

Vaccination against urinary tract infections caused by *E. coli*

Hartwig Wilhelm Bauer, Hira Shams, Ricarda Michaela Bauer

Urological Clinic Maximilian, Urological Clinic LM University München, München, Germany
Email: mail@praxis-bauer.de

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ABSTRACT

Uropathogenic *Escherichia coli* (UPEC) causes uncomplicated urinary tract infection (UTI) depicts a prevalent and potentially uncompromising infectious disease. In this analysis, we explained the functions of an immunoproteomics concept to vaccine development that has been successfully employed to recognize vaccine targets in other pathogenic bacteria. Pyelonephritis strains *E. coli* CFT073 are used for outer membrane isolation mimics urinary tract environment in which iron limitation, osmotic stress, human urine, and exposure to uroepithelial cells are included. During experiments of UTI, the antigens that induce the humoral immune response is identified, two-dimensional gel electrophoresis are employed for the isolation of outer membrane protein and probed using pooled antisera from 20 CBA/J mice chronically infected with *E. coli* CFT073. 23 total outer membrane antigens, in which a unique iron compound receptor is included, are reacted with antisera and were identified by mass spectrometry. These antigens comprises of proteins with known functions in UPEC pathogenesis such as, ChuA, IroN, IreA, Iha, IutA, and FliC. These all information and data elaborate that these factors are associated with virulence during UTI are directed by antibody response. We also represents that the genes encoding ChuA, IroN, hypothetical protein c2482, and IutA are significantly more prevalent among UPEC strains than among fecal-commensal *E. coli* isolates. Therefore we concluded that, the outer membrane antigens are identified in this study are conserved, could be reflective part for the UTI vaccine generated to induce protective immunity against UPEC infections.

Keywords: Pyelonephritis; Outer Membrane Antigen; Iron Compound Receptors

1. INTRODUCTION

Urinary tract infection (UTI) is a widespread infectious

disease having possibly uncompromising complexities. Almost 11.3 million communities suffered from UTIs come off in United States each year, with an annual cost of \$1.6 billion [1]. If UTI left untreated, they can lead to more extreme conditions including acute pyelonephritis, bacteremia, and renal scarring. Furthermore, increasing rates of antimicrobial resistance among uropathogens will likely complicate future treatment of these infections. [2,3] Consequently, there is an urgent public health need to develop an efficacious vaccine to prevent UTI.

The most common etiological agent of UTIs accounts for greater than 80% of these infections is Uropathogenic *Escherichia coli* (UPEC) [4]. Most of the virulence determinants expedite the ability of UPEC to colonize the urinary tract and exert degenerative changes in cell, including Type 1 fimbriae, [5] P fimbriae, [6] Dr adhesins, [7] hemolysin, [8,9] cytotoxic necrotizing Factor 1, [10] flagella, [11] capsule polysaccharide, [12] lipopolysaccharide O antigen, [13] and TonB-dependent iron transport systems [14].

currently, the perseverance of the *in vivo* transcriptome—the set of all RNA molecule including mRNA, tRNA, rRNA and other non coding RNAs, of UPEC more accentuated the significance of adhesion and iron salvage during UTI, because genes associated in these processes were exceeding in the number of receptors on the surface of target cells, making the cells more sensitive to the agent during experimental infection. [15] Most of the factors associated with virulence have been experimented as vaccine target due to the medical and economic impact of UPEC and UTI. For instance, immunization with FimH—the type 1 fimbrial adhesin, specifically decreased bladder colonization in C3H/J mice [16] and explained shield in a primate model of UTI. [17] Furthermore, a subunit vaccine using PapG—the P fimbrial adhesin, complexed with its periplasmic chaperone, PapD, specifically shielded primates from histological indications of pyelonephritis, [18] as periplasmic chaperone used to enhance functional secretion of proteins in *E. coli*.

The success rate is found to be very limited by using

hemolysin, [19] Dr Fimbriae, [20] and the siderophore receptor IroN [21] in some experiments for the induction of protective immunity against uropathogenic *E. coli*. Uropathogens mixture that is deactivated by heat treatment and mucosal immunization specifically decreased urinary tract infection events among in women. This is occurred recent clinical trials of Phase II [22]. Although these vaccine preparations are not considered to be the best for providing long term protection. So, it is necessary to identify some other antigens, which is mainly employed for the production of vaccine against UPEC.

Former endeavors for vaccine production against UPEC were relied initially on some specific virulence factors or whole cells, genomic and proteomic process provide an extensive approach in order to design vaccine. Reverse vaccinology, the most advance technique used for screening of genome of serogroup B *Neisseria meningitidis* that exhibit a conserved surface exposed antigens among *Neisseria meningitidis* strains [23]. Antibody response that is activated in immunized animals by the antigens, were employed successfully to produce a universal multivalent vaccine against UPEC. Meanwhile, the immunoproteomics methods that is the isolation and mass spectrometric identification of MHC (major histocompatibility complex) binding peptides, purification and identification of protein antigens binding specific antibodies (or other affinity reagents, and comparative immunoproteomics to identify proteins and pathways modulated by a specific infectious organism, disease or toxin. These methods are linked with the bacterial proteomes screening by using sera from infected individuals, are mainly employed for antigen identification in pathogens particularly *Campylobacter jejuni* [24], *Anaplasma marginale*, [25] *Bartonella henselae*, [26] and *Klebsiella pneumonia* [27]. The addition of novel proteins and non virulence factors as candidate for immunization is the main advantage of these genomics and proteomics techniques that is proteins which are particularly eliminated from the strategies used for traditional vaccine design.

Innate and adaptive immune mechanisms are involved for immune response to UTI. Toll-like receptors (TLRs) are a class of proteins that play a key role in the innate immune system [28,29]. They are single, membrane-spanning, non-catalytic receptors that recognize structurally conserved molecules derived from microbes. Once these microbes have breached physical barriers such as the skin or intestinal tract mucosa, they are recognized by TLRs, which activate immune cell responses. [30] neutrophil infiltration is considered to be the primary mechanism of the innate immune response to control UTI [31]. Because, the mice have lack of neutrophils contains degenerated capability to eradicate UPEC infection compared to neutrophil-replete animals. Neverthe-

less, adaptive immune responses also integrate to immunity against UPEC. High susceptibility to infection with UPEC are found in severe combined immunodeficient mice that implicating B and T cell mediated immunity take part in eradication of bacteria particularly. As a result, the natural infections with UPEC are elicited by many aspects of immune response. And this clue declared that the hummoral response induced by the vaccine is actually protected from uncomplicated UTI.

Because an antibody response is probably a particular component of the adaptive immune response to UPEC, [32,33] ideal vaccine targets should be surface exposed and accessible to circulating immunoglobulins. Surface exposed proteins in *E. coli* are attached in the outer membrane. So the group of anticipated vaccine candidates is depicted on the outer membrane protein (OMPs) of UPEC. Meanwhile it is necessary that ideal vaccine candidates must be particular to pathogenic *E. coli* to refrain cross reactivity with commensal strains. In this analysis, powerful vaccine targets in UPEC are being identified by immunoproteomic methods. *E. coli* CFT073 grown under certain conditions and OMPs has been purified that mimics the urinary tract with antisera from the chronically infected mice that shows 23 antigenic OMPs are found that elicit the strong immune response during infection. Several OMPs are UPEC specific and novel iron-induced protein as well. This indicated that these antigenic OMPs represents novel identified targets for the production of multivalent vaccine against UTI agents (**Figures 1 and 2**).

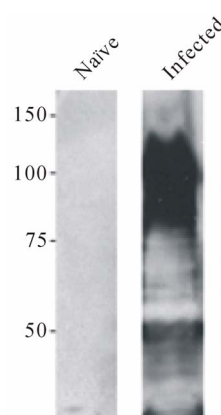


Figure 1. Specificity of antisera from chronically infected mice: Specificity of antisera generated against *E. coli* CFT073. Western blots of a CFT073 whole-cell lysate electrophoresed on a 12% polyacrylamide gel and probed with nonimmune sera from naïve CBA/J mice (left) or antisera from chronically infected mice (right) are shown. Molecular mass standards are shown in kDa.

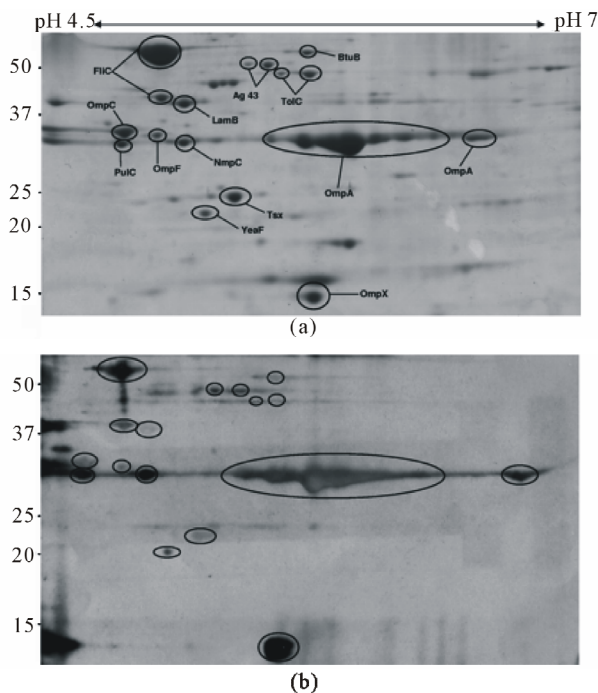


Figure 2. Outer membrane antigens from UPEC are antigenic: Identification of antigenic OMPs of *E. coli* CFT073. (a) Colloidal Coomassie-stained 2D-PAGE gel of outer membrane fractions isolated from CFT073 cultured in rich medium; (b) Western blot of 2D-PAGE gel probed with pooled antisera from chronically infected CBA/J mice. Proteins annotated in A are seroreactive and were identified by mass spectrometry. Molecular mass standards are shown in kDa.

2. DISCUSSION

OMP antigens that are produced during UPEC infection is being recognized by immunoproteomic approach and depicts the first wide screened for vaccine targets for this UPEC. *In vivo* studies shows that 23 outer membrane antigens from UPEC are expressed and capable of eliciting the humoral response by using immunoreactive antisera from chronically infected mice. Hypothetical protein c2482 has been identified in this screen. This novel antigen expressed in iron limitation under certain conditions. Moreover, it is also discussed that the genes encoding at least four of these OMPs, *chuA*, c2482, *iroN*, and *iutA* are the most predominate genes among UPEC strains as compare to fecal commensal UPEC strains. This shows that these conserved antigenic OMPs may be useful and great target for a vaccine against UPEC.

There has been wide investigation for the deterrent of UTI by evaluating the virulence factors of UPEC as protected immunogens for provoking adaptive immune response. Though, the disparate trait of UPEC isolates proposes that increased vaccine targets will be attained to secure protection against a widespread series of strains.

Therefore, there is main focus on identifying conserved outer membrane antigens of UPEC that may be used

in vaccine against UTI.

In addition to Porins and adhesins—a beta barrel protein that forms channels that allow certain cellular material to pass through cell membranes. Any type of material, below a certain size, may pass through the porins, which join together to create cylindrical passageways that allow various molecules to passively move from one side of a membrane to another, screen recognized a new outer membrane of antigen of UPEC. c2482, a speculated iron receptor, was identified during iron limitation conditions, proliferation in human urine and association with the cells of bladder epithelial. These recommendations conclude the anticipated outer membrane localization of this protein and accommodate indirect affirmation of its engrossment in iron redemption as well. Current studies show the high expression of c2482 during murine UTI [15] and this protein is considered to be the most eliciting proteins during the growth of CFT073 in human urine [34], proposing that this immunogen is a promising vaccine target.

This analysis also suggests during experimental UTI *E. coli* CFT073 expresses proteins. However, the transcript level gene expression is discussed by transcriptome of murine UTI *in vivo* during infection, [15] the recent outcomes assist and boost former research by observing *in vivo* protein expression. As a result, the 23 seroreactive OMPs identified are expressed in the UPEC outer membrane during infection. This data coincide with the former work, as 17 of these 23 seroreactive OMPs were among the top 30% of CFT073 transcripts detected *in vivo*. Indeed, 11 of these 17 OMPs were upregulated at least twofold during experimental UTI. Besides, these discoveries aid in understanding of the pathogenesis of UPEC as well.

The most outstanding benefit of this recent analysis study over other immunoproteomics analyses is the involvement of several optimum cultured conditions, has designed to mimic natural environment of the infectious agent. While these different conditions only modestly extend the number of antigens recognized, the approach nevertheless enhances our confidence that few major outer membrane antigens were excluded from the screen. This method also declared different iron-related antigens, *ChuA*, *Iha*, *IroN*, c2482, and *IreA*, which were identified during growth under only three culture conditions: iron limitation, human urine, and exposure to bladder epithelial cells. As these environments likely elicit iron loss, it is not astounding that they also activate the expression of additional proteins involved in iron acquisition. A recent study from our laboratory concealing these outcomes, showing the induction of these five OMPs during culture in human urine as well as their repression during growth in iron-replete medium [34]. It is interesting that the genes encoding three of the iron-related antigens, *chuA*,

iroN, and *c2482*, were also found to be both conserved and UPEC specific by dot blot hybridization. Given the well-established role of iron acquisition in pathogenesis, we speculate that UPEC, compared to commensal *E. coli*, expresses a greater range of iron receptors in response to iron-limiting environments.

Fimbriae—the anchored outer membrane surface structures were thoroughly missing by western blots that predict the infected mouse antisera used in the screen consists of fimbriae-specific antibodies. In the sera of primates infected with UPEC, [18] anti-P fimbria immunoglobulin G (IgG) was discovered undoubtedly. Nevertheless, fimbrial proteins mostly need some more steps for solubilization [35] and during preparation they are clipped easily from the surface. These proteins are probably absent from 2D gel electrophoresis. Additionally, an absolute 2D-polyacrylamide gel electrophoresis analysis of the *E. coli* outer membrane proteome also deficit the apprehension of fimbriae, [36] this finding shows that this is the most expected outcome.

The identification of important outer membrane antigens expressed by *E. coli* CFT073 during UTI with limitations of several congenital stipulations with this technique is considered to be the major breakthrough. OMPs are seroreactive that are less abundantly found, could have been below the limits of detection of 2D-PAGE colloidal Coomassie staining and mass spectrometry analyses. IgG response induced by primary antigens during infection are detected as well. Secretory IgG is considered to play a pivotal role in the clearance of UTI [37] as UPEC is a mucosal pathogen but these limitations do not affect harshly on our breakthrough.

Many OMPs recognized as being powerful protective antigens have roles in the virulence of UPEC and other pathogens. *E. coli* strains lacking the heme/hemoglobin receptor *ChuA* [14], the aerobactin receptor *IutA*, [14] the salmochelin receptor *IroN*, [38] the iron-responsive element *IreA*, [39] or the iron-regulated adhesin *Iha* [40] were particularly outcompeted by the wild-type strain in a mouse model of UTI, determining the significance of iron possession to the fitness of this pathogen *in vivo*. Moreover, isogenic mutants lacking the major flagellum subunit *FliC* were similarly outcompeted by wild-type *E. coli* CFT073 [41]. Therefore, this data provide evidence that a humoral response is generated against these virulence-associated factors during murine infection. Furthermore, while a role in pathogenesis is likely not a requirement for a vaccine target, it may be beneficial, as neutralizing antibodies may block critical functions of such targets upon infection.

In recent studies the outer membrane antigen are identifies contain additional features that assist in assumed roles as vaccine candidate. β -barrel structures are formed by several outer membrane antigens including the iron

compound receptors. Although, most of the β -barrel proteins will be plunged in the membrane, extracellular loops accommodate surface exposed regions due to the neutrophils, as they are the most pivotal constituent of immune response to UTI, [31] opsonizing antibodies against such surface-exposed proteins may be essential to enhance phagocytosis at the site of infection. Furthermore, some of the OMPs are identified in this screen are associated in cellular processes that are typical for *in vivo* bacterial growth like iron redemption. Bacterial clearance from urinary tract can also be possible by ceasing the function of these proteins through neutralizing antibodies. Eventually, four of the antigens that are *ChuA*, *c2482*, *IutA*, and *IroN* are conserved among most UPEC strains and mostly absent in some fecal commensal-*E. coli* strains. Some genetic assortments at the amino acid level are detected in each of these antigens, among sequenced strains of pathogenic *E. coli*. Each antigen is 90 to 100% similar between strains. This concludes as these proteins may produce protection against a wide series of pathogenic strains, if attained as vaccine target, which is least cross react with normal flora.

3. CONCLUSION

Uncomplicated UTI that is secured by a vaccine would have outstanding public health advantages. The information and data existed in this study depicts a first step towards the development of such a broadly protective vaccine against UPEC. Some of the outer membrane antigens have not observed yet as vaccine targets for UPEC. Moreover, *c2482*, a unique antigen recognized in this screen shows not only a novel vaccine target but a newly recognized OMP as well that could function as an iron compound receptor. Although, lots of examinations are required to enhance these observations and investigate immunization with these antigens before an efficient UTI vaccine can be cultivated.

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