Anti-PcrV antibody strategies against virulent Pseudomonas aeruginosa

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Keywords: antibody, PcrV, Pseudomonas aeruginosa, type III secretion system, V-antigen

Abbreviations: CF, cystic fibrosis; Fab, fragment antigen binding; Fc, fragment crystallizable region; immunoglobulin G, IgG; mAb, monoclonal antibody; MDR, multidrug resistant; MDRP, multidrug resistant *Pseudomonas aeruginosa*; *P. aeruginosa*, *Pseudomonas aeruginosa*; TTSS, type III secretion system; TTS, type III secretory; VAP, ventilator-associated pneumonia

Pseudomonas aeruginosa is an opportunistic bacterial pathogen that causes fatal acute lung infections in critically ill individuals. Its pathogenesis is associated with bacterial virulence conferred by the type III secretion system (TTSS), through which P. aeruginosa causes necrosis of the lung epithelium and disseminates into the circulation, resulting in bacteremia, sepsis, and mortality. TTSS allows P. aeruginosa to directly translocate cytotoxins into eukaryotic cells, inducing cell death. The P. aeruginosa V-antigen PcrV, a homolog of the Yersinia V-antigen LcrV, is an indispensable contributor to TTS toxin translocation. Vaccination against PcrV ensures the survival of challenged mice and decreases lung inflammation and injury. Both the rabbit polyclonal anti-PcrV antibody and the murine monoclonal anti-PcrV antibody, mAb166, inhibit TTS toxin translocation. mAb166 lgG was cloned, and a molecular engineered humanized anti-PcrV IgG antigenbinding fragment, KB001, was developed for clinical use. KB001 is currently undergoing Phase-II clinical trials for ventilator-associated pneumonia in France and chronic pneumonia in cystic fibrosis in USA. In these studies, KB001 has demonstrated its safety, a favorable pharmacokinetic profile, and promising potential as a nonantibiotic strategy to reduce airway inflammation and damage in P. aeruginosa pneumonia.

Introduction

Pseudomonas aeruginosa is an opportunistic bacterial pathogen that causes fatal acute lung infections in critically ill individuals.¹⁻⁴ Its pathogenesis is frequently associated with the development of septic shock and multiple organ failure, because certain strains of *P. aeruginosa* have the ability to cause necrosis of the lung epithelium and disseminate into the circulation.^{5,6} Damage to the lung epithelium is associated with the expression of toxins that are directly translocated into eukaryotic cells through the type III

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secretion system (TTSS).^{7,8} The P. aeruginosa V-antigen PcrV, a homolog of the Yersinia V-antigen LcrV, is an indispensable contributor to the process of TTS toxin translocation. Vaccination against PcrV ensures the survival of challenged mice and decreases lung inflammation and injury (Table 1).9 Both the rabbit polyclonal anti-PcrV antibody and the murine monoclonal anti-PcrV antibody, mAb166, inhibit the TTS toxin translocation.^{10,11} Till date, the therapeutic effects of anti-PcrV antibodies are the subject of most published studies on P. aeruginosa infections in animal models. Previously, we cloned mAb166 from a hybridoma, and humanized this monoclonal antibody for potential clinical use.¹² This humanized anti-PcrV IgG antigen-binding (Fab') fragment, KB001, is currently undergoing Phase-II clinical trials for ventilator-associated pneumonia (VAP) in France and chronic pneumonia in cystic fibrosis (CF) in USA (Table 1).^{13,14} In this review, we summarize the development and characterization of anti-PcrV antibodies, including the early outcomes of their Phase-II clinical trials. Figure 1

TTSS and P. Aeruginosa PcrV

Understanding the precise mechanism of acute lung injury caused by *P. aeruginosa* is key to identifying new therapeutic targets. Reportedly, the ability of this bacterium to cause epithelial injury, disseminate into the circulation, and avoid host innate immune responses is due to TTSS.^{7,8,15,16} Although most toxins produced by bacteria are secreted into the surrounding extracellular environment via classical type I or II secretion systems, recent studies in gram-negative bacteria have identified a specific group of toxins that are injected directly into adjacent host cells. This protein secretion mechanism is termed TTSS. TTSS is found in a wide variety of pathogenic strains of gram-negative bacteria, including *Salmonella*, *Shigella*, and *Yersinia*.^{17,18} Figure 2

TTSS was shown to contribute to the virulence of *P. aeruginosa*.^{7,8} In addition, many *in-vitro* clinical reports have recently shown that patients infected with strains of *P. aeruginosa* expressing TTSS have a higher risk of mortality than those infected with strains not expressing TTSS.^{15,19-22} In our *in vitro* cell culture experiments, *P. aeruginosa* strains exhibited cytotoxicity if they



Figure 1. The type III secretory apparatus of *Pseudomonas aeruginosa*: A basal component, a **needle structure**, and a translocon. The type III secretory (TTS) apparatus also known as the *injectisome* comprises many protein components. The basal component comprises an outer ring PscC, periplasmic joint PscJ, ATPase PscN, and more, and is the mechanism by which TTS toxins pass through the bacterial cell membrane. The needle structure comprises approximately 140 PscF proteins. The translocon comprises PcrV, PopB, and PopD, and is the mechanism by which TTS toxins translocate through the eukaryotic cell membrane. PcrV forms a cap structure on the tip of the secretory needle rod.

produced TTS products.^{23,24} In our animal models of P. aeruginosa pneumonia, airspace instillation of cytotoxic P. aeruginosa strains caused consistent alveolar epithelial injury, progressive bacteremia, and septic shock in sheep, rabbits, rats, and mice.²⁵⁻²⁹ In contrast, airspace instillation of noncytotoxic strains unable to produce TTS toxins did not cause a systemic inflammatory response or septic shock, despite a potent inflammatory response in the lung.¹⁶ Therefore, blocking TTSS-associated virulence is key to improving acute lung injury and mortality caused by cytotoxic P. aeruginosa. However, neutralizing TTS toxins using a specific antibody is ineffective, because these toxins are translocated from the bacterial cytosol directly into the cytosol of target eukaryocytes through TTSS without entering the extracellular environment. Therefore, it is necessary to target the toxin secretion and/or translocation system to disrupt toxin translocation and abrogate virulence. PcrV, a translocational component of the P. aeruginosa TTSS, is reportedly a homolog of the Yersinia V-antigen LcrV.^{9,30} Immunoprotective effects of Yersinia V-antigen have been reported in animal models of pneumonic plague.³¹ In our previous study, active immunization with Escherichia coli recombinant PcrV was protective in mice infected with lethal doses of P. aeruginosa and the mortality of the infected mice decreased following the administration of an antibody to PcrV.¹⁰ Figure 3



Figure 2. The mAb166 blocking epitope of PcrV. PcrV consists of 294 amino acids. The C-terminal center region between amino acids 144–257, consisting of 114 amino acids and located between 2 coiled-coil regions, is the blocking epitope for the monoclonal anti-PcrV IgG antibody, mAb166. This blocking epitope overlaps with both the channel-forming region for erythrocyte hemolysis and the blocking region of the homologous *Yersinia* LcrV, based on a sequence alignment between PcrV and LcrV. mAb166 binds to this epitope and blocks the translocation of type III secretory toxins. Amino acid 204 corresponds to the hypervariable region of *Yersinia* LcrV.

The Molecular Mechanism of The TTS Toxin Translocation

The process of toxin translocation across the eukaryotic plasma membrane and directly into the cytosol via TTSS has been extensively studied in *Yersinia* over the last 25 years.^{32,33} The translocator proteins LcrV, YopB, and YopD are involved in the process of translocation via the *Yersinia* TTSS, and *lcrV*-null mutant *Yersinia* is unable to translocate TTS toxins.^{34,35} The homologous translocator proteins PcrV, PopB, and PopD are involved in toxin translocation in *P. aeruginosa*.³⁶ Isogenic mutant, *P. aeruginosa*, lacking *pcrV* and PcrV expression cannot translocate toxins and is unable to affect eukaryotic cells.¹⁰ Figure 4

The supramolecular structure of the TTS apparatus, also known as the *injectisome*, has been well studied over the last 15 years.³⁷⁻³⁹ The *injectisome* is evolutionarily associated with bacterial flagella, and comprises:

- 1. a basal body, anchored in the bacterial outer and inner membranes;
- 2. a needle structure, extending from the outer surface of the membrane; and
- 3. translocon proteins, including a needle cap structure and proteins that form pores in the plasma membrane of the target eukaryotic cell.

In *Yersinia*, the needle has a helical structure and comprises oligomers of more than

100 YscF subunits, and LcrV, which forms a distinct mushroomlike structure at the needle tip acts as a bridge during the translocation of TTSS effectors between the YscF secretion needle and the translocational pore, composed of YopB and YopD, on the eukaryotic cell membrane.⁴⁰ An oligomeric donut-like LcrV-protein assembly structure, with approximate internal and external diameters of 3-4 and 8-10 nm, respectively, was observed in vitro, and in the atomic model of the tip complex, the N-terminal globular domain of LcrV forms the base, the central globular domain forms the head, and the tip complex itself is likely formed by pentameric LcrV.⁴¹⁻⁴⁵ For translocating TTS, the formation of a translocational pore on the eukaryotic plasma membrane, involving a cap protein such as LcrV or PcrV, was demonstrated.⁴⁶⁻⁴⁹ In *Yersinia*, the pore associated with LcrV was approximately 3 nm in diameter, whereas in *P. aeruginosa*, the homologous PcrV protein induces pores of approximately 2 nm in diameter.⁴¹ The regions important for translocational pore size were localized to amino acids 127-195 in LcrV and 106-173 in PcrV.^{9,11,35} The size of the translocational pore correlates with its ability to translocate Yop effectors into host cells, suggesting that LcrV is a size-determining structural component of the



Figure 3. Predicted tertiary structure of PcrV. Based on the structural information for LcrV (Swissplot IR6F), the tertiary structure of PAO1 PcrV was predicted by the protein structure prediction server RaptorX.⁵⁸ PcrV, consisting of 294 amino acids, forms a central shaft structure, with coiled-coil double strands and 2 globular domains at either end. The C-terminal globular domain includes amino acid 204, corresponding to the hypervariable region of *Yersinia* LcrV, and amino acids 144–257, recognized as the mAb166 blocking epitope.

Yop translocon.⁴¹ The mechanism of translocational blockade by specific antibodies against LcrV or PcrV remains poorly elucidated. However, specific antibodies probably interfere with the toxin passage through the translocational pore formed by LcrV or PcrV, because the binding sites of such antibodies overlap with the pore-forming regions of LcrV and PcrV.⁹

The Anti-PcrV Polyclonal Antibody

Immunization with PcrV

The *in-vitro* inhibition of translocation by anti-PcrV IgG suggested that anti-PcrV or immunization with PcrV may protect against the lethal effects of cytotoxic *P. aeruginosa*.¹⁰ Mice were immunized with endotoxin-free rPcrV or the recombinant TTS toxin ExoU, and subsequently challenged via airspace instillations of a lethal dose of the cytotoxic *P. aeruginosa* strain PA103. When survival was measured, both vaccines protected the mice to some degree but animals immunized with rPcrV had significantly fewer bacteria in their lungs. Next, to determine whether prophylactic intervention was possible, mice were passively



Figure 4. PcrV forms a ring structure at the tip of the secretion needle. A. Monomeric PcrV. B. Five PcrV aligned in a circle. C. Top view of the ring structure formed by pentameric PcrV.^{43,44}

immunized with preimmune rabbit IgG or rabbit anti-PcrV IgG or anti-ExoU IgG prior to airspace instillation of PA103. Anti-PcrV IgG provided complete protection to a lethal infection, whereas anti-ExoU IgG provided partial protection, which was significantly different from the administration of control IgG.¹⁰ If PcrV is accessible for neutralization, then concomitant administration of the bacterial inoculum with anti-PcrV IgG should completely protect against lung injury and lethality. Then, either anti-PcrV IgG or control IgGs were mixed with the inoculum (at a 10-fold higher dose than the lethal inoculum) prior to instilling the bacteria into the lung and the survival was examined. Only anti-PcrV IgG was protective against this extreme infection.

In the next series of experiments, studies were performed in mice with lethal pulmonary *P. aeruginosa* infections to examine the therapeutic effects of rabbit polyclonal anti-PcrV IgG on survival after this infection.⁵⁰ Nonspecific control IgG, anti-PcrV IgG, or anti-ExoU IgG were administered intravenously after airspace instillation of a lethal dose of PA103. All mice that received anti-PcrV IgG 4 h after airspace instillation survived, but fewer than 20% of mice that received either control IgG or anti-ExoU IgG 1 or 4 h after infection survived only for more than 3 days. Anti-PcrV IgG significantly decreased inflammation and lung injury, even when administered 4 h after infection, whereas increased recruitment of neutrophils and destruction of the alveolar structure were observed in the lungs of mice treated with nonspecific control IgG.⁵⁰

The mechanism of anti-PcrV blockade

In vitro study, anti-PcrV IgG demonstrated no bactericidal activity and did not affect bacterial growth.⁵⁰ It did not affect the other toxic exoproducts such as elastase, exotoxin A, alkaline protease, and pigments. We still don't know whether anti-PcrV blockade affect the biofilm production or not while it was reported that biofilms and type III secretion are not mutually exclusive in *P. aeruginosa.*⁵¹ In the co-culture of eukaryotic cells such as macrophages or lung epithelial cells with cytotoxic *P. aeruginosa*, anti-PcrV IgG prevented the eukaryotic cells from bacterial intoxication. Therefore, to directly test whether anti-PcrV IgG can inhibit type III translocation, the TTS toxin translocation was quantified by measuring cyclic AMP levels in macrophages infected with an isogenic mutant of *P. aeruginosa* that secretes only ExoY, adenylate cyclase type III secretory toxin. As a result, neutralization of PcrV by the antibody significantly decreased ExoY translocation and prevents or inhibits the intoxication of eukaryotic cells. In addition, anti-PcrV IgG restored phagocytic function of macrophages.¹⁰

Use of the anti-PcrV polyclonal antibody against sepsis

Studies on the pathogenesis of septic shock following *P. aeru-ginosa* pneumonia have shown that airspace instillation of cytotoxic PA103 into rabbits caused consistent alveolar epithelial injury, progressive bacteremia, and septic shock.¹⁶ However, instilling a noncytotoxic, isogenic mutant strain (PA103 ΔUT), unable to produce TTS toxins, did not cause either a systemic inflammatory response or septic shock, despite a potent inflammatory response in the lung.¹⁶ These results suggest that injury to the alveolar epithelium caused by TTSS-associated virulence allows the release of proinflammatory mediators into the circulation that are primarily responsible for septic shock. This demonstrates the importance of the compartmentalization of inflammatory mediators in the lung and the crucial role of bacterial cytotoxins in causing alveolar epithelial damage in the pathogenesis of acute septic shock in *P. aeruginosa* pneumonia.

To analyze the therapeutic effects of anti-PcrV IgG in a lung infection associated with sepsis, a rabbit model in which acute septic shock was induced by airspace instillation of *P. aeruginosa* was utilized.⁵⁰ Rabbits receiving airspace instillation of PA103 consistently exhibit septic physiology within 8 h.⁵⁰ By

Table 1. Reports on antibodies against Pseudomonas aeruginosa PcrV

Authors	Antibody	Species	Target	Country	Year	Ref
Sawa T, et al.	Rabbit polyclonal IgG	mice	pneumonia	USA	1999	10
Shime N, et al.	Rabbit polyclonal IgG & F(ab)'	rabbit	sepsis	USA	2001	50
Frank DW, et al.	Murine monoclonal IgG (mAb166)	mice	pneumonia	USA	2002	11
Faure K, et al.	Murine monoclonal IgG (mAb166), and Fab	rat	pneumonia	USA	2003	54
Neely AN, et al.	Rabbit polyclonal IgG	mice	burned	USA	2005	51
Imamura K, et al.	Rabbit polyclonal IgG	mice	chronic pneumonia	Japan	2007	52
Baer M, et al.	Humanized Fab' (KB001)	mice	pneumonia	USA	2008	12
Moriyama K, et al.	Rabbit polyclonal IgG (affinity purified)	mice	pneumonia (immunocompromised)	USA	2009	53
Song Y, et al.	Murine monoclonal (mAb166)	mice	pneumonia	USA	2012	55
François B, et al.	Humanized Fab' (KB001)	human	Ventilator-associated pneumonia	France	2012	13
Milla CE, et al.	Humanized Fab' (KB001)	human	cystic fibrosis	USA	2013	14
Wang Q, et al.	Monoclonal (mAb166)	mice	pneumonia with MDRP	China	2014	56

monitoring parameters of septic shock, including hemodynamics, metabolic acidosis, bacteremia, and the production of inflammatory cytokines, the therapeutic effects of the anti-PcrV IgG were analyzed in detail. In these experiments, either anti-PcrV IgG or control IgG was administered intravenously or intratracheally 1 h after airspace instillation of PA103. Rabbits receiving control IgG developed septic shock regardless of the route of administration. These rabbits exhibited a decrease in cardiac output to less than 70% of the baseline; mean arterial pressures decreased to approximately 60% of the baseline, and they developed severe metabolic acidosis. The rabbits that received control IgG intratracheally died from septic shock and severe hypoxemia at 8 h. Lung edema, as assessed by the lung water:dry weight ratio, was severe and progressive lung epithelial injury, quantified by the efflux of an alveolar protein tracer, was detected in rabbits that received control IgG. Significant increases in bacteremia and plasma tumor necrosis factor (TNF)- α concentrations were also observed in rabbits that received control IgG. In contrast, rabbits receiving anti-PcrV IgG intravenously or intratracheally exhibited stable hemodynamics and did not develop metabolic acidosis. Lung damage was also significantly less severe in these rabbits, particularly in those given IgG intratracheally. Bacteremia was significantly reduced from 10- to 100-fold compared with that in control IgG groups and developed later in the course of infection. The plasma TNF-a concentration was also significantly lower (10-20 pg/ml), compared with that in the control IgG groups (>100 pg/ml). Finally, the number of bacteria cultured from the instilled lung was also significantly decreased by intravenous anti-PcrV IgG. These findings indicate that anti-PcrV IgG not only has significant therapeutic effects against acute lung injury but also decreases systemic bacteremia and TNF-a concentrations and prevents septic shock.

Anti-PcrV F(ab')₂ protects animals comparably to whole anti-PcrV immunoglobulin G

To further explore the mechanism of the effect of anti-PcrV IgG, the administration of a $F(ab')_2$ fragment of anti-PcrV IgG was examined in infected animals.⁵⁰ One hundred micrograms of anti-PcrV $F(ab')_2$ administered intravenously 1 h after airspace instillation of PA103 resulted in an 80% survival rate at 1 week.

An examination of the histology of the lungs of animals that received anti-PcrV $F(ab')_2$ therapy revealed significantly decreased inflammation and lung injury, similar to that evident after anti-PcrV IgG therapy. Next, the effects of anti-PcrV F $(ab')_2$ were examined in a rabbit model of septic shock. Anti-PcrV $F(ab')_2$ (3 mg/kg) was administered intratracheally 1 h after instillation of PA103⁵⁰. This treatment provided statistically significant protection from lung damage, bacteremia, and septic shock. These effects were comparable to those achieved with whole anti-PcrV IgG molecules.

Use of the anti-PcrV polyclonal antibody in burned mice

Because P. aeruginosa is a major cause of infection in burned patients, passive anti-PcrV treatment was examined in burned mice that received a fatal P. aeruginosa challenge.⁵² Mice were given a third-degree burn, challenged with a lethal dose of P. aeruginosa, and treated intraperitoneally with either anti-PcrV IgG or control IgG. Protection against 3 different P. aeruginosa isolates was tested. Genotyping by PCR and phenotyping by immunoblotting showed all P. aeruginosa strains to all be of the invasive TTS toxin phenotype: ExoS⁺ and/or ExoT⁺, ExoU⁻, and ExoY⁺. Against all strains, the anti-PcrV treatment yielded significantly better survival than the control IgG treatment. Therefore, passive anti-PcrV immunization could protect burned mice against a fatal challenge with P. aeruginosa of an invasive TTS phenotype. These findings suggest that immunotherapy represents a possible treatment for a P. aeruginosa infection in burned hosts.

Use of the anti-PcrV polyclonal antibody against chronic pneumonia

P. aeruginosa is one of the most important pathogens in patients with chronic airway conditions, such as CF and diffuse panbronchiolitis. The effects of anti-PcrV IgG were evaluated in a mouse model of chronic *P. aeruginosa* infection.⁵³ In this model, a chronic airway infection was induced in mice by placing a plastic catheter infected with *P. aeruginosa* 6073, a human corneal isolate, in the main bronchus of the mouse. The analysis of bronchoalveolar lavage fluid (BALF) revealed that total cell numbers and neutrophil ratios in anti-PcrV IgG-treated groups were

lower than those in the control group. In addition, macrophage inflammatory protein-2, TNF- α , and interleukin (IL)-1 β concentrations in the BALF were lower in anti-PcrV IgG-treated groups than in that in controls. These data suggest that anti-PcrV IgG reduces the inflammatory reaction caused by chronic *P. aeruginosa* respiratory infection and may be useful in treating respiratory diseases. In this experiment, the changes in anti-PcrV IgG titer in mouse plasma were evaluated. For 3 weeks after administration of anti-PcrV IgG (100 µg/mouse), a significant concentration was maintained. Pro-inflammatory cytokine levels were significantly lower in the anti-PcrV IgG twice-treated group, but not in the once-treated group; therefore, pro-inflammatory cytokines appear to be beneficial in maintaining higher serum anti-PcrV titer.

Use of affinity purified antibody and truncated vaccines in neutropenic mice

Rabbit-derived polyclonal anti-PcrV IgG was affinity purified using rPcrV covalently conjugated to sepharose beads.⁵⁴ In the case of passive immunization, the administration of affinity purified anti-PcrV IgG against either $PcrV_{1-294}$ or $PcrV_{139-258}$ showed a significantly higher efficacy against lethal *P. aeruginosa* infection than that of unpurified anti-PcrV IgG. The increased efficacy of affinity purified anti-PcrV IgG implies that more potent anti-PcrV strategies are possible. The results of this study are crucial for the development of both an effective PcrV vaccine for active immunization and an appropriate-blocking anti-PcrV antibody against *P. aeruginosa* infection in humans.

Monoclonal Anti-PcrV Antibody

Development of an anti-PcrV monoclonal blocking antibody

Because polyclonal anti-PcrV antibodies inhibit the delivery of TTS toxins and enhance the clearance of bacteria during acute lung infections, the next issue was whether a specific monoclonal antibody against PcrV could demonstrate the same blocking capacity as that of rabbit anti-PcrV IgG. To generate blocking monoclonal clones against PcrV, approximately 80 clones from a hybridoma library derived from the spleen cells of a PcrV-vaccinated mouse were tested in a protection assay against TTSSmediated intoxication.¹¹ Of the initial 80 cell lines, only 2 hybridomas, mAb166 and mAb179, demonstrated protective activity. mAb166 (IgG2bK) and mAb179 (IgG1K) were purified and tested for protection in an acute lung infection model. Groups of mice that received mAb166 had smaller decrements in their body temperatures, and all infected mice survived.

Anti-PcrV monoclonal antibody mAb166 and the blocking epitope

Monoclonal antibodies were tested for their ability to affect the survival of mice infected with cytotoxic PA103.¹¹ The survival of each group of animals was monitored for 1 week. Only mAb166 demonstrated significant protection in this assay. The effects of mAb166 on survival were further characterized by testing different doses in co-instillation experiments. The addition of 0.5 µg or 1 µg of mAb166 IgG to the bacterial inoculum (at a 4-fold lethal dose) appeared to have a slight effect on survival within the first 48 h of infection. Higher doses of antibody, 10 and 50 µg, resulted in 80% and 90% survival, respectively. In another series of experiments, the ability of mAb166 to passively protect against PA103 infection was compared with that of polyclonal rabbit IgG.¹¹ When the antibodies were administered by an intraperitoneal injection 1 h before bacterial lung instillation (at a 4-fold lethal dose), mAb166 and rabbit polyclonal anti-PcrV were protective at a dose of 100 µg. Different doses of PA103 were prepared and mixed with a fixed amount of mAb166 IgG for co-instillation. Fifty micrograms of mAb166 protected 100% of the animals against infection with 4-, 8-, and 10-fold lethal doses of strain PA103. Seventy percent of the animals were protected at a 20-fold lethal dose of PA103 with 50 µg of mAb166. Survival was diminished to nominal levels as the bacterial instillation was increased to 30-fold the lethal dose. Although only 20% of the animals survived this severe infection, co-instillation with mAb166 seemed to delay mortality by 1-2 days.

Next, to determine whether the protective properties of mAb166 were dependent on its functional fragment crystallizable (Fc) region, Fab fragments from mAb166 were purified. Purified fragments (10 μ g) were premixed and co-instilled with a 4-fold lethal dose of strain PA103, and survival was assessed.¹¹ The administration of mAb166 Fab fragments resulted in a high level of protection (80%), comparable to the effect of whole IgG. These data indicate that the Fc portion of both the polyclonal and monoclonal antibodies is unnecessary to mediate protection *in vivo*.

Full-length PcrV comprises 294 amino acids. To define the amino acid residues bound by mAb166, truncated rPcrV tagged with glutathione S-transferase was constructed, and mAb166 binding with truncated rPcrV was tested by immunoblotting or the dot blot technique.¹¹ mAb166 failed to bind to expression clones encoding the N terminus of PcrV or to one internal deletion of PcrV, but it did bind to a construct encoding the C terminus of PcrV. The smallest epitope recognized by mAb166 is located between amino acids 144–257 of PcrV.

The therapeutic effects of the monoclonal anti-PcrV antibody mAb166

The therapeutic effects of mAb166 on acute lung injury induced by *P. aeruginosa* were analyzed in a rat model.⁵⁵ In this model, lung injury was induced by instilling the *P. aeruginosa* strain PA103 directly into the left lungs of anesthetized rats. One hour after bacterial instillation, either rabbit polyclonal anti-PcrV IgG, murine monoclonal anti-PcrV IgG mAb166, or mAb166 Fab fragments (4 mg/kg, respectively) were administered intratracheally directly into the lungs. The degree of alveolar epithelial injury, amount of lung edema, decrease in oxygenation, and extent of lung inflammation were histologically evaluated as independent parameters of acute lung injury. Rats that received phosphate-buffered saline alone 1 h after bacterial instillation showed a significant increase in lung epithelial injury and lung edema after 4 h. Arterial blood pressure gradually decreased to 80 mmHg over the experimental period. Arterial blood oxygenation remained significantly decreased. Severe metabolic acidosis developed over 4 h. Rats that received either rabbit polyclonal anti-PcrV IgG, murine monoclonal anti-PcrV IgG mAb166, or mAb166 Fab fragments intratracheally showed significant improvement of alveolar epithelial injury and lung edema 4 h after bacterial instillation. The protective effect of mAb166 Fab fragments on lung epithelial injury was the most significant among the 3 antibodies, whereas rabbit polyclonal and murine monoclonal anti-PcrV IgGs were better at improving lung edema than mAb166 Fab fragments. Hypotension did not develop in any group. Arterial oxygenation in the 3 treated groups was significantly improved compared with that in the untreated rats. Although mild metabolic acidosis did develop in rats that received rabbit polyclonal anti-PcrV IgG and mAb166 Fab fragments, those that received mAb166 did not become acidotic. In this series of experiments, lung histology was compared between rats treated with mAb166 and those treated with control IgG. Whereas rats that received control IgG 1 h after bacterial instillation exhibited severe neutrophil recruitment and destruction of alveolar structures; those that received mAb166 had almost no neutrophils in their airspaces and normal alveolar structure was preserved. Therefore, the therapeutic administration of mAb166 showed comparable effects to that of rabbit polyclonal anti-PcrV IgG in preventing acute lung injury and subsequent systemic distress.

Recently, Song *et al.*⁵⁶ examined whether the combined administration of mAb166 and an antibiotic could further improve the survival of *P. aeruginosa*-infected mice. Three clinically relevant antibiotics (ciprofloxacin, tobramycin, and ceftazidime) were used in this study. The administration of mAb166 with an antibiotic significantly improved the survival of mice infected with 3 times the lethal dose (LD₉₀) of PA103 compared with that of either antibiotic or antibody alone. This synergistic effect was because of enhanced bactericidal effects and protection against lung injury, which prevented bacterial dissemination to other organs. Therefore, combined mAb166 and antibiotic administration provides a new, more effective strategy against *P. aeruginosa* airway infection, particularly when large numbers of highly virulent strains are present.

Wang *et al.*⁵⁷ recently investigated the level of TTSS expression in clinically isolated multidrug resistant *P. aeruginosa* (MDRP) strains and the effects of anti-PcrV antibody on an MDR isolate-induced acute lung injury. The level of TTSS expression was quantified in 53 isolates, comprising 25 MDR strains and 28 susceptible strains. They also investigated the effect of anti-PcrV antibody using a murine model created by instilling an MDRP strain intratracheally into the left lung through and showed that the level of TTSS expression in MDRP strains is comparable to that of susceptible strains. Anti-PcrV ensured the survival of challenged mice, reduced bacterial numbers and attenuated lung inflammation and injury. Therefore, these results suggest that anti-PcrV may be a potentially effective strategy against an MDRP-induced acute lung injury.

A Humanized Anti-PcrV Monoclonal Antibody

Cloning anti-PcrV antibody genes

We cloned the blocking murine monoclonal anti-PcrV antibody mAb166 and generated a murine-human chimeric monoclonal antibody to PcrV, chimera166, which is protective against P. aeruginosa when co-instilled with the bacterial inoculum or intraperitoneally transferred into mice. As a first step in characterizing the structural features of mAb166 underlying its interactions with PcrV, we sequenced both the heavy- and light-chain variable regions of mAb166 IgG mRNA. The predicted amino acid sequence of the κ light-chain is shown in Table 2. This κ variable region is a Class II mouse κ variable region. Although its sequence is not identical to any germline-variable region present in the data bank, the DNA sequence of the 5'-untranslated region and V-region in the k light-chain showed highest homology to the germline IGKV12-41 (IGKV Subgroup 12; Accession Number: AJ235953). The DNA sequence in the J-region was identical to the germline IGKJ2 (Accession Number: V00777). The predicted amino acid sequence of the mAb166 heavy-chain region is shown in Table 2. The DNA sequence of the 5'-untranslated region and V-region segments in the heavy-chain containing complementarity determining regions (CDRs) 1 and 2 is identical, with the exception of 2 amino acids in the Frame 3 region, to the germline IGHV2S2 (IGHV Subgroup 2, VH101; Accession Number: J00502). The V-region sequence also showed the same level of homology with the reported pseudogene IGHV2S5 (Accession Number: M21165). The DNA sequence in the Jregion was identical to the germline IGHJ4 (Accession Number: V00770). The unique CDR3 sequence includes an Arg-Gly-Asp sequence known to function as a recognition sequence for adhesion receptors in many adhesive proteins, including fibrinogen, fibronectin, von Willebrand factor, and vitronectin.

An engineered human anti-PcrV antibody Fab fragment

An engineered human antibody Fab fragment that binds to the *P. aeruginosa* PcrV protein with a high affinity has been identified, and it has potent *in-vitro* neutralization activity against TTSS.¹² Instillation of a single dose of Fab into the lungs of mice provided protection against a lethal pulmonary challenge with *P. aeruginosa*, leading to a substantial reduction in viable bacterial cell counts in the lungs. These results demonstrate that blocking TTSS with a Fab fragment that lacks antibody Fc-mediated effector functions is sufficient for the effective clearance of a pulmonary *P. aeruginosa* infection.

Clinical trial for ventilator-associated pneumonia (VAP)

In France, a multicenter, Phase-IIa clinical trial of the recombinant, PEGylated, engineered, human Fab' anti-PcrV fragment KB001 involving 10 intensive care units was undertaken.¹³ The purpose of the study was to determine the safety and pharmacokinetics of KB001, and its ability to prevent *P. aeruginosa* VAP. In this trial, 39 mechanically ventilated patients, who were colonized by *P. aeruginosa* but not infected, were randomized at a 1:1:1 ratio to receive a single intravenous 3 mg/kg or 10 mg/kg

Table 2. Primary sec	quences of the Anti-PcrV	monoclonal antibody, n	1Ab166
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Chain	Туре	Region	Amino Acid Sequence	KB001 modification
heavy chain	lgG2b	Sig-pep	MAVLGLLFCLVTFPSCVLS	
		FR1	QVQLKQSGPGLVQPSQSLSITCTVSGFSLT	
		CDR1	SYGVH	
		FR2	WVRQSPGKGLEWLG	
		CDR2	VIWSGGDTDYNAAFIS	
		FR3	RLSISKDNSKSQLFFKMNSLRATDTAIYYCAR	
		CDR3	NRGDIYYDFTYAMDY	NRGDIYYDFTYAXDZ
(X = M/F, Z = I/S/Q)				
		FR4	WGQGTSVTVSS	
		CH1	AKTTPPSVYPLAPGCGDTTGSSVTLGCLVKGYFPESVTVT	
			WNSGSLSSSVHTFPALLQSGLYTMSSSVTVPSST	
			WPSQTVTCSVAHPASSTTVDKKLEPSGPISTINPCPP	
			CKECHKCPAPNLEGGPSVFIFPPNIKDVLMISLTPKVT	
			CVVVDVSEDDPDVQISWFVNNVEVHTAQTQTHREDYN	
			STIRVVSTLPIQHQDWMSGKEFKCKVNNKDLPSPIERTIS	
			KIKGLVRAPQVYILPPPAEQLSRKDVSLTCLVVGFNPGDISV	
			EWTSNGHTEENYKDTAPVLDSDGSYFIYSKLNMKTSKWE	
			KTDSFSCNVRHEGLKNYYLKKTISRSPGK	
light chain	к	Sig-pep	MSVLTQVLALLLLWLTGARC	
		FR1	DIQMTQSPASLSASVGETVTITC	
		CDR1	RASGNIQNYLA	
		FR2	WYQQTQGKSPQLLVY	
		CDR2	SAKTLAD	
		FR3	GVPSRFSGSGSGTQYSLKINSLQPEDFGSYYC	
		CDR3	QHFWSTPYT	QQFWXTPYT ($X = S/G$)
		FR4	FGGGTKLEIKR	FGGGTKLTVLR
		CL	ADAAPTVSIFPPSSEQLTSGGASVVCFLNNFYPKDINVKW	
			KIDGSERQNGVLNSWTDQDSKDSTYSMSSTLTLTKDEYE	
			RHNSYTCEATHKTSTSPIVKSFNRNEC	

infusion of KB001 or a placebo. KB001 was well tolerated and not immunogenic. The respective maximum serum concentrations of the 3- and 10-mg/kg groups were $5-8 \times 10^4$ and $12-28 \times 10^4$ ng/ml, with mean elimination half-lives of 8.1 and 9.3 days, respectively. KB001 was detected in the endotracheal aspirates of all patients receiving it as early as day 1 and up to 28 days. Patients who received either 3 or 10 mg/kg of KB001 developed P. aeruginosa pneumonia less frequently (33% and 31%, respectively) than placebo recipients (60%). In this study, KB001 demonstrated its safety, a favorable pharmacokinetic profile, and a promising potential for reducing the incidence of P. aeruginosa pneumonia in mechanically ventilated patients colonized with the bacterium in intensive care. The administration of 3 mg/kg of KB001 appears to be promising to keep the effective serum titer which is high enough to neutralize PcrV of P. aeruginosa in the bronchial secretes at least for a month and prevent the ventilated patients from development of pneumonia.

Clinical trial for chronic pneumonia in cystic fibrosis

In USA, the safety and pharmacokinetic and pharmacodynamic properties of KB001 were evaluated in CF patients with a chronic *P. aeruginosa* infection.¹⁴ In this study, 27 CF patients with a sputum *P. aeruginosa* density $>10^5$ CFU/g received a single intravenous dose of KB001 (3 mg/kg or 10 mg/kg) or a placebo. After administration, KB001 demonstrated a mean serum half-life of 11.9 days. At day 28, there was a trend toward a dose-dependent reduction in sputum myeloperoxidase, IL-1 β , and IL-8, and there were significant overall differences in the change in sputum neutrophil elastase and neutrophil counts favoring the KB001 10 mg/kg group versus the control group, suggesting that targeting the *P. aeruginosa* TTSS with KB001 as a nonantibiotic strategy reduces airway inflammation and damage in CF patients with a chronic *P. aeruginosa* infection.

Summary

The observations described in this review support the hypothesis that anti-PcrV strategies have potential as nonantibiotic immunostrategies for cytotoxic *P. aeruginosa* infections. Clinical trials of the humanized anti-PcrV IgG Fab' fragment, KB001, which was molecularly engineered from the monoclonal anti-PcrV IgG mAb166 are currently ongoing for patients with VAP and CF. In preliminary reports from these clinical trials, KB001 has demonstrated its safety, a favorable pharmacokinetic profile, and a promising potential as a nonantibiotic strategy to reduce airway inflammation and damage in *P. aeruginosa* pneumonia.¹⁴ Further analyses of the role of PcrV in TTSS and the mechanism of its antibody mediated disruption are critical for developing improved anti-V-antigen strategies. Up to date, there is no report about the effects of anti-PcrV on *Staphylococcus aureus* and other potentially drug resistant pathogens in polymicrobial environment. Therefore, in further clinical trials, an evaluation of the efficacy of antibody mediated therapies including the analysis of anti-PcrV in the polymicrobial environment which is frequently observed in the cystic fibrosis airways, will also strengthen these strategies.

Antibodies to other bacterial V-antigen homologs may protect hosts from other infections such as diarrheal enterocolitis and sepsis caused by *Vibrio parahaemolyticus* and *Aeromonas* sp.⁹ This strategy could also be used to protect fish and shrimp from massive mortality caused by *A. salmonicida* or *V. harveyi* infections at fish and shrimp farms. V-antigenassociated virulence is linked to lethal infections caused by pathogenic gram-negative bacteria at fish and shrimp farms, in potential plagues caused by bioterrorism, and in opportunistic infections in humans.

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Disclosure of Potential Conflicts of Interest

With regard to the content of the manuscript, Teiji Sawa has a patent fee from the Regent of the University of California, USA.

Acknowledgments

The authors would like to thank Dr. Jeanine P. Wiener-Kronish, Anesthetist-in-Chief; Henry Isaiah Dorr, Professor of Anesthesia; and Harvard Medical School for critical support of the authors' research.

Funding

This work was supported by grants from a Grant-in-Aid for Scientific Research (KAKENHI No. 24390403) and The Ministry of Education, Culture, Sports, Science and Technology, Japan to Teiji Sawa.

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