine candidates. However, no in vivo challenge study was performed on various types of porins. [35]. There are two variable regions in OmpC and OmpF porin proteins useful for a typhoid fever diagnostic test based on antigen - antibody interaction [41]. OmpF generates only bactericidal antibodies after boosting, suggesting that the long-lasting antiOmpF antibodies recognize mainly epitopes not exposed on the bacterial surface [40]. Marked IgG responses in mice immunized with OmpF support the immunogenic nature of recombinant OmpF suggestive of its application as a good immunogen for vaccine studies.

OmpL

Outer membrane porin L (OmpL also called YshA) was predicted as a 230-residue TMBB, which contained 10 trans membrane strands and an N-terminal signal sequence, recombinant OmpL localizes to the outer membrane when expressed in *Escherichia coli* [42]. Studies demonstrated that OmpL an outer membrane protein of *S. Typhimurium*, is likely to adopt a β barrel conformation consisting of 12 antiparallel β strands connected by 6 long, flexible, surface-exposed loops and 5 short periplasmic turns. OmpL was used as a candidate protein not only because of its location on the bacterial surface but also because of its abundant expression [42,43]. Analysis of the amino acid sequence of the OmpL indicates OmpL is a widely distributed and conserved outer membrane protein among different *Salmonella* serovars which raises the possibility that OmpL could be used in diagnostics or as part of a subunit or conjugate vaccine. Large number of studies with bacterial outer membrane molecules as candidate vaccines has shown considerable promise [11,12]. OmpL may induce strong protective immunity in the host and remain attractive vaccine targets. Researchers suggest that OmpL immunization is protective in murine infection, reducing bacterial loads and increasing survival time after infection, which confirms and extends that OmpL-based vaccine, may be a potential vaccine candidate against multiple *Salmonella Typhimurium* infection. When mice immunized with rOmpL were challenged with lethal doses of *S. Typhimurium*, 100% of them were protected against salmonellosis. However, the task of producing and testing OmpL in subunit vaccine trials remains a challenging objective. The ELISA results indicated that the

protein elicits a significant humoral response. OmpL is a potential vaccine candidate against *Salmonella* infections and also promises to be a potent adjuvant [43].

Omp A

Outer membrane protein A (OmpA) is a major heat-modifiable OMP in *Escherichia coli* and is one of the best characterized OMPs. Originally purified in 1977, the molecular mass of OmpA was demonstrated to be 33 kDa, and since then various studies have identified its molecular mass to range from 28 to 36 kDa depending on the temperature and conditions to which it is subjected prior to SDS-PAGE [44]. OmpA proteins are characterized by an N-terminal domain that forms an eight-stranded, anti-parallel β barrel, which is embedded in the outer membrane and the C-terminal domain is globular and located in the periplasmic space. Thus, the protein spans the membrane many times with the protein interacts with lipid present in outer membrane non-covalently in *Chlamydia*, a gram negative bacteria [45]. So, only a part of the protein, N-terminal is exposed for binding of phages and other properties while other portion, C-terminal is not essential for its localisation or for its function as phage receptor.

OmpA has both structural and ion-permeable porin roles, with its ionic pore controlled by a salt-influenced electrostatic gating mechanism that allows bacterial survival during osmotic stress [46] OmpA is abundant in the membrane with estimates of approximately 100,000 copies per cell [47]. Because OmpA homologues are highly immunogenic, in certain Gram-negative bacteria, OmpA homologues can serve as potential targets for vaccine development. *Salmonella* spp. are severe enteric pathogens in many animal species. Although, OmpA is not so critical to pathogenesis of enteric or systemic salmonellosis, *Salmonella* spp. OmpA is highly immunogenic as determined by detection of antibodies in mice infected with attenuated *Salmonella* and humans with confirmed cases of typhoid [48] Vaccination of BALB/c mice with formalin-killed *Salmonella* enhanced resistance against challenge, and survivors had high anti-OmpA antibodies [49] which suggests *Salmonella* OmpA may exist in more than one conformational form. *Salmonella* OmpA is immune-stimulatory as demonstrated by stimulation of IFN-g production and enhanced expression of MHC and co-stimulatory molecules in dendritic cells and/or T cells and may play a role in modulation of the immune response in salmonellosis [50,51]. OmpA is surface exposed and highly immunogenic stimulating bactericidal antibodies in the presence of complement [52].

Hence, studies showed that Omp A function in bacterial structure, physiology, and also adaptation to environmental stresses, whereas in disease, they can serve as virulence factors causing adhesion, invasion, and damage of host tissue or evasion of host defenses resulting in clinical disease or death. Various OmpA homologues also serve as targets for host innate and adaptive immune responses resulting in increased host resistance against infection. As per studies about pathogenic and immunogenic roles of the OmpA family of proteins, their involvement in natural and recombinant-derived vaccines will likely increase.

Porins are part of outer membrane proteins in Gram-negative bacteria stimulating immune system [35,53,54]. Researchers showed that denatured forms of recombinant OmpC and OmpF of *S. enterica* ssp Typhi are immunogenic proteins in murine models [35]. The major porins in *Salmonella* viz OmpC, OmpF, OmpD are widely distributed and conserved among different *Salmonella* serovars which raises the possibility that they could be used in diagnostics or as part of a subunit or conjugate vaccine. Various studies have shown that a crude preparation of outer membrane proteins (OMPs) of *Salmonella* evokes strong immune response and induces a protective immunity against infection caused by diverse Gram-negative bacteria. Therefore, from the studies on the above different Outer membrane proteins (OMPs) discussed it can be concluded that OMPs of *Salmonella* seem to have strong immunogenic potential and also have been implicated as possible candidates for conferring protection against typhoid. Studies have also shown that some recombinant OMPs also may evoke strong immune response in animals and some of these proteins have very strong potential for the development of a subunit vaccine against *Salmonella*. Thus, outer membrane proteins (OMPs) could be a candidate antigen for diagnostics and vaccination and may act as promising potential vaccine candidates against *Salmonella*.

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