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Epitope Mapping of the Novel Anti-Human CCR9 Monoclonal Antibody (C₉Mab-11) by 2×Alanine Scanning

Yu Isoda,^{1,*} Tomohiro Tanaka,^{1,*} Hiroyuki Suzuki,² Teizo Asano,¹ Kaishi Kitamura,² Yuma Kudo,² Ryo Ejima,² Kazuki Ozawa,² Takeo Yoshikawa,³ Mika K. Kaneko,¹ and Yukinari Kato¹⁻³

We recently developed a novel anti-human C-C chemokine receptor 9 (hCCR9) monoclonal antibody (mAb), C₉Mab-11, which is applicable to flow cytometry, western blotting, and enzyme-linked immunosorbent assay (ELISA). This study aims to identify the binding epitope of C₉Mab-11 by using 1× and 2×alanine (or glycine) substituted-hCCR9 peptides (1× and 2×Ala-scan) by ELISA. According to the 1×Ala-scan analysis, the response of C₉Mab-11 was diminished against M13A of the hCCR9 peptide, but was not eliminated. In the 2×Ala-scan analysis, the reactions were abolished in the substitution of P11A-N12A, N12A-M13A, and M13A-A14G of hCCR9 N-terminal peptides. The results indicate that the binding epitope of C₉Mab-11 includes Pro11, Asn12, Met13, and Ala14 of hCCR9, with the region around Met13 being particularly important. The successful identification of the C₉Mab-11 epitope might be useful for the future pathophysiological analysis of hCCR9.

Keywords: human CCR9, monoclonal antibody, epitope, ELISA, 2×Ala-scan

Introduction

THE C-C CHEMOKINE RECEPTOR 9 (CCR9) is a member of the G protein-coupled receptors (GPCRs), which are highly expressed in T cells, thymocytes, B cells, plasmacytoid dendritic cells (pDCs), and intestinal cells.¹ Chemokine ligand 25 (CCL25), formerly thymus-expressed chemokine (TECK), regulates the activation and infiltration of immature T cells by binding to CCR9.^{2,3} CCL25 is the only known ligand for CCR9 and is mainly secreted from the epithelial cells of the thymus and small intestine. In addition, CCR9+ T cells are gut-homing T cells because the CCR9 and CCL25 interactions potently recruit T cells to intestinal tissues.⁴⁻⁶ CCR9 on immune cells guides the spatial organization and cellular communications of immune cells within tissues.

The CCR9/CCL25 axis is involved in the development of various inflammatory diseases. The increase in CCR9+ monocytes was confirmed in the synovium of patients with rheumatoid arthritis.⁷ In the ovalbumin-induced allergic inflammation model, the recruitment of eosinophils and T cells was impaired at the peribronchial and perivenular levels in CCR9-deficient mice.⁸ In hepatitis, CCR9+ macrophages

trigger acute liver damage by collaborating with helper T1 (Th1) cells.⁹ However, CCR9 axis inhibition avoided pDCs capture in the small intestine, enhanced pDCs infiltration into the liver, and protected against liver injury.¹⁰ Therefore, managing the CCR9/CCL25 axis could be a target for immunotherapy in numerous inflammation-associated disorders.

CCR9 is highly expressed in T cell acute lymphoblastic leukemia (T-ALL), contributing to the progression of T-ALL.¹¹ Furthermore, CCR9 expression has been confirmed with high probability in melanomas that have metastasized to the intestine.^{12,13} Thus, CCR9 is considered a therapeutic target for cancer. In the T-ALL-bearing mouse model, an anti-human CCR9 monoclonal antibody (hCCR9 mAb) (92R) and humanized 92R (Srb1) inhibited tumor growth and prolonged the survival of mice.^{14,15} A clinical study of the CCR9 antagonist CCX282-B against intestinal bowel disease was conducted.^{16,17} However, there are still no approved drugs despite CCR9 being an attractive target.

Previously, we developed various anti-mouse and anti-human GPCR mAbs,¹⁸⁻²⁵ including an anti-hCCR9 mAb (clone C₉Mab-11).²⁶ We further characterized the binding

Departments of ¹Antibody Drug Development, ²Molecular Pharmacology, and ³Pharmacology, Tohoku University Graduate School of Medicine, Sendai, Japan.

*These authors contributed equally to this study.

epitopes of these mAbs.^{27–29} The binding epitope of CX₆Mab-1, an anti-mouse CXCR6 mAb, was identified by enzyme-linked immunosorbent assay (ELISA) using the two alanine scanning method (2×Ala-scan).²⁹ We demonstrated that the 2×Ala-scan was an effective method for epitope mapping of mAbs.

In this study, we determined the epitope of C₉Mab-11 using the 1× and 2×Ala-scan strategies.

Materials and Methods

Peptides

The hCCR9 (Accession No. NM_031200) peptide (4-TDFTSPIPNMADDYGSEST-₂₂), one-alanine (or glycine) residue-substituted peptide (Supplementary Table S1), and two alanine (or glycine) residue-substituted peptides (Table 1) were synthesized using PEPscreen (Sigma-Aldrich Corp., St. Louis, MO).

Enzyme-linked immunosorbent assay

Synthesized hCCR9 peptides were immobilized on Nunc Maxisorp 96-well immunoplates (Thermo Fisher Scientific, Inc., Waltham, MA) at a 1 μg/mL concentration for 30 minutes at 37°C. After washing with phosphate-buffered saline containing 0.05% Tween20 (PBST; Nacalai Tesque, Inc., Kyoto, Japan), the wells were blocked with 1% bovine serum albumin-containing PBST for 30 minutes at 37°C. The plates were incubated with 1 μg/mL of C₉Mab-11, followed by peroxidase-conjugated anti-mouse immunoglobulin (1:2000 diluted; Agilent Technologies Inc., Santa Clara, CA). Enzymatic reactions were performed using the ELISA POD Substrate TMB Kit (Nacalai Tesque, Inc.). The optical density was measured at 655 nm using an iMark microplate reader (Bio-Rad Laboratories, Inc., Berkeley, CA).

TABLE 1. IDENTIFICATION OF THE C₉MAB-11 EPITOPE USING 2×ALANINE-SUBSTITUTED hCCR9 PEPTIDES

Peptides	Sequences	C ₉ Mab-11
WT	TDFTSPIPNMADDYGSEST	+++
T4A–D5A	AAFTSPIPNMADDYGSEST	+++
D5A–F6A	TAATSPIPNMADDYGSEST	+++
F6A–T7A	TDAASPIPNMADDYGSEST	+++
T7A–S8A	TDFAAIPNMADDYGSEST	+++
S8A–P9A	TDFTAAIPNMADDYGSEST	+++
P9A–I10A	TDFTSAAPNMADDYGSEST	+++
I10A–P11A	TDFTSPAANMADDYGSEST	+++
P11A–N12A	TDFTSPIAAMADDYGSEST	-
N12A–M13A	TDFTSPIPAAADDYGSEST	-
M13A–A14G	TDFTSPIPNAGDDYGSEST	-
A14G–D15A	TDFTSPIPNMGADYGSEST	+++
D15A–D16A	TDFTSPIPNMAAAYGSEST	+++
D16A–Y17A	TDFTSPIPNMADAAGSEST	+++
Y17A–G18A	TDFTSPIPNMADDAASEST	+++
G18A–S19A	TDFTSPIPNMADDYAAEST	+++
S19A–E20A	TDFTSPIPNMADDYGAEST	+++
E20A–S21A	TDFTSPIPNMADDYGSAAEST	+++
S21A–T22A	TDFTSPIPNMADDYGSEEA	+++

+++ , OD₆₅₅ ≥ 0.3; -, OD₆₅₅ < 0.1.

hCCR9, human C-C chemokine receptor 9; WT, wild-type.

Results and Discussion

Epitope mapping of C₉Mab-11 with alanine-substituted hCCR9 peptides

We previously established an anti-hCCR9 mAb (clone C₉Mab-11) by immunizing the hCCR9 N-terminal peptide (4-TDFTSPIPNMADDYGSEST-₂₂) + C-terminal cysteine residue with keyhole limpet hemocyanin.²⁶ To investigate the binding epitope of C₉Mab-11, we first synthesized one-alanine- (or glycine-) substituted peptides of hCCR9, referred to as the 1×Ala-scan method (Supplementary Table S1). C₉Mab-11 reacted with all one-alanine- (or glycine-) substituted and wild-type (WT) hCCR9 peptides. However, compared with other substituted peptides, the reaction of C₉Mab-11 against M13A was attenuated (Supplementary Fig. S1).

Furthermore, we synthesized 18 peptides, substituting two amino acid sequences with two alanine (or glycine) residues (2×Ala-scan method). For instance, a peptide (T4A–D5A) indicates the alanine substitution of the fourth Thr and the fifth Asp of the hCCR9 peptide (Table 1). As shown in Figure 1A, C₉Mab-11 exhibited reactions with T4A–D5A, D5A–F6A, F6A–T7A, T7A–S8A, S8A–P9A, P9A–I10A, I10A–P11A, A14G–D15A, D15A–D16A, D16A–Y17A, Y17A–G18A, G18A–S19A, S19A–E20A, E20A–S21A, S21A–T22A, and WT of hCCR9 peptides. However, C₉Mab-11 did not react with P11A–N12A, N12A–M13A, and M13A–A14G (Fig. 1A), indicating that Pro11, Asn12, Met13, and Ala14 are included in the critical epitope of C₉Mab-11. The results are summarized in Table 1. Figure 1B describes a schematic illustration of hCCR9 and the estimated binding epitope of C₉Mab-11.

The 2×Ala-scan is a useful method for identifying the epitope of mAbs. Considering the results of the 1×Ala-scan (Supplementary Fig. S1), we speculate that amino acids in the region centered on Met13 of hCCR9 are pivotal epitope for C₉Mab-11 binding. Anti-hCCR9 mAbs, 91R (mouse IgG_{2b}), and 92R (mouse IgG_{2a}) generated by immunization of plasmid bearing-hCCR9 cDNA using the gene gun method have been reported to possess antitumor effects in mouse models.^{14,15,30} The hotspot of epitopes of 91R and 92R are ₁₁.PNMADD₁₆ of hCCR9,¹⁴ consistent with the epitope of C₉Mab-11 in this study (Fig. 1). Since the subclass of C₉Mab-11 is also mouse IgG_{2a},²⁶ C₉Mab-11 is expected to have an antitumor effect on CCR9-expressing cancers. In addition to antitumor activity, chimeric antigen receptor-T cells that target hCCR9 have high antitumor activity against T-ALL.⁵¹ C₉Mab-11 could contribute to the development of novel immunotherapy modalities.

Author Disclosure Statement

No competing financial interests exist.

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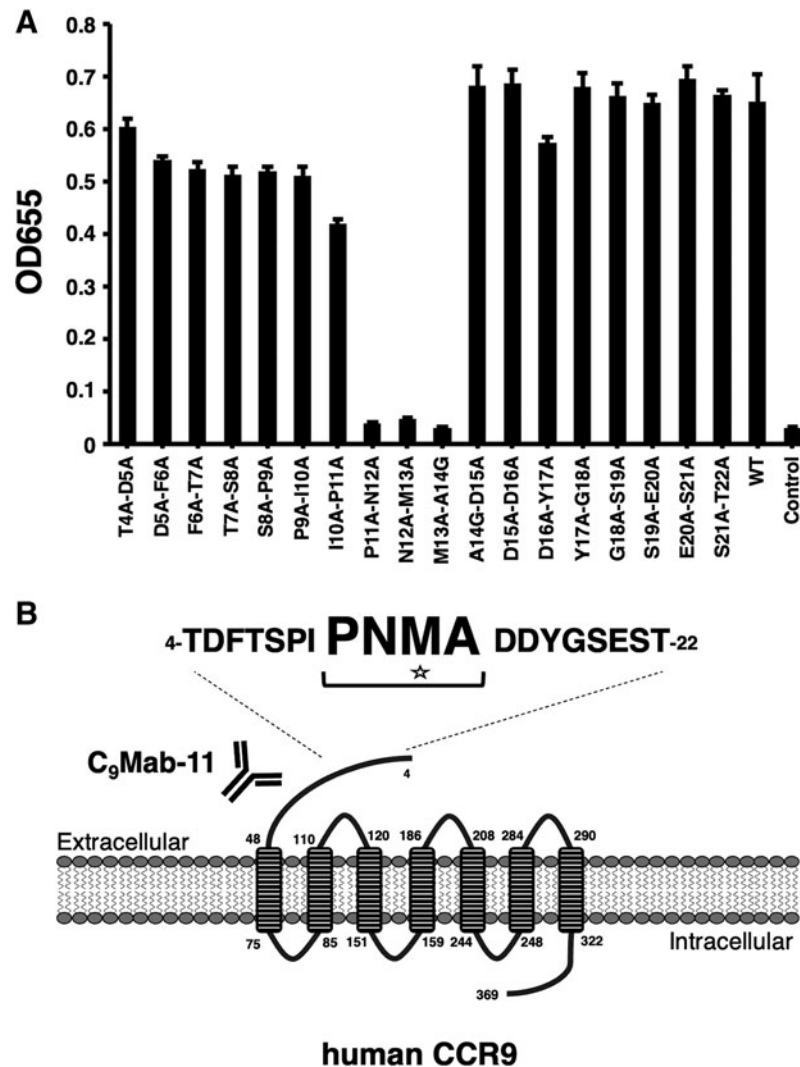


FIG. 1. Epitope identification of the C₉Mab-11 by ELISA using the 2×Ala-scan method. **(A)** The 2×alanine-substituted and WT of hCCR9 peptides were immobilized on immunoplates. The plates were incubated with C₉Mab-11 (1 μg/mL), followed by peroxidase-conjugated anti-mouse immunoglobulins. **(B)** Schematic illustration of hCCR9 and the critical epitope of C₉Mab-11. The C₉Mab-11 epitope involves Pro11, Asn12, Met13, and Ala14 of hCCR9. Star describes the estimated epitope from Supplementary Figure S1. 2×Ala-scan, 2×alanine scanning; ELISA, enzyme-linked immunosorbent assay; hCCR9, human C-C chemokine receptor 9; WT, wild-type.

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Supplementary Material

Supplementary Table S1

Supplementary Figure S1

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Address correspondence to:

Yukinari Kato
Department of Molecular Pharmacology
Tohoku University Graduate School of Medicine
2-1, Seiryomachi, Aoba-ku
Sendai 980-8575
Japan

E-mail: yukinari.kato.e6@tohoku.ac.jp

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