Preparation of a Nanobody Specific to Dectin 1 and Its Anti-inflammatory Effects on Fungal Keratitis

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Objective: To prepare a nanobody specific to dectin 1 and verify its specificity and anti-inflammatory effects on Aspergillus fumigatus keratitis.

Methods: The nanobody was selected from a high-quality shark-antibody library constructed with phage-display technology. The nanobody was developed in the expression systems of Escherichia coli. Indirect ELISA was used to determine the specificity of the nanobody to recombinant dectin 1 protein. The potential of the nanobody to be recognized and expressed on the surfaces of cells and corneas was detected by immunofluorescence, and its anti-inflammatory effect on A. fumigatus keratitis was further verified. After infection with A. fumigatus, eyes of C57BL/6 mice were treated with nanobodies. Human corneal epithelial cells (HCECs) were pretreated with nanobodies and then incubated with A. fumigatus. Clinical scores and slit-lamp photography were used to assess disease response in mouse corneas. RT-PCR and ELISA were used to evaluate mRNA and protein expression of IL1β and IL6 in both mouse corneas and HCECs.

Results: The nanobody was successfully expressed through microbial system and showed specific high-affinity binding to recombinant dectin 1. Furthermore, it exhibited specific binding to dectin 1 expressed on the surfaces of cells and recognized dectin 1 in mouse corneas. Importantly, it reduced clinical scores of A. fumigatus keratitis in mice compared with a PBS-treatment group. In addition, it decreased mRNA and protein expression of IL1β and IL6 in infected corneas and HCECs stimulated with A. fumigatus.

Conclusion: These results suggest that this nanobody can bring about anti-inflammatory effects. This highlights the potential of these nanobodies as innovative therapeutic agents in A. fumigatus.

Keywords: nanobody, Aspergillus fumigatus, dectin 1, inflammation, cornea

Introduction

Fungal keratitis is a severe blinding eye infection caused by pathogenic fungi and has a high incidence in developing countries.1-4 Corneal trauma due to vegetative material provides an opportunity for fungal infection of the cornea.5 With the excessive use of broad-spectrum antibiotics, irregular use of glucocorticoids, and wearing of contact lenses, the incidence of fungal keratitis has increased in recent years.6 Aspergillus fumigatus is one of the most common pathogenic fungi in fungal keratitis, resulting in inflammatory responses triggered by pattern-recognition receptors (PRRs) of innate immunity.7 A significant PRR, dectin 1 participates in inflammatory responses against fungal infection in fungal keratitis.8 Although innate immunity plays a significant role in the early stage of fungal infection, it can cause excessive inflammatory reactions.9 Excessive inflammatory reactions are detrimental to tissue repair and fungi clearance in fungal keratitis.10,11 A recent study has shown that suppression of excessive inflammatory responses may improve prognosis.12 Therefore, it is necessary to find a new drug that can act against dectin 1 to relieve the severe inflammatory response in fungal keratitis.
Nanobodies originate naturally from the sera of camelids and shark species, and are part of the variable region of the heavy chain. Small proteins with molecular weight of only 15 kDa, nanobodies display considerable ability to bind antigens not accessible to conventional antibodies with the long third complementarity determining region (CDR3). They are the smallest antigen-binding fragment that retains affinity and specificity in recognizing antigens. Nanobodies can be expressed in microorganism systems through genetic engineering technology, which can reduce production costs. Nanobodies have the advantages of low toxicity, favorable permeability, and high specificity. Previous studies have shown nanobodies to have promising potential as diagnostic and therapeutic tools in many diseases related to infection and inflammation. However, no study has investigated the application of nanobodies in the treatment of fungal keratitis. In this study, a nanobody specific to dectin 1 was prepared and its recognition by dectin 1 in different models verified. The inhibitory effect of the nanobody on the inflammatory response during *A. fumigatus* infection was investigated. All the results implied that this nanobody could be used as a new candidate for the treatment of fungal keratitis.

**Methods**

**Affinity Selection of Phages Specific to Dectin 1 by Phage Display**

The nanobody (IgNAR V) phage-display library was generously provided by the Ocean University of China (Qingdao, China). Phages were generated by infection of the stock library with the M13K07 helper phage, and three rounds of biopanning were carried out as per standard protocols. In the first round, an immunotube was coated with 4 mL recombinant dectin 1 (100 µg/mL dissolved in PBS) overnight at 4°C and then blocked with 4 mL of MPBS (5% skimmed-milk powder in PBS) for 2 hours at room temperature. Generated phage particles were added and incubated at room temperature for 1 hour in a vertical rotator and 1 hour without rotation. After being washed ten times with PBS, the bound phages were eluted with 1 mL 100 mM TEA for 10 minutes at room temperature and neutralized with 1 M Tris–HCl (pH 7.4). The phage particles were used to infect *Escherichia coli* TG1 for the next round of panning. The second and third rounds of biopanning followed the same procedure, except the coating concentration was changed to 10 µg/mL and 1 µg/mL, and phages were washed ten times with PBS and ten times with PBST (PBS containing 0.05% Tween 20).

**Isolation of Specific Clone Gene**

Polyclonal phage ELISA was used to monitor the selection of positive phages throughout the biopanning. A 96-well plate was coated with 10 µg/mL recombinant dectin 1 (100 µL/well) overnight at 4°C and blocked with MPBS for 2 hours at room temperature. After being washed three times with PBST, 100 µL phages from each round of selection containing approximately 10^9 CFU/mL of phage particles was added to the wells and incubated for 1 hour at room temperature. Then, the plate was washed three times with PBST and further incubated with 100 µL 1:2,000 dilution of HRP-anti-M13 mAb (GE Healthcare) at room temperature for another hour. After another three washes with PBST, 100 µL tetramethylbenzidine substrate solution was added and incubated in the dark for 15 minutes. The reaction was finally stopped by the addition of 50 µL 1M H_2SO_4 and absorbance at 450 nm measured with a microplate reader.

Monoclonal phage ELISA was carried out to determine individual colonies’ ability to recognize dectin 1 specifically. In sum, 95 colonies were randomly picked from the third round of selection and inoculated in 200 µL 2xTY medium (supplemented with 100 µg/mL ampicillin and 20 g/L glucose) into separate wells of the cell-culture microplate. The plate was incubated overnight at 37°C with shaking at 200 rpm. A fresh 2xTY microplate containing 200 µL 2xTY per well was inoculated with 2 µL monolayer plate cultures and incubated for 2 hour at 37°C. After coinfection with helper phage (10^9 per well), the latter plate was centrifuged at 1,000 rpm for 10 minutes at room temperature and bacterial pellets resuspended in 200 µL 2xTY (supplemented with 100 µg/mL ampicillin, 50 µg/mL kanamycin, and 1 g/L glucose), and incubated overnight at 30°C. The supernatant was detected by ELISA after 1,000 rpm centrifugation for 10 minutes on the next day.

Positive colonies were checked again with monoclonal phage ELISA. The positive phage with the highest absorbance was used to performed a colony PCR (forward primer 5\’--ATA ATA AGG AAT TCC ATG GCT CGA GTG GAC CAA ACA CCG, reverse primer 5\’--ATA ATC AAG CTT GCG GCC GCA TTC ACA GTC ACG ACA GTG CCA CCT C)
using PCR Master Mix DNA polymerase (Tsingke Biotech). Finally, sequencing of the phage clone was determined by Tsingke Biotech.

**Expression, Purification, and Identification of Nanobodies Specific to Dectin 1**

The gene of the nanobody was digested with *BamHI* (NEB) and *NotI* (NEB) and ligated with a similarly digested pET-28a vector by T4 DNA ligase (NEB). After verification by colony PCR (forward primer 5’–CGG GAT CCATGG CTC GAG TGG ACC AAA CA, reverse primer 5’–ATA GTT TAG CCG CCG CAT TCA CAG TCA CAG TGC) using the Master Mix DNA polymerase, the recombinant plasmid was transformed into competent *E. coli* Transetta-DE3 (Tsingke Biotech) cells by heat shock. Expression of the nanobody specific to dectin 1 was induced by 0.5 mmol/L isopropyl-β-D-thiogalactopyranoside (IPTG) in 16°C for 20 hours. Induction cells were collected and broken up by an ultrasonic cell crusher (JY88-II ultrasonic cell disruptor, Xinzhi Biological Technology), and the protein in the inclusion body was renatured by lysis buffer (8 M urea, 50 mM Tris-HCl, 300 mM NaCl, pH 8). Then, the recombinant nanobody was purified under gradient-washing using a nickel-ion column (GE Healthcare) and the elution product dialyzed in the solutions listed in Table 1. SDS-PAGE 12% was used to analyze the expression and purification of the nanobody. The concentration of purified nanobodies was determined by the Bradford method. A typical indirect ELISA was carried out to determine the specificity of the nanobody to recombinant dectin 1 protein.

**Cell Culture and *A. fumigatus* Stimulation**

Human corneal epithelium cells (HCECs) were generously provided by the Ocular Surface Laboratory of Zhongshan Ophthalmic Center, Guangzhou, Guangdong, China. RAW 264.7 cells were purchased from the Chinese Academy of Sciences in Shanghai, China. The study on HCECs was approved by the Ethics Review Committee of the Affiliated Hospital of Qingdao University, Qingdao, Shandong, China (QYFYWZLL-26567). The standard *A. fumigatus* strain 3.0772 was purchased from the China General Microbiological Culture Collection Center in Beijing. HCECs were cultured in DMEM (Gibco) containing 10% FBS (Gibco, San Diego, CA, USA) at 37°C in 5% CO₂. RAW 264.7 cells were cultured in high-glucose DMEM supplemented with 12% FBS and incubated at 37°C in 5% CO₂. The inactive *A. fumigatus* hyphae (to a final concentration of 5×10⁶ CFU/mL) were prepared as per Xia et al and added to the cell culture for stimulation of dectin 1.

**Immunofluorescence Staining of Cells**

HCECs or RAW 264.7 (500 μL) cells at a concentration of 2×10⁵/mL were seeded onto 24-well plates with poly-L-lysine–coated slips. When cultured to nearly 80% confluence, the cells were stimulated with 30 μL *A. fumigatus* hyphae (5×10⁶ CFU/mL) for 16 hours. Then, cells were washed three times with PBST, coated with 4% paraformaldehyde (Solarbio) for 10 minutes, and blocked with 1:10 diluted goat serum (Solarbio) for 30 minutes at room temperature after another three washes with PBST. Purified recombinant nanobodies (100 μL, 1 mg/mL) were added and incubated with cells at 4°C overnight. PBS or aspecific nanobodies of equal volume were used as blank and negative controls, respectively. Subsequently, the HCECs were incubated with 100 μL 1:1,000 mouse antishark VNAR antibodies and 100 μL 1:1,000 FITC-conjugated goat antihuman IgG antibodies (Abcam) at 37°C for 1 hour successively. The RAW 264.7 cells were incubated with 100 μL 1:1,000 rabbit anti–His tag antibodies and 100 μL 1:1,000 FITC-conjugated goat antirabbit IgG antibodies (Abcam) at 37°C for 1 hour successively. Then, PBST was used to wash the cells three times. Cell nuclei were stained with 100 μL 1:100 DAPI for 10 minutes and washed three times with PBST. Fluorescent signals were finally detected with a Zeiss Axiovert microscope at 200× magnification.

**Table 1 Components of dialysate solution**

<table>
<thead>
<tr>
<th>Ingredients</th>
<th>Buffer 1</th>
<th>Buffer 2</th>
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<tbody>
<tr>
<td>Tris 50 mM</td>
<td>50 mM Tris, 150 mM NaCl, 2 mM EDTA, 4 mM GSH, 0.4 mM GSSG, 0.4 M L-arginine, 10% glycerin (pH 8)</td>
<td>50 mM Tris, 150 mM NaCl, 10% glycerin (pH 8)</td>
</tr>
<tr>
<td>NACl 150 mM</td>
<td>50 mM Tris, 150 mM NaCl, 2 mM EDTA, 4 mM GSH, 0.4 mM GSSG, 0.4 M L-arginine, 10% glycerin (pH 8)</td>
<td>50 mM Tris, 150 mM NaCl, 10% glycerin (pH 8)</td>
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Anti-inflammatory Effect of Nanobodies in HCEC Model

HCECs (1.5 mL) at a concentration of $3 \times 10^5$/mL were seeded into 12-well plates and cultured until 70%–80% confluence, then pretreated with 100 μL (1 mg/mL) of nanobodies specific to dectin 1 dissolved in PBS (Solarbio) for 2 hours. After stimulation with inactive *A. fumigatus* hyphae ($5 \times 10^6$ CFU/mL) for 8 or 16 hours, the cells and supernatant were collected for RT-PCR and ELISA tests, respectively. Protein levels of IL1β and IL6 (n=6/group/time) were assessed using human ELISA kits (ELISA Max Deluxe, BioLegend) according to the protocols. Absorbance at 450 nm was detected with a reference wavelength of 570 nm (VersaMax plate reader, Molecular Devices). mRNA levels of IL1β and IL6 were assessed by RT-PCR (n=6/group/time). Total RNA of the cells was extracted using a RNA-extraction reagent (RNAiso Plus, Solarbio) and quantified by NanoDrop (Thermo Fisher Scientific). mRNA expression of β-actin, one of the housekeeping genes, was used as a reference. HiScript III RT SuperMix for qPCR (Vazyme) was used to reverse RNA into complementary cDNA. Quantitative RT-PCR (primers shown in Table 2) was performed using ChamQ SYBR Color qPCR Master Mix (Vazyme). Values were recorded to verify the fold-change ratios of proinflammatory cytokines. Cells pretreated with PBS and nonspecific nanobodies were used as blank and negative controls, respectively.

Specificity and Anti-inflammatory Effect of Nanobodies in Mouse Model

C57BL/6 mice (female, 8 weeks old) were purchased from Pengyue Laboratory Animal (Jinan, China) and treated in accordance with the ARVO statement for the Use of Animals in Ophthalmic and Visual Research. All experimental protocols were performed in accordance with the National Institution of Health’s *Guidelines for the Care and Use of Laboratory Animals* and every effort was made to minimize both the number of animals and their suffering. The study was approved by the Ethics Review Committee of the Affiliated Hospital of Qingdao University, Qingdao, Shandong, China (QFYFWZLL-26567). All corneas were separately inspected under slit-lamp microscopy before experimental use. After anesthetisation by intraperitoneal injection with 8% choral hydrate, central corneal epithelia of mouse eyes were removed within 2 mm diameter. A 10 μL aliquot containing $10^8$ CFU/mL of *A. fumigatus* was coated on the ocular surface and a soft contact lens placed. Subsequently, the eyelids of the mice were sutured. Then, the eye of each mouse was injected with 5 μL specific nanobodies (to a final concentration of 5 mg/mL) or sterile PBS subconjunctivally at 1 day postinfection and with 5 μL specific nanobodies (to a final concentration of 5 mg/mL) or PBS three times daily from day 2 postinfection. Epithelia of the control corneas were removed without *A. fumigatus* stimulation. No treatment was administered to normal corneas.

Mouse eyeballs (n=3/group/time) were embedded and frozen. The corneas of the normal group and *A. fumigatus*-infected mice were sectioned to 10 μm. Corneal lesions (10 μm) were acquired and then fixed in acetone. After being blocked with goat serum (1:100), sections were incubated with 30 μL nanobodies, followed by 30 μL monoclonal rabbit anti–His-tag antibodies and FITC-conjugated goat antirabbit IgG antibodies (1:1,000, Abcam) at 37°C for 1 hour.

### Table 2 Nucleotide sequences of human primers for rT-PCR

<table>
<thead>
<tr>
<th>Gene</th>
<th>GenBank identifier</th>
<th>Primer sequence (5’–3’)</th>
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<tr>
<td>hβ-GAPDH</td>
<td>NM_001101.5</td>
<td>F: TGG CAC CCA GCA CAA TGA A</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: CTA AGT CAT AGT CCG CCT AGA AGC A</td>
</tr>
<tr>
<td>hIL1β</td>
<td>NM_000576.2</td>
<td>F: GCT GAT GGC CCT AAA CAG ATG AA</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: TCC ATG GCC ACA ACA ACT GAC</td>
</tr>
<tr>
<td>hIL6</td>
<td>NM_000600.4</td>
<td>F: AAG CCA GAG CTG TGC AGA TGA GTA</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: TGT CCT GCA GCC ACT GTG TC</td>
</tr>
<tr>
<td>Mβ-actin</td>
<td>NM_007393.5</td>
<td>F: GAT TAC TGC TCT GGC TCC TAG C</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: GAC TCA TCG TAC TGC TGC TTG C</td>
</tr>
<tr>
<td>MIL1β</td>
<td>NM_008361.4</td>
<td>F: CGC AGC AGC ACA TCA ACA AGA GC</td>
</tr>
<tr>
<td></td>
<td></td>
<td>R: TGT CCT CAT CCT GGA AGG TCC ACG</td>
</tr>
<tr>
<td>MIL6</td>
<td>NM_001314054.1</td>
<td>F: TGA TGG ATG CTA CCA AAC TGG A</td>
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<tr>
<td></td>
<td></td>
<td>R: TGT GAC TCC AGC TTA TCT CTG GG</td>
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successively. Cell nuclei were stained with DAPI. Images were photographed with a fluorescence microscope (Zeiss Axio Vert, 200×).

There were six mice in each group (n=6/time/group). Based on the observation under a slit lamp at 1, 3, and 5 days postinfection, the severity of keratitis was evaluated by a clinical score that was the sum of the three aspects of cornea — opacity density, opacity area, and surface regularity — each of which had a grade of 0 to 4. Severity was classed as normal (0), mild (1–5), moderate (6–9), and severe (10–12). Taking a normal cornea as an example, a live cornea would be given a score of 0 for each class, thus yielding a total score of 0. Mouse corneas removed with a scalpel and microscissor at the indicated times after treatment were prepared for RT-PCR and ELISA, respectively.

Cytotoxicity Assessment of Nanobodies
HCECs (100 µL, 3×10^4/mL) were suspended and seeded into a 96-well plate and cultured at 37°C for 24 hours. Nanobodies specific to dectin 1 diluted in PBS for various concentrations were added to each well. After incubation for another 24 hours, 10 µL CCK8 solution (MCE) was added to each well. The plates were incubated for 2 hours and absorbance was measured by a microplate reader (Solarbio) at 450 nm. Each sample had six replicates.

Statistical Analysis
Differences in clinical scores between two groups at each time point were analyzed by Mann–Whitney U test. Statistical significance between two groups was evaluated by two-tailed Student’s t-test, and one-way ANOVA was used to make comparisons among three or more groups. Multiple comparisons among groups were performed using Bonferroni’s test. GraphPad Prism 7.0 and SPSS 23.0 were used for statistical analyses, with values presented as means ± SD. P<0.05 was considered statistically significant. All experiments were repeated at least three times.

Results
Screening of Nanobody Gene Specific to Dectin 1
Phages displaying nanobodies specific to dectin 1 were panned for three rounds using immunotubes coated with different concentrations of recombinant dectin 1 protein. Polyclonal phage ELISA results showed that the phages specific to dectin 1 were enriched as panning progressed. After the first round of panning, the value of OD_{450} values showed no obvious difference from the negative-control group, but a significant increase in binding ability was observed from the first to the third rounds (Figure 1A). A total of 95 phages were randomly picked from the third round panning and then analyzed by monoclonal phage ELISA, and 26 showed high absorbance (at least three times that of control wells, Figure 1C). Four phages with the highest absorbance were selected to repeat the phage ELISA in order to avoid false-positive reactivity. The absorbance of these was at least three times that of the negative control (Figure 1B). The phage with the strongest antigen-specific binding capacity was selected for colony PCR to amplify the nanobody gene. A gene band about 500 bp appeared in the agarose gel (Figure 2).

Expression and Purification of Nanobodies
Sequencing results showed that nanobodies were shark-derived with a complete CDR3 (Figure 3A). Therefore, this gene was used as the gene sequence for specific nanobody expression. The gene of nanobody specific to dectin 1 was recombined with the plasmid after restriction by T4 DNA ligase to construct a new protein-expression vector. A solid plate with kanamycin was used to select a positive expression vector (Figure 3B). PCR and nucleic acid electrophoresis were used to verify the construction of plasmid-expression vectors. Nucleic acid electrophoresis in a 1% agarose gel showed a bright band around 350 bp (Figure 3C).

The expression vector was transformed into competent E. coli Transetta-DE3 by heat shock. Expression of the recombinant nanobodies was induced by 0.5 mmol/L IPTG at 16°C overnight. After induction with 0.5 mmol/L IPTG, the 17 kDa protein band was present in the precipitation of E. coli and broken up by the ultrasonic cell crusher, as shown on 12% SDS-PAGE stained with Coomassie (Figure 4A). Nanobodies were purified by the nickel-ion column under different concentrations of imidazole. When passing through the column, the His-tag carried by the nanobodies
Figure 1 Screening of special nanobody gene against dectin 1. (A) Polyclonal phage ELISA results. The phages generated from the original library were used as negative controls, and affinity of the panned phages showed a significant increase from the first round to the third round of panning. (B) Monoclonal phage ELISA results of four phages. NC, negative control. (C) Monoclonal phage ELISA results. 1–95, clones selected from the third round of panning; 96, negative control.

Figure 2 Agarose-gel electrophoresis of the positive clones. M, DNA marker DL2000; lanes 1–4, positive clones; lane 5, negative control.
Figure 3 Construction of expression vector of dectin 1–specific nanobodies. (A) The 119 amino–acid sequence of the nanobody. The area marked in red is the CDR3. (B) Nucleic acid electrophoresis of the target gene (M, DNA marker DL2000). Lanes 1–3, target gene. (C) Screening of positive clones on solid plates with kanamycin.

Figure 4 Expression and purification of soluble nanobodies. (A) M, protein marker; 1, bacterium protein before induction; 2, 16°C supernatant; 3, 16°C precipitation. (B) M, Protein marker; 1, upper sample; 2, effluent; 3–7, 20, 50, 100 mM, 200 mM, 500 mM imidazole-elution component; 8: 16°C precipitation. (C) M, protein marker; 1–5, protein after dialysis.
interacted with nickel ions. Imidazole of low concentration was used to wash away impurities, and imidazole of high concentration eluted the target protein (Figure 4B). In order to remove imidazole and other impurities components left, the eluted outflow solution was dialyzed to obtain the final purified nanobodies. SDS-PAGE 12% was used to analyze the purification of the nanobodies (Figure 4C). Bright protein bands appeared around 17 kDa. The total protein content of nanobody was determined by the Bradford method. After measurement, the yield of nanobody specific to dectin 1 was 5 mg/L.

Affinity Determination of Nanobodies Specific to Dectin 1
The specificity of nanobodies for binding to dectin 1 were determined by ELISA. Dectin 1 nanobodies at final concentrations of 5, 2.5, 1, and 0.5 mg/mL were added to recombinant dectin 1–coated microplates. According to the results, OD$_{450}$ of nanobodies specific to dectin 1 at a concentration of 5 mg/mL was about 3. OD$_{450}$ decreased with decreased nanobody concentration. OD$_{450}$ of 2.5 mg/mL is slightly lower than that of 5 mg/mL. OD$_{450}$ of 1 mg/mL decreased significantly, and for 0.5 mg/mL was still three times that of the negative control. OD$_{450}$ of negative nanobodies was not significantly different from that of blank controls (Figure 5). To further verify the ability of nanobodies to bind to dectin 1 expressed in cells, we selected HCECs and RAW 264.7 cells for immunofluorescence staining. Compared with the blank control group and negative-nanobody group, stronger fluorescence was seen in the membrane and cytoplasm of nanobodies specific to dectin 1. Furthermore, fluorescence was stronger still in cells stimulated by A. fumigatus hyphae (Figure 6A and B).

Anti-inflammatory Effects of Nanobodies Specific to Dectin 1 in A. fumigatus Keratitis
In order to determine whether nanobodies specific to dectin 1 could inhibit the expression of proinflammatory cytokines, RT-PCR and ELISA were used to detect the expression of IL1β and IL6. Pretreatment with nanobodies 2 hours prior to A. fumigatus stimulation markedly downregulated mRNA and protein levels of IL1β and IL6 in HCECs (Figure 7A–D). The differences were statistically significant ($P<0.001$). A mouse model of A. fumigatus keratitis was established and

![Figure 5](https://www.DOI.org/10.2147/IJN.S319874)

**Figure 5** Specificity determination of nanobody specific to dectin 1. Binding activity and specificity of the were analyzed by indirect ELISA, 1–4: 5, 2.5, 1, and 0.5 mg/mL of nanobody, 5, other negative used as negative control; 6, PBS used as blank control.
Figure 6 Immunofluorescence analysis of binding activity of nanobody specificities. (A) HCECs with or without A. fumigatus stimulation were incubated with nanobodies and detected with FITC–streptavidin (green fluorescence). Negative nanobodies and PBS were used as negative controls. DAPI was used for nuclear staining (blue fluorescence). (B) RAW 264.7 cells stimulated by A. fumigatus were incubated with nanobodies. Negative nanobodies were used as negative controls.
clinical scores used to evaluate the degree of inflammation. As shown in Figure 8A and B, the nanobodies alleviated the inflammatory response and reduced clinical scores. Immunofluorescence staining showed that the fluorescence in the corneas of the normal group was weak, but stronger in the corneal epithelium 3 days after infection (Figure 8C). The nanobodies decreased cytokine levels in infected corneas. (Figure 8D–G). This difference was statistically significant (P<0.05).

Effects of Nanobodies on Proliferation of HCECs
CCK8 analysis was used to evaluate the effect of nanobodies on the proliferation of HCECs. HCECs were seeded onto 96-well cell-culture plates and treated with different concentrations of nanobodies specific to dectin 1 for 24 hours. The nanobodies did not affect the viability of HCECs (Figure 9).

Discussion
Fungal keratitis is a serious corneal disease that can cause visual impairment.26–28 It is caused by infection by pathogenic fungi and usually elicits ulceration or even perforation of the infected cornea.29–31 The lack of high-efficiency, low-toxicity, and high-permeability drugs for corneal fungal infections means poor prognosis.32–34 Nanobodies that naturally lack light chains are unique antibodies first found in the sera of sharks and camels.35 Because they possess many desirable attributes, such as small molecules, low immunogenicity, strong penetration, and high binding affinity, different types nanobodies have become novel diagnosis and treatment methods for tumors and infectious diseases.36–38 We constructed a nanobody specific to dectin 1, which is an important PRR in fungal infection. We also verified that this nanobody can bind to the dectin 1 antigen and relieve the inflammatory response in fungal keratitis.39,40

As stated in other studies, panning of a nanobody-gene library with phage display is a practical method of preparing antibodies. Phage display is a feasible method for selection of a particular domain of an antigen.41,42 Through the construction of a large-capacity natural nanobody-gene library, a variety of nanobodies can be prepared without
Figure 8 Anti-inflammatory effect of nanobody specific to dectin 1 in mouse *A. fumigatus* keratitis. (A) Photographs of nanobody- and PBS-treated corneas taken with a slit lamp at 3 and 5 days post-infection. Nanobody treatment relieved the severity of fungal keratitis in mice. Magnification 25×. (B) Significant reductions in clinical scores were observed at 3 and 5 days in nanobody-treated corneas (n=6) compared with PBS-treated corneas (n=6). (C) Immunofluorescence images taken under fluorescence microscopy. The fluorescence in the cornea of the normal group was weak, and the fluorescence in the corneal epithelium was stronger 3 days after infection. Blue, nuclear staining (DAPI); green, dectin 1 staining. Magnification 200×. (D–G) IL1β and IL6 expression in PBS- and nanobody-treatment groups.
immunization. For instance, nanobodies specific to LPS can be panned out through natural gene libraries derived from Camelid. In our study, after three rounds’ selection, the gene sequence of a nanobody specific to dectin 1 was selected. The gene was expressed in the inclusion body of *E. coli* with a high yield. The 12% SDS-PAGE result showed it was specific to dectin 1 at 17 kDa. This consistent with results of previous studies showing nanobody molecular weight of about 15 kDa. The specificity of our nanobody was further assessed by indirect ELISA and immunofluorescence staining. Indirect ELISA results suggested that it had strong binding ability to dectin 1 protein. Due to their small dimensions and permeability, nanobodies have been widely used as probes for immunoimaging and diagnostic assays. For instance, in one study PEDV-infected Vero cells were immunofluorescence-stained with S7 nanobodies. Compared with control Vero cells, those infected with PEDV were well stained by the S7 nanobody. Our immunofluorescence results showed expression of dectin 1 on HCECs and RAW 264.7 cells. These nanobodies can combine with the dectin 1 expressed in the mouse cornea. This further illustrated that dectin 1 antigens and nanobodies can interact with high affinity. These results provided a foundation for our research on anti-inflammatory effect of nanobodies in HCECs and a mouse model.

Extensive research has demonstrated that nanobodies play a significant role in several inflammatory diseases. In related experiments, nanobodies specific to TLR4 treatment significantly decreased mRNA and protein levels of proinflammatory cytokines in NR8383 cells. Consistently with that research, our in vitro tests showed that compared with negative nanobody treatment, specific nanobody treatment significantly suppressed mRNA and protein expression of IL1β and IL6 in HCECs stimulated by *A. fumigatus*. Nanobody specifically inhibits P2X7 on immune cells was designed to play a anti-inflammatory reaction which is the first nanobody that blocks the channel protein. In one study, systematic injection of nanobodies into mice blocked P2X7 on T cells and macrophages in vivo and ameliorated experimental glomerulonephritis and allergic contact dermatitis. Research has shown that orally administered *Lactococcus lactis* secreting an anti-TNF nanobody had anti-inflammatory efficacy in chronic colitis. Injecting nanobodies against TLR4 into mice can successfully reduce excessive inflammatory reactions in sepsis without obvious side effects. We found that specific nanobody treatment alleviated inflammation and improved the outcome of fungal keratitis. Clinical scores and photographs taken with a slit lamp showed less corneal opacity in nanobody-treated corneas than PBS-treated ones, and nanobody treatment reduced the expression of inflammatory factors in fungal keratitis in

![Figure 9](https://doi.org/10.2147/IJN.S338974)
mice. These results show that this nanobody can restrain excessive inflammation induced by *A. fumigatus* in HCECs and corneas. These nanobodies can have great clinical application potential in the treatment of *A. fumigatus* infection by alleviating excessive inflammatory responses. In addition, cytotoxicity assessment in HCECs was performed to verify the low toxicity of the nanobodies.

**Conclusion**

Nanobodies specific to dectin 1 were successfully expressed through a microbial system with a high yield, interacted with dectin 1 antigen with high affinity, and had an anti-inflammatory effect in a mouse fungal keratitis model. All the results implied that these nanobodies could be used as potential therapeutic agent for the treatment of fungal keratitis.

**Ethics**

Our experiments were carried out in accordance with the Basel Declaration and the ethical guidelines of the International Council for Laboratory Animal Science with the approval of an experimental animal ethics committee (QYFYWZLL-26567).

**Acknowledgments**

This work was supported by the National Natural Science Foundation of China (81870632), Natural Science Foundation of Shandong Province (ZR2017BH025), Shandong Province Key Research and Development Plan (2019GSF107022), Taishan Scholar Project of Shandong Province (tsqn201812151), and Youth National Natural Science Foundation of China (81500695).

**Disclosure**

Xing Liu and Jianxin Sui are joint first authors. The authors declare no conflict of interest.

**References**


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https://doi.org/10.2147/IJN.S338974

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