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Epitope Mapping of Anti-mouse CCR3 Monoclonal Antibodies using Flow Cytometry

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Abstract: The CC chemokine receptor 3 (CCR3) is a receptor for CC chemokines, including CCL5/RANTES, CCL7/MCP-3, and CCL11/eotaxin. CCR3 is expressed on the surface of eosinophils, basophils, a subset of Th2 lymphocytes, mast cells, and airway epithelial cells. CCR3 and its ligands are involved in airway hyperresponsiveness in allergic asthma, ocular allergy, and cancers. Therefore, CCR3 is an attractive target for those therapies. Previously, anti-mouse CCR3 (mCCR3) monoclonal antibodies (mAbs), C₃Mab-3 (rat IgG_{2a}, kappa), and C₃Mab-4 (rat IgG_{2a}, kappa) were developed using the Cell-Based Immunization and Screening (CBIS) method. In this study, the binding epitope of these mAbs was investigated using flow cytometry. The CCR3 extracellular domain-substituted mutant analysis showed that C₃Mab-3, C₃Mab-4, and a commercially available mAb (J073E5) recognized the N-terminal region (amino acids 1–38) of mCCR3. Next, the alanine scanning was conducted in C₃Mab-3 binding, whereas Ala2, Phe3, and Thr5 are essential to C₃Mab-4 binding, and Ala2 and Phe3 are crucial to J073E5 binding. These results reveal the involvement of the N-terminus of mCCR3 in the recognition of C₃Mab-3, C₃Mab-4, and J073E5.

Keywords: mouse CCR3; monoclonal antibody; epitope mapping; alanine scanning; flow cytometry

1. Introduction

Chemokines are a family of small cytokines secreted by cells and play essential roles in cell migration, inflammation, and immune responses by binding to chemokine receptors [1-4]. The CC chemokine receptor 3 (CCR3) is a receptor for CC chemokines, including CCL5/RANTES, CCL7/MCP-3, and CCL11/eotaxin [5-7]. CCR3 is expressed on the surface of eosinophils, basophils, a subset of Th2 lymphocytes, mast cells, and airway epithelial cells [8-13]. CCR3 is a family of G protein-coupled receptors (GPCRs) that transduce extracellular signals to intracellular signaling molecules [14]. The CCR3 signaling pathway is critical in eosinophil migration [15,16]. It has been reported that CCR3 and its ligands can cause airway hyperresponsiveness in a murine allergic asthma model [17-20], contributing to the ocular allergy [21]. Moreover, elevated eotaxin expression has been observed in colorectal cancer [22], breast cancer [23], and oral squamous cell carcinomas [24]. Therefore, CCR3 and its ligands are the therapeutic targets for allergic diseases and cancers [7,25].

GPCR has seven transmembrane domains, four extracellular regions, including an N-terminal region (residues 1–38), and three extracellular loops (ECL1; amino acids [aa] 96–111, ECL2; aa 176–207, and ECL3; aa 269–285). Previously, monoclonal antibodies

(mAbs) have been developed against GPCRs, including an anti-mouse CCR2 mAb [26], an anti-human CCR2 mAb [27], an anti-mouse CCR3 (mCCR3) mAbs [28-30], an anti-mouse CCR4 mAb [31], an anti-mouse CCR8 mAbs [32-34], an anti-human CCR9 mAb [35], and an anti-mouse CXCR6 mAb [36]. The binding epitopes of anti-CCR2, CCR4, CCR9, and CXCR6 mAbs, which were established using the peptide immunization method, were determined by enzyme-linked immunosorbent assay (ELISA) [37-39]. However, anti-mCCR3 mAbs, C₃Mab-3 (rat IgG_{2a}, kappa) [30], and C₃Mab-4 (rat IgG_{2a}, kappa) were established by the Cell-Based Immunization and Screening (CBIS) method. Therefore, ELISA could not be applied to their epitope mapping.

In this study, epitope mapping of the anti-mCCR3 mAbs was conducted by flow cytometry using the extracellular region substitution and the alanine scanning methods to clarify the features of C₃Mab-3 and C₃Mab-4.

2. Materials and Methods

2.1. Cell lines

Chinese hamster ovary (CHO)-K1 cell was obtained from the America Type Culture Collection (ATCC, Manassas, VA, USA). The CHO/mCCR3 cells were produced in our previous study [28]. The chimera and the point mutant plasmids were transfected into CHO-K1 cells using the Neon Transfection System (Thermo Fisher Scientific Inc., Waltham, MA, USA). Stable transfectants were selected using a cell sorter (SH800; Sony Biotechnology Inc., Tokyo, Japan). Cells were cultured in Roswell Park Memorial Institute (RPMI) 1640 medium (Nacalai Tesque, Inc., Kyoto, Japan) supplemented with 10% heat-inactivated fetal bovine serum (FBS) (Thermo Fisher Scientific Inc.), 100 units/mL of penicillin, 100 μ g/mL streptomycin, and 0.25 μ g/mL amphotericin B (Nacalai Tesque, Inc.) at 37 °C in a humidified atmosphere containing 5% CO₂. The stable transfectants were cultivated in a medium containing 0.5 mg/mL Zeocin (InvivoGen, San Diego, CA, USA).

2.2. Plasmid construction

Synthesized DNA (Eurofins Genomics KK, Tokyo, Japan) encoding mCCR3 (Accession No.: NM_009914.4) [28-30] and mouse CCR8 (mCCR8; Accession No.: NM_007720.2) [32-34] were subcloned into a pCAG-Ble vector (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan), respectively. Chimeric mutants mCCR8 (mCCR3p1-38), mCCR8 (mCCR3p96-111), mCCR8 (mCCR3p176-207), and mCCR8 (mCCR3p269-285) were produced with a RAP [40,41] and a MAP tag [42,43] at their C-terminus using the HotStar HiFidelity polymerase kit (Qiagen Inc., Hilden, Germany). Alanine (glycin) substitutions in the mCCR3 N-terminal region were conducted using QuikChange Lightning Site-Directed Mutagenesis Kits (Agilent Technologies Inc., Santa Clara, CA, USA). PCR fragments bearing the desired mutations were inserted into the pCAG-Ble vector (FUJIFILM Wako Pure Chemical Corporation) using the In-Fusion HD Cloning Kit (TaKaRa Bio, Inc., Shiga, Japan).

2.3. Antibodies

C₃Mab-3 [30] and C₃Mab-7 [29] were described previously. C₃Mab-4 was also established together with C₃Mab-3. An anti-mCCR3 mAb (clone J073E5) was purchased from BioLegend (San Diego, CA, USA). A secondary Alexa Fluor 488-conjugated anti-rat IgG was purchased from Cell Signaling Technology, Inc. (Danvers, MA, USA) as well.

2.4. Flow cytometry

Cells were harvested after brief exposure to 0.25% trypsin/1 mM ethylenediaminetetraacetic acid (Nacalai Tesque, Inc.). After washing with 0.1% bovine serum albumin in phosphate-buffered saline, cells were treated with primary mAbs (1 μ g/ml) for 30 min at 4 °C and subsequently with Alexa Fluor 488-conjugated anti-rat IgG (1:1000; Cell Signaling Technology, Inc., Danvers, MA, USA). Fluorescence data were obtained using the EC800 Cell Analyzer (Sony Biotechnology Inc.).

3. Results

3.1. Determination of the epitope of anti-mCCR3 mAbs by flow cytometry using chimeric proteins.

C₃Mab-3 and C₃Mab-4 were established using the CBIS method, and are applicable to flow cytometry, but not to ELISA. To investigate the binding epitope of C₃Mab-3 and C₃Mab-4, we focused on four extracellular regions of mCCR3 including the N-terminal region (aa 1-38), ECL1 (aa 96-111), ECL2 (aa 176-207), and ECL3 (aa 269-285). The four extracellular regions of mCCR3 were substituted into the corresponding regions of mCCR8, which possesses a similar amino acid structure to mCCR3. As shown in Fig. 1, mCCR8 (mCCR3p1-38), mCCR8 (mCCR3p96-111), mCCR8 (mCCR3p176-207), and mCCR8 (mCCR3p269-285) were generated. The chimeric proteins were transiently expressed on CHO-K1 cells, and their reactivities to C₃Mab-3, C₃Mab-4, and commercially available J073E5 reacted with mCCR8 (mCCR3p1-38) and mCCR3. In contrast, they did not react with mCCR8 (mCCR3p96-111), mCCR8 (mCCR3p176-207), and mCCR8 (mCCR3p269-285). These results show that the N-terminal region of mCCR3 is recognized by C₃Mab-4, and J073E5.



Figure 1. Schematic illustration of mCCR8 and mCCR3 chimeric proteins. The four extracellular regions of mCCR3, including the N-terminal region (residues 1–38), ECL1 (residues 96–111), ECL2 (residues 176–207), and ECL4 (residues 269–285) were substituted into the corresponding regions of mCCR8. ECL, extracellular loop.



Figure 2. Determination of the epitope of anti-mCCR3 mAbs by flow cytometry using chimeric proteins. C₃Mab-3 (1 μ g/mL) (A), C₃Mab-4 (1 μ g/mL) (B), and J073E5 (1 μ g/mL) (C) were treated with CHO-K1 cells which were transiently expressed chimeric proteins for 30 min at 4°C, followed by the addition of Alexa488-conjugated anti-rat IgG. Red lines show the cells with anti-mCCR3 mAb treatment, and black lines show cells without anti-mCCR3 mAbs treatment as a negative control.

3.2. Determination of the C₃Mab-3 epitope by flow cytometry using alanine scanning.

Next, alanine scanning was conducted in the N-terminal region except for Cys28. Thirty-six alanine substitution mutants of mCCR3 were constructed, and the mutant proteins were transiently expressed on CHO-K1 cells. The reactivity against C₃Mab-3, C₃Mab-4, and J073E5 was assessed using flow cytometry. As shown in Fig. 3A, C₃Mab-3 did not react with the four mutants (A2G, F3A, N4A, and T5A). In contrast, C₃Mab-3 reacted with the other 32 mutants. These results showed that four residues (Ala2, Phe3, Asn4, and Thr5) of mCCR3 are important for C3Mab-3 binding (Fig. 3B). C3Mab-4 did not react with three mutants (A2G, F3A, and T5A) but reacted with others (Fig. 4A), indicating that three residues (Ala2, Phe3, and Thr5) of mCCR3 are important for C₃Mab-4 binding (Fig. 4B). J073E5 did not react with two mutants (A2G and F3A) but reacted with others (Fig. 5A), indicating that two residues (Ala2 and Phe3) of mCCR3 are important for J073E5 binding (Fig. 5B). The cell surface expression of mCCR3 mutants on CHO-K1 cells was confirmed using anti-mCCR3 mAbs, C3Mab-7. It has already been confirmed that Phe15 and Glu16 are essential for C₃Mab-7 binding (manuscript submitted). We could confirm the cell surface expression of four mutants (A2G, F3A, N4A, and T5A) of mCCR3 using C3Mab-7 (Fig. 6).



Figure 3. Determination of the C₃Mab-3 epitope by flow cytometry using alanine scanning. (A) C₃Mab-3 (1 μ g/mL) was treated with CHO-K1 cells which were transiently expressed mutant proteins for 30 min at 4°C, followed by the addition of Alexa488-conjugated anti-rat IgG. Red lines show the cells with C₃Mab-3 treatment, and black lines show cells without Ab treatment as a negative control. (B) The C₃Mab-3 epitope for mCCR3 involves Ala2, Phe3, Asn4, and Thr5 of mCCR3.



Figure 4. Determination of the C₃Mab-4 epitope by flow cytometry using alanine scanning. (A) C₃Mab-4 (1 μ g/mL) was treated with CHO-K1 cells which were transiently expressed mutant proteins for 30 min at 4°C, followed by the addition of Alexa488-conjugated anti-rat IgG. Red lines show the cells with C₃Mab-4 treatment, and black lines show cells without Ab treatment as a negative control. (B) The C₃Mab-4 epitope for mCCR3 involves Ala2, Phe3, and Thr5 of mCCR3.



Figure 5. Determination of the J073E5 epitope by flow cytometry using alanine scanning. (A) J073E5 (1 μ g/mL) was treated with CHO-K1 cells which were transiently expressed mutant proteins for 30 min at 4°C, followed by the addition of Alexa488-conjugated anti-rat IgG. Red lines show the cells with J073E5 treatment, and black lines show cells without Ab treatment as a negative control. (B) The J073E5 epitope for mCCR3 involves Ala2, and Phe3 of mCCR3.



Fluorescence intensity

Figure 6. Cell surface expression of mCCR3 mutants on CHO-K1 cells by flow cytometry. C₃Mab-7 (1 μ g/mL) was treated with CHO-K1 cells which were transiently expressed mutant proteins for 30 min at 4°C, followed by the addition of Alexa488-conjugated anti-rat IgG. Red lines show the cells with C₃Mab-7 treatment, and black lines show cells without Ab treatment as a negative control.

4. Discussion

We have established various mAbs against membrane proteins using the CBIS method. Because the mAbs sometimes recognize the conformational epitope, they can be applied to flow cytometry but not to Western blotting and ELISA. Two anti-mCCR3 mAbs examined in this study, C₃Mab-3 and C₃Mab-4, were established using the CBIS method [30]. We first attempted to identify their epitope using synthetic peptides using ELISA. However, they did not recognize the synthetic peptides, including the mCCR3 N-terminal region (p1–19), which contains the epitope determined using flow cytometry (Fig. 3 and 4). These results suggest that the residues participate in the formation of conformational epitope and/or undergo the post-translational modification on the cell surface. Furthermore, we could not exclude the possibility of the first Met as their epitopes. In the case of C₃Mab-3 and C₃Mab-4 epitopes, Asn4 and Thr5 are involved in the recognition. Although Asn and Thr are known to be *N*- and *O*- glycosylated respectively, there is no report on the glycosylation of Asn4 and Thr5 of mCCR3. Further studies were required to analyze the involvement of the posttranslational modification of these residues in the recognition by C₃Mab-3 and C₃Mab-4.

Other anti-mCCR3 mAbs, C₃Mab-6, and C₃Mab-7, were also developed by mCCR3 N-terminal peptide immunization [29]. It was found that they could recognize the synthetic peptide of the mCCR3 N-terminal region (p1-19) using ELISA and cell surface expressed mCCR3 by flow cytometry. Furthermore, Phe3, Asn4, Thr5, Asp6, Glu7, Lys9, Thr10, and Glu13 of mCCR3 were determined as C₃Mab-6 epitope, whereas Phe15 and Glu16 as C₃Mab-7 epitope (manuscript submitted). These results indicate that C₃Mab-6 and C₃Mab-7 recognize both the naked N-terminal peptide and cell surface expressed mCCR3. Further studies are essential to understand the difference of mCCR3 recognitions between C₃Mab-3/C₃Mab-4 and C₃Mab-6/C₃Mab-7.

It has been reported that a CCR3 ligand, CCL11/eotaxin binds to the N-terminal region of CCR3 [44,45]. Therefore, our established anti-mCCR3 mAbs could compete with the ligand binding to mCCR3 and have neutralizing activity. Shen *et al.* reported that anti-CCR3 mAb could significantly suppress airway eosinophilia and mucus overproduction in asthmatic mice; therefore, the blockage of the CCR3 axis may be an attractive strategy for asthma therapy [46]. In future studies, we would like to examine the neutralizing activities of these anti-mCCR3 mAbs.

Author Contributions: NT, GL, TT, TY, and TA performed the experiments. MKK and YK designed the experiments. NT, TA, HS, and YK analyzed the data. TA, HS, and YK wrote the manuscript. All authors read and approved the final manuscript and agreed to be accountable for all aspects of the research in ensuring that the accuracy or integrity of any part of the work is appropriately investigated and resolved.

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Conflicts of Interest: The authors declare no conflict of interest involving this article.

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