



Epitope Mapping of Anti-Mouse CCR3 Monoclonal Antibodies (C₃Mab-6 and C₃Mab-7)

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One of G protein-coupled receptors, CC chemokine receptor 3 (CCR3), is expressed in eosinophils, basophils, a subset of Th2 lymphocytes, mast cells, and airway epithelial cells. CCR3 levels in the serum of colorectal cancer patients are significantly higher than in control groups. Moreover, CCR3 is essential for recruiting eosinophils into the lung. Therefore, CCR3 is considered both a therapeutic target for colorectal cancer and allergic diseases. Previously, we established anti-mouse CCR3 (mCCR3) monoclonal antibodies (mAbs), C₃Mab-6 (rat IgG₁, kappa) and C₃Mab-7 (rat IgG₁, kappa), by immunizing a rat with an N-terminal peptide of mCCR3. These mAbs can be used in flow cytometry and enzyme-linked immunosorbent assays. In this study, we performed the epitope mapping of C₃Mab-6 and C₃Mab-7 using alanine scanning. The reactivity between these mAbs and point mutants of mCCR3 were analyzed using flow cytometry. The results indicated that Phe3, Asn4, Thr5, Asp6, Glu7, Lys9, Thr10, and Glu13 of mCCR3 are essential for C₃Mab-6 binding, whereas Phe15 and Glu16 are essential for C₃Mab-7 binding.

Keywords: mouse CCR3, monoclonal antibody, epitope, flow cytometry, alanine scanning

Introduction

THE CC CHEMOKINE RECEPTOR 3 (CCR3), also known as CD193, is a member of the G protein-coupled receptor.¹ CCR3 is a seven-transmembrane domain receptor with four extracellular domains, and the N-terminal region of CCR3 is located in extracellular, whereas the C-terminal region is located in intracellular. CCR3 is expressed on the surface of eosinophils, basophils, a subset of Th2 lymphocytes, mast cells, and airway epithelial cells.²⁻⁶ Some CC chemokines, such as eotaxin-1, eotaxin-2, eotaxin-3, MCP-3, and RANTES, are known to be ligands for CCR3.⁷⁻⁹

CCR3 and its ligands play an important role in recruiting eosinophils into the lung, inducing airway hyper-responsiveness in a murine model of allergic asthma.¹⁰⁻¹³ CCR3 also contributes to ocular allergy.¹⁴ Moreover, CCR3 levels in the serum of colorectal cancer patients are significantly higher than in control groups.^{15,16} High expression levels of eotaxins occur in some tumors such as colorectal

cancer,¹⁷ breast cancer,¹⁸ and oral squamous cell carcinoma.¹⁹ Therefore, CCR3 is regarded as the therapeutic target for both allergic diseases and cancers.^{9,15,20-22}

Monoclonal antibodies (mAbs) are used for the treatment of inflammatory diseases and cancers.²³⁻²⁶ Many therapeutic mAbs possess neutralizing activity through the blockade between the targets and their ligands.²⁷ The epitope identification of mAbs is important to elucidate the pharmacological function of mAbs, and is essential to avoid unexpected cross-reactivity. Previously, we produced novel anti-mouse CCR3 (mCCR3) mAbs, C₃Mab-6 (rat IgG₁, kappa), and C₃Mab-7 (rat IgG₁, kappa) by immunizing a rat with the mCCR3 N-terminal peptide.²⁸ C₃Mab-6 and C₃Mab-7 reacted not only with mCCR3-overexpressed Chinese hamster ovary-K1 (CHO/mCCR3) cells, but also P388 (mouse lymphoid neoplasm) and J774-1 (mouse macrophage-like) cells, which express mCCR3 endogenously in flow cytometry. However, the critical amino acids of mCCR3 for the binding of C₃Mab-6 and C₃Mab-7 have not been determined.

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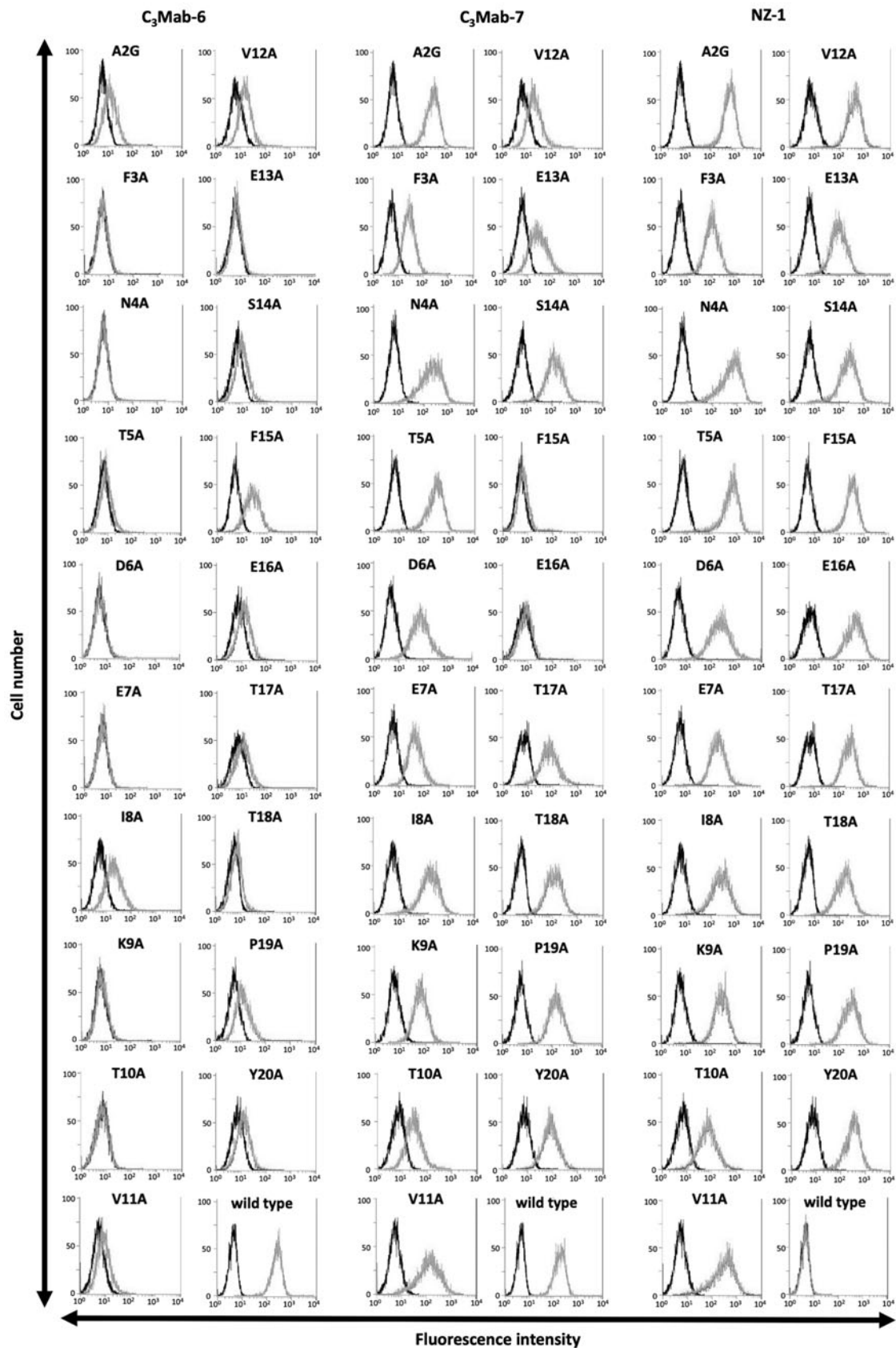


FIG. 1. Determination of the C₃Mab-6 and C₃Mab-7 epitope for mCCR3 by flow cytometry. C₃Mab-6 (1 µg/mL; gray line), C₃Mab-7 (1 µg/mL; gray line), or NZ-1 (1 µg/mL; gray line) were treated with mCCR3 point mutants-overexpressed CHO-K1 cells for 30 minutes at 4°C, followed by the addition of the secondary antibody. The black line represents the negative control. CHO-K1, Chinese hamster ovary-K1; mCCR3, mouse CC chemokine receptor 3.

In this study, we determined the binding epitope of C₃Mab-6 and C₃Mab-7 to mCCR3 using alanine scanning by flow cytometry.^{29–34}

Materials and Methods

Plasmids of mCCR3 point mutants

The pEX-A2J2 plasmid subcloned mCCR3 (accession No. NM_009914.4) was purchased from Eurofins Genomics KK (Tokyo, Japan).^{28,35–37} Substitutions of amino acids to alanine in the mCCR3 sequence were conducted by QuikChange Lightning Site-Directed Mutagenesis Kits (Agilent Technologies, Inc., Santa Clara, CA). PCR fragments bearing the desired mutations were inserted into the pCAG-Ble vector (FUJIFILM Wako Pure Chemical Corporation, Osaka, Japan) and a PA16 tag^{38–41} was added at the N-terminus using the In-Fusion HD Cloning Kit (Takara Bio, Inc., Shiga, Japan). The alanine (glycine) substitution mutants (A2G, F3A, N4A, T5A, D6A,

E7A, I8A, K9A, T10A, V11A, V12A, E13A, S14A, F15A, E16A, T17A, T18A, P19A, and Y20A) were generated using QuikChange Lightning Site-Directed Mutagenesis Kits (Agilent Technologies Inc., Santa Clara, CA, USA).

Cell lines

CHO-K1 cells were obtained from the America Type Culture Collection (ATCC, Manassas, VA). The CHO/mCCR3 cells were produced in our previous study.³⁷ mCCR3 point mutant plasmids were transfected into CHO-K1 cells using the Neon Transfection System (Thermo Fisher Scientific, Inc., Waltham, MA). Stable transfectants were selected using a cell sorter (SH800; Sony Corp., Tokyo, Japan). CHO/mCCR3 cell and stable transfectants were cultured in RPMI 1640 medium (Nacalai Tesque, Inc., Kyoto, Japan) supplemented with 10% heat-inactivated fetal bovine serum (Thermo Fisher Scientific, Inc.), 100 U/mL of penicillin, 100 μg/mL streptomycin, and

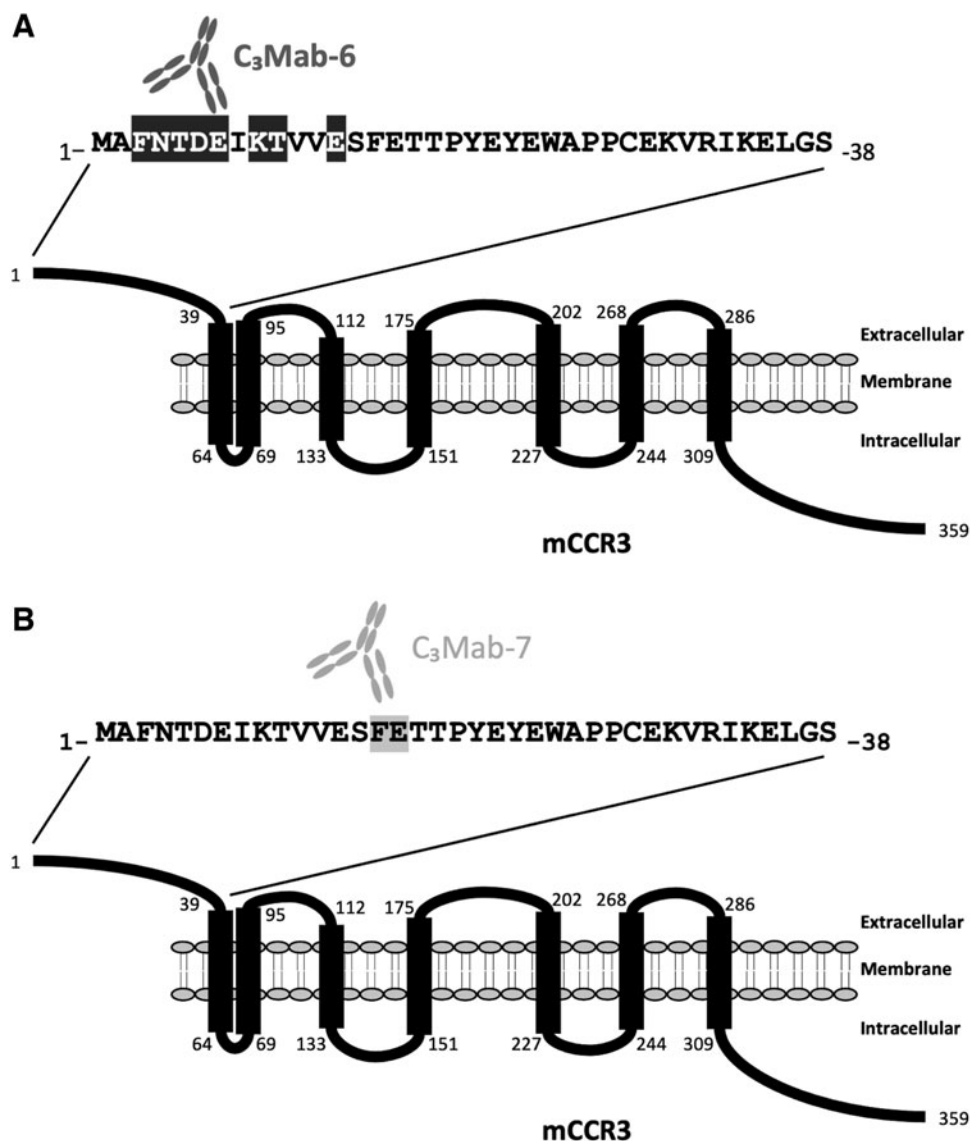


FIG. 2. Schematic illustration of C₃Mab-6 and C₃Mab-7 epitope for mCCR3. (A) C₃Mab-6 epitope for mCCR3 involves Phe3, Asn4, Thr5, Asp6, Glu7, Lys9, Thr10, and Glu13 of mCCR3. (B) C₃Mab-7 epitope for mCCR3 involves Phe15 and Glu16 of mCCR3.

0.25 µg/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 µg/mL Zeocin (InvivoGen, San Diego, CA).

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 C₃Mab-6, C₃Mab-7, or NZ-1 (1 µg/mL) for 30 minutes at 4°C and subsequently with Alexa Fluor 488-conjugated anti-rat IgG (1:1000; Cell Signaling Technology, Inc., Danvers, MA). Fluorescence data were collected using EC800 Cell Analyzer (Sony Corp.).

Results

To investigate the epitope of C₃Mab-6 and C₃Mab-7 for mCCR3, we conducted alanine scanning. We constructed 19 alanine substitution mutants of mCCR3. Mutant proteins were stably expressed on CHO-K1 cells, and the reactivities against C₃Mab-6, C₃Mab-7, or anti-PA16 tag mAb (NZ-1) were analyzed using flow cytometry.

First, we investigated the cell surface expression of mCCR3 mutants on CHO-K1 cells using NZ-1. The reactivities of NZ-1 against each mCCR3 mutant were confirmed in all the mutants (Fig. 1, right). Since PA16 tag was not conjugated in the wild-type mCCR3, NZ-1 did not react with it (Fig. 1, right). Next, we examined the reactivities of C₃Mab-6 and C₃Mab-7 against mCCR3 mutants. C₃Mab-6 reacted with 11 mutants (A2G, I8A, V11A, V12A, S14A, F15A, E16A, T17A, T18A, P19A, and Y20A) and wild-type mCCR3 (Fig. 1, left).

In contrast, C₃Mab-6 did not react with 8 mutants (F3A, N4A, T5A, D6A, E7A, K9A, T10A, and E13A) (Fig. 1, left). C₃Mab-7 reacted with 17 mutants (A2G, F3A, N4A, T5A, D6A, E7A, I8A, K9A, T10A, V11A, V12A, E13A, S14A, T17A, T18A, P19A, and Y20A) and wild-type mCCR3, whereas 2 mutants (F15A and E16A) lost their reactivity to C₃Mab-7 (Fig. 1, middle). These results indicated that eight amino acids of mCCR3 (Phe3, Asn4, Thr5, Asp6, Glu7, Lys9, Thr10, and Glu13) are critical for C₃Mab-6 binding (Fig. 2A), whereas two amino acids (Phe15 and Glu16) are critical for that on C₃Mab-7 (Fig. 2B).

Discussion

Therapeutic mAbs possesses biological activities, including neutralizing activity against the physiological ligands.²⁷ The epitope determination in the target protein is necessary to understand the properties of mAbs. Previously, we established two anti-mCCR3 mAbs, C₃Mab-6 and C₃Mab-7, which were established by immunizing the N-terminal peptide of mCCR3 (₁-MAFNTDEIKTVVESFETTP-₁₉).²⁸ Therefore, the binding epitope of C₃Mab-6 and C₃Mab-7 are included within the residues.

In this study, we performed the epitope mapping of C₃Mab-6 and C₃Mab-7 using alanine scanning of the mCCR3 N-terminal domain. The results showed that C₃Mab-6 and C₃Mab-7 have different binding regions of mCCR3 (Fig. 1). The C₃Mab-6 epitope is mainly located in the first half of the N-terminal region, whereas that of C₃Mab-7 is located in another half of the N-terminal region (Fig. 2). Structural analyses showed that eotaxin binds to the

N-terminus of human CCR3 (residues 8–23).^{42,43} Therefore, it is expected that both C₃Mab-6 and C₃Mab-7 may compete with eotaxin for binding to mCCR3.

Zhu *et al.* showed that the dissociation constants (*K_D*) of CCR3's binding to eotaxin-1, -2, and -3 are 2.1, 9.7, and 1.2 × 10⁻⁹ M, respectively.⁴⁴ We previously showed that C₃Mab-6 and C₃Mab-7 bound to CHO/mCCR3 with *K_D* of 8.7 × 10⁻⁹ M and 3.7 × 10⁻⁹ M, respectively.²⁸ The *K_D* values of C₃Mab-6 and C₃Mab-7 to CHO/mCCR3 were comparable with that of eotaxin. Therefore, we expect that C₃Mab-6 and C₃Mab-7 treatments would block eotaxin's binding to CHO/mCCR3.

In future studies, the neutralizing activities of C₃Mab-6 and C₃Mab-7 should be further assessed. Moreover, these mAbs will be useful for depleting mCCR3-expressing eosinophils and basophils, and targeting mCCR3-positive cancers.

Author Disclosure Statement

No competing financial interests exist.

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