DESCRIPTION

Title of Invention: ANTI-HUMAN XCR1 ANTIBODIES

Technical Field

The present invention relates to an antibody that binds to human XCR1.

Background Art

Chemokine is a collective term for basic heparin-binding proteins that have effects on leukocyte chemotaxis and leukocyte activation. Based on a comparison of the primary structures of various chemokines, the chemokines are classified into CXC, CC, C, and CX3C subfamilies according to the positions of conserved cysteine residues. XCL1 (also referred to as lymphotactin (Ltn) or lymphotactin α (Ltn-α)) and XCL2 (also referred to as lymphotactin β (Ltn-β)) are chemokines classified into the subfamily C described above. XCR1 (also referred to as GPR5, SCM-1α, or ATAC) is a G protein-coupled chemokine receptor, which specifically binds to XCL1 and XCL2.

Expression of XCR1 in various human tissues has been examined at the mRNA level. Expression of XCR1 is reportedly high in placenta, but low in spleen and thymus gland (NON PATENT LITERATURE 1). Further, XCR1 is mainly expressed in dendritic cells. In mice, XCR1 is highly expressed, particularly in CD8α+ dendritic cells (NON PATENT LITERATURE 2; and NON PATENT LITERATURE 3). The CD8α+ dendritic cells are normally present in secondary lymphoid tissues such as spleen and lymph nodes, and are known to perform "cross-presentation," which serves an important role in reactions against infection and immunological responses to tumor cells. XCR1 is also known to be highly expressed in human CD141+ dendritic cells, which are considered to be homologues of mouse CD8α+ dendritic cells (NON PATENT LITERATURE 4).

Antigen taken up from the outside of cells into antigen-presenting cells is usually degraded into peptide,

presented on class II major histocompatibility antigen (MHC class II), and recognized by CD4+ T-cells. In contrast, there is a case where the antigen taken up from the outside of cells is presented on class I major histocompatibility antigen (MHC class I) via a pathway different from the usual pathway described above. This antigen presentation process is referred to as "cross-presentation." In this process, the antigen presented on MHC class I is recognized by the CD8+ T-cells, and then differentiated into cytotoxic T-cells (CTL) that play a role in phylaxis and the elimination of tumor cells in the host (Non Patent Literature 5).

Migration of various immune-related cells occurs during inflammation reaction. In particular, migration of dendritic cells to a local inflammatory site occurs for phagocytosis of antigens. Chemokines and chemokine receptors play important roles in causing such migration of dendritic cells. After migration to a local inflammatory site, the dendritic cells present antigens to T-cells, and activate T-cells. Subsequently, the information is transmitted from T-cells to many more immune-related cells, amplifying the immune reaction (Non Patent Literature 6).

Among antigen presenting cells, the dendritic cells have particularly excellent antigen-presenting ability, and play a very important role in the activation of the T-cells. It has been suggested that because T-cells are involved in the development and exacerbation of various immune diseases including autoimmune diseases, to control dendritic cells is to control the activation of T-cells, which may lead to the amelioration of various immune diseases (Non Patent Literature 6; and Non Patent Literature 7).

Further, it has been shown that a rabbit-derived polyclonal antibody against human XCR1 has an effect of inhibiting XCL-induced migration of normal oral keratinocytes and oral cancer cells (Non Patent Literature 8).

Citation List

Non Patent Literature

NPL 1: Yoshida T, Imai T, Kakizaki M, Nishimura M, Takagi S, Yoshie O. "Identification of Single C motif-1/lymphotactin receptor XCR1," J. Biol. Chem. 273: 16551-16554 (1998)

NPL 2: Crozat K, Guiton R, Contreras V, Feuillet V, Dutertre CA, Ventre E, Vu Manh TP, Baranek T, Storset AK, Marvel J, Boudinot P, Hosmalin A, Schwartz-Cornil I, Dalod M "The XC chemokine receptor 1 is a conserved selective marker of mammalian cells homologous to mouse CD8α+ dendritic cells," J Exp Med, 207: 1283-1292 (2010)

NPL 3: Dorner BG, Dorner MB, Zhou X, Opitz C, Mora A, Güttler S, Hutloff A, Mages HW, Ranke K, Schaefer M, Jack RS, Henn V, Kroczek RA "Selective expression of the chemokine receptor XCR1 on cross-presenting dendritic cells determines cooperation with CD8+ T-cells," Immunity, 31: 823-833 (2009)

NPL 4: Bachem A, Güttler S, Hartung E, Ebstein F, Schaefer M, Tannert A, Salama A, Movassaghi K, Opitz C, Mages HW, Henn V, Kloetzel PM, Gurka S, Kroczek RA, "Superior antigen cross-presentation and XCR1 expression define human $CD11c^{+}CD141^{+}$ cells as homologues of mouse $CD8^{+}$ dendritic cells," J Exp Med, 207: 1273-1281 (2010)

NPL 5: Kurts C, Robinson BW, Knolle PA, "Cross-priming in health and disease," Nat Rev Immunol, 10: 403-414 (2010)

NPL 6: Kurts C, Robinson BW, Knolle PA, "Cross-priming in health and disease," Nat Rev Immunol, 10: 403-414 (2010)

NPL 7: Waldner H, "The role of innate immune responses in autoimmune disease development," Autoimmun, Rev 8: 400-404 (2009)

NPL 8: Khurram SA, Whawell SA, Bingle L, Murdoch C, McCabe BM, Farthing PM, "Functional expression of the chemokine receptor XCR1 on oral epithelial cells," J Pathol, 221: 153-63 (2010)

Summary of Invention

Technical Problem

Knowledge that dendritic cells are involved in the development, exacerbation, and the like of immune diseases has been thus far accumulated using disease animal models. However, at present, neither an effective treatment method nor a prevention method has been developed for many immune diseases. Further, although an anti-human XCR1 antibody having an effect of inhibiting cell migration is known (Khurram SA, Whawell SA, Bingle L, Murdoch C, McCabe BM, Farthing PM, "Functional expression of the chemokine receptor XCR1 on oral epithelial cells," J Pathol, 221: 153-63 (2010)), because such an antibody is a rabbit-derived polyclonal antibody, it is unlikely to be immediately clinically applicable as a pharmaceutical product. In addition, the above document does not suggest that such an antibody inhibits cell migration of dendritic cells, and it is impossible to even predict that such an antibody will be effective in the treatment or prevention of immune diseases.

An object of the present invention is to provide a monoclonal antibody that selectively binds to human XCR1; preferably, a monoclonal antibody that selectively binds to human XCR1 and inhibits cell migration; further preferably an antibody that is effective in the treatment or prevention of immune diseases, in particular, immune diseases of the skin, based on the above-described effect.

Solution to Problem

The present inventors conducted intensive studies in an attempt to solve the above problem. As a result, they developed antibodies that bind to human XCR1, and found that such antibodies have an effect of inhibiting cell migration as well as a significant effect in the treatment or prevention of immune diseases, such as immune diseases of the skin, associated with migration of dendritic cells.

Hereinafter, in the present specification, the above-described antibodies are sometimes simply referred to as the "antibodies," "antibodies of the present invention," or "anti-

human XCR1 antibodies."

Advantageous Effects of Invention

The antibodies of the present invention bind to human XCR1. The antibodies of the present invention include an antibody that inhibits binding between human XCR1 and human XCL1. Such an antibody has potential as an active ingredient to be added to a human XCR1-human XCL1 binding inhibitor.

The antibodies of the present invention also include an antibody that inhibits cell migration, particularly that of dendritic cells. Such an antibody has potential as an active ingredient to be added to a cell migration inhibitor, particularly a dendritic cell migration inhibitor. Further, the antibodies of the present invention also include an antibody that specifically recognizes BDCA3 (also referred to as CD141) positive cells. Therefore, a pharmaceutical composition comprising the antibodies of the present invention has potential as a therapeutic agent for the treatment of immune diseases associated with cell migration, particularly dendritic cell migration. In particular, the pharmaceutical composition has potential as a therapeutic agent for the treatment of immune diseases of the skin such as delayed-type hypersensitivity, psoriasis, parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (e.g., pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, systemic scleroderma, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin diseases associated with collagenosis (e.g., systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin diseases associated with Addison's disease, skin diseases associated with graft-versus-host disease (GVHD), eczema, and urticaria.

In addition to these immune diseases of the skin, the antibodies of the present invention also has potential as therapeutic agents for the treatment of immune diseases such as

diabetes mellitus type 1, glomerulonephritis, autoimmune hepatitis, multiple sclerosis, ankylosing spondylitis, thyroiditis, graft rejection, Crohn’s disease, rheumatoid arthritis, inflammatory bowel disease, anterior uveitis, Wegener's granulomatosis, or Behçet’s disease.

Brief Description of Drawings

[Fig. 1] Fig. 1 shows the results of FACS analysis of the reactivity of mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) to human XCR1-EGFP-expressing B300.19 cells.

[Fig. 2] Fig. 2 shows the analysis results of a chemotaxis assay of the neutralizing activity of the mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) on human lymphotactin-induced migration of human XCR1-EGFP-expressing B300.19 cells.

[Fig. 3] Fig. 3 shows the results of FACS analysis of the reactivity of humanized anti-human XCR1 antibodies (HK1L2 and HK5L5) to human XCR1-EGFP-expressing B300.19 cells, in parallel with the reactivity of their parent mouse anti-human XCR1 antibody (5G7) and its chimeric antibody.

[Fig. 4] Fig. 4 shows the results of FACS analysis of the reactivity of mouse anti-human XCR1 antibody (5G7) and humanized anti-human XCR1 antibodies (HK1L2 and HK5L5) to human BDCA3+ dendritic cells.

[Fig. 5] Fig. 5 shows the analysis results of a chemotaxis assay of the neutralizing activity of the humanized anti-human XCR1 antibodies (HK1L2 and HK5L5) on human lymphotactin-induced migration of human XCR1-EGFP-expressing B300.19 cells, in parallel with the neutralizing activity of their parent mouse anti-human XCR1 antibody (5G7) and its chimeric antibody.

[Fig. 6] Fig. 6 shows the analysis results of a transendothelial migration assay of the neutralizing activity of the humanized anti-human XCR1 antibodies (HK1L2 and HK5L5) and a chimeric antibody on human lymphotactin-induced migration of human BDCA3+ dendritic cells, in parallel with an isotype control

antibody (human IgG2, κ).

[Fig. 7] Fig. 7 shows the comparison of amino acid sequences of heavy chain CDRs 1 to 3 and light chain CDRs 1 to 3 of the antibodies of the present invention.
The figure also shows the generalized amino acid sequences of heavy chain CDRs 1 to 3 and light chain CDRs 1 to 3.

[Fig. 8] Fig. 8 shows a pharmacological effect of the mouse anti-human XCR1 antibody (5G7) of the present invention on a mouse model of delayed-type contact dermatitis (DTH).
Figs. 8A and 8B respectively show the results obtained by comparing the degree of ear swelling (mm) 24 hours and 48 hours after induction by DNFB between the mouse anti-human XCR1 antibody (5G7) of the present invention and the control antibody.

[Fig. 9] Fig. 9 shows binding specificity of the mouse anti-human XCR1 antibody (5G7) of the present invention to various human chemokine receptors. The abscissa axis of the graph in the figure indicates the fluorescence intensity of phycoerythrin (PE).

[Fig. 10] Fig. 10 shows amino acid sequences of human XCR1 to which the antibodies of the present invention bind.

[Fig. 11] Fig. 11 shows cytotoxity to human XCR1 expressing cells, using the antibodies of present invention.

[Fig. 12] Fig. 12 shows the analysis of the result of cytotoxic T lymphocyte assay of the mouse anti-human XCR1 antibody (5G7) of the present invention.

[Fig. 13] Fig. 13 shows reactivity of mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) of the present invention to the chimeric human/mouse XCR1-expressing cells.

[Fig. 14] Fig. 14 shows the analysis of result of mapping of mouse anti-human XCR1 antibodies (2H6, and 5G7)-binding sites on human XCR1 extracellular domains by peptide ELISA.

[Fig. 15] Fig. 15 shows the analysis of the result of mapping of anti-human XCR1 polyclonal antibody-binding sites on human XCR1 extracellular domains by using alanine mutants.

[Fig. 16] Fig. 16 shows the analysis of the result of mapping of mouse anti-human XCR1 antibody (2H6) -binding sites on human XCR1 extracellular domains by using alanine mutants.

[Fig. 17] Fig. 17 shows the analysis of the result of mapping of mouse anti-human XCR1 antibody (5G7) -binding sites on human XCR1 extracellular domains by using alanine mutants.

[Fig. 18] Fig. 18 shows the analysis of the result of mapping of mouse anti-human XCR1 antibody (11H2)-binding sites on human EXR1 extracellular domains by using alanine mutants.

[Fig. 19] Fig. 19 shows the analysis of the result of mapping of humanized anti-human XCR1 antibody (HK1L2)-binding sites on human XCR1 extracellular domains by using alanine mutants.

[Fig. 20] Fig. 20 shows the analysis of the result of mapping of humanized anti-human XCR1 antibody (HK5L5)-binding sites on human XCR1 extracellular domains by using alanine mutants.

[Fig. 21] Fig. 21 shows the analysis of the result of the competition among mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) for binding to human XCR1-expressing cells.

[Fig. 22] Fig. 22 shows binding specificity of the mouse anti-human XCR1 monoclonal antibody (5G7) and commercial goat anti-human XCR1 polyclonal antibody to various human chemokine receptors. The abscissa axis of the graph in the figure indicates the fluorescence intensity of phycoerythrin (PE).

[Fig. 23] Fig. 23 shows binding specificity of the mouse anti-human XCR1 monoclonal antibody (5G7) and humanized anti-human XCR1 monoclonal antibodies (HK1L2 and HK5L5) to various human chemokine receptors. The abscissa axis of the graph in the figure indicates the fluorescence intensity of phycoerythrin (PE).

[Fig. 24] Fig. 24 shows a pharmacological effect of the mouse anti-human XCR1 antibody (5G7) of the present invention on a mouse model of delayed-type contact dermatitis (DTH) induced by Mycobacterium butyricum.

[Fig. 25] Fig. 25 shows a pharmacological effect of the mouse anti-human XCR1 antibody (5G7) of the present invention on a mouse model of multiple sclerosis (MS) by experimental autoimmune encephalomyelitis (EAE).

[Fig. 26] Fig.26 shows the analysis of the result of the competitive ligand binding assay of mouse anti-human XCR1 antibodies of the present invention.

Description of Embodiments

Various techniques used to practice the present invention are easily and reliably enabled for a person skilled in the art based on known documents and the like, except for those techniques whose sources are clearly identified herein. For example, in regard to genetic engineering and molecular biological techniques, reference may be made to documents such as Sambrook and Russell, "Molecular Cloning: A Laboratory Manual," Cold Spring Harbor Laboratory Press, New York, (2001); and Ausubel, F M et al., "Current Protocols in Molecular Biology," John Wiley & Sons, New York, NY.

Further, in regard to antibody engineering techniques, reference may be made to documents such as Kabat et al., "Sequences of Proteins of Immunological Interest," U.S. Department of Health and Human Services, (1983); and Konterman and Dübel, "Antibody Engineering," Springer.

Explanation of the Terms

The term "nucleic acid" encompasses, for example, ribonucleotides, deoxyribonucleotides, and their modified forms. The nucleic acid may be either single- or double-stranded, and either polynucleotide or oligonucleotide.

The term "protein" refers to a compound in which two or more amino acids are linked by peptide bonds.

The term "monoclonal antibody" refers to an antibody obtained from a population of substantially homogeneous antibodies. In other words, the individual antibodies included in

the population are identical except for naturally occurring mutations that may be present in minor amounts. Monoclonal antibodies are highly specific, and directed to a single antigenic site. Further, in contrast to polyclonal antibody preparations comprising different antibodies directed to different determinants (epitopes), each monoclonal antibody is directed to a single determinant on the antigen. In addition to their specificities, monoclonal antibodies are also advantageous in that they can be synthesized without contamination by other antibodies. The modifier "monoclonal" refers to a characteristic of an antibody obtained from a population of substantially homogeneous antibodies, and should not be interpreted to mean that antibodies must be produced by any specific method.

For example, a monoclonal antibody that should be used in accordance with the present invention can be prepared by the hybridoma method first described by Köhler G and Milstein C, "Continuous cultures of fused cells secreting antibody of predefined specificity," Nature, 256: 495-7 (1975), or by a recombinant DNA method (see U.S. Patent No. 4816567).

Further, "monoclonal antibodies" can be isolated from phage antibody library by using a technique, for example, described by Clackson T, Hoogenboom HR, Griffiths AD, and Winter G, "Making antibody fragments using phage display libraries," Nature, 352: 624-8 (1991); or Marks JD, Hoogenboom HR, and Bonnert TP, McCafferty J, Griffiths AD, Winter G, "By-passing immunization: Human antibodies from V-gene libraries displayed on phage," J Mol Biol, 222: 581-97 (1991).

The "identity" between amino acid sequences or nucleotide sequences refers to the degree of identical amino acid sequences or nucleotide sequences between two or more comparable amino acid sequences or nucleotide sequences. Accordingly, when the identity between two amino acid sequences or nucleotide sequences is high, the identity or similarity of these sequences is high. The level of identity between amino acid sequences or nucleotide sequences is determined, for example, using FASTA,

which is a sequence analysis tool, based on default parameters.

Alternatively, it can be determined using the algorithm BLAST by Karlin and Altschul (Karlin S, Altschul SF, "Methods for assessing the statistical significance of molecular sequence features by using general scoring schemes," Proc Natl Acad Sci USA, 87: 2264-2268 (1990); and Karlin S, Altschul SF, "Applications and statistics for multiple high-scoring segments in molecular sequences," Proc Natl Acad Sci USA, 90: 5873-7 (1993)). Programs such as BLASTN and BLASTX based on the above-described BLAST algorithm have been developed (Altschul SF, Gish W, Miller W, Myers EW, Lipman DJ, "Basic local alignment search tool," J Mol Biol, 215: 403-10 (1990)). For example, BLASTN may be used when analyzing the nucleotide sequence, by setting, for example, the score to 100 and the word length to 12, as parameters.

In addition, BLASTX may be used when analyzing the amino acid sequence, by setting, for example, the score to 50 and the word length to 3, as parameters.

When BLAST and Gapped BLAST programs are used, default parameters of each program may be used. Specific techniques of these analysis methods are known. Reference may be made to the website of the National Center of Biotechnology Information (NCBI) (http://www.ncbi.nlm.nih.gov/).

Anti-Human XCR1 Antibody

The antibodies of the present invention are isolated antibodies.

The antibodies of the present invention bind to human XCR1. The amino acid sequence of human XCR1 is an amino acid sequence shown by NCBI Reference Sequence: NP_001019815.1 or NP_005274.1. In regard to these amino acid sequences, reference may be made to the NCBI websites (respectively, http://www.ncbi.nlm.nih.gov/protein/NP_001019815.1 and http://www.ncbi.nlm.nih.gov/protein/NP_005274.1).

A specific antibody of a first embodiment of the present invention is an antibody comprising a heavy chain variable region comprising
a heavy chain CDR 1 described in (A) or (a) below,
a heavy chain CDR 2 described in (B) or (b) below, and
a heavy chain CDR 3 described in (C) or (c) below; and
a light chain variable region comprising
a light chain CDR 1 described in (D) or (d) below,
a light chain CDR 2 described in (E) or (e) below, and
a light chain CDR 3 described in (F) or (f) below.

(A) A heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 53,
(B) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 54,
(C) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 55;
(D) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 56,
(E) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 57, and
(F) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 58.

(a) A heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 41,
(b) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 42,
(c) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 43;
(d) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 44,
(e) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 45, and
(f) a light chain CDR 3 consisting of the amino acid sequence of

SEQ ID NO: 46.

The term "CDR" defined in relation to the antibodies of the present invention is an abbreviation for Complementarity Determining Region, and is also referred to as a complementarity determining region. CDRs are present in the variable region of immunoglobulin, and are deeply involved in the specific binding of an antibody to an antigen. Further, "light chain CDR" refers to a CDR that is present in the variable region of the light chains of immunoglobulin, and "heavy chain CDR" refers to a CDR that is present in the variable region of the heavy chains of immunoglobulin.

In addition, "variable region" refers to a region that includes the above-described CDR 1 to CDR 3 (hereinafter simply referred to as "CDRs 1 to 3"). Although the order of arrangement of the CDRs 1 to 3 is not particularly limited, preferably, CDR 1, CDR 2, and CDR 3 are arranged in that order or in the opposite order from N-terminus to C-terminus in a sequential manner or via other amino acid sequences called framework regions (FRs). Further, the "heavy chain variable region" is a region where the above-described heavy chain CDRs 1 to 3 are located, and the ”light chain variable region” is a region where the above-described light chain CDRs 1 to 3 are located.

As described above, the region other than the above-described CDRs 1 to 3 in the each variable region is called a framework region (FR). In particular, the region between the N-terminus and CDR 1 in each variable region is defined as FR 1, the region between CDR 1 and CDR 2 is defined as FR 2, the region between CDR 2 and CDR 3 is defined as FR 3, and the region between CDR3 and the C-terminus in each variable region is defined as FR 4.

The FRs also have a function as linker sequences for linking the CDRs 1 to 3 that are particularly important as the antigen recognition sequences. The FRs are the regions that contribute to the formation of the three-dimensional structure of

the entire variable region.

A preferable antibody of the first embodiment according to the present invention is an antibody comprising
a heavy chain variable region comprising
a heavy chain CDR 1 of (g) below, (m) below, or (a) above,
a heavy chain CDR 2 of (h) below, (n) below, or (b) above, and
a heavy chain CDR 3 of (i) below, (o) below, or (c) above; and
a light chain variable region comprising
a light chain CDR 1 of (j) below, (p) below, or (d) above,
a light chain CDR 2 of (k) below, (q) below, or (e) above, and
a light chain CDR 3 of (l) below, (r) below, or (f) above.

(g) A heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 17,
(h) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 18,
(i) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 19;
(j) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 20,
(k) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 21,
(l) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 22;

(m) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 29,
(n) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 30,
(o) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 31;
(p) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 32,
(q) a light chain CDR 2 consisting of the amino acid sequence of

SEQ ID NO: 33, and
(r) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 34.

The heavy chain CDR 3 described in (i) and the heavy chain CDR 3 described in (o) comprise identical amino acid sequences.

An antibody of a second embodiment of the present invention is an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (A)-(C) above or
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (a)-(c) above; and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (D)-(F) above or
a light chain variable region comprising the light chain CDRs 1 to 3 described in (d)-(f) above.

A more preferable antibody of the second embodiment is an antibody comprising any one of
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (g)-(i) above,
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (m)-(o) above, and
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (a)-(c) above; and
any one of
a light chain variable region comprising the light chain CDRs 1 to 3 described in (j)-(l) above,
a light chain variable region comprising the light chain CDRs 1 to 3 described in (p)-(r) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (d)-(f) above.

An antibody of a third embodiment of the present

invention is an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (A)-(C) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (D)-(F) above, or
an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (a)-(c) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (d)-(f) above.

A more preferable antibody of the third embodiment is an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (g)-(i) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (j)-(l) above;
an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (m)-(o) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (p)-(r) above; or
an antibody comprising
a heavy chain variable region comprising the heavy chain CDRs 1 to 3 described in (a)-(c) above, and
a light chain variable region comprising the light chain CDRs 1 to 3 described in (d)-(f) above.

The molecular structures of the antibodies of the present invention are not limited to that of immunoglobulin insofar as the antibodies have the above-described heavy and light chain variable regions. Examples of specific structures include molecular structures of $F(ab')_2$ that does not comprise the Fc region; Fab formed by papain digestion of immunoglobulin and composed of CH1 and CL domains as well as the heavy and light

chain variable regions; Fv that does not comprise the immunoglobulin constant region; and scFv, which is a single-chain Fv antibody.

The antibodies of the present invention may also be multivalent, in which the above molecular structures are combined. Such a multivalent antibody is formed by a technique of accumulating scFv constructs, as in an scFv-Fc construct formed by the combination of the Fc region and scFv construct described above; and a construct called a minibody, formed by the combination of CH3 domain of the constant region and the scFv construct described above. The term "multivalent" refers to the presence of multiple antigen-binding sites. In regard to the antibodies of the present invention, the term is used in the same meaning as the presence of multiple sites that bind to human XCR1.

The antibodies of the present invention may also have a human constant region in addition to the above-described heavy and light chain variable regions.

In immunoglobulin, the "constant region" of the heavy chains comprises domains called CH1, CH2, and CH3; and the "constant region" of the light chains comprises a domain called CL.

As described above, when the antibodies of the present invention have the constant region, it is preferable that the heavy chain variable region is linked to at least one of the CH1, CH2, and CH3 domains, and that the light chain variable region is linked to CL. Further, the heavy chain variable region is preferably directly linked to CH1.

The constant region of the antibodies of the present invention is a constant region derived from human immunoglobulin, preferably, a constant region derived from human immunoglobulin IgG. The subtype of human immunoglobulin IgG is not particularly limited, and may be suitably selected, for example, according to whether to impart ADCC activity, CDC activity, and the like described below to the antibodies.

The term "ADCC activity" is an abbreviation for

Antibody-Dependent Cellular Cytotoxicity activity. It is an activity in which cells such as NK cells expressing receptors specific for the antibody Fc region bind to the antibodies and damage cells present in the vicinity of the antibodies. Additionally, the term "CDC activity" is an abbreviation for Complement-Dependent Cytotoxicity activity. In the case of humans, the subtype of IgG having a high ADCC and/or CDC activity is IgG1, and the subtype of IgG having a low ADCC and/or CDC activity is IgG2 or IgG4.

Amino acid residues in the Fc region of the antibodies of the present invention may be mutated in order to induce a change in ADCC and/or CDC activity. Mutations to be introduced are not particularly limited, and known mutations may be introduced. For example, the following mutations may be introduced into the constant region of IgG1 for the purpose of increasing ADCC activity: S239D, I332E, S239D/I332E, S239D/I332E/A330L, and the like (Lazar GA, Dang W, Karki S, Vafa O, Peng JS, Hyun L, Chan C, Chung HS, Eivazi A, Yoder SC, Vielmetter J, Carmichael DF, Hayes RJ, Dahiyat BI, "Engineered antibody Fc variants with enhanced effector function," Proc Natl Acad Sci USA, 103: 4005-10 (2006)); and S298A, K334A, S298A/K334A, S298A/E333A/K334A, etc., (Shields RL, Namenuk AK, Hong K, Meng YG, Rae J, Briggs J, Xie D, Lai J, Stadlen A, Li B, Fox JA, Presta LG, "High resolution mapping of the binding site on human IgG1 for Fc gamma RI, Fc gamma RII, Fc gamma RIII, and FcRn and design of IgG1 variants with improved binding to the Fc gamma R," J Biol Chem, 276: 6591-604 (2001)).

Examples of mutations that increase CDC activity include S267E, H268F, S324T, S267E/H268F, S267E/S324T, H268F/S324T, S267E/H268F/S324T (Moore GL, Chen H, Karki S, Lazar GA, "Engineered Fc variant antibodies with enhanced ability to recruit complement and mediate effector functions," MAbs, 2:181-9 (2010)).

Additionally, for the purpose of lowering ADCC activity, known mutations may be introduced; for example, V234A/G237A (Cole

MS, Anasetti C, Tso JY, "Human IgG2 variants of chimeric anti-CD3 are nonmitogenic to T cells," J Immunol, 159:3613-21 (1997)), H268Q/V309L/A330S/P331S (An Z, Forrest G, Moore R, Cukan M, Haytko P, Huang L, Vitelli S, Zhao JZ, Lu P, Hua J, Gibson CR, Harvey BR, Montgomery D, Zaller D, Wang F, Strohl W, "IgG2m4, an engineered antibody isotype with reduced Fc function," MAbs, 1:572-9 (2009)), and the like.

The numbering of the above-described amino acids to be mutated is in accordance with the Eu numbering (see Sequences of proteins of immunological interest, NIH Publication No. 91-3242).

Chimeric Antibody

Among the antibodies of the present invention, an antibody in which the heavy and light chain variable regions comprise amino acid sequences derived from non-human species and the constant region comprises amino acid sequences derived from human is defined as a "chimeric antibody."

A first embodiment of the chimeric antibody of the present invention is a chimeric antibody comprising a heavy chain consisting of the amino acid sequence of SEQ ID NO: 13 and a light chain of SEQ ID NO: 14.

As shown in Table 5, the amino acid sequence of SEQ ID NO: 13 comprises the heavy chain CDRs 1 to 3 of SEQ ID NOs: 17 to 19 among the heavy chain CDRs 1 to 3 described above in the heavy chain variable region. Further, as shown in Table 5, the amino acid sequence of SEQ ID NO: 14 comprises the light chain CDRs 1 to 3 of SEQ ID NOs: 20 to 22 among the light chain CDRs 1 to 3 described above in the light chain variable region.

The chimeric antibody of the present invention comprises variants caused by mutations in the heavy chain consisting of the amino acid sequence of SEQ ID NO: 13 and/or the light chain consisting of the amino acid sequence of SEQ ID NO: 14, insofar as such mutations do not abolish the binding ability of the chimeric antibody to human XCR1.

Such variants in the heavy and light chains are

preferably obtained by introducing mutations into at least any one of FR 1 to FR 4 (hereinafter simply referred to as "FRs 1 to 4") of the variable region, or at least one site in the constant region of the respective amino acid sequences of SEQ ID NOs: 13 and 14.

The specific number of mutations introduced into the heavy and light chains is not particularly limited. Mutations are usually introduced to obtain a variant having 85% or higher identity, preferably 90% or higher identity, more preferably 95% or higher identity, and most preferably 99% or higher identity with the amino acid sequence before mutation.

The term "mutation" used herein includes substitution, deletion, insertion, and the like. A known method without specific limitation can be employed as a specific method for introducing mutations. For example, in the case of substitution, conservative substitution may be employed. The term "conservative substitution" refers to a substitution of an amino acid residue with another amino acid residue having a similar side chain.

For example, a substitution between amino acid residues with basic side chains such as lysine, arginine, and histidine corresponds to a conservative substitution. In addition, the following substitutions between the amino acid residues also correspond to conservative substitutions: substitutions between amino acid residues with acid side chains such as aspartic acid and glutamic acid; substitutions between amino acid residues with non-charged polar side chains such as glycine, asparagine, glutamine, serine, threonine, tyrosine, and cysteine; substitutions between amino acid residues with non-polar side chains such as alanine, valine, leucine, isoleucine, proline, phenylalanine, methionine, and tryptophan; substitutions between amino acid residues with β-branched side chains such as threonine, valine, and isoleucine; and substitutions between amino acid residues with aromatic side chains such as tyrosine, phenylalanine, tryptophan, and histidine.

Humanized Antibodies

Among the antibodies of the present invention, the antibody comprising the above-described CDRs 1 to 3 in the heavy and light chain variable regions, in which the FRs 1-4 comprise a human-derived amino acid sequence or a variant thereof, is defined as a "humanized antibody."

Such FRs comprising a human-derived amino acid sequence are not particularly limited, and may be determined based on a known technique.

Examples of such FRs include fully human framework regions or sub-regions, with FRs derived from human germline sequences being preferable. Reference may be suitably made to, for example, the NCBI website, which shows a list of currently known sequences of FRs as examples of fully human framework regions or sub-regions.

Non-limiting examples of the sequences of the human heavy chain variable region include VH1-18, VH1-2, VH1-24, VH1-3, VH1-45, VH1-46, VH1-58, VH1-69, VH1-8, VH2-26, VH2-5, VH2-70, VH3-11, VH3-13, VH3-15, VH3-16, VH3-20, VH3-21, VH3-23, VH3-30, VH3-33, VH3-35, VH3-38, VH3-43, VH3-48, VH3-49, VH3-53, VH3-64, VH3-66, VH3-7, VH3-72, VH3-73, VH3-74, VH3-9, VH4-28, VH4-31, VH4-34, VH4-39, VH4-4, VH4-59, VH4-61, VH5-51, VH6-1, and VH7-81.

Non-limiting examples of the sequences of the human light chain variable region include VL1-11, VL1-13, VL1-16, VL1-17, VL1-18, VL1-19, VL1-2, VL1-20, VL1-22, VL1-3, VL-4, VL1-5, VL1-7, VL1-9, VL2-1, VL2-11, VL2-13, VL2-14, VL2-15, VL2-17, VL2-19, VL2-6, VL2-7, VL-8, VL3-2, VL3-3, VL3-4, VL4-1, VL4-2, VL4-3, VL4-4, VL4-6, VL5-1, VL5-2, VL5-4, and VL5-6.

Fully human FRs are selected from these functional germline genes. Each of these FRs is usually different because of the modification of a limited number of amino acids. These FRs may be used in a combination with the CDRs described in the present specification. Non-limiting additional examples of human FRs to be used in combination with the above-described CDRs include KOL, NEWM, REI, EU, TUR, TEI, LAY, and POM. In regard to

the examples of these human FRs, reference may be made to the following documents: Kabat, et al., "Sequences of Proteins of Immunological Interest," US Department of Health and Human Services, NIH (1991) USA; Wu TT, Kabat EA, "An analysis of the sequences of the variable regions of Bence Jones proteins and myeloma light chains and their implications for antibody complementarity," J Exp Med, 132: 211-50 (1970); and the like.

A first embodiment of the humanized antibody of the present invention is a humanized antibody comprising a heavy chain variable region comprising the amino acid sequence of either SEQ ID NO: 60 or SEQ ID NO: 64, and a light chain variable region of either SEQ ID NO: 68 or SEQ ID NO: 72.

A more preferable embodiment is a humanized antibody comprising a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 60 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 68, or a humanized antibody comprising a heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 64 and a light chain variable region comprising the amino acid sequence of SEQ ID NO: 72.

As respectively shown in Tables 11-1 and 12-1, the amino acid sequences of SEQ ID NO: 60 and SEQ ID NO: 64 comprise the heavy chain CDRs 1 to 3 of SEQ ID NOs: 17 to 19 among the above-described heavy chain CDRs 1 to 3 in the heavy chain variable region. As respectively shown in Tables 13-1 and 14-1, the amino acid sequences of SEQ ID NO: 68 and SEQ ID NO: 72 comprise the light chain CDRs 1 to 3 of SEQ ID NOs: 20-22 among the above-described light chain CDRs 1 to 3 in the light chain variable region.

The humanized antibody of the present invention comprises variants caused by mutations in the heavy chain variable region comprising the amino acid sequence of SEQ ID NO: 60 or 64 and/or the light chain variable region comprising the amino acid sequence of SEQ ID NO: 68 or 72, insofar as such

mutation do not abolish the binding ability to human XCR1. Such variants in the heavy and light chain variable regions are preferably obtained by introducing mutations into the respective FRs 1 to 4.

The specific number of mutations into the heavy and light chain variable regions is not particularly limited. Mutations are usually introduced to obtain a variant having 85% or higher identity, preferably 90% or higher identity, more preferably 95% or higher identity, and most preferably 99% or higher identity with the amino acid sequence before mutation.

The term "mutation" used herein includes substitution, deletion, insertion, and the like. As is the case with the chimeric antibody described above, conservative substitution and the like may be employed as a specific method for introducing mutations.

The second embodiment of the humanized antibody of the present invention includes an antibody comprising a human constant region. Examples thereof include a humanized antibody comprising a heavy chain comprising the amino acid sequence of either SEQ ID NO: 59 or 63, and a light chain comprising the amino acid sequence of either SEQ ID NO: 67 or 71.

A more preferable embodiment is a humanized antibody comprising a heavy chain comprising the amino acid sequence of SEQ ID NO: 59 and a light chain comprising the amino acid sequence of SEQ ID NO: 67, or a humanized antibody comprising a heavy chain comprising the amino acid sequence of SEQ ID NO: 63 and a light chain comprising the amino acid sequence of SEQ ID NO: 71.

As shown in Table 11-1, the amino acid sequence of SEQ ID NO: 59 comprises an amino acid sequence corresponding to the heavy chain variable region of SEQ ID NO: 60, and therefore comprises the heavy chain CDRs 1 to 3 of SEQ ID NO: 17-19 among the heavy chain CDRs 1 to 3 described above. Further, as shown in Table 13-1, the amino acid sequence of SEQ ID NO: 67 comprises an

amino acid sequence corresponding to the light chain variable region of SEQ ID NO: 68, and therefore comprises the light chain CDRs 1 to 3 of SEQ ID NO: 20-22 among the light chain CDRs 1 to 3 described above.

The heavy and/or light chain described above comprises variants caused by mutations insofar as such mutations do not abolish the binding ability to human XCR1. Such variants in the heavy and light chains are preferably obtained by introducing mutations into the FRs 1 to 4 or the constant region.

The specific number of mutations into the heavy and light chains is not particularly limited. Mutations are usually introduced to obtain a variant having 85% or higher identity, preferably 90% or higher identity, more preferably 95% or higher identity, and most preferably 99% or higher identity with the amino acid sequence before mutation.

The term "mutation" used herein includes substitution, deletion, insertion, and the like. As is the case with the chimeric antibody described above, conservative substitution and the like may be employed as a specific method for introducing mutations.

Function of the Antibodies

The antibodies of the present invention bind to human XCR1. The meaning of the term "bind" used herein encompasses, at least, binding through hydrophobic bonds and the like as seen in the case of an interaction between proteins. In other words, antibodies that bind to human XCR1 at least by hydrophobic binding are sufficient as the antibodies of the present invention. Further, the antibodies of the present invention and human XCR1 may or may not be dissociated after binding.

The antibodies of the present invention preferably specifically bind to human XCR1. The term "specific binding" as used herein refers to specific binding to human XCR1, meaning that the antibodies preferentially bind to human XCR1 when human XCR1 is present concurrently with molecules other than human XCR1,

in particular, molecules having a structure similar to that of human XCR1, such as a homologue of human XCR1 or an orthologue of human XCR1.

That the antibodies of the present invention specifically bind to human XCR1 does not mean that the ability of binding to the above-described homologue or orthologue of human XCR1 is excluded.

The degree of binding of the antibodies of the present invention to human XCR1 can be evaluated by a reaction rate constant such as a Kd, Koff, or Kon value. A Kd value is a value obtained by dividing a Koff value by a Kon value.

The reaction rate constant between the antibodies of the present invention and human XCR1 is not particularly limited.

The antibodies of the present invention bind to the extracellular domain of human XCR1. Specifically, the antibodies bind to one or more of amino acid regions 1 to 31, 90 to 103, 168 to 190, and 251 to 267, which corresponds to the extracellular domain region of the amino acid sequence (SEQ ID NO: 91; Fig. 10) of the above-described NCBI Reference Sequence: NP_001019815.1 or NP_005274.1.

More preferably, in the amino acid sequence of SEQ ID NO: 91, the antibodies bind to at least three amino acids selected from the group consisting of the 8^{th}, 11^{th}, 12^{th}, 13^{th}, 14^{th}, 16^{th}, 17^{th}, 22^{nd}, 23^{rd}, 176^{th}, and 177^{th} amino acids.

Said “at least three amino acids” includes, for example, three or more amino acids, four or more amino acids, five or more amino acids, six or more amino acids, seven or more amino acids, eight or more amino acids, nine or more amino acids, ten or more amino acids, or eleven amino acids.

Note that the “epitope” in the present invention is also referred to as an “antigenic determinant,” and includes “linear epitope” and “discontinuous epitope.” A “linear epitope” is an epitope that is recognized by antibodies by the primary structure of the amino acid sequence, rather than by its

conformational structure. A “discontinuous epitope” is an epitope that is recognized by antibodies by the conformational structure of the amino acid sequence, based on the higher-order structure

Note that a person skilled in the art can determine the epitope of the antibodies of the present invention by suitably modifying the methods described in the Examples of the present invention. For example, the epitope can be determined by synthesizing a protein or peptide consisting of a desirable amino acid sequence that falls within the extracellular domain of the amino acid sequence of human XCR1 using a known method, and confirming the binding between the obtained protein or peptide and the antibody by a known method. Alternatively, the epitope can be determined by preparing a mutant by introducing an appropriate mutation to desired amino acids in the amino acid sequence of human XCR1 by a known method, and confirming whether the binding between the prepared mutant and the antibody is reduced.

As described above, because the antibodies of the present invention bind to human XCR1, the antibodies of the present invention also include an antibody that inhibits binding between human XCR1 and human XCL1. Human XCL1 is also referred to as human lymphotactin (Ltn) or human lymphotactin α (Ltn-α). Such an inhibitory activity is sometimes referred to as "neutralizing activity" induced by the antibodies of the present invention. Because human XCR1 is present on the cellular surface as a receptor protein *in vivo*, inhibition of binding between human XCR1 and XCL1 by the antibodies of the present invention is preferably performed on the cellular surface. It does not matter whether the antibodies of the present invention have an inhibitory activity against binding between human XCR1 and XCL2 insofar as the antibodies have activity to at least inhibit binding between human XCR1 and XCL1. Accordingly, the antibodies of the present invention also include an antibody that inhibits binding not only between human XCR1 and XCL1, but also between human XCR1 and XCL2.

Examples of preferred cells include cells associated with an immune system activated by the binding between human XCR1 and human XCL1, with dendritic cells being particularly preferable. In particular, as shown by the later-described examples, because the antibodies of the present invention specifically recognize BDCA3+ dendritic cells, which are dendritic cells expressing a significant amount of human XCR1 proteins, it is preferable that the antibodies have an effect of inhibiting binding between human XCR1 and human XCL1 on BDCA3+ dendritic cells.

Binding between human XCR1 and human XCL1 is inhibited by the antibodies of the present invention. Non-limiting examples of forms of such inhibition include:

(1) The antibodies of the present invention bind to XCR1 at a site to which human XCL1 originally should bind, causing a steric obstruction to binding to human XCL1, and resulting in the inhibition of binding between human XCR1 and human XCL1.

(2) The antibodies of the present invention bind to human XCR1, causing a change in the three-dimensional structure of human XCR1, which consequently causes a change in the structure of human XCR1 to which human XCL1 should bind, thus resulting in the inhibition of binding between human XCR1 and human XCL1.

(3) The antibodies of the present invention bind to XCR1, causing an internalization of the receptor, which leads to the inhibition of binding between human XCR1 and human XCL1.

The inhibitory activity of the antibodies of the present invention against binding between human XCR1 and human XCL1 is evaluated based on IC_{50} or IC_{90} values. These values can be obtained, for example, by performing a competitive inhibition experiment or the like of binding of human XCL1 to human XCR1, using cells those express human XCR1 proteins in the presence of the antibodies of the present invention. A known method may be

employed as a specific method of such a competitive inhibition experiment.

The antibodies of the present invention include an antibody that has an effect of inhibiting cell migration. The term "cell migration" refers to the phenomenon in which cells actively migrate as a result of external stimuli given to the cells and stimulus-induced activation of the intracellular signal transduction mechanism. Effects produced by the active cell migration vary depending on the functions of the cells. For example, in the case of cell migration of dendritic cells, such cell migration is a phenomenon that serves as one of the mechanisms in the immune system. In the present invention, inhibitory activity against cell migration is sometimes referred to as "neutralizing activity."

As described above, because the antibodies of the present invention suitably inhibit binding between human XCR1 and human XCL1 in dendritic cells, particularly BDCA3+ dendritic cells, the antibodies particularly preferably inhibit migration of dendritic cells, particularly BDCA3+ dendritic cells.

Human XCR1 is a seven-transmembrane G protein-coupled receptor. When human XCL1 binds to human XCR1, the three-dimensional structure of human XCR1 changes; and, as a result, a G protein coupled to the intracellular domain of human XCR1 is released, and a signal is transduced into the cells.

G proteins is prevented from release by the antibodies of the present invention inhibiting the binding between human XCR1 and XCL1 in accordance with the above-described forms (1), (2) or the like. As a result, no signal is transduced, thereby inhibiting the phenomena of cell migration.

Alternatively, the phenomena of cell migration may be inhibited as a result of a mechanism in which binding of the antibodies of the present invention to human XCR1 strengthens the bond between human XCR1 and G protein coupled to the intracellular domain of human XCR1, the release of G proteins

consequently does not occur, thereby inhibiting intracellular signal transduction.

The inhibitory activity of the antibodies of the present invention against cell migration of human cells is evaluated based on an IC_{50} or IC_{90} value. Specific values are not particularly limited. For example, an IC_{50} value is usually about 0.36 nM or less, preferably about 0.27 nM or less, and more preferably about 0.16 nM or less. For example, an IC_{90} value is usually about 2.38 nM or less, preferably about 1.52 nM or less, and more preferably about 0.86 nM or less.

The antibodies of the present invention include, as an embodiment, an antibody that has an effect of decreasing cytotoxic T lymphocyte (CTL) activity. The mechanism of decreasing the CTL activity is, for example, the antibodies of the present invention inhibiting the interaction between human XCR1 and human XCL1 in dendritic cells. Among the dendritic cells, the above-described BDCA3+ dendritic cells are preferable.

Method for Preparing the Antibodies of the Present Invention

The antibodies of the present invention can be prepared by a method comprising the following three steps, although it is not limited thereto.

(i) Step 1 of introducing a vector into the host to transform the host, the vector comprising a nucleic acid comprising a nucleotide sequence encoding the antibodies of the present invention;

(ii) Step 2 of culturing the transformed host obtained in step 1 and collecting a fraction containing antibodies that bind to human XCR1; and

(iii) Step 3 of isolating or purifying the above antibodies from the fraction obtained in step 2.

Step 1

The nucleic acid used in step 1 is a nucleic acid that encodes the antibodies of the present invention. The nucleotide sequence of the above nucleic acid can be determined using the *in silico* technique based on the amino acid sequence information of the antibodies of the present invention. At that time, it is preferable to determine the nucleotide sequence with reference to the codon frequency in the host employed in step 2. Specific examples of nucleotide sequences include the nucleotide sequence of SEQ ID NO: 3, 4, 7, 8, 11, 12, 15, 16, 61, 62, 65, 66, 69, 70, 73, or 74; or a variant thereof.

The above variant is preferably generated by introducing mutations (deletion, substitution, insertion, or the like) in the FR or constant region of the antibodies.

The specific number of mutations introduced into the variant is not particularly limited. Mutations are usually introduced to obtain a variant having 85% or higher identity, preferably 90% or higher identity, more preferably 95% or higher identity, and most preferably 99% or higher identity with the amino acid sequence before mutation.

Further, the above nucleic acid may comprise a nucleotide sequence that encodes a secretion signal peptide at the 5'-terminus. A specific nucleotide sequence encoding a secretion signal peptide is preferably a nucleotide sequence that effectively functions as a secretion signal peptide in the host cells employed in step 2. The term "secretion signal peptide" refers to a peptide comprising an amino acid sequence that acts as a recognition sequence for introducing proteins or peptides produced in the host into a pathway for secretion of the proteins or peptides to the outside of the host.

Examples of nucleotide sequences encoding a secretion signal peptide include:
ATGGGATTCAGCAGGATCTTTCTCTTCCTCCTGTCAGTAACTACAGGTGTCCACTCC (SEQ ID NO: 75),
ATGAAGTTGCCTGTTAGGCTGTTGGTGCTGCTGTTCTGGTTTCCTGCTTCCAACACT (SEQ ID NO: 76),

ATGGAATGGTCATGGGTCTTTCTGTTCTTTCTGAGTGTCACAACCGGGGTGCATAGC (SEQ ID NO: 77),

ATGGAATGGTCTTGGGTCTTTCTGTTCTTTCTGTCCGTCACTACCGGGGTCCACTCT (SEQ ID NO: 78),

ATGTCCGTGCCTACTCAGGTGCTGGGGCTGCTGCTGCTGTGGCTGACCGATGCTCGTTGC (SEQ ID NO: 79), and

ATGTCCGTGCCTACTCAGGTGCTGGGGCTGCTGCTGCTGTGGCTGACCGATGCTCGTTGT (SEQ ID NO: 80).

The vector used in step 1 comprises at least one of the above nucleic acids.

Such a vector may be one of the following vectors:

(I) a vector comprising a nucleic acid comprising a nucleotide sequence encoding at least one member selected from the group consisting of heavy chains, heavy chain variable region, and heavy chain CDRs 1 to 3 of the antibodies of the present invention;

(II) a vector comprising a nucleic acid comprising a nucleotide sequence encoding at least one member selected from the group consisting of light chains, light chain variable region, and light chain CDRs 1 to 3 of the antibodies of the present invention; or

(III) a vector comprising a nucleic acid comprising a nucleotide sequence encoding at least one member selected from the group consisting of heavy chains, heavy chain variable region, and heavy chain CDRs 1 to 3 of the antibodies of the present invention, and a nucleic acid comprising a nucleotide sequence encoding at least one member selected from the group consisting of light chains, light chain variable region, and light chain CDRs 1 to 3 the antibodies of the present invention.

The above vector may be a gene expression vector. The "gene expression vector" is a vector having a function to cause expression of the nucleotide sequence of the above nucleic acid. The gene expression vector may contain a promoter sequence, enhancer sequence, repressor sequence, insulator sequence, and the like to control the expression of the nucleotide sequence.

These sequences are not particularly limited insofar as they function in the above-described host.

The host used in step 1 is not particularly limited insofar as the above gene is expressed. Examples thereof include insect cells, eukaryotic cells, and mammalian cells. Of these cells, HEK cells, CHO cells, NS0 cells or SP2/O cells, which are mammalian cells, are particularly preferable in terms of more efficient expression of the nucleotide sequence that encodes antibodies.

A technique for introducing the above vector into the host in step 1 is not particularly limited. A known technique may be used. The vectors shown in (I) to (III) above may be introduced singly or in a combination of two or more into the host.

A host with the above vector can be produced by such a technique. The vector may be maintained as is in the host, or in such a manner that the nucleic acid comprising the nucleotide sequence encoding antibodies in the vector is incorporated into the genome of the host. The prepared host may be maintained using a known technique, and can be stored for a long period of time, if necessary.

Step 2

Step 2 is a step of culturing the above-described host obtained in step 1 and collecting a fraction containing the antibodies of the present invention, which bind to human XCR1. Culturing the host maintaining the above-described vector allows the host to express the nucleotide sequence encoding the antibodies of the present invention based on the nucleic acid in the vector, resulting in the production of the antibodies of the present invention. The produced antibodies are stored in the host or in the medium used for culturing the host.

In step 2, a known method may be employed as a technique for collecting a fraction containing the antibodies of the present invention. For example, for collecting a fraction

containing the antibodies of the present invention from the host, the host is disrupted by physical or chemical means, and the solution obtained by disruption is subjected to solid-liquid separation treatment, thereby obtaining a liquid fraction. The obtained liquid fraction may be used as the fraction containing the antibodies of the present invention.

On the other hand, for collecting a fraction containing the antibodies of the present invention from the medium used for culturing the host, the medium, i.e., the culture solution of the host obtained in step 1, is subjected to solid-liquid separation treatment, thereby obtaining a liquid fraction. The obtained liquid fraction may be used as the fraction containing the antibodies of the present invention.

In view of simplification of the isolation or purification step in the subsequent step 3, it is preferable to collect a fraction containing the antibodies of the present invention from the culture solution of the host.

The medium used for cultivation in step 2 is not particularly limited insofar as the medium allows the host to express the nucleotide sequence encoding the antibodies of the present invention, thereby producing the antibodies of the present invention. However, when collecting a fraction containing the antibodies of the present invention from the culture solution of the host as described above, it is preferable to employ serum-free medium in view of simplifying the isolation or purification step as much as possible in the subsequent step 3.

In regard to various conditions employed during cultivation of the host, such as container, temperature, time, host concentration in the medium, and culture conditions, the conditions used in a known method for producing antibodies may be employed.

Step 3

Step 3 is a step of isolating or purifying the antibodies of the present invention, which bind to human XCR1,

from the fraction obtained in step 2. The method for isolating and purifying the antibodies of the present invention is not particularly limited. A generally used method for isolating or purifying protein is widely applicable.

Medicinal Use of the Antibodies of the Present Invention

(1) Use as Therapeutic Agents for Immune Diseases

As described above, the antibodies of the present invention have an effect of inhibiting the phenomena of cell migration of dendritic cells associated with the immune system. Based on this effect, the antibodies of the present invention, in particular, the humanized antibody, have potential as an active ingredient of a pharmaceutical composition that is clinically applicable to human.

Diseases to which the antibodies of the present invention are applicable are explained below.

Applicable Diseases (Immune Diseases)

XCR1 is highly expressed in CD141+ dendritic cells in the case of humans, and in CD8α+ dendritic cells in the case of mice. These dendritic cells activate T-cells using the above-described antigen presentation method called cross-presentation (Bachem A, Güttler S, Hartung E, Ebstein F, Schaefer M, Tannert A, Salama A, Movassaghi K, Opitz C, Mages HW, Henn V, Kloetzel PM, Gurka S, Kroczek RA, "Superior antigen cross-presentation and XCR1 expression define human $CD11c^{+}CD141^{+}$ cells as homologues of mouse $CD8^{+}$ dendritic cells," J Exp Med, 207: 1273-1281 (2010)).

Further, because the source of production of XCL1, which is a ligand for human XCR1, comprises T-cells, in particular, CD8+ T-cells, the chemokine system in which XCL1-XCR1 is involved controls dendritic cell-induced activation of CD8+ T-cells (Crozat K, Guiton R, Contreras V, Feuillet V, Dutertre CA, Ventre E, Vu Manh TP, Baranek T, Storset AK, Marvel J, Boudinot P, Hosmalin A, Schwartz-Cornil I, Dalod M, "The XC chemokine

receptor 1 is a conserved selective marker of mammalian cells homologous to mouse CD8α+ dendritic cells," J Exp Med, 207: 1283-1292 (2010); and Dorner BG, Dorner MB, Zhou X, Opitz C, Mora A, Güttler S, Hutloff A, Mages HW, Ranke K, Schaefer M, Jack RS, Henn V, Kroczek RA, "Selective expression of the chemokine receptor XCR1 on cross-presenting dendritic cells determines cooperation with CD8+ T-cells," Immunity, 31: 823-833 (2009)).

As described above, the antibodies of the present invention include, as an embodiment, an antibody that exhibits an effect of inhibiting binding between human XCL1 and human XCR1 in dendritic cells, in particular, BDCA3+ dendritic cells. Accordingly, the antibodies of the present invention has potential as therapeutic agents for the treatment of immune diseases in which T-cells that are activated by migration of the dendritic cells are involved. In particular, the antibodies have potential as therapeutic agents for the treatment of diseases associated with the control of the activation of CD8+ T-cells.

As described above, the antibodies of the present invention include, as an embodiment, an antibody that exhibits an effect of decreasing CTL activity. CTL has a mechanism to activate the immune system by attacking cells or tissues. Various immunological diseases are known to have accelerated CTL activity; therefore, the antibodies of the present invention have potential as a therapeutic agent for the treatment of immunological diseases by decreasing CTL activity.

Non-limiting examples of such diseases include diabetes mellitus type 1, psoriasis, glomerulonephritis, autoimmune hepatitis, multiple sclerosis, ankylosing spondylitis, thyroiditis, graft rejection, delayed-type hypersensitivity, Crohn’s disease, dermatomyositis, polymyositis, inclusion body myositis, rheumatoid arthritis, inflammatory bowel disease, anterior uveitis, Wegener's granulomatosis, graft-versus-host disease, and Behçet’s disease (Kurts C, Robinson BW, Knolle PA, "Cross-priming in health and disease," Nat Rev Immunol, 10: 403-

414 (2010); Kehren J, Desvignes C, Krasteva M, Ducluzeau MT, Assossou O, Horand F, Hahne M, Kägi D, Kaiserlian D, Nicolas JF. "Cytotoxicity is mandatory for CD8(+) T cell-mediated contact hypersensitivity," J Exp. Med. 189: 779-786 (1999); Middel P, Thelen P, Blaschke S, Polzien F, Reich K, Blaschke V, Wrede A, Hummel KM, Gunawan B, Radzun HJ, "Expression of the T-cell chemoattractant chemokine lymphotactin in Crohn's disease," Am J Pathol, 159: 1751-1761 (2001); Sugihara T, Sekine C, Nakae T, Kohyama K, Harigai M, Iwakura Y, Matsumoto Y, Miyasaka N, Kohsaka H, "A new murine model to define the critical pathologic and therapeutic mediators of polymyositis," Arthritis Rheum, 56: 1304-1314 (2007); Wang CR, Liu MF, Huang YH, Chen HC, "Up-regulation of XCR1 expression in rheumatoid joints," Rheumatology (Oxford) 43: 569-573 (2004); Muroi E, Ogawa F, Shimizu K, Komura K, Hasegawa M, Fujimoto M, Sato S, "Elevation of serum lymphotactin levels in patients with systemic sclerosis," J Rheumatol, 35: 834-838 (2008); Torrence AE, Brabb T, Viney JL, Bielefeldt-Ohmann H, Treuting P, Seamons A, Drivdahl R, Zeng W, Maggio-Price L, "Serum biomarkers in a mouse model of bacterial-induced inflammatory bowel disease," Inflamm Bowel Dis, 14: 480-490 (2008); Yeh PT, Lin FA, Lin CP, Yang CM, Chen MS, Yang CH, "Expressions of lymphotactin and its receptor, XCR, in Lewis rats with experimental autoimmune anterior uveitis," Graefes Arch Clin Exp Ophthalmol, 248: 1737-1747 (2010); Blaschke S, Brandt P, Wessels JT, Müller GA, "Expression and function of the C-class chemokine lymphotactin (XCL1) in Wegener's granulomatosis," J Rheumatol, 36: 2491-2500 (2009); Asuka H, Okazaki Y, Kawakami Y, Hirakata M, Inoko H, Ikeda Y, Kuwana M, "Autoreactive CD8+ cytotoxic T lymphocytes to major histocompatibility complex class I chain-related gene A in patients with Behçet's disease," Arthritis Rheum, 50: 3658-3662 (2004); Serody JS, Burkett SE, Panoskaltsis-Mortari A, Ng-Cashin J, McMahon E, Matsushima GK, Lira SA, Cook DN, Blazar BR, "T-lymphocyte production of macrophage inflammatory protein-1alpha is critical to the recruitment of CD8(+) T cells to the liver, lung, and spleen

during graft-versus-host disease," Blood, 96: 2973-2980 (2000); Sugihara T, Sekine C, Nakae T, Kohyama K, Harigai M, Iwakura Y, Matsumoto Y, Miyasaka N, Kohsaka H, "A new murine model to define the clinical pathologic and therapeutic mediators of polymyositis," Arthritis & Rheumatism. 56: 1304-1314 (2007)); and Dalakas MC, "Review: An update on inflammatory and autoimmune myopathies," Neuropathol Appl Neurobiol, 37: 226-242 (2011).

It was also revealed that the antibody (anti-human XCR1 mouse monoclonal antibody (5G7)) of the present invention significantly inhibits the DTH reaction in the later-described experiment that used a mouse model of delayed-type hypersensitivity (hereinafter sometimes referred to as "DTH"). As described above, delayed-type hypersensitivity is a disease known as one of the immune diseases in which CD8+ T-cells that are activated by migration of the dendritic cells are involved. The fact that the antibodies of the present invention are effective in the treatment of delayed-type hypersensitivity provides evidence that the antibodies of the present invention have activity of inhibiting cell migration, in particular, dendritic cell migration, because the antibodies of the present invention affect CD8+ T-cells.

Further, in addition to delayed-type hypersensitivity, atopic dermatitis and contact dermatitis are also known as immune diseases of the skin in which the DTH reaction is involved (Fabrizi G, Romano A, Vultaggio P, Bellegrandi S, Paganelli R, Venuti A, "Heterogeneity of atopic dermatitis defined by the immune response to inhalant and food allergy," Eur J Dermatol, 9: 380-384 (1999); and Fonacier LS, Dreskin SC, Leung DYM, "Allergic skin diseases," J Allergy Clin Immunol, 125: S138-149 (2010)).

Based on the above, the antibodies of the present invention have potential as therapeutic agents for the treatment of immune diseases of the skin such as atopic dermatitis or contact dermatitis.

It has been pointed out that the activation of CD8+ T-cells may also be involved in the DTH reaction (Mody CH, Pain III R, Jackson C, Chen G-H, Toews GB, "CD8 Cells play a critical role in delayed-type hypersensitivity to intact Cryptococcus neoformans," J Immunol, 152: 3970-3979 (1994), etc.).

Invasion of CD8+ T-cells into the epidermis is observed in psoriasis, which is an autoimmune skin disease affecting a large number of patients, in particular, in chronic psoriasis lesions. These cells are considered to be the main effector cells that cause psoriasis lesions (Gudjonsson JE, Johnston A, Sigmundsdottir H, Valdimarsson H, "Immunopathogenic mechanisms in psoriasis," Clin Exp Immunol, 135: 1-8 (2004)).

Based on the above, the antibodies of the present invention have potential as therapeutic agents for the treatment of immune diseases of the skin in which the activation of CD8+ T-cells is involved.

In addition to delayed-type hypersensitivity, atopic dermatitis, and contact dermatitis, non-limiting examples of immune diseases of the skin in which CD8+ T-cells are involved also include dermatomyositis, polymyositis, inclusion body myositis, psoriasis, parapsoriasis, autoimmune blistering diseases (e.g., pemphigus, pemphigoid, and acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin diseases associated with collagenosis (e.g., systemic lupus erythematosus, Sjögren syndrome, and mixed connective tissue disease), skin diseases associated with Addison's disease, skin diseases associated with graft-versus-host disease (GVHD), eczema, and urticaria.

Herein below, the relationship between the antibodies of the present invention and various immune diseases (multiple sclerosis, human type 1 diabetes mellitus, glomerulonephritis, autoimmune hepatitis, thyroiditis, graft-versus-host disease, dermatomyositis, polymyositis, and inclusion body myositis) is

described. Diseases to which the antibodies of the present invention are effective are not limited to the following specific diseases.

Multiple Sclerosis

Invasion of CD8+ T-cells in addition to CD4+ T-cells into the central lesions in multiple sclerosis in humans has been recently reported. Further, it has been reported that, in the experiment using mice, implantation of CD8+ T-cells activated by antigen derived from the myelin sheath in central nerves induce experimental autoimmune encephalomeningitis, which is a model of human multiple sclerosis. Compared to the conventional model, the above-mentioned model more closely mimics the pathology of human multiple sclerosis (repeated exacerbations and remissions, significant demyelination, invasion of many CD8+ T-cells and macrophages/microglial cells in demyelinated lesions). As described above, it has been suggested that CD8+ T-cells play an important role in human multiple sclerosis and its mouse model (Friese MA, Fugger L, "Autoreactive CD8+ cells in multiple sclerosis: a new target for therapy?" Brain, 128: 1747-1763 (2005)).

Accordingly, the antibodies of the present invention that control the activation of CD8+ T-cells has potential as therapeutic agents for the treatment of multiple sclerosis.

Human Type 1 Diabetes Mellitus

Non-obese diabetic (NOD) mice representing a model of human type 1 diabetes mellitus have shown that depletion of CD8+ T-cells results in inhibition of the onset of diabetes mellitus (Wang B, Gonzales A, Benoist C, Mathis D, "The role CD8+ T-cells in the initiation of insulin-dependent diabetes mellitus," Eur J Immunol, 26: 1762-1769 (1996)). This suggests that CD8+ T-cells are also involved in the development of the pathology of diabetes mellitus type 1.

Accordingly, the antibodies of the present invention

that control the activation of CD8+ T-cells has potential as therapeutic agents for the treatment of human type 1 diabetes mellitus.

Glomerulonephritis

In a mouse model of glomerulonephritis, it has been shown that CD8+ T-cells are involved in the process of the formation of renal lesions (Heymann F, Meyer-Schwesinger C, Hamilton-Williams EE, Hammerich L, Panzer W, Kaden S, Quaggin SE, Floege J, Gröne H-J, Kurts C, "Kidney dendritic cell activation is required for progression of renal disease on a mouse model of glomerular injury," J Clin Invest, 119: 1286-1297 (2009)). Invasion of many CD8+ T-cells into the kidney is observed in patients with severe autoimmune lupus nephritis. The correlation between the number of these CD8+ T-cells and an increase in the renal activity score and the serum creatinine level, which indicate aggravation of the renal function, has been reported (Couzi L, Merville P, Deminière C, Moreau J-F, Combe C, Pellegrin J-L, Viallard J-F, Blanco P, "Predominance of CD8+ T lymphocytes among periglomerular infiltrating cells and link to the prognosis of class III and class IV lupus nephritis," Arthritis Rheum, 56: 2362-2370 (2007)). As described above, CD8+ T-cells are considered to be involved in the onset of autoimmune glomerulonephritis or progression of the pathology thereof in human and mouse models.

Accordingly, the antibodies of the present invention, which control the activation of CD8+ T-cells, have potential as therapeutic agents for the treatment of glomerulonephritis.

Autoimmune Hepatitis

It has been suggested that infection with hepatitis C virus (HCV) is involved in the process of the development of autoimmune hepatitis. It has also been suggested that CD8+ CTLs induced with respect to HCV are involved in the development of autoimmune hepatitis by eliminating HCV and damaging the infected

liver cells (Kammer AR, van der Burg SH, Grabscheid B, Hunziker IP, Kwappenberg KMC, Reichen J, Melief CJM, Cerny A, "Molecular mimicity of human cytochrome P450 by hepatitis C virus at the level of cytotoxic T cell recognition," J Exp Med, 190: 169-175 (1999)).

Accordingly, the antibodies of the present invention, which control the activation of CD8+ T-cells, have potential as therapeutic agents for the treatment of autoimmune hepatitis.

Thyroiditis

CD8+ CTLs are known to be involved in the development of experimental autoimmune thyroiditis (EAT), which is a mouse model of human thyroiditis (for example, Hashimoto's disease). It has been reported that the mice in the model show lesions similar to human thyroiditis (antithyroglobulin antibodies are found in the peripheral blood, and the invasion of CD8+ T-cells and CD4+ T-cells into the thyroid gland is observed). As described above, it has been suggested that CD8+ T-cells are involved in the development of thyroiditis in human and mouse models (Brazillet M-P, Batteux F, Abehsira-Amar O, Nicoletti F, Charreire J, "Induction of experimental autoimmune thyroiditis by heat-denatured porcine thyroglobulin: a Tc1-mediated disease," Eur J Immunol, 29: 1342-1352 (1999)).

Accordingly, the antibodies of the present invention, which control the activation of CD8+ T-cells, have potential as therapeutic agents for the treatment of human thyroiditis.

Rheumatoid arthritis

As described in the Examples below, 5G7, which is one of the antibodies of the present invention, exhibits a significant effect in treating rheumatoid arthritis in the experiment of DTH induced by *Mycobacterium butyricum*. Therefore, the antibodies of the present invention have potential as a therapeutic agent for the treatment of rheumatoid arthritis.

Graft Rejection

CD8+ T-cells play an important role in the graft rejection after human organ transplantation. A graft is rejected by CD8+ T-cells in the host that recognizes MHC class I being expressed in the cells in the graft. Further, invasion of many CD8+ T-cells into the kidney has been reported in renal transplant patients experiencing rejection. As described above, it has been suggested that CD8+ T-cells also play a central role in the graft rejection after human organ transplantation (Bueno V, Pestana JOM, "The role of CD8+ T-cells during allograft rejection," Braz J Med Biol Res, 35: 1247-1258 (2002)).

Accordingly, the antibodies of the present invention, which control the activation of CD8+ T-cells, have potential as therapeutic agents for the treatment of graft-versus-host disease.

Dermatomyositis, Polymyositis, and Inclusion Body Myositis

When lymphocytes that invade the lesion site of patients with dermatomyositis and polymyositis were established as cell lines, CD8+ T-cell lines showed cytotoxicity against their own cultured muscle cells. This indicates that muscle cell damage in the patients with the above-described myositis is caused by CD8+ T-cells with antigen-specific cytotoxicity (Hohlfeld R, Engel AG, "Coculture with autologous myotubes of cytotoxic T cells isolated from muscle in inflammatory myopathies," Ann Neurol, 29: 498-507 (1991)). Further, invasion of CD8+ T-cells into the lesion site has been observed in the patients with inclusion body myositis (Dalakas MC, "Review: An update on inflammatory and autoimmune myopathies," Neuropathol Appl Neurobiol, 37: 226-242 (2011)). Accordingly, the antibodies of the present invention, which control the activation of CD8+ T-cells, have potential as therapeutic agents for the treatment of dermatomyositis, polymyositis, or inclusion body myositis.

As described above, because the antibodies of the present invention have potential as therapeutic agents for the

treatment of immune diseases, in particular, immune diseases of the skin, the present invention provides a pharmaceutical composition comprising the antibodies of the present invention.

Such a pharmaceutical composition has potential as a therapeutic agent for the treatment of immune disease, for the purpose of treating immune diseases, in particular, immune diseases of the skin.

The term "treatment" used herein means attainment of desired pharmacological and/or physiological effects. The effects include an effect of partially or completely curing disease and/or adverse effects caused by the disease (pathologies and symptoms). The above effects also include an effect of inhibiting or delaying the progression of the disease and/or adverse effects caused by the disease (pathologies and symptoms); an effect of alleviating pathologies and symptoms (i.e., ameliorating the disease or symptoms, or causing reversal of the progression of symptoms); and an effect of preventing recurrence of the disease. The above effects also include an effect of partially or completely preventing the onset of the disease and/or adverse effects caused by the disease (pathologies and symptoms) in the individuals who may possess a predisposition to the disease and/or adverse effects caused by the disease (pathologies and symptoms) but who have not been diagnosed as having the predisposition. Accordingly, the term "treatment" also means "relief," "prevention of recurrence," and "prevention of disease."

In the present invention, a pharmaceutical composition comprising the antibodies of the present invention can be suitably used for the treatment of human immune diseases, in particular, immune diseases of the skin. It is understood that the above pharmaceutical composition is capable of providing, for example, an effect of partially or completely curing various symptoms of immune diseases; an effect of partially or completely inhibiting various symptoms of immune diseases (i.e., inhibiting or delaying the progression); an effect of alleviating various

symptoms of immune diseases (i.e., ameliorating the disease or symptoms, or causing reversal of the progression of symptoms); or an effect of preventing recurrence of various symptoms of immune diseases.

Specific examples of target diseases are as described above, with immune diseases of the skin being preferable.

The content of the antibodies of the present invention in the above pharmaceutical composition is not particularly limited insofar as the pharmaceutical composition comprises an effective amount of the antibodies of the present invention. The content can be suitably determined, for example, in such a manner the antibodies of the present invention are contained in the pharmaceutical composition in an amount of 0.001 to 99.99 wt% relative to 100 wt% of the composition, by taking into account the type of the target immune disease, dosage form, administration method, and the like.

The term "effective amount" used herein refers to an amount that allows the antibodies of the present invention to demonstrate an effect of inhibiting cell migration of dendritic cells, or an amount that allows the antibodies to demonstrate the above-described desired pharmacological and/or physiological effects (treatment effect for immune diseases).

Pharmaceutically acceptable carriers or additives may be added in combination with the antibodies of the present invention to the pharmaceutical composition. The term "pharmaceutically acceptable carriers or additives" used herein refers to optional carriers, diluents, excipients, suspending agents, lubricants, adjuvants, vehicles, delivery systems, emulsifiers, disintegrants, absorbents, preservatives, surfactants, colorants, fragrances, or sweeteners. Known carriers or additives may be used.

Non-limiting examples of dosage forms of the pharmaceutical composition include tablets, syrups, liniments, injections, and infusions, with injections or infusions being

preferable. Such injections and infusions may be in aqueous, non-aqueous, or suspension form. Additionally, the pharmaceutical composition may have a dosage form that is prepared just before administration.

The pharmaceutical composition of the present invention, specifically, a therapeutic agent for an immune disease, has potential in methods of treating an immune disease, comprising a step of administering the composition to a human subject with an immune disease, in particular, an immune disease of the skin. As described above, the pharmaceutical composition also has potential in methods of preventing an immune disease, comprising administering the composition to a human subject who has not developed pathologies or symptoms of an immune disease, in particular, an immune disease of the skin, but who may possess a predisposition to the immune disease.

The dosage amount and administration method of the pharmaceutical composition (therapeutic agent for immune diseases) can be suitably determined within a range of 0.001 to 100 mg/kg/day, according to the type of immune disease, the human subject's sex, race, age, and general condition, the severity of the disease, and the like.

The antibodies of the present invention may be administered at the above-described dosage once a day, or in divided dosage several times per day. Further, in the range that the antibodies have a treatment effect on the above-described diseases, the administration interval may be every day, every other day, every week, every other week, every 2 to 3 weeks, every month, or every 2 to 3 months. Non-limiting examples of administration methods include oral, intramuscular, intravenous, intraarterial, intrathecal, intradermal, intraperitoneal, intranasal, intrapulmonary, intraocular, intravaginal, intracervical, intrarectal, and subcutaneous administrations.

(2) Application as Immunotoxin

The antibodies of the present invention may have been

conjugated to cytotoxic molecules. Because such antibodies bind to human XCR1 protein that is expressed in a significant amount in dendritic cells associated with the immune system, the antibodies may be used as immunotoxins that target dendritic cells.

The term "cytotoxic molecules" used herein refers to molecules that demonstrate effects, such as apoptosis and/or necrosis, which cause the death of cells.

Examples of such molecules include saporin, ricin, Pseudomonas exotoxin, diphtheria toxin, and chemotherapeutic agents. Binding between the antibody and a toxic substance may be performed by a method used for the preparation of conventional immunotoxins.

(3) Other Applications of the Antibodies of the Present Invention

Because the antibodies of the present invention also include, as an embodiment, an antibody that binds to XCR1 that is expressed in a significant amount in dendritic cells, the antibodies has potential in a method for detecting dendritic cells. In this case, it is preferable to label the antibodies of the present invention for the use. The term "label" used herein refers to binding the antibodies to labeled molecules such as fluorescent molecules, luminescent molecules, chromogenic molecules and radioisotope molecules.

The binding pattern is not limited insofar as the bond is not dissociated in a detection step. A known method may be employed as a specific detection method. For example, a flow cytometry technique may be employed.

Further, the antibodies of the present invention may also be suitably applicable in methods of isolating and/or removing dendritic cells after the detection of dendritic cells. Known methods may also be employed for these methods. For example, a known cell-sorting device may be suitably used in a combination with a flow cytometry technique.

The present invention relates to the antibodies explained above, and widely encompasses the inventions of the embodiments described below.

Item 1

An antibody binding to human XCR1, wherein the antibody binds to linear or discontinuous epitopes which comprise at least three amino acids selected from the group consisting of the 8th, 11th, 12th, 13th, 14th, 16th, 17th, 22nd, 23rd, 176th, and 177th amino acids in the amino acid sequence of SEQ ID NO: 91.

Item 2

The antibody according to above item 1 , wherein the antibody is:

the antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (g) to (i) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (j) to (l) below;

the antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (m) to (o) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (p) to (r) below; or

the antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (a) to (c) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (d) to (f) below:

(a) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 41,

(b) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 42,

(c) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 43;

(d) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 44,

(e) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 45, and

(f) a light chain CDR 3 consisting of the amino acid sequence of

SEQ ID NO: 46;
(g) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 17,
(h) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 18,
(i) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 19;
(j) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 20,
(k) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 21,
(l) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 22;
(m) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 29,
(n) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 30,
(o) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 31;
(p) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 32,
(q) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 33, and
(r) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 34.

Item 3

The antibody according to above item 1 or 2, wherein the antibody comprises a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60 or 64, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 68 or 72.

Item 4

The antibody according to any one of above items 1 to 3,

wherein the antibody comprises a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 68.

Item 5

The antibody according to any one of above items 1 to 3, wherein the antibody comprises a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 64, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 72.

Item 6

The antibody according to any one of above items 1 to 5, wherein the antibody comprises a human constant region.

Item 7

The antibody according to any one of above items 1 to 6, wherein the antibody comprises a heavy chain comprising an amino acid sequence of SEQ ID NO: 59, and a light chain comprising an amino acid sequence of SEQ ID NO: 67.

Item 8

The antibody according to any one of above items 1 to 6, wherein the antibody comprises a heavy chain comprising an amino acid sequence of SEQ ID NO: 63, and a light chain comprising an amino acid sequence of SEQ ID NO: 71.

Item 9

The antibody according to any one of above items 1 to 8 comprising an Fc region, wherein the Fc region is mutated to induce a change in ADCC activity.

Item 10

The antibody according to above item 9, wherein the Fc

region is mutated to lower ADCC activity.

Item 11

The antibody according to any one of above items 1 to 10, wherein the antibody is conjugated to a cytotoxic molecule.

Item 12

The antibody according to any one of above items 1 to 11, wherein the antibody inhibits interaction between human XCR1 and human XCL1.

Item 13

The antibody according to any one of above items 1 to 12, wherein the antibody inhibits cell migration of dendritic cells.

Item 14

The antibody according to any one of above items 1 to 13, wherein the antibody suppresses the activity of cytotoxic T lymphocytes.

Item 15

A pharmaceutical composition comprising the antibody according to any one of above items 1 to 14 and a pharmaceutically acceptable carrier or additive.

Item 16

The pharmaceutical composition according to above item 15, wherein the pharmaceutical composition is a therapeutic agent for an immune disease.

Item 17

The pharmaceutical composition according to above item 16, wherein the immune disease is an immune disease of the skin.

Item 18

The pharmaceutical composition according to above item 17, wherein the immune disease of the skin is psoriasis, parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin disease associated with collagenosis (systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin disease associated with Addison's disease, skin disease associated with graft-versus-host disease (GVHD), eczema, or urticaria.

Item 19

The pharmaceutical composition according to above item 17, wherein the immune disease of the skin is psoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, or inclusion body myositis.

Item 20

The pharmaceutical composition according to above item 17~~19~~, wherein the immune disease of the skin is atopic dermatitis or contact dermatitis.

Item 21

The pharmaceutical composition according to above item 16, wherein the immune disease is thyroiditis, rheumatoid arthritis, type 1 diabetes, or multiple sclerosis.

Item 22

A nucleic acid comprising a nucleotide sequence encoding the antibody according to any one of above items 1 to 14.

Item 23

A method of treating an immune disease comprising administering an effective amount of the antibody according to any one of above items 1 to 14 or the pharmaceutical composition according to above item 15 to a human affected by an immune disease.

Item 24

The method according to above item 23, wherein the immune disease is an immune disease of the skin.

Item 25

The method according to above item 24, wherein the immune disease of the skin is psoriasis, parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin disease associated with collagenosis (systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin disease associated with Addison's disease, skin disease associated with graft-versus-host disease (GVHD), eczema, or urticaria.

Item 26

The method according to above item 24, wherein the immune disease of the skin is psoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, or inclusion body myositis.

Item 27

The method according to above item 23, wherein the immune disease is thyroiditis, rheumatoid arthritis, type 1 diabetes, or multiple sclerosis.

The present invention also encompasses the embodiments

described below

Item 1-A

An antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (g) to (i) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (j) to (l) below;

an antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (m) to (o) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (p) to (r) below; or

an antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (a) to (c) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (d) to (f) below:

(a) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 41,

(b) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 42,

(c) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 43;

(d) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 44,

(e) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 45, and

(f) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 46;

(g) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 17,

(h) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 18,

(i) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 19;

(j) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 20,

(k) a light chain CDR 2 consisting of the amino acid sequence of

SEQ ID NO: 21,

(l) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 22;

(m) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 29,

(n) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 30,

(o) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 31;

(p) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 32,

(q) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 33, and

(r) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 34.

Item 2-A

The antibody according to above item 1-A, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60 or 64, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 68 or 72.

Item 3-A

The antibody according to above item 1-A or 2-A, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 68.

Item 4-A

The antibody according to above item 1-A or 2-A, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 64, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 72.

Item 5-A

The antibody according to any one of above items 1-A to 4-A, comprising a human constant region.

Item 6-A

The antibody according to any one of above items 1-A to 5-A, comprising a heavy chain comprising an amino acid sequence of SEQ ID NO: 59, and a light chain comprising an amino acid sequence of SEQ ID NO: 67.

Item 7-A

The antibody according to any one of above items 1-A to 5-A, comprising a heavy chain comprising an amino acid sequence of SEQ ID NO: 63, and a light chain comprising an amino acid sequence of SEQ ID NO: 71.

Item 8-A

The antibody according to any one of above items 1-A to 7-A comprising an Fc region, wherein the Fc region is mutated to induce a change in ADCC activity.

Item 9-A

The antibody according to above item 8-A, wherein the Fc region is mutated to lower ADCC activity.

Item 10-A

The antibody according to any one of above items 1-A to 9-A, wherein the antibody inhibits interaction between human XCR1 and human XCL1.

Item 11-A

The antibody according to any one of above items 1-A to 10-A, wherein the antibody inhibits cell migration of dendritic cells.

Item 12-A

A pharmaceutical composition comprising the antibody according to any one of above items 1-A to 11-A and a pharmaceutically acceptable carrier or additive.

Item 13-A

The pharmaceutical composition according to above item 12-A, wherein the pharmaceutical composition is a therapeutic agent for an immune disease.

Item 14-A

The pharmaceutical composition according to above item 13-A, wherein the immune disease is an immune disease of the skin.

Item 15-A

The pharmaceutical composition according to above item 14-A, wherein the immune disease of the skin is psoriasis, parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin disease associated with collagenosis (systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin disease associated with Addison's disease, skin disease associated with graft-versus-host disease (GVHD), eczema, or urticaria.

Item 16-A

The pharmaceutical composition according to above item14-A, wherein the immune disease of the skin is psoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, or inclusion body myositis.

Item 17-A

The pharmaceutical composition according to above item

14-A, wherein the immune disease of the skin is atopic dermatitis or contact dermatitis.

Item 18-A

A nucleic acid comprising a nucleotide sequence encoding the antibody according to any one of above items 1-A to 11-A.

Item 19-A

An immune disease treatment method comprising a step of administering an effective amount of the antibody according to any one of above items 1-A to 11-A to a human affected by an immune disease.

The present invention further encompasses the embodiments described below.

Item 1-B

An antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (g) to (i) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (j) to (l) below;

an antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (m) to (o) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (p) to (r) below; or

an antibody comprising a heavy chain variable region comprising heavy chain CDRs 1 to 3 described in (a) to (c) below and a light chain variable region comprising light chain CDRs 1 to 3 described in (d) to (f) below:

(a) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 41,

(b) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 42,

(c) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 43;

(d) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 44,

(e) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 45, and

(f) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 46;

(g) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 17,

(h) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 18,

(i) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 19;

(j) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 20,

(k) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 21,

(l) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 22;

(m) a heavy chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 29,

(n) a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 30,

(o) a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 31;

(p) a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO: 32,

(q) a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO: 33, and

(r) a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO: 34.

Item 2-B

The antibody according to above item 1-B, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60 or 64, and a light chain variable region comprising

an amino acid sequence of SEQ ID NO: 68 or 72.

Item 3-B

The antibody according to above item 1-B or 2-B, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 60, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 68.

Item 4-B

The antibody according to above item 1-B or 2-B, comprising a heavy chain variable region comprising an amino acid sequence of SEQ ID NO: 64, and a light chain variable region comprising an amino acid sequence of SEQ ID NO: 72.

Item 5-B

The antibody according to any one of above items 1-B to 4-B, comprising a human constant region.

Item 6-B

The antibody according to any one of above items 1-B to 5-B, comprising a heavy chain comprising an amino acid sequence of SEQ ID NO: 59, and a light chain comprising an amino acid sequence of SEQ ID NO: 67.

Item 7-B

The antibody according to any one of above items 1-B to 5-B, comprising a heavy chain comprising an amino acid sequence of SEQ ID NO: 63, and a light chain comprising an amino acid sequence of SEQ ID NO: 71.

Item 8-B

The antibody according to any one of above items 1-B to 7-B comprising an Fc region, wherein the Fc region is mutated to induce a change in ADCC activity.

Item 9-B

The antibody according to above item 8-B, wherein the Fc region is mutated to lower ADCC activity.

Item 10-B

The antibody according to any one of above items 1-B to 9-B, wherein the antibody is conjugated to a cytotoxic molecule.

Item 11-B

The antibody according to any one of above items 1-B to 10-B, wherein the antibody inhibits interaction between human XCR1 and human XCL1.

Item 12-B

The antibody according to any one of above items 1-B to 11-B, wherein the antibody inhibits cell migration of dendritic cells.

Item 13-B

A pharmaceutical composition comprising the antibody according to any one of above items 1-B to 12-B and a pharmaceutically acceptable carrier or additive.

Item 14-B

The pharmaceutical composition according to above item 13-B, wherein the pharmaceutical composition is a therapeutic agent for an immune disease.

Item 15-B

The pharmaceutical composition according to above item 14-B, wherein the immune disease is an immune disease of the skin.

Item 16-B

The pharmaceutical composition according to above item 15-B, wherein the immune disease of the skin is psoriasis,

parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin disease associated with collagenosis (systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin disease associated with Addison's disease, skin disease associated with graft-versus-host disease (GVHD), eczema, or urticaria.

Item 17-B

The pharmaceutical composition according to above item15-B, wherein the immune disease of the skin is psoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, or inclusion body myositis.

Item 18-B

The pharmaceutical composition according to above item 15-B, wherein the immune disease of the skin is atopic dermatitis or contact dermatitis.

Item 19-B

A nucleic acid comprising a nucleotide sequence encoding the antibody according to any one of above items 1-B to 12-B.

Item 20-B

A method of treating an immune disease comprising administering an effective amount of the antibody according to any one of above items 1-B to 12-B or the pharmaceutical composition according to above item 13-B to a human affected by an immune disease.

Item 21-B

The method according to above item 20-B, wherein the immune disease is an immune disease of the skin.

Item 22-B

The method according to above item 21-B, wherein the immune disease of the skin is psoriasis, parapsoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, inclusion body myositis, autoimmune blistering disease (pemphigus, pemphigoid, or acquired epidermolysis bullosa), pustulosis, herpes gestationis, linear IgA bullous dermatosis, alopecia areata, vitiligo vulgaris, skin disease associated with collagenosis (systemic lupus erythematosus, Sjögren syndrome, or mixed connective tissue disease), skin disease associated with Addison's disease, skin disease associated with graft-versus-host disease (GVHD), eczema, or urticaria.

Item 23-B

The method according to above item 21-B, wherein the immune disease of the skin is psoriasis, atopic dermatitis, contact dermatitis, dermatomyositis, polymyositis, or inclusion body myositis.

Item 24-B

The method according to above item 21-B, wherein the immune disease of the skin is atopic dermatitis or contact dermatitis.

Examples

Herein below, the present invention is described in more detail based on Examples. Needless to say, the present invention is not limited to the Examples.

Example 1

(1) Preparation of Mouse Anti-Human XCR1 Monoclonal Antibodies

To obtain monoclonal antibodies against human XCR1,

membrane fraction of human XCR1-expressing B300.19 cells was immunized to XCR1 knockout mice. The membrane fraction was prepared as the following procedure: first, human XCR1-expressing B300.19 cells, being suspended in a Ho buffer (0.25 M Sucrose, 10 mM Hepes (pH 7.4), 1 mM EGTA, 0.5 mM $MgCl_2$, 1x Complete mini EDTA-free (Roche Applied Science)), were disrupted (800 psi, for 30 minutes on ice) by a nitrogen gas cell disruption vessel (Parr Instrument Company) and then centrifuged (2,000 g, 10 minutes). The supernatant was collected and re-centrifuged (100,000 g, 30 minutes). The pellet was suspended in a 50 mM Hepes (pH 7.4) buffer, and designated as a membrane fraction.

160 μg or 260 μg of this membrane fraction was mixed with equal volume of GERBU adjuvant (GERBU Biotechnik GmbH), and then injected subcutaneously into the footpads of the XCR1 knockout mice (Deltagen). Five or six additional injections were then administered every other week. Three or four days after the final immunization, the mice were sacrificed, and the peripheral lymph-node cells were fused with P3U1 myeloma cells at a 2:1 or 5:1 ratio in the presence of GenomeONE-CF (Ishihara Sangyo Kaisha, Ltd.). The fused cells were then cultured in 96-well plastic plates.

FACS analysis was performed for primary screening. Parent CHO cells and human XCR1-EGFP-expressing CHO cells were mixed at a 1:1 ratio, and suspended in a FACS buffer (1 mM EDTA, 1% FBS-containing PBS^- (Sigma)). The cells were incubated for 20 minutes on ice with culture supernatants from each hybridoma. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted at 1:100 in the FACS buffer). The cells were washed with the FACS buffer three times, and then suspended in the FACS buffer. The fluorescence intensity was measured using a FACSCanto II Cell analyzer (BD Bioscience). As results, supernatants collected from three wells showed high reactivity to human XCR1-EGFP-expressing CHO cells.

A standard limiting dilution method was used to obtain clones from these three positive wells (2H6, 5G7, and 11H2). The reactivity of each clone was confirmed by the FACS analysis described above.

Subsequently, an *in vitro* chemotaxis assay was performed to evaluate the neutralizing activity of these three clones on human lymphotactin-induced migration of human XCR1-expressing BaF3 cells or B300.19 cells. The chemotaxis assay was performed in 24-well transwell culture supports (pore 3 μm, Costar, #3399) or 96-well transwell culture plates (MultiScreen, pore 5 μm, Millipore, #MAMIC 5S10).

In the case of the 24-well transwell culture supports, human XCR1-expressing BaF3 cells (1 x 10^6 cells) were suspended in a mixture of 50 μL of a chemotaxis buffer (RPMI1 640 medium (Invitrogen) containing 0.5% BSA, 0.5% FBS, and 20 mM HEPES (pH 7.4)) and 50 μL of each culture supernatant, and incubated at room temperature for 30 minutes. Subsequently, recombinant human lymphotactin (Genzyme, #2695) dissolved in the chemotaxis buffer at a concentration of 1 μg/mL was added to the lower wells at 600 μL/well, and the incubated cells were added to the upper wells. After 4 hours of incubation in a 5% CO_2 incubator at 37°C, the transwells were centrifuged at 1,350 rpm for 5 minutes, and migrated cells were collected into the lower wells. The collected cells were fixed with paraformaldehyde (final concentration: 1%), and 30 μL of each sample was applied to the FACSCanto II cell analyzer to count the number of the cells.

In the case of the 96-well transwell culture plates, human XCR1-expressing B300.19 cells (2 x 10^5 cells) were suspended in a mixture of 25 μL of a chemotaxis buffer (RPMI1 640 (Invitrogen) containing 0.5% BSA, 0.5% FBS, and 20 mM HEPES (pH 7.4), and 50 μM 2-mercaptoethanol) and 50 μL of each culture supernatant, and incubated at room temperature for 30 minutes. Subsequently, recombinant human lymphotactin (Genzyme, #2695) dissolved in chemotaxis buffer at a concentration of 1 μg/mL was added to the lower wells at 150 μL/well, and the incubated cells were added to

the upper wells. After 4 hours of incubation in a 5% CO_2 incubator at 37°C, the transwells were centrifuged at 1,350 rpm for 5 minutes, and migrated cells were collected in the lower wells. 30 µL of each sample was applied to the FACSCanto II cell analyzer to count the number of the cells.

The culture supernatants produced by three hybridoma clones (2H6, 5G7, and 11H2) demonstrated neutralizing activity against human lymphotactin-induced migration of human XCR1-expressing BaF3 cells and B300.19 cells.

(2) Reactivity of Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2) to Human XCR1-Expressing Cells

In order to evaluate the reactivity and neutralizing activity of purified antibodies from these three clones, the antibodies were purified with recombinant protein A (GE Healthcare, #17-5080-01) from culture supernatants of each clone. The isotype of each clone was determined using monoclonal antibody isotyping kit (Serotec, #MMT1). 2H6 and 5G7 were IgG2b, κ and 11H2 was IgG2a, κ.

The reactivity of the purified antibodies to human XCR1 was evaluated by FACS analysis. Parent B300.19 cells and human XCR1-EGFP-expressing B300.19 cells were mixed at a 1:1 ratio and suspended in a FACS buffer (1% FBS-containing PBS^- (Sigma)). The cells were blocked for 10 minutes on ice with the FACS buffer containing 100 µg/mL of human immunoglobulin. The cells were then incubated for 20 minutes on ice with the purified antibodies (2H6, 5G7, and 11H2) at various concentrations from 0 to 10 µg/mL or with mouse isotype control antibody, IgG2a (eBioscience, #14-4724-82) or IgG2b (eBioscience, #14-4732-82), at a concentration of 10 µg/mL. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted at 1:50 in the FACS buffer). The cells were washed with the FACS buffer three times, and then suspended in the FACS buffer. The fluorescence intensity was measured by a FACSCanto II

cell analyzer.

These three purified antibodies (2H6, 5G7, and 11H2) showed the reactivity to human XCR1-EGFP-expressing B300.19 cells, but not to parent B300.19 cells (Fig. 1). In contrast, the mouse isotype control antibody did not react to either human XCR1-EGFP-expressing B300.19 cells, or to parent cells (data not shown).

(3) Neutralizing Activity of Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2) against Human Lymphotactin-Induced Migration of Human XCR1-Expressing Cells

The neutralizing activity of purified antibodies from these clones was evaluated by *in vitro* chemotaxis assay. The chemotaxis assay was performed using 96-well transwell culture plates (MultiScreen, pore 5 μm, Millipore, #MAMIC 5S10). Human XCR1-expressing B300.19 cells (2 x 10^5 cells) were suspended in a 75 μL of chemotaxis buffer (RPMI 1640 medium (Invitrogen) containing 0.5% BSA, 0.5% FBS, 20 mM HEPES (pH 7.4), and 50 μM 2-mercaptoethanol) containing each of the purified antibodies (2H6, 5G7, and 11H2) at various concentrations from 0 to 10 μg/mL; and incubated at room temperature for 30 minutes. Further, recombinant human lymphotactin (R&D, #695-LT/CF) was dissolved in the chemotaxis buffer at a final concentration of 1 μg/mL, in which the purified antibodies were dissolved at various concentrations from 0 to 10 μg/mL. The mixture of lymphotactin and purified antibodies were added to the lower wells at 150 μL/well, and incubated at room temperature for 30 minutes. 30 minutes later, the incubated cells were added to the upper wells, and incubated in a 5% CO_2 incubator at 37°C for 4 hours. Subsequently, 30 μL of each sample was applied to the FACSCanto II cell analyzer to count the number of the cells. 2H6, 5G7, and 11H2 mAbs completely inhibited cell migration at a concentration of about 3 μg/mL. Fig. 2 shows the typical pattern of the concentration-dependent inhibition. IC_{50} and IC_{90} values were calculated from three independent experiments. Table 1 shows these values as the mean ± standard error.

Table 1

IC_{50} and IC_{90} Values of 2H6, 5G7, and 11H2 by Chemotaxis Assay

	2H6	5G7	11H2
IC_{50} (nM)	1.28 ± 0.172	0.63 ± 0.139	0.99 ± 0.168
IC_{90} (nM)	7.60 ± 2.331	2.52 ± 0.645	6.32 ± 1.830

(4) Sequence Analysis of Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2)

A polynucleotide comprising a gene sequence encoding the heavy and light chains of the clones (2H6, 5G7, and 11H2) were amplified by 5'-RACE (5'-rapid amplification of cDNA ends) method. The total RNA was prepared from the hybridoma of these three clones using TRIZOL (Invitrogen) and treated with DNase (QIAGEN, RNase free DNase set). Double-stranded cDNA was prepared from the total RNA, using a cDNA synthesis kit (TAKARA). 5' adaptor obtained by annealing ad29S; ACATCACTCCGT (SEQ ID NO: 81) and as29AS; ACGGAGTGATGTCCGTCGACGTATCTCTGCGTTGATACTTCAGCGTAGCT (SEQ ID NO: 82) was added to the cDNA. The obtained cDNA was amplified using

5' forward primer

(5'-PCR4 primer, AGCTACGCTGAAGTATCAACGCAGAG: SEQ ID NO: 83)

and

3' reverse primer

(AGGACAGGGGTTGATTGTTGA: SEQ ID NO: 84, or

CTCAAGTTTTTTGTCCACCGTGGTGC: SEQ ID NO: 85

was used to amplify IgG2b heavy chain;

CTCAATTTTCTTGTCCACCTTGGTGC: SEQ ID NO: 86, or

GCCAGTGGATAGACTGATG: SEQ ID NO: 87

was used to amplify IgG2a heavy chain; and

CTCATTCCTGTTGAAGCTCTTGACAAT: SEQ ID NO: 88,

GATGGATACAGTTGGTGCAGC: SEQ ID NO: 89, or

CAGATCCTCAGCCTCCACTCTGCT: SEQ ID NO: 90

was used to amplify Igκ light chain). The amplified cDNA was inserted into pCR2.1 vector (Invitrogen). The gene sequences were analyzed using ABI3130XL. Tables 2-1 to 4-2 show amino acid

sequences encoded by the gene sequences identified by the analysis.

Table 2-1

Amino Acid Sequences of Mouse Anti-Human XCR1 Antibody (2H6)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 1)	QAYLQQSGAELVRPGASVKMSCKASGYTFSSHNMHWIKQTLRQGLE WIGAIYPGKGNTSYNQKFKGKATLTVDKSSSTAYMQLSSLTSEDSA VYFCARWGSVVGDWYFDVWGTGTTVTVSS
Light chain variable region (SEQ ID NO: 2)	DVVVTQTPLSLPVSLGDQASISCRSSQSLVHSNGNTYLHWYLQKPG QSPKLLIYRVSNRFSGVPDRFSGSGLGRDFTLKISRVEAEDLGVYF CSQSTFVPWTFGGGTKLEIK

Table 2-2

Nucleic Acid Sequences of Mouse Anti-Human XCR1 Antibody (2H6)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 3)	CAGGCTTATCTACAGCAGTCTGGGGCTGAACTGGTGAGGCCTGGGG CCTCAGTGAAGATGTCCTGCAAGGCTTCTGGCTACACATTTAGCAG TCACAATATGCACTGGATAAAGCAGACACTTAGACAGGGCCTGGAA TGGATAGGAGCTATTTATCCAGGAAAAGGTAATACTTCCTACAATC AGAAGTTCAAGGGCAAGGCCACACTGACTGTAGACAAATCCTCCAG CACAGCCTACATGCAGCTCAGCAGCCTGACATCTGAAGACTCTGCG GTCTATTTCTGTGCAAGATGGGGTTCGGTAGTAGGAGACTGGTACT TCGATGTCTGGGGCACAGGGACCACGGTCACCGTCTCTTCA
Light chain variable region (SEQ ID NO: 4)	GATGTTGTGGTGACCCAAACTCCACTCTCCCTGCCTGTCAGTCTTG GAGATCAAGCCTCCATCTCTTGCAGATCTAGTCAGAGCCTTGTACA CAGTAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGC CAGTCTCCAAAGCTCCTGATCTACAGAGTTTCCAATCGATTTTCTG GGGTCCCAGACAGGTTCAGTGGCAGTGGATTAGGGAGAGATTTCAC ACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGAGTTTATTTC TGCTCTCAAAGTACATTTGTTCCGTGGACGTTCGGTGGAGGCACCA AGCTGGAAATCAAA

Table 3-1

Amino Acid Sequences of Mouse Anti-Human XCR1 Antibody (5G7)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 5)	QAYLQQSGAELVRPGASVKMSCKASGYTFTSHNLHWVKQTPRQGLQ WIGAIYPGNGNTAYNQKFKGKATLTVDKSSSTAYMQLSSLTSDDSA VYFCARWGSVVGDWYFDVWGTGTTVTVSS
Light chain	DVVMTQTPLSLPVTLGNQASIFCRSSLGLVHRNGNTYLHWYLQKPG

variable region (SEQ ID NO: 6)	QSPKLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYF CSQSTHVPWTFGGGTKLEIK

Table 3-2

Nucleic Acid Sequences of Mouse Anti-Human XCR1 Antibody (5G7)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 7)	CAGGCTTATCTTCAGCAGTCTGGGGCTGAACTGGTGAGGCCTGGGG CCTCAGTGAAGATGTCCTGCAAGGCTTCTGGCTACACATTCACCAG TCACAATTTGCACTGGGTAAAGCAGACACCTAGACAGGGCCTGCAA TGGATTGGAGCTATTTATCCAGGAAATGGTAATACTGCCTACAATC AGAAGTTCAAGGGCAAGGCCACGCTGACTGTAGACAAATCCTCCAG TACAGCCTACATGCAGCTCAGCAGCCTGACATCTGATGACTCTGCG GTCTACTTCTGTGCAAGATGGGGTTCGGTTGTAGGAGACTGGTACT TCGACGTCTGGGGCACAGGGACCACGGTCACCGTCTCCTCA
Light chain variable region (SEQ ID NO: 8)	GATGTTGTGATGACCCAAACTCCACTCTCCCTGCCTGTCACTCTTG GAAATCAAGCCTCCATTTTTTGTAGATCTAGTCTGGGCCTTGTACA CAGAAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGC CAGTCTCCAAAGCTCCTGATCTACAAAGTTTCCCACCGATTTTCTG GGGTCCCAGACAGGTTCAGTGGCAGTGGCTCAGGGACAGATTTCAC ACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGGGTTTATTTC TGCTCTCAAAGTACCCATGTTCCGTGGACGTTCGGTGGAGGCACCA AGCTGGAAATCAAA

Table 4-1

Amino Acid Sequences of Mouse Anti-Human XCR1 Antibody (11H2)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 9)	EVQLQQSGPVLVKPGASVKMSCKASGYTFTDYYVNWVKQSHGASLE WIGVSNPKNGDKSYNQKFKGKATLTVDKSSSTAYMELNSLTSEDSA VYYCARGLYYAGTYGYFDVWGTGTTVTVSS
Light chain variable region (SEQ ID NO: 10)	DIQMTQATSSLSASLGDRVTISCRASQDISNYLNWYQQKPDGTVKL LIYYTSRLHSGVPSRFRGSGSGTDFSLTISNLEQEDIATYFCQQGK TLPRTLGGGTKLEIK

Table 4-2

Nucleic Acid Sequences of Mouse Anti-Human XCR1 Antibody (11H2)

Name	Sequence
Heavy chain variable region (SEQ ID NO: 11)	GAGGTCCAGCTTCAACAGTCTGGACCTGTGCTGGTGAAGCCTGGGG CTTCAGTGAAGATGTCCTGTAAGGCTTCTGGATACACATTCACTGA CTACTATGTGAACTGGGTGAAACAGAGCCATGGAGCGAGCCTTGAG TGGATTGGAGTTAGTAATCCTAAGAACGGTGATAAAAGTTACAACC AGAAGTTCAAGGGCAAGGCCACATTGACTGTTGACAAGTCCTCCAG

	TACAGCCTACATGGAGCTCAACAGCCTGACATCTGAGGACTCTGCT GTCTATTACTGTGCAAGAGGGCTTTACTACGCTGGTACCTACGGGT ACTTCGATGTCTGGGGCACGGGGACCACGGTCACCGTCTCCTCA
Light chain variable region (SEQ ID NO: 12)	GATATCCAGATGACACAGGCTACATCCTCCCTGTCTGCCTCTCTGG GAGACAGAGTCACCATCAGTTGTAGGGCAAGTCAGGACATTAGCAA TTATTTAAACTGGTATCAGCAGAAGCCAGATGGAACTGTTAAACTC CTGATCTACTACACATCAAGATTACACTCAGGTGTCCCATCAAGGT TCAGAGGCAGTGGGTCTGGGACAGATTTCTCTCTCACCATTAGCAA CCTGGAGCAAGAAGATATTGCCACTTATTTTTGCCAACAGGGTAAA ACGCTTCCTCGGACGCTCGGTGGAGGCACCAAGCTGGAAATCAAA

Example 2

(1) Preparation of Chimeric Anti-Human XCR1 Antibody and Humanized Anti-Human XCR1 Antibodies

5G7, which demonstrated the highest neutralizing activity among 2H6, 5G7, and 11H2, was used to produce a chimeric antibody and humanized antibodies.

The chimeric antibody was prepared by combining, by overlapping extension PCR, the gene sequence of the 5G7 heavy chain variable region and the gene sequence of the human IgG2 constant region into which V234A/G237A mutation was inserted for heavy chain and the sequence of the 5G7 light chain variable region and the gene sequence of human Igκ constant region, and by inserting the resulting sequence into expression vectors (pEE6.4 or pEE12.4). Tables 5 and 6 respectively show amino acid sequences and nucleotide sequences of the specific chimeric antibody.

Table 5

Amino Acid Sequences of Chimeric Anti-Human XCR1 Antibody

Name	Sequence (The variable region is indicated in **bold**, and CDRs in the variable region are underlined.)
Heavy chain (SEQ ID NO: 13)	**QAYLQQSGAELVRPGASVKMSCKASGYTFTSHNLHWVKQTPRQGLQ WIGAIYPGNGNTAYNQKFKGKATLTVDKSSSTAYMQLSSLTSDDSA VYFCARWGSVVGDWYFDVWGTGTTVTVSS**ASTKGPSVFPLAPCSRS TSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY SLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTVERKCCVECPPC PAPPAAAPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFN WYVDGVEVHNAKTKPREEQFNSTFRVVSVLTVVHQDWLNGKEYKCK VSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCL

	VKGFYPSDIAVEWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDK SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
Light chain (SEQ ID NO: 14)	**DVVMTQTPLSLPVTLGNQASIFCRSSLGLVHRNGNTYLHWYLQKPG QSPKLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDLGVYF CSQSTHVPWTFGGGTKLEIK**RTVAAPSVFIFPPSDEQLKSGTASVV CLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTL TLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC

Table 6

Nucleic Acid Sequences of Chimeric Anti-Human XCR1 Antibody

Name	Sequence (The variable region is indicated in **bold.**)
Heavy chain (SEQ ID NO: 15)	**CAGGCTTATCTTCAGCAGTCTGGGGCTGAACTGGTGAGGCCTGGGG CCTCAGTGAAGATGTCCTGCAAGGCTTCTGGCTACACATTCACCAG TCACAATTTGCACTGGGTAAAGCAGACACCTAGACAGGGCCTGCAA TGGATTGGAGCTATTTATCCAGGAAATGGTAATACTGCCTACAATC AGAAGTTCAAGGGCAAGGCCACGCTGACTGTAGACAAATCCTCCAG TACAGCCTACATGCAGCTCAGCAGCCTGACATCTGATGACTCTGCG GTCTACTTCTGTGCAAGATGGGGTTCGGTTGTAGGAGACTGGTACT TCGACGTCTGGGGCACAGGGACCACGGTCACCGTCTCCTCA**GCTAG CACCAAGGGCCCATCGGTCTTCCCCCTGGCGCCCTGCTCCAGGAGC ACCTCCGAGAGCACAGCGGCCCTGGGCTGCCTGGTCAAGGACTACT TCCCCGAACCGGTGACGGTGTCGTGGAACTCAGGCGCTCTGACCAG CGGCGTGCACACCTTCCCGGCTGTCCTACAGTCCTCAGGACTCTAC TCCCTCAGCAGCGTGGTGACCGTGCCCTCCAGCAACTTCGGCACCC AGACCTACACCTGCAACGTAGATCACAAGCCCAGCAACACCAAGGT GGACAAGACAGTTGAGCGCAAATGTTGTGTCGAGTGCCCACCGTGC CCAGCACCACCTGCCGCAGCCCCGTCAGTCTTCCTGTTCCCCCCAA AACCCAAGGACACCCTCATGATCTCCCGGACCCCTGAGGTCACGTG CGTGGTGGTGGACGTGAGCCACGAAGACCCCGAGGTCCAGTTCAAC TGGTACGTGGACGGCGTGGAGGTGCATAATGCCAAGACAAAGCCAC GGGAGGAGCAGTTCAACAGCACGTTCCGTGTGGTCAGCGTCCTCAC CGTCGTGCACCAGGACTGGCTGAACGGCAAGGAGTACAAGTGCAAG GTCTCCAACAAAGGCCTCCCAGCCCCCATCGAGAAAACCATCTCCA AAACCAAAGGGCAGCCCCGAGAACCACAGGTGTACACCCTGCCCCC ATCCCGGGAGGAGATGACCAAGAACCAGGTCAGCCTGACCTGCCTG GTCAAAGGCTTCTACCCCAGCGACATCGCCGTGGAGTGGGAGAGCA ATGGGCAGCCGGAGAACAACTACAAGACCACACCTCCCATGCTGGA CTCCGACGGCTCCTTCTTCCTCTACAGCAAGCTCACCGTGGACAAG AGCAGGTGGCAGCAGGGGAACGTCTTCTCATGCTCCGTGATGCATG AGGCTCTGCACAACCACTACACACAGAAGAGCCTCTCCCTGTCTCC GGGTAAATGA
Light chain (SEQ ID NO: 16)	**GATGTTGTGATGACCCAAACTCCACTCTCCCTGCCTGTCACTCTTG GAAATCAAGCCTCCATTTTTTGTAGATCTAGTCTGGGCCTTGTACA CAGAAATGGAAACACCTATTTACATTGGTACCTGCAGAAGCCAGGC CAGTCTCCAAAGCTCCTGATCTACAAAGTTTCCCACCGATTTTCTG GGGTCCCAGACAGGTTCAGTGGCAGTGGCTCAGGGACAGATTTCAC ACTCAAGATCAGCAGAGTGGAGGCTGAGGATCTGGGGGTTTATTTC**

	TGCTCTCAAAGTACCCATGTTCCGTGGACGTTCGGTGGAGGCACCA AGCTGGAAATCAAACGTACGGTGGCTGCACCATCTGTCTTCATCTT CCCGCCATCTGATGAGCAGTTGAAATCTGGAACTGCCTCTGTTGTG TGCCTGCTGAATAACTTCTATCCCAGAGAGGCCAAAGTACAGTGGA AGGTGGATAACGCCCTCCAATCGGGTAACTCCCAGGAGAGTGTCAC AGAGCAGGACAGCAAGGACAGCACCTACAGCCTCAGCAGCACCCTG ACGCTGAGCAAAGCAGACTACGAGAAACACAAAGTCTACGCCTGCG AAGTCACCCATCAGGGCCTGAGCTCGCCCGTCACAAAGAGCTTCAA CAGGGGAGAGTGTTAG

The antibody was humanized by grafting the complementary determining region of mouse antibody 5G7 into the human antibody variable region. The complementary determining region was determined according to the Kabat numbering system and a method for identifying the complementary determining region (for example, Kabat et al., (1991) Sequences of Proteins of Immunological Interest: US Department of Health and Human Services, NIH, USA). Further, the complementary determining regions of 2H6 and 11H2 were also determined in a similar manner. Tables 7-1 to 9-2 show amino acid sequences and nucleotide sequences of the complementary determining regions of these three clones.

Table 7-1

Amino Acid Sequences of Complementary Determining Region of 5G7

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 17)	SHNLH
Heavy chain CDR 2 (SEQ ID NO: 18)	AIYPGNGNTAYNQKFKG
Heavy chain CDR 3 (SEQ ID NO: 19)	WGSVVGDWYFDV
Light chain CDR 1 (SEQ ID NO: 20)	RSSLGLVHRNGNTYLH
Light chain CDR 2 (SEQ ID NO: 21)	KVSHRFS
Light chain CDR 3 (SEQ ID NO: 22)	SQSTHVPWT

Table 7-2

Nucleic Acid Sequences of Complementary Determining Region of 5G7

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 23)	AGTCACAATTTGCAC
Heavy chain CDR 2 (SEQ ID NO: 24)	GCTATTTATCCAGGAAATGGTAATACTGCCTACAATCAGAAGTT CAAGGGC
Heavy chain CDR 3 (SEQ ID NO: 25)	TGGGGTTCGGTTGTAGGAGACTGGTACTTCGACGTC
Light chain CDR 1 (SEQ ID NO: 26)	AGATCTAGTCTGGGCCTTGTACACAGAAATGGAAACACCTATTT ACAT
Light chain CDR 2 (SEQ ID NO: 27)	AAAGTTTCCCACCGATTTTCT
Light chain CDR 3 (SEQ ID NO: 28)	TCTCAAAGTACCCATGTTCCGTGGACG

Table 8-1

Amino Acid Sequences of Complementary Determining Region of 2H6

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 29)	SHNMH
Heavy chain CDR 2 (SEQ ID NO: 30)	AIYPGKGNTSYNQKFKG
Heavy chain CDR 3 (SEQ ID NO: 31)	WGSVVGDWYFDV
Light chain CDR 1 (SEQ ID NO: 32)	RSSQSLVHSNGNTYLH
Light chain CDR 2 (SEQ ID NO: 33)	RVSNRFS
Light chain CDR 3 (SEQ ID NO: 34)	SQSTFVPWT

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Table 8-2

Nucleic Acid Sequences of Complementary Determining Region of 2H6

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 35)	AGTCACAATATGCAC
Heavy chain CDR 2 (SEQ ID NO: 36)	GCTATTTATCCAGGAAAAGGTAATACTTCCTACAATCAGAAGTT CAAGGGC
Heavy chain CDR 3 (SEQ ID NO: 37)	TGGGGTTCGGTAGTAGGAGACTGGTACTTCGATGTC
Light chain CDR 1	AGATCTAGTCAGAGCCTTGTACACAGTAATGGAAACACCTATTT

(SEQ ID NO: 38)	ACAT
Light chain CDR 2 (SEQ ID NO: 39)	AGAGTTTCCAATCGATTTTCT
Light chain CDR 3 (SEQ ID NO: 40)	TCTCAAAGTACATTTGTTCCGTGGACG

Table 9-1

Amino Acid Sequences of Complementary Determining Region of 11H2

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 41)	DYYVN
Heavy chain CDR 2 (SEQ ID NO: 42)	VSNPKNGDKSYNQKFKG
Heavy chain CDR 3 (SEQ ID NO: 43)	GLYYAGTYGYFDV
Light chain CDR 1 (SEQ ID NO: 44)	RASQDISNYLN
Light chain CDR 2 (SEQ ID NO: 45)	YTSRLHS
Light chain CDR 3 (SEQ ID NO: 46)	QQGKTLPRT

Table 9-2

Nucleic Acid Sequences of Complementary Determining Region of 11H2

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 47)	GACTACTATGTGAAC
Heavy chain CDR 2 (SEQ ID NO: 48)	GTTAGTAATCCTAAGAACGGTGATAAAAGTTACAACCAGAAGTTCAAGGGC
Heavy chain CDR 3 (SEQ ID NO: 49)	GGGCTTTACTACGCTGGTACCTACGGGTACTTCGATGTC
Light chain CDR 1 (SEQ ID NO: 50)	AGGGCAAGTCAGGACATTAGCAATTATTTAAAC
Light chain CDR 2 (SEQ ID NO: 51)	TACACATCAAGATTACACTCA
Light chain CDR 3 (SEQ ID NO: 52)	CAACAGGGTAAAACGCTTCCTCGGACG

As is clear from Tables 7-1 and 8-1, the identity of

the amino acid sequences of the CDRs between 5G7 and 2H6 is high; in particular, the heavy chain CDR 3 amino acid sequences were completely identical. Accordingly, in regard to 5G7 and 2H6, the amino acid sequences can be generalized as shown in Table 10 below. Additionally, Fig. 7 shows the comparison of amino acid sequences of the CDRs 1 to 3 of these clones.

Table 10

Generalized Amino Acid Sequences of Complementary Determining Regions of 5G7 and 2H6

Name	Sequence
Heavy chain CDR 1 (SEQ ID NO: 53)	SHNXH
Heavy chain CDR 2 (SEQ ID NO: 54)	AIYPGXGNTXYNQKFKG
Heavy chain CDR 3 (SEQ ID NO: 55)	WGSVVGDWYFDV
Light chain CDR 1 (SEQ ID NO: 56)	RSSXXLVHXNGNTYLH
Light chain CDR 2 (SEQ ID NO: 57)	XVSXRFS
Light chain CDR 3 (SEQ ID NO: 58)	SQSTXVPWT

The "X" in the table may be any of the following: alanine (Ala: A), arginine (Arg: R), asparagine (Asn: N), aspartic acid (Asp: D), cysteine (Cys: C), glutamine (Gln: Q), glutamic acid (Glu: E), glycine (Gly: G), histidine (His: H), isoleucine (Ile: I), leucine (Leu: L), lysine (Lys: K), methionine (Met: M), phenylalanine (Phe: F), proline (Pro: P), serine (Ser: S), threonine (Thr: T), tryptophan (Trp: W), tyrosine (Tyr: Y), and valine (Val: V).

The FRs of a human antibody with high identity to the FR of 5G7 were selected as the FRs of the humanized antibodies. Subsequently, the amino acids in the FRs, which interact with the CDRs of 5G7, were predicted using the 3D model of the resulting

antibody, and grafted with the CDRs. The human IgG2 constant region into which V234A/G237A mutation was inserted was used as the constant region. HK1 and HK5 were designed as the humanized antibody heavy chains, and L2 and L5 were designed as the humanized antibody light chains. Tables 11-1 to 14-2 show amino acid sequences and nucleotide sequences of the specific humanized antibodies.

Table 11-1

Amino Acid Sequences of Humanized Anti-Human XCR1 Antibody Heavy Chain (HK1)

Name	Sequence (The variable region is indicated in **bold**, and CDRs in the variable region are underlined.)
Heavy chain (SEQ ID NO: 59)	**QVQLVQSGAEVKKPGASVKVSCKASGYTFTSHNLHWVRQAPGQRLE WMGAIYPGNGNTAYNQKFKGRVTITRDTSASTAYMELSSLRSEDTA VYYCARWGSVVGDWYFDVWGQGTLVTVSS**ASTKGPSVFPLAPCSRS TSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY SLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTVERKCCVECPPC PAPPAAAPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFN WYVDGVEVHNAKTKPREEQFNSTFRVVSVLTVVHQDWLNGKEYKCK VSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCL VKGFYPSDIAVEWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDK SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
Heavy chain variable region (SEQ ID NO: 60)	**QVQLVQSGAEVKKPGASVKVSCKASGYTFTSHNLHWVRQAPGQRLE WMGAIYPGNGNTAYNQKFKGRVTITRDTSASTAYMELSSLRSEDTA VYYCARWGSVVGDWYFDVWGQGTLVTVSS**

Table 11-2

Nucleic Acid Sequences of Humanized Anti-Human XCR1 Antibody Heavy Chain (HK1)

Name	Sequence (The variable region is indicated in **bold**, and CDRs in the variable region are underlined.)
Heavy chain (SEQ ID NO: 61)	**CAGGTGCAGCTGGTGCAGTCTGGAGCCGAAGTGAAGAAACCAGGGG CCTCTGTCAAGGTGAGTTGCAAGGCCTCCGGTTACACTTTCACCTC CCACAACCTGCATTGGGTGAGACAGGCTCCTGGACAGCGACTGGAG TGGATGGGAGCAATCTACCCAGGCAACGGAAATACTGCCTATAATC AGAAGTTTAAAGGCAGGGTGACAATTACTCGGGACACTTCCGCAAG**

	CACCGCCTACATGGAGCTGTCCAGCCTGAGGAGTGAAGATACCGCT GTGTACTATTGTGCACGATGGGGATCCGTGGTCGGAGACTGGTATT TCGATGTGTGGGGGCAGGGTACCCTGGTCACAGTGTCTAGTGCCTC CACAAAGGGCCCCAGCGTGTTTCCACTGGCTCCCTGCTCTAGGAGT ACATCAGAGTCCACTGCCGCTCTGGGATGTCTGGTGAAGGACTATT TCCCTGAACCAGTCACCGTGAGTTGGAACTCAGGGGCTCTGACATC TGGTGTCCACACTTTTCCTGCAGTGCTGCAGTCATCCGGCCTGTAC TCCCTGAGCTCTGTGGTCACAGTCCCAAGTTCAAATTTCGGAACCC AGACATATACTTGCAACGTGGACCATAAGCCCAGCAATACCAAGGT CGATAAAACAGTGGAGCGAAAGTGCTGTGTCGAATGCCCACCTTGT CCAGCTCCACCAGCAGCAGCTCCTTCTGTGTTCCTGTTTCCTCCAA AGCCAAAAGACACTCTGATGATCAGCCGGACCCCCGAGGTCACATG TGTGGTCGTGGACGTGTCTCACGAGGATCCTGAAGTCCAGTTTAAC TGGTACGTGGATGGGGTCGAAGTGCATAATGCAAAGACAAAACCAC GAGAGGAACAGTTCAACTCTACATTTCGTGTCGTGAGTGTGCTGAC TGTCGTGCACCAGGATTGGCTGAACGGCAAGGAGTATAAGTGCAAA GTGTCCAATAAGGGACTGCCCGCCCCTATCGAGAAAACTATTAGCA AGACCAAAGGCCAGCCTAGAGAACCACAGGTGTACACCCTGCCCCC TAGTCGCGAGGAAATGACTAAGAACCAGGTCTCACTGACCTGTCTG GTGAAAGGGTTCTATCCCAGCGACATTGCCGTGGAGTGGGAATCTA ATGGTCAGCCTGAGAACAATTACAAGACCACACCACCCATGCTGGA CTCCGATGGGAGCTTCTTTCTGTATTCAAAGCTGACCGTGGATAAA TCCAGGTGGCAGCAGGGTAATGTCTTTAGCTGCTCTGTGATGCACG AAGCCCTGCACAACCATTACACTCAGAAGTCCCTGTCCCTGTCACC TGGAAAGTGA
Heavy chain variable region (SEQ ID NO: 62)	**CAGGTGCAGCTGGTGCAGTCTGGAGCCGAAGTGAAGAAACCAGGGG CCTCTGTCAAGGTGAGTTGCAAGGCCTCCGGTTACACTTTCACCTC CCACAACCTGCATTGGGTGAGACAGGCTCCTGGACAGCGACTGGAG TGGATGGGAGCAATCTACCCAGGCAACGGAAATACTGCCTATAATC AGAAGTTTAAAGGCAGGGTGACAATTACTCGGGACACTTCCGCAAG CACCGCCTACATGGAGCTGTCCAGCCTGAGGAGTGAAGATACCGCT GTGTACTATTGTGCACGATGGGGATCCGTGGTCGGAGACTGGTATT TCGATGTGTGGGGGCAGGGTACCCTGGTCACAGTGTCTAGT**

Table 12-1
Amino Acid Sequences of Humanized Anti-Human XCR1 Antibody Heavy Chain (HK5)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Heavy chain (SEQ ID NO: 63)	**QVQLVQSGAEVKKPGASVKVSCKASGYTFTSHNLHWVRQAPGQGLE WMGAIYPGNGNTAYNQKFKGRVTMTRDTSTSTVYMELSSLRSEDTA**

	VYYCARWGSVVGDWYFDVWGQGTLVTVSSASTKGPSVFPLAPCSRS TSESTAALGCLVKDYFPEPVTVSWNSGALTSGVHTFPAVLQSSGLY SLSSVVTVPSSNFGTQTYTCNVDHKPSNTKVDKTVERKCCVECPPC PAPPAAAPSVFLFPPKPKDTLMISRTPEVTCVVVDVSHEDPEVQFN WYVDGVEVHNAKTKPREEQFNSTFRVVSVLTVVHQDWLNGKEYKCK VSNKGLPAPIEKTISKTKGQPREPQVYTLPPSREEMTKNQVSLTCL VKGFYPSDIAVEWESNGQPENNYKTTPPMLDSDGSFFLYSKLTVDK SRWQQGNVFSCSVMHEALHNHYTQKSLSLSPGK
Heavy chain variable region (SEQ ID NO: 64)	**QVQLVQSGAEVKKPGASVKVSCKASGYTFTSHNLHWVRQAPGQGLE WMGAIYPGNGNTAYNQKFKGRVTMTRDTSTSTVYMELSSLRSEDTA VYYCARWGSVVGDWYFDVWGQGTLVTVSS**

Table 12-2

Nucleic Acid Sequences of Humanized Anti-Human XCR1 Antibody Heavy Chain (HK5)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Heavy chain (SEQ ID NO: 65)	**CAGGTGCAGCTGGTGCAGTCTGGGGCCGAAGTGAAGAAACCAGGGG CTTCTGTCAAGGTGAGTTGCAAAGCATCAGGTTACACTTTCACCTC CCACAACCTGCATTGGGTGCGACAGGCTCCTGGACAGGGACTGGAG TGGATGGGAGCAATCTACCCAGGGAACGGTAATACCGCTTATAATC AGAAGTTTAAAGGCAGGGTCACAATGACTCGGGACACCTCCACAAG CACTGTGTACATGGAGCTGTCCAGCCTGCGAAGTGAAGATACAGCA GTGTACTATTGTGCACGTTGGGGATCCGTGGTCGGTGACTGGTATT TCGATGTGTGGGGCCAGGGAACCCTGGTCACAGTGTCTAGT**GCTTC CACTAAGGGGCCCAGCGTGTTTCCACTGGCACCCTGCTCTCGGAGT ACTTCAGAGTCCACCGCCGCTCTGGGCTGTCTGGTGAAGGACTATT TCCCTGAACCAGTCACAGTGAGTTGGAACTCAGGCGCACTGACTTC TGGAGTCCACACCTTTCCTGCCGTGCTGCAGTCATCCGGCCTGTAC TCCCTGAGCTCTGTGGTCACTGTCCCAAGTTCAAATTTCGGAACCC AGACATATACTTGCAACGTGGACCATAAGCCCAGCAATACAAAGGT CGATAAAACTGTGGAGAGAAAGTGCTGTGTGGAATGCCCACCTTGT CCAGCACCACCAGCAGCAGCTCCTTCTGTGTTCCTGTTTCCTCCAA AGCCAAAAGACACACTGATGATCAGCCGCACACCCGAGGTCACTTG TGTGGTCGTGGACGTGTCTCACGAGGATCCTGAAGTCCAGTTTAAC TGGTACGTGGATGGCGTCGAAGTGCATAATGCCAAGACCAAACCAA GAGAGGAACAGTTCAACTCTACTTTTCGCGTCGTGAGTGTGCTGAC CGTCGTGCACCAGGATTGGCTGAACGGCAAGGAGTATAAGTGCAAA GTGTCCAATAAGGGACTGCCCGCTCCTATCGAGAAAACCATTAGCA AGACAAAAGGACAGCCTAGGGAACCACAGGTGTACACCCTGCCCCC TAGTCGGGAGGAAATGACCAAGAACCAGGTCTCACTGACATGTCTG

	GTGAAAGGGTTCTATCCCAGCGACATTGCCGTGGAGTGGGAATCTA ATGGTCAGCCTGAGAACAATTACAAGACCACACCACCCATGCTGGA CTCCGATGGCAGCTTCTTTCTGTATTCAAAGCTGACCGTGGATAAA TCCAGGTGGCAGCAGGGAAATGTCTTTAGCTGCTCTGTGATGCACG AAGCACTGCATAATCACTACACTCAGAAGAGCCTGTCCCTGTCACC TGGTAAATGA
Heavy chain variable region (SEQ ID NO: 66)	**CAGGTGCAGCTGGTGCAGTCTGGGGCCGAAGTGAAGAAACCAGGGG CTTCTGTCAAGGTGAGTTGCAAAGCATCAGGTTACACTTTCACCTC CCACAACCTGCATTGGGTGCGACAGGCTCCTGGACAGGGACTGGAG TGGATGGGAGCAATCTACCCAGGGAACGGTAATACCGCTTATAATC AGAAGTTTAAAGGCAGGGTCACAATGACTCGGGACACCTCCACAAG CACTGTGTACATGGAGCTGTCCAGCCTGCGAAGTGAAGATACAGCA GTGTACTATTGTGCACGTTGGGGATCCGTGGTCGGTGACTGGTATT TCGATGTGTGGGGCCAGGGAACCCTGGTCACAGTGTCTAGT**

Table 13-1

Amino Acid Sequences of Humanized Anti-Human XCR1 Antibody Light Chain (L2)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Light chain (SEQ ID NO: 67)	**DVVMTQSPLSLPVTLGQPASISCRSSLGLVHRNGNTYLHWFQQRPG QSPRLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGVYY CSQSTHVPWTFGQGTKVEIK**RTVAAPSVFIFPPSDEQLKSGTASVV CLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTL TLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC
Light chain variable region (SEQ ID NO: 68)	**DVVMTQSPLSLPVTLGQPASISCRSSLGLVHRNGNTYLHWFQQRPG QSPRLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGVYY CSQSTHVPWTFGQGTKVEIK**

Table 13-2

Nucleic Acid Sequences of Humanized Anti-Human XCR1 Antibody Light Chain (L2)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Light chain (SEQ ID NO: 69)	**GATGTCGTGATGACCCAGTCTCCTCTGAGCCTGCCTGTGACTCTGG GCCAGCCAGCATCAATCTCCTGCCGATCCAGCCTGGGACTGGTGCA CCGTAACGGGAATACCTACCTGCATTGGTTCCAGCAGAGGCCTGGT CAGAGTCCCCGGCTGCTGATCTATAAGGTGTCTCACAGATTCAGTG GCGTCCCAGACCGCTTTAGCGGCTCTGGAAGTGGGACTGATTTCAC**

	CCTGAAAATTTCCCGAGTGGAGGCAGAAGACGTGGGAGTCTACTAT TGCTCACAGTCCACACATGTGCCCTGGACTTTTGGTCAGGGCACCA AGGTCGAGATCAAACGCACCGTGGCCGCTCCTAGCGTCTTCATTTT TCCCCCTTCTGACGAACAGCTGAAGTCAGGAACAGCTTCCGTGGTC TGTCTGCTGAACAATTTTTACCCCAGAGAGGCAAAGGTGCAGTGGA AAGTCGATAACGCCCTGCAGAGCGGCAACTCCCAGGAGAGTGTGAC AGAACAGGACTCAAAGGATTCCACTTATAGCCTGTCTAGTACCCTG ACACTGTCTAAAGCTGATTACGAGAAGCACAAAGTGTATGCATGTG AAGTCACCCACCAGGGGCTGTCATCACCCGTCACCAAGTCCTTTAA TAGAGGGGAGTGTTGA
Light chain variable region (SEQ ID NO: 70)	**GATGTCGTGATGACCCAGTCTCCTCTGAGCCTGCCTGTGACTCTGG GCCAGCCAGCATCAATCTCCTGCCGATCCAGCCTGGGACTGGTGCA CCGTAACGGGAATACCTACCTGCATTGGTTCCAGCAGAGGCCTGGT CAGAGTCCCCGGCTGCTGATCTATAAGGTGTCTCACAGATTCAGTG GCGTCCCAGACCGCTTTAGCGGCTCTGGAAGTGGGACTGATTTCAC CCTGAAAATTTCCCGAGTGGAGGCAGAAGACGTGGGAGTCTACTAT TGCTCACAGTCCACACATGTGCCCTGGACTTTTGGTCAGGGCACCA AGGTCGAGATCAAA**

Table 14-1

Amino Acid Sequences of Humanized Anti-Human XCR1 Antibody Light Chain (L5)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Light chain (SEQ ID NO: 71)	**DIVMTQTPLSLPVTPGQPASISCRSSLGLVHRNGNTYLHWYLQKPG QSPQLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGVYY CSQSTHVPWTFGQGTKVEIK**RTVAAPSVFIFPPSDEQLKSGTASVV CLLNNFYPREAKVQWKVDNALQSGNSQESVTEQDSKDSTYSLSSTL TLSKADYEKHKVYACEVTHQGLSSPVTKSFNRGEC
Light chain Variable Region (SEQ ID NO: 72)	**DIVMTQTPLSLPVTPGQPASISCRSSLGLVHRNGNTYLHWYLQKPG QSPQLLIYKVSHRFSGVPDRFSGSGSGTDFTLKISRVEAEDVGVYY CSQSTHVPWTFGQGTKVEIK**

Table 14-2

Nucleic Acid Sequences of Humanized Anti-Human XCR1 Antibody Light Chain (L5)

Name	Sequence (The variable region is indicated in **bold,** and CDRs in the variable region are underlined.)
Light chain	**GATATTGTGATGACTCAGACTCCACTGAGCCTGCCAGTGACTCCCG**

(SEQ ID NO: 73)	**GCCAGCCTGCATCAATCTCCTGCAGATCCAGCCTGGGACTGGTGCA CCGCAACGGGAATACCTACCTGCATTGGTATCTGCAGAAGCCTGGT CAGAGTCCCCAGCTGCTGATCTACAAAGTGTCTCACAGGTTCAGTG GCGTCCCCGACCGGTTTAGCGGCTCTGGAAGTGGGACTGATTTCAC CCTGAAGATTTCCCGAGTGGAGGCCGAAGACGTGGGCGTCTACTAT TGCTCACAGTCCACACATGTGCCTTGGACTTTTGGTCAGGGCACCA AGGTCGAGATCAAA**AGGACCGTGGCCGCTCCAAGCGTCTTCATTTT TCCCCCTTCTGACGAACAGCTGAAGTCAGGAACAGCTTCCGTGGTC TGTCTGCTGAACAATTTCTACCCCAGAGAGGCAAAGGTGCAGTGGA AAGTCGATAACGCCCTGCAGAGCGGCAACTCCCAGGAGAGTGTGAC AGAACAGGACTCAAAGGATTCCACTTATAGCCTGTCTAGTACCCTG ACACTGTCTAAAGCTGATTACGAGAAGCACAAAGTGTATGCATGTG AAGTCACACACCAGGGTCTGAGTTCCCCCGTCACCAAATCCTTTAA TCGTGGAGAGTGCTGA
Light chain Variable Region (SEQ ID NO: 74)	**GATATTGTGATGACTCAGACTCCACTGAGCCTGCCAGTGACTCCCG GCCAGCCTGCATCAATCTCCTGCAGATCCAGCCTGGGACTGGTGCA CCGCAACGGGAATACCTACCTGCATTGGTATCTGCAGAAGCCTGGT CAGAGTCCCCAGCTGCTGATCTACAAAGTGTCTCACAGGTTCAGTG GCGTCCCCGACCGGTTTAGCGGCTCTGGAAGTGGGACTGATTTCAC CCTGAAGATTTCCCGAGTGGAGGCCGAAGACGTGGGCGTCTACTAT TGCTCACAGTCCACACATGTGCCTTGGACTTTTGGTCAGGGCACCA AGGTCGAGATCAAA**

Gene sequences of these humanized antibodies were entirely synthesized by GenScript USA Inc., and inserted into expression vectors (pEE6.4 or pEE12.4 purchased from Lonza). To produce antibodies, the expression vectors were transfected into HEK293E cells (Invitrogen) using Lipofectamine 2000 according to the instructions for Lipofectamine 2000 (Invitrogen). The supernatants were collected and purified using Protein A (GE Healthcare). The neutralizing activity was evaluated using these purified humanized antibodies.

Humanized antibodies having neutralizing activity against human lymphotactin-induced migration of human XCR1-expressing cells were identified by performing *in vitro* chemotaxis assays using human XCR1-expressing B300.19 cells. The chemotaxis assay was performed as described above using 96-well transwell culture plates (MultiScreen, pore 5 μm, Millipore, #MAMIC 5S10 or Corning #3387 or #3388). However, in the case of using Corning

transwell culture plates, the amount of the recombinant human lymphotactin and purified antibodies to be added to the lower wells is 235 µL per well.

Among humanized antibodies having neutralizing activity, two types of the following antibodies, HK1L2 and HK5L5, were evaluated in further detail.

(2) Reactivity of Humanized Anti-Human XCR1 Antibodies (HK1L2 and HK5L5) to Human XCR1-Expressing Cells

FACS analysis was performed using these two humanized antibodies (HK1L2 and HK5L5), the parent antibody 5G7, and the chimeric antibody. Parent B300.19 cells and human XCR1-EGFP-expressing B300.19 cells were mixed at a 1:1 ratio and suspended in a FACS buffer (1% FBS-containing PBS^- (Sigma)). The cells were incubated for 20 minutes on ice with the purified antibodies at various concentrations from 0 to 10 µg/mL. The cells were washed three times with the FACS buffer, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151: used for cells which had been stained with parent antibody 5G7, diluted at 1:100 in the FACS buffer) or with PE-labeled anti-human IgG polyclonal antibody (Jackson, #709-116-149: used for cells which had been stained with chimeric antibody or humanized antibodies (HK1L2 and HK5L5), diluted at 1:100 in FACS buffer. The cells were washed three times with FACS buffer, and then suspended in FACS buffer. The fluorescence intensity was measured using a FACSCanto II cell analyzer (BD Bioscience).

Humanized antibodies (HK1L2 and HK5L5) showed concentration-dependent reactivity to human XCR1-EGFP-expressing B300.19 cells. Parent antibody 5G7 and chimeric antibody showed substantially the same reactivity (Fig. 3).

The reactivity of humanized antibodies (HK1L2 and HK5L5) to human XCR1 was further examined by FACS analysis using human peripheral blood mononuclear cells. Because human XCR1 gene is known to be expressed in BDCA3+ dendritic cells, which is a

minor population in human peripheral blood mononuclear cells, first, the dendritic cells were concentrated from human peripheral blood mononuclear cells and used for FACS analysis. Human peripheral blood mononuclear cells were isolated from the blood of healthy human subjects using Ficoll-Paque (GE Healthcare, #17-1440-02). CD3, CD14, CD19, and CD56 positive cells from human peripheral blood mononuclear cells were labeled with CD3, CD14, CD19, CD56 antibody microbeads (Miltenyi, #130-050-101, #130-050-201, #130-050-301, #130-050-401), and depleted using auto-MACS (Miltenyi). Thereby, human dendritic cells were concentrated. The concentrated dendritic cells were blocked for 10 minutes on ice with a FACS buffer (1% FBS-containing PBS^- (Sigma)) containing 1% rat serum, 1% mouse serum, 100 μg/mL human immunoglobulin. The cells were then stained for 30 minutes on ice separately using PE-labeled 5G7, HK1L2, HK5L5, and isotype control antibody mouse IgG2b, κ (eBioscience, #14-4732-82) or human IgG2, κ (Sigma, #I5404) with FITC-labeled anti-BDCA3 antibody (Miltenyi, #130-090-513), APC-labeled anti-CD123 antibody (Miltenyi, #130-090-901), APC-Cy7-labeled anti-HLA-DR antibody (BioLegend, #307617), and Alexa700-labeled anti-CD3, CD14, CD19, CD56 antibodies (BioLegend, #300324, #301822, #302225, and #318316). The cells were washed three times with the FACS buffer, and then suspended in the FACS buffer. The fluorescence intensity was measured using a FACSCanto II cell analyzer.

As is the case with parent antibody 5G7, the humanized antibodies (HK1L2 and HK5L5) selectively reacted to BDCA3+ dendritic cells expressing human XCR1 (Fig. 4).

(3) Neutralizing Activity of Humanized Anti-Human XCR1 Antibodies (HK1L2 and HK5L5) on Human Lymphotactin-Induced Migration of Human XCR1-Expressing Cells

The neutralizing activity of these humanized antibodies was evaluated in parallel with parent antibody 5G7 and a chimeric antibody by *in vitro* chemotaxis assay as described above.

In comparison with parent antibody 5G7, both humanized

antibodies maintained the neutralizing activity. Fig. 5 shows the typical pattern of concentration-dependent inhibition. IC_{50} and IC_{90} values were calculated from three independent experiments. Table 16 shows these values as the mean ± standard error.

Table 16

Neutralizing Activity of Humanized Antibodies (HK1L2 and HK5L5) in Chemotaxis Assay

	Mouse mAb	Chimeric mAb
	Mean ± S.E.	Mean ± S.E.
IC_{50} (nM)	0.23 ± 0.120	0.27 ± 0.101
IC_{90} (nM)	1.31 ± 0.452	1.52 ± 0.755

HK1L2				
	Assay-1	-2	-3	Mean ± S.E.
IC_{50} (nM)	0.10	0.25	0.12	0.16 ± 0.081
IC_{90} (nM)	0.66	1.01	0.92	0.86 ± 0.182
HK5L5				
	Assay-1	-2	-3	Mean ± S.E
IC_{50} (nM)	0.52	0.36	0.20	0.36 ± 0.160
IC_{90} (nM)	2.30	3.15	1.70	2.38 ± 0.729

Next, the neutralizing activity of humanized antibodies (HK1L2 and HK5L5) was further examined by transendothelial migration assay that used human dendritic cells instead of the human XCR1-expressing B300.19 cells. The transendothelial migration assay was performed using 24-well transwell culture supports (pore 5 μm, Costar, #3421). First, ECV304 cells were suspended in 10% FBS-containing Medium 199 Earle's medium (Invitrogen), and seeded into the upper chamber of the transwell at 2 x 10^5 cells per well, followed by incubation in a 5% CO_2 incubator at 37°C for 3 days. On the day of an assay, ECV304 cells were washed with assay buffer (a mixture of Medium 199 Earle's medium and RPMI 1640 medium at a 1:1 ratio, to which 0.5% BSA and 20 mM HEPES (pH 7.4) were added). Recombinant human lymphotactin dissolved in the assay buffer at a concentration of 1 μg/mL, to which the chimeric antibody, HK1L2, HK5L5, or isotype

control antibody human IgG2, κ (Sigma) was added at a concentration of 10 μg/mL, was added to the lower wells at 600 μL/well. Human dendritic cells were concentrated as described above, suspended in the assay buffer to which the chimeric antibody, the humanized antibodies (HK1L2 and HK5L5) and isotype control antibody human IgG2, κ (Sigma) were added at a concentration of 10 μg/mL, and added to the upper wells containing ECV304 cells. After incubation for 4 hours in a 5% CO_2 incubator at 37°C, the cells in the transwell were centrifuged at 1,350 rpm for 5 minutes, and migrated cells were collected. The collected cells were stained for 30 minutes on ice, using cell lineage markers, FITC-labeled anti-BDCA3 antibody (Miltenyi, #130-090-513), PE-labeled anti-BDCA1 antibody (BioLegend, #331517), APC-labeled anti-CD123 antibody (Miltenyi, #130-090-901), and APC-labeled anti-HLA-DR antibody (BioLegend, #307617). 170 μL of each sample was then applied to a FACSCanto II cell analyzer (BD Bioscience) to count the number of cells.

Both humanized antibodies inhibited migration of BDCA3+ dendritic cells, as is the case with the chimeric antibody (Fig. 6).

Example 3

Pharmacological Effect of Mouse Anti-XCR1 Antibody

Pharmacological effect of anti-human XCR1 mouse monoclonal antibody (5G7) prepared in Example 2 above was confirmed using a mouse model of delayed-type contact dermatitis (DTH).

(1) Effect of Mouse Anti-human XCR1 Antibody on Ear Swelling of DNFB (Dinitrofluorobenzene)-Sensitized Mice

Experimental Method

1. Sample Mice

Human XCR1 knock-in mice (mice whose XCR1 gene has been

replaced with human XCR1 gene) on C57BL/6 background between the ages of 7 weeks and 12 weeks were used for the experiment.

2. Method for Preparing DNFB for Sensitization and DNFB for Induction

DNFB for sensitization and induction was prepared by mixing DNFB to a 4:1 mixture of acetone and olive oil to obtain a concentration of 0.5%. Further, a 4:1 mixture of acetone and olive oil was used as a control solution for induction.

3. Method for Administering DNFB

The abdominal hair of the mice was shaved to expose the skin, and 50 µL of 0.5% DNFB for sensitization was applied thereto. On the following day, 50 µL of 0.5% DNFB was applied again to the same site. 4 days after the application, 25 µL of 0.5% DNFB for induction was applied to the front side of the right ear of the mice. At the same time, as a control, 25 µL of the control solution obtained by mixing acetone and olive oil at a 4:1 ratio was applied to the front side of the left ear of the mice.

4. Method for Administering Antibodies

Anti-human XCR1 mouse monoclonal antibody (5G7) and its control antibody, i.e., mouse IgG (Jackson Laboratory), were prepared in PBS to a final concentration of 2 mg/mL. The day when the first sensitization was conducted was defined as Day 0. Each of the above antibodies was intraperitoneally administered into the mice in an amount of 250 µL/mouse (500 ug/mouse) on Day -1, Day 1, and Day 4.

5. Method for Measuring the Ear Swelling of DNFB-Sensitized Mouse Model

On the first day and the following day, the mice were sensitized by applying 50 µL of 0.5% DNFB to the exposed skin of the abdomen. 4 days after the sensitization, the ear thickness

was measured using a caliper. After measurement, 25 μL of 0.5% DNFB was applied to the front side of the right ear of the mice for induction. Further, as a control, 25 μL of the control solution formed by mixing acetone and olive oil at a 4:1 ratio was applied to the left ear of the mice. The ear thickness was measured 24 hours and 48 hours after induction. The swelling was determined by converting measured values by the following formula.

Formula

Ear thickness changed by DNFB (swelling: mm) = ([A]—[B])-([C]—[D])

[A]: thickness of right ear after induction (mm)

[B]: thickness of right ear before induction (mm)

[C]: thickness of left ear after application of control solution (mm)

[D]: thickness of left ear before application of control solution (mm)

Experimental Results and Analysis

Fig. 8 clearly shows a significant suppression of ear swelling 24 after induction by DNFB in the mice administered with anti-human XCR1 mouse monoclonal antibody (5G7), compared to the mice administered with the control antibody (Fig. 8A). The effect also showed significant suppression in a similar manner 48 hours after induction by DNFB (Fig. 8B).

Although the antibody was systemically administered intraperitoneally, the swelling was suppressed in the ear induced by DNFB. Therefore, it is presumed that the antibody transferred from the abdominal cavity into the blood and, along with the blood flow, reached an inflammatory site or a lymph node, where the antibody demonstrates the effect of suppressing ear swelling.

This suggests that the antibodies of the present invention have a specific effect on the inflammatory site in a site-specific manner.

Example 4

Reactivity of Mouse Anti-Human XCR1 Monoclonal Antibody (5G7) to Various Human Chemokine Receptors

The reactivity of mouse anti-human XCR1 monoclonal antibody (5G7) to various human chemokine receptors was evaluated by FACS analysis. Parent B300.19 cells and human chemokine receptor-EGFP-expressing B300.19 cells (XCR1, CXCR1, CXCR3, CXCR4, CXCR5, CXCR6, CCR1, CCR2, CCR3, CCR4, CCR5, CCR6, CCR7, CCR8, CCR9, CCR11, or CX3CR1, and) were suspended in a FACS buffer (PBS⁻ (Sigma) containing 1 % fetal bovine serum). The cells were blocked for 20 minutes on ice with a blocking buffer (a FACS buffer containing 100 μg/mL of human immunoglobulin). The cells were then incubated for 30 minutes on ice with the blocking buffer containing 5G7 or mouse isotype control antibody IgG2b (eBioscience, #14-4732-82) at a concentration of 10 μg/mL. The cells were washed three times with the FACS buffer, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted at 1:50 in the blocking buffer). The cells were washed with the FACS buffer three times, and then suspended in the FACS buffer. The fluorescence intensity was measured using a FACSCanto II cell analyzer.

Anti-human XCR1 antibody 5G7 showed a high reactivity to human XCR1-EGFP-expressing B300.19 cells. Further, anti-human XCR1 antibody 5G7 showed a very low reactivity to human CX3CR1-EGFP-expressing B300.19 cells, and no reactivity to other human chemokine receptors-EGFP-expressing cells (Fig. 9). On the other hand, mouse isotype control antibody did not show reactivity to any B300.19 cells.

Example 5

Cytotoxicity of anti-human XCR1 antibodies using saporin-conjugated Fab anti-mouse IgG secondary antibody to human XCR1 expressing cells

In order to demonstrate the cytotoxic activity of anti-human XCR1 antibodies to XCR1-expressing cells, cytotoxicity of mouse anti-human XCR1 mAbs to cells on which human XCR1 is exogenously expressing was examined by using saporin-conjugated Fab anti-mouse IgG secondary antibody.

2 x 10^3 cells of B300.19 parent cells or human XCR1-EGFP-expressing B300.19 cells in 80 µL of RPMI1640 (Invitrogen, #11875-093) containing 10% fetal bovine serum (Cell Culture Bioscience, #171012), 100 µg/ml of kanamycin sulfate (Invitrogen, #15160-054) and 50 µM 2-Mercaptoethanol (2-ME, Invitrogen, #21985-023) were added into each well of a 96 well plate. Mouse anti-human XCR1 antibodies (2H6, 5G7, or 11H2), mouse isotype control antibodies, IgG2a, κ (eBioscience, #16-4724-85) or IgG2b, κ (eBioscience, #16-4732-85), were diluted with RPMI1640 containing 10% fetal bovine serum, 100 µg/ml of kanamycin sulfate and 50 µM 2-ME and 10 µl of the diluted antibodies were added to the cells at various concentrations from 0 to 0.17 µg/ml. The cells were then incubated in a 5% CO_2 incubator at 37°C for 20 min. Then, saporin-conjugated Fab anti-mouse IgG (Advanced Targeting Systems, #IT-48) was diluted to 10 µg/ml with RPMI1640 containing 10% fetal bovine serum, 100 µg/ml of kanamycin sulfate and 50 µM 2-ME, and 10 µl of the diluted saporin-conjugated Fab anti-mouse IgG was added to each well at the final concentration of 1 µg/ml. The cells were then incubated in a 5% CO_2 incubator at 37°C for 72 hrs. Then the number of cells in each well was measured by using Cell Count Reagent SF (Nacalai tesque, 07553-15 or -44). The reagent was added to each well and the cells were incubated in a 5% CO_2 incubator at 37°C for 2 or 3 hrs. OD_{450} was then measured with a plate reader (Arvo, PerkinElmer).

Mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) with saporin-conjugated secondary antibody showed growth suppression of human XCR1-EGFP-expressing B300.19 cells (Fig. 11). The IC_{50} values of 2H6, 5G7, and 11H2 calculated by Graphpad Prism software were 0.141 nM, 0.017 nM, and 0.155 nM, respectively. On

the other hand, these antibodies with saporin-conjugated secondary antibody did not show cell growth suppression of parent B300.19 cells. Control antibodies with saporin-conjugated secondary antibody did not suppress cell growth of human XCR1-EGFP-expressing B300.19 cells nor parent B300.19 cells (Fig. 11). These findings indicate that mouse anti-human XCR1 antibodies, 2H6, 5G7, and 11H2, were internalized with saporin-conjugated secondary antibody and acted as an immunotoxin.

Example 6

Effects of 5G7 Mab on cytotoxitic T lymphocyte assay *in vivo*

In order to investigate the inhibitory activity of 5G7 Mab on the CTL function, CTL assay was performed.

The engineered hXCR1-knocked-in mice, in which human XCR1 is expressed instead of mouse XCR1, were immunized subcutaneously with ovalbumin (200 µg/head) emulsified with CFA on day 0. The 5G7 Mab or the control mouse IgG (Jackson Laboratory), was intraperitoneally injected at a dose of 500 ug/head on day -1, day 2 and day 5. Six days later, splenocytes from naïve C57BL/6 mice were incubated for 30 min at 37°C with or without 10 µg/ml $OVA_{257-264}$ peptide (SIINFEKL; MBL).

These peptide-pulsed target and non-target cell populations were labeled with 2.5 and 0.25 µM CFSE (Invitrogen Life Technologies), respectively, then mixed at a 1:1 ratio, and injected intravenously into the immunized mice.

One day after the injection of CFSE-labeled splenocytes, the target cell-killing activity was evaluated using the ratio of CFSE-positive populations in the spleen as follows.

The CFSE-positive cells in the spleen in the immunized mice were detected by flow cytometry, and the CTL activity of each mouse was calculated with the ratio of $CFSE^{high}$ cells and $CFSE^{low}$ cells as follows: CTL activity = (% of $CFSE^{high}$/% of $CFSE^{low}$).

Then, the relative CTL activity was calculated as follows: Relative CTL activity = (CTL activity in each immunized mouse)/(CTL activity in control mouse).

The results showed that the relative CTL activity in the mice treated with 5G7 Mab showed lower relative CTL activity as compared to that in the mice treated with the control IgG (Fig. 12).

The data indicated the suppression of the *in vivo* CTL activity by the treatment with anti-XCR1 antibody, and suggested that the treatment with anti-XCR1 antibodies may be beneficial for immune diseases, such as graft rejection, GVHD and tissue injury in autoimmune diseases.

Example 7

Reactivity of Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2) to the Chimeric Human/Mouse XCR1-Expressing Cells

To determine epitopes of human XCR1 recognized by mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2), reactivity of these antibodies to chimeric human/mouse XCR1-expressing cells was evaluated.

Because mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) reacted to human XCR1 but not to mouse XCR1, a panel of human/mouse XCR1 chimeric receptors was prepared. In this panel, each extracellular domain of human XCR1 was replaced by the homologous region of mouse XCR1, and vice versa. Expression vectors of this panel were constructed using an overlapping extension polymerase chain reaction (PCR) method. Each chimeric receptor-EGFP was expressed in TK-1 cells and mAb reactivity was determined by FACS analysis. Parent TK-1 cells, human XCR1-EGFP-, mouse XCR1-EGFP-, or chimeric XCR1-EGFP-expressing TK-1 cells were suspended in a FACS buffer (PBS^- (Sigma) containing 1% fetal bovine serum). The cells were blocked for 10 minutes on ice with a FACS buffer containing 100 μg/mL of human immunoglobulin. The cells were then incubated for 20 minutes on ice with the anti-human XCR1 antibodies (2H6, 5G7, or 11H2) at various concentrations from 0 to 10 μg/mL, mouse isotype control antibodies, IgG2a (eBioscience, #14-4724-82) or IgG2b (eBioscience, #14-4732-82), at a concentration of 10 μg/mL, or a

FACS buffer without antibody. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted at 1:50 in the FACS buffer) or PE-labeled anti-human XCR1 polyclonal antibody (R&D, #FAB857P, diluted at 2:5 in the FACS buffer, used for cells that had been incubated with the FACS buffer without antibody). The cells were washed with the FACS buffer three times, and then suspended in the FACS buffer. The fluorescence intensity was measured by a FACSCanto II cell analyzer.

Mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) showed the reactivity to human XCR1-EGFP-expressing TK-1 cells, but not to parent TK-1 cells or mouse XCR1-EGFP-expressing TK-1 cells (Fig. 13; the origins of the four extracellular domains were designated by four-letter codes (e.g., HHHH is wild-type human XCR1, Hmmm has human N-terminal extracellular domain and mouse first, second, and third extracellular loops, etc.)). These three antibodies showed reactivity to chimeric XCR1s, having the human XCR1 N-terminus, -EGFP-expressing TK-1 cells. Reactivity to chimera receptor, mmHm, was also examined in another experiment, and reactivity was not observed (data not shown).

In contrast, the mouse isotype control antibodies did not show reactivity to any TK-1 cells (data not shown).

Example 8

Mapping of Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2)-Binding Sites on the Extracellular Domains of Human XCR1 by Peptide ELISA

To define the contact residues of anti-human XCR1 antibodies (2H6, 5G7, and 11H2) on human XCR1 extracellular domains, peptide scan analysis was performed using sets of 12-mer peptides covering the extracellular domains of human XCR1.

Two sets of peptides with biotin and spacer GSGS at N-terminal were synthesized by Sigma. The first set of 13 peptides

contained all possible 12-mers from the human XCR1 N terminus, each offset by 2 amino acids. The second set of 13 peptides contained all possible 12-mer from the human XCR1 extracellular loops, each offset by 3 amino acids. Peptides were initially reconstituted in 100% dimethyl sulfoxide and subsequently diluted in 30% dimethyl sulfoxide solution to give a final concentration of 50 µg/mL for direct ELISA.

Streptavidin-coated microtiter plates (Perkin Elmer) were coated with 50 µg/mL of peptide per well in a volume of 50 µL, and incubated at room temperature for 1 hour. The peptide solution was removed and PBS^- containing 4% Block-Ace was added to each well and incubated overnight at 4°C. Each well was washed three times with an ELISA wash buffer (0.02% Tween20 in PBS-). Anti-human XCR1 antibodies (2H6, 5G7, or 11H2) were added to each well in amount of 10 µg/mL, and incubated for 6 hours at room temperature. Each well was washed three times with the ELISA wash buffer. Horseradish peroxidase-conjugated donkey anti-mouse IgG antibody (Jackson, #715-035-150), diluted 1:5,000 in the ELISA wash buffer, was added to each well and incubated for 1 hour at room temperature. Each well was washed three times with the ELISA wash buffer. TMBZ (3,3',5,5' tetramethyl benzidine; Sigma) was added to each well and incubated at room temperature. The reaction was stopped with 2N H_2SO_4, and A_{450nm} was measured by Arvo plate reader (PerkinElmer).

Anti-human XCR1 antibodies 2H6 and 5G7 showed strong binding to one peptide containing ^{7}PESTTFFYYDLQ^{18} (SEQ ID NO: 96), and weak binding to ^{11}TFFYYDLQSQPC^{22} (SEQ ID NO: 110) (Fig. 14). 5G7 also showed weak binding to three non-sequential peptides containing ^{19}SQPCENQAWVFA^{30} (SEQ ID NO: 101), ^{172}SSGCDYSELTWY^{183} (SEQ ID NO: 110), and ^{175}CDYSELTWYLTS^{186} (SEQ ID NO: 111). On the other hand, 11H2 showed no reactivity to these peptides (data not shown).

Example 9

Mapping of Binding Residues of Mouse Anti-Human XCR1 Antibodies

(2H6, 5G7, and 11H2) and Humanized Anti-Human XCR1 Antibodies (HK1L2 and HK5L5) on Human XCR1 Extracellular Domains by Using Alanine Mutants

To determine the critical residues of human XCR1 recognized by mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) and humanized anti-human XCR1 antibodies (HK1L2 and HK5L5), alanine substitution assay was performed.

A panel of alanine substitution mutants of human XCR1 was prepared. In this panel, each amino acid in ^{7}PESTTFFYYDLQSQPCENQAWVFA^{30} (SEQ ID NO: 118) and ^{175}CDYSELTWYLTS^{186} (SEQ ID NO: 119) of human XCR1 extracellular regions were replaced by alanine. Expression vectors for alanine substitution mutants were constructed by using site-directed mutagenesis. Each mutant was expressed on B300.19 cells, and antibody reactivity was determined by FACS analysis. Parent B300.19 cells and human XCR1-EGFP- or each alanine mutant human XCR1-EGFP-expressing B300.19 cells were mixed at a 1:1 ratio and suspended in a FACS buffer (PBS^{-} (Sigma) containing 1% fetal bovine serum). The cells were blocked for 10 minutes on ice with a FACS buffer containing 10% rat serum. The cells were then incubated for 20 minutes on ice with mouse anti-human XCR1 antibodies (2H6, 5G7, or 11H2), humanized antibodies (HK1L2 or HK5L5), mouse isotype control antibodies, IgG2a (eBioscience, #14-4724-82) or IgG2b (eBioscience, #14-4732-82), or human isotype control antibody IgG2 (Sigma, #I5404), at a concentration of 10 μg/mL; or incubated with a FACS buffer without antibody. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted at 1:50 in the FACS buffer, used for cells that had been incubated with mouse antibodies), PE-labeled anti-human IgG polyclonal antibody (Jackson, #709-116-149, diluted at 1:50 in the FACS buffer, used for cells which had been incubated with humanized antibodies or human control IgG), or PE-labeled anti-human XCR1 polyclonal

antibody (R&D, #FAB857P, diluted at 2:5 in the FACS buffer, used for cells that had been incubated with the FACS buffer without antibody). The cells were washed three times with the FACS buffer, and then suspended in the FACS buffer. The fluorescence intensity was measured using a FACSCanto II cell analyzer (BD Bioscience).

Each alanine mutant was detected by PE-labeled anti-human XCR1 polyclonal antibody, except for C175A mutant (Fig. 15). Because expression amounts of each alanine mutants on cell surface were varied among these mutants as shown in Fig. 15, reactivity of antibodies to each alanine mutant was evaluated by a relative PE mean value (mAb/pAb), calculated as per the following procedure. At first, a relative PE mean value for each antibody was calculated by setting the PE mean value, which was obtained by staining of human XCR1-EGFP-expressing B300.19 cells (wild type) using each antibody, as 1.0. The relative PE mean values (mAb/pAb) were then calculated by the following equation: each relative PE mean values for mouse anti-human XCR1 antibodies (2H6, 5G7, or 11H2) or humanized antibodies (HK1L2 or HK5L5) was divided by the relative PE mean values for PE-labeled anti-human XCR1 polyclonal antibody. The results showed that 2H6 (Fig. 16), 5G7 (Fig. 17), HK1L2 (Fig. 19), and HK5L5 (Fig. 20) showed lower reactivity to many alanine mutants in which each residue in N-terminus or 2nd loop was replaced with alanine. In particular, no reactivity or weak reactivity to Y14A, D16A, and L17A mutants were observed. Additionally, reactivity to E8A, F13A, C22A, and Y177A were lower among these mutants. Taken together, these results indicate that 2H6, 5G7, HK1L2, and HK5L5 recognize E8, F13, Y14, D16, L17, C22 and Y177 on human XCR1 extracellular domain. 11H2 (Fig. 18) showed similar reactivity to other mAbs except for F13A and D16A, indicating that 11H2 binds to E8, Y14, L17, C22, and Y177.

Example 10

Competition among Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2), Recognizing Similar Epitopes, for Binding to Human XCR1-

Expressing Cells

To determine whether anti-human XCR1 antibodies, recognizing similar epitopes, compete with each other for binding to human XCR1, a competition assay was performed.

The competition assay was performed as per the following procedure. Parent B300.19 cells and human XCR1-EGFP-expressing B300.19 cells were mixed at a 1:1 ratio and suspended in a FACS buffer (PBS^- (Sigma) containing 1% fetal bovine serum). The cells were blocked for 10 minutes on ice with the FACS buffer containing 10% rat serum. The cells were then incubated with mouse anti-human XCR1 antibodies (2H6, 5G7, or 11H2), mouse isotype control antibodies, IgG2a (eBioscience, #16-4724-85) or IgG2b (eBioscience, #16-4732-85), at various concentrations from 0 to 10 µg/mL in the FACS buffer for 20 minutes on ice. The cells were then incubated with biotinylated mouse anti-human XCR1 antibody (5G7) at a concentration of 0.3 µg/mL in the FACS buffer for 20 minutes on ice. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled streptavidin (BD Pharmingen, #554061, diluted with the FACS buffer at a dilution factor of 1:50). The cells were washed three times with the FACS buffer, and then suspended in the FACS buffer. Fluorescence intensity was measured using a FACSCanto II cell analyzer (BD Bioscience).

Binding of biotinylated mouse anti-human XCR1 antibody (5G7) to human XCR1-EGFP-expressing B300.19 cells was competed with unlabeled 5G7 itself, unlabeled 2H6 and 11H2, recognizing similar epitopes on human XCR1 (Fig. 21). On the other hand, control antibodies did not compete with biotinylated antibody (5G7) for binding to human XCR1-EGFP-expressing B300.19 cells.

Example 11

Reactivity of Mouse Anti-Human XCR1 Monoclonal Antibody, 5G7, and Humanized Anti-Human XCR1 Monoclonal Antibodies, HK1L2 and HK5L5, to Various Human Chemokine Receptors

The reactivity of mouse anti-human XCR1 monoclonal antibody, 5G7 and humanized anti-human XCR1 monoclonal antibodies, HK1L2 and HK5L5 to various human chemokine receptors were evaluated by FACS analysis.

Parent B300.19 cells and human chemokine receptor-EGFP-expressing B300.19 cells (XCR1, CXCR1, CXCR3, CXCR4, CXCR5, CXCR6, CCR1, CCR2B, CCR3, CCR4, CCR5, CCR6, CCR7, CCR8, CCR9, CCR11, or CX3CR1) were suspended in a FACS buffer (PBS^- (Sigma) containing 1% fetal bovine serum) at a concentration of 1 x 10^6 cells/mL and aliquots of 100 µl were dispensed into wells of a 96 well round bottom plate. The cells were then centrifuged, and supernatants were discarded. Mouse anti-human XCR1 mAb, 5G7, mouse isotype control antibody IgG2b (eBioscience, #14-4732-82), humanized anti-human XCR1 monoclonal antibodies, HK1L2 and HK5L5, and control human IgG (Mitsubishi, #128-26053-9) were diluted with the FACS buffer at a concentration of 5 µg/mL. PE-labeled goat anti-human XCR1 polyclonal antibodies (R&D, #FAB857P, and LifeSpan BioScience, #LS-C76885) were diluted with the FACS buffer at the dilution factors of 2:5 and 1:5, respectively. Fifty µL of the diluted antibodies were added to each well, and the cells were incubated for 20 minutes on ice. The cells were then washed three times with the FACS buffer. PE-labeled anti-mouse IgG polyclonal antibody (Jackson, #715-116-151, diluted with the FACS buffer at a dilution factor of 1:50) was added to the cells that had been incubated with 5G7 or mouse isotype control antibody. PE-labeled anti-human IgG polyclonal antibody (Jackson, #709-116-149, diluted with the FACS buffer at a dilution factor of 1:50) was added to the cells that had been incubated with HK1L2, HK5L5 or human control IgG. The FACS buffer was added to the cells that had been incubated with anti-hXCR1 polyclonal antibodies. The cells were then incubated for 20 minutes on ice. The cells were washed with the FACS buffer three times, and then suspended in the FACS buffer. Fluorescence intensity was measured using a FACSCanto II cell analyzer, and expressed as a delta PE mean value. The delta PE mean value was

calculated by subtracting background PE mean value from each PE mean value, which was obtained by staining each cell line with each antibody.

Mouse anti-human XCR1 antibody, 5G7 selectively reacted to human XCR1-EGFP-expressing B300.19 cells except for human CX3CR1-EGFP-expressing cells (Fig. 22). On the other hand, goat anti-human XCR1 polyclonal antibodies reacted to various human chemokine receptor-EGFP-expressing cells in addition to human XCR1-EGFP-expressing cells (Fig. 22). Humanized anti-human XCR1 antibodies, HK1L2 and HK5L5 showed reduced reactivity to human CX3CR1-EGFP-expressing cells in spite of their high reactivity to human XCR1-EGFP-expressing cells (Fig. 23).

Example 12

Effect of 5G7 Mab on Mycobacterium butyricum-induced DTH response

It is known that a delayed-type hypersensitivity(DTH) response is one of the main mechanisms causing autoimmune diseases such as thyroiditis, rheumatoid arthritis and type 1 diabetes when this response is directed against self-antigens (Actor, J.K. and Ampel, N.M. (December 2009) Hypersensitivity: T Lymphocyte-mediated (Type IV). In: Encyclopedia of Life Sciences (ELS). John Wiley & Sons, Ltd: Chichester). T cell-Dendritic cell interaction is critical for DTH responses. Thus the inhibition of T cell-DC interaction is believed to be useful to treat those diseases. We investigated the effect of the anti-human XCR1, 5G7 Mab, on a model of DTH reaction, Mycobacterium (M.) butyricum-induced DTH response, in human XCR1 knocked-in mice (Mihara, M. et al, Immunology Letters 2002, 84: 223-229; Mohan K et al, Eur. J. Immunol. 2005, 35: 1702-1711) .

(Methods)

The engineered hXCR1-knoked-in mice, in which human XCR1 is expressed instead of mouse XCR1, were immunized subcutaneously with heat-killed M. butyricum (100 μg/head) with mineral oil on

day 0. A 5G7 Mab or a control mouse IgG (Jackson Laboratory), were intraperitoneally injected at the dose of 500 μg/head on day 1, day 3, day 7 and day 9. 10 days after the immunization with M. butyricum, the mice were challenged with M. butyricum suspended in mineral oil on the right footpad (20 μg/foot, M. butyricum challenge), and mineral oil on alone left footpad (control challenge). One day after the challenge injection, the DTH response was evaluated by measuring the footpad thickness of each footpad. The footpad swelling was calculated according to the following formula.

Footpad swelling = ([A]-[B])-([C]-[D])

[A] = thickness of right footpad after M. butyricum challenge

[B] = thickness of right footpad before M. butyricum challenge

[C] = thickness of left footpad after control challenge

[D] = thickness of left footpad before control challenge

(Results)

The result showed that the M. butyricum-induced DTH response in mice treated with 5G7 Mab showed significantly lower DTH response compared to the mice treated with the control IgG (Fig. 24).

(Conclusion)

The data showed the efficacy of anti-XCR1 antibody treatment in the DTH response. It is suggested that the use of anti-XCR1 antibodies may be beneficial in the treatment of DTH-driven autoimmune diseases such as thyroiditis, rheumatoid arthritis and type 1 diabetes.

Example 13

Effect of 5G7 Mab on MOG 37-50 peptide mediated EAE

Multiple sclerosis (MS) is a chronic demyelinating disease of the human central nervous system (CNS) which can be characterized clinically by a remitting-relapsing or a chronic progressive

course. The most intensively studied animal model of MS, experimental autoimmune encephalomyelitis (EAE), classically leads to deficit in motor functions. Many reports showed that T cells play crucial roles in the pathogenesis of MS and EAE. Therefore, we performed an EAE model experiment to investigate the inhibitory activity of 5G7 Mab on the pathogenesis of MS.

(Experimental Method)

1. Sample Mice

Human XCR1 knock-in mice (7-12 weeks old), in which human XCR1 is expressed instead of mouse XCR1 on C57BL/6 background, were used for the experiment.

2. Induction of EAE

The induction of EAE was performed according to the method reported in the journal Eur. J. Immunol. 2005, 35: 76-85, in which the probable role of CD8+ T cells was indicated in the EAE development. Briefly, the human XCR1 knock-in mice were injected subcutaneously with 200 μg of myelin oligodendrocyte glycoprotein 37-50 peptide (MOG 37-50) emulsified in Freund's complete adjuvant (CFA) containing 20 mg/ml of Mycobacterium tuberculosis H37Ra. 200 ng of pertussis toxin was administered intravenously on days 0 and 2, post-immunization.

3. Method for Administering Antibodies

Anti-human XCR1 mouse monoclonal antibody (5G7) and its control antibody, i.e., mouse IgG (Jackson Laboratory), were prepared in PBS to a final concentration of 2 mg/mL. Each of the above antibodies was intravenously administered into the mice with the volume of 250 μl/mouse (500 μg/mouse) on day 7, day 10, day 14 and day 17.

4. Scoring of the pathology of this model

Clinical symptom of EAE was monitored from the day of the immunization, and was scored on a scale of 0-5, based on the

following criteria:

grade 0: no disease, grade 0.5: mild-tail paralysis, grade 1: tail paralysis, grade 2: uneven gait, grade 2.5: one paralyzed rear leg, grade 3: rear limb paralysis, grade 4: paralyzed front and rear legs: and grade 5: moribund or death.

(Experimental result and conclusion)

The obtained clinical scores of the mice administered with 5G7 Mab showed lower levels than those in the mice administered with the control IgG (Fig. 25). The data indicated that the treatment with anti-XCR1 antibody showed a certain level of suppression in the EAE development, and suggested that the treatment with anti-XCR1 antibodies may be beneficial for MS in human.

Example 14

Inhibition of Human XCL1 Binding to Human XCR1-Expressing Cells with Mouse Anti-Human XCR1 Antibodies (2H6, 5G7, and 11H2)

To determine whether mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2) inhibit human XCL1 binding to human XCR1, a competitive ligand binding assay was performed.

First, the binding of human XCL1-SSS-His(10) to human XCR1-EGFP-expressing BaF3 cells were determined according to the following procedure . Parent BaF3 cells and human XCR1-EGFP-expressing BaF3 cells were mixed at a 1:1 ratio, and suspended in a FACS buffer (1% FBS-containing PBS^- (Sigma)). The cells were incubated for 30 minutes on ice with an increasing concentration of human XCL1-SSS-His(10) in the presence or absence of 2.5 μM soluble XCL1 (R&D, #695-LT-025/CF) in the FACS buffer. Next, the cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with anti-6x His tag antibody (BETHYL, #A190-114A, diluted at 1:100 in the FACS buffer). The cells were again washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-rabbit IgG antibody (Jackson, #711-166-152, diluted at 1:50 in FACS buffer). Next, the cells were once again washed three times with the FACS buffer, and then suspended in the FACS buffer. The fluorescence intensity was measured

using a FACSCanto II cell analyzer (BD Bioscience). Specific binding was determined by subtracting the non-specific binding (in the presence of 2.5 μM soluble XCL1) from the total binding (in the absence of 2.5 μM soluble XCL1).

The competitive ligand binding assay was performed according to the following procedure. Parent BaF3 cells and human XCR1-EGFP-expressing BaF3 cells were mixed at a 1:1 ratio, and suspended in a FACS buffer (1% FBS-containing PBS^- (Sigma)). The cells were blocked for 10 minutes on ice with a FACS buffer containing 10% rat serum. The cells were then incubated for 20 minutes on ice with mouse anti-human XCR1 antibodies (2H6, 5G7, or 11H2), mouse isotype control antibodies, IgG2a (eBioscience, #16-4724-85), or IgG2b (eBioscience, #16-4732-85) at various concentrations from 0 to 150 μg/mL. Next the cells were incubated for 30 minutes on ice with human XCL1-SSS-His(10) at a saturating concentration of 0.12 μg/mL. The cells were washed with the FACS buffer three times, and then incubated for 20 minutes on ice with anti-6x His tag antibody (BETHYL, #A190-114A, diluted at 1:100 in FACS buffer). The cells were again washed with the FACS buffer three times, and then incubated for 20 minutes on ice with PE-labeled anti-rabbit IgG antibody (Jackson, #711-166-152, diluted at 1:50 in FACS buffer). Next the cells were once again washed three times with the FACS buffer, and then suspended in FACS buffer. The fluorescence intensity was measured using a FACSCanto II cell analyzer (BD Bioscience).

Human XCL1 binding to human XCR1-EGFP-expressing BaF3 cells was inhibited with mouse anti-human XCR1 antibodies (2H6, 5G7, and 11H2), and the IC_{50} of the antibodies was 37.0, 6.9, and 23.8 nM, respectively. On the other hand, control antibodies did not inhibit human XCL1 binding to human XCR-EGFP-expressing BaF3 cells.

CLAIMS

1. An antibody that binds to human XCR1, wherein the antibody comprises (i) a heavy chain variable region comprising a heavy chain complementarity determining region (CDR) 1 consisting of the amino acid sequence of SEQ ID NO:17, a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:18, and a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:19; and (ii) a light chain variable region comprising a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO:20, a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:21, and a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:22.

2. The antibody according to claim 1, wherein the heavy chain variable region comprises the amino acid sequence of SEQ ID NO: 60 or 64, and the light chain variable region comprises the amino acid sequence of SEQ ID NO: 68 or 72.

3. The antibody according to claim 1, wherein the heavy chain variable region comprises the amino acid sequence of SEQ ID NO: 60, and the light chain variable region comprises the amino acid sequence of SEQ ID NO: 68.

4. The antibody according to claim 1, wherein the heavy chain variable region comprises the amino acid sequence of SEQ ID NO: 64, and the light chain variable region comprises the amino acid sequence of SEQ ID NO: 72.

5. The antibody according to claim 1, wherein the antibody comprises a human constant region.

6. The antibody according to claim 1, wherein the antibody comprises a heavy chain comprising the amino acid sequence of SEQ ID NO: 59, and a light chain comprising the amino acid sequence

of SEQ ID NO: 67.

7. The antibody according to claim 1, wherein the antibody comprises a heavy chain comprising the amino acid sequence of SEQ ID NO: 63, and a light chain comprising the amino acid sequence of SEQ ID NO: 71.

8. The antibody according to claim 1, comprising an Fc region, wherein the Fc region is mutated to induce a change in antibody-dependent cellular cytotoxicity (ADCC) activity.

9. The antibody according to claim 8, wherein the Fc region is mutated to lower ADCC activity.

10. The antibody according to claim 1, wherein the antibody is conjugated to a cytotoxic molecule.

11. The antibody according to claim 1, wherein the antibody inhibits interaction between human XCR1 and human XCL1.

12. The antibody according to claim 1, wherein the antibody inhibits cell migration of dendritic cells.

13. The antibody according to claim 1, wherein the antibody suppresses the activity of cytotoxic T lymphocytes.

14. A pharmaceutical composition comprising the antibody according to claim 1 and a pharmaceutically acceptable carrier or additive.

15. The antibody according to claim 1, wherein the antibody is a humanized antibody.

16. An antibody that binds to human XCR1, wherein the antibody comprises (i) a heavy chain variable region comprising a heavy

chain complementarity determining region (CDR) 1 consisting of the amino acid sequence of SEQ ID NO:41, a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:42, and a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:43; and (ii) a light chain variable region comprising a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO:44, a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:45, and a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:46.

17. A pharmaceutical composition comprising the antibody according to claim 16 and a pharmaceutically acceptable carrier or additive.

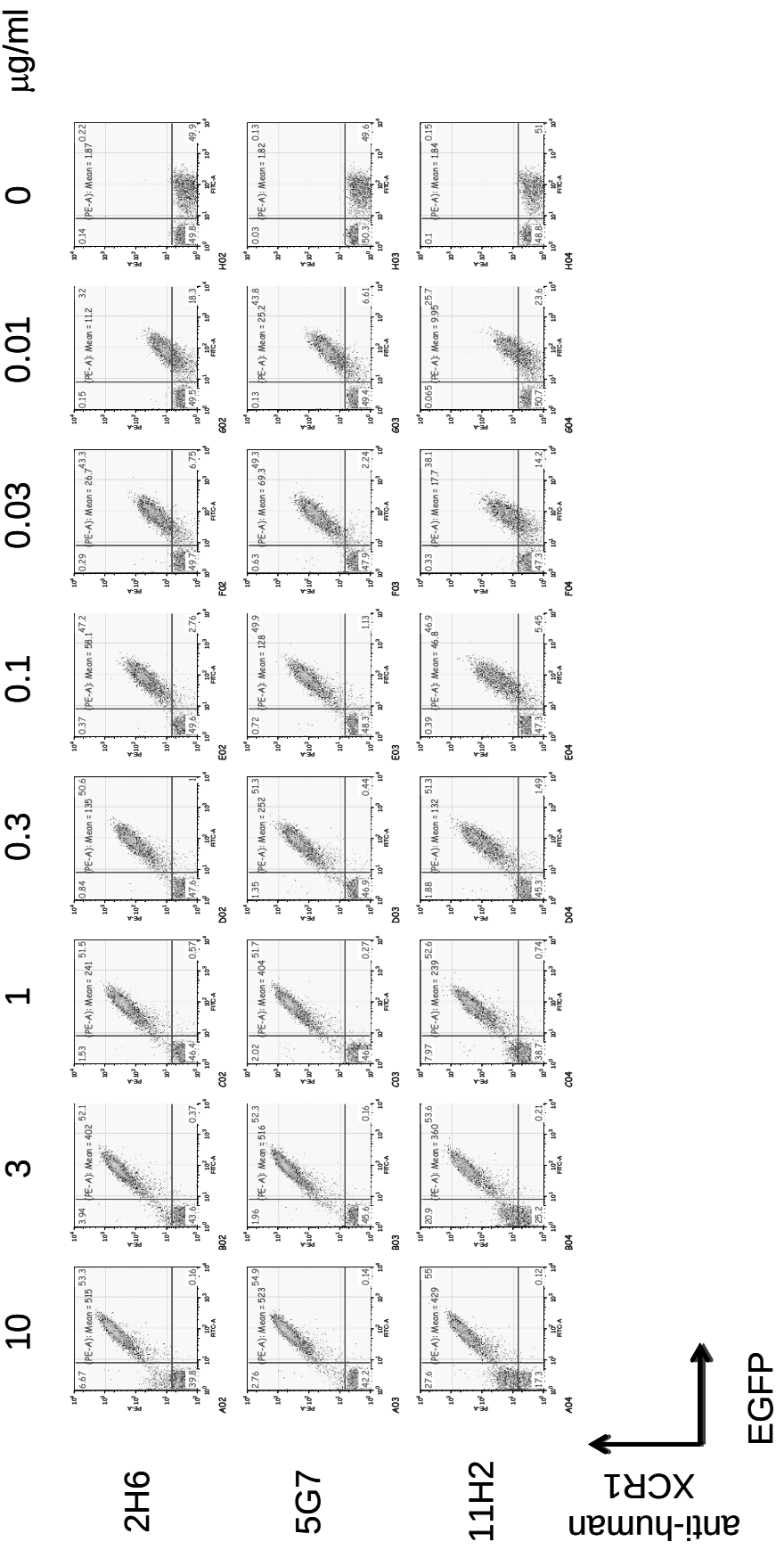
18. The antibody according to claim 16, wherein the antibody is a humanized antibody.

19. An antibody that binds to human XCR1, wherein the antibody comprises (i) a heavy chain variable region comprising a heavy chain complementarity determining region (CDR) 1 consisting of the amino acid sequence of SEQ ID NO:29, a heavy chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:30, and a heavy chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:31; and (ii) a light chain variable region comprising a light chain CDR 1 consisting of the amino acid sequence of SEQ ID NO:32, a light chain CDR 2 consisting of the amino acid sequence of SEQ ID NO:33, and a light chain CDR 3 consisting of the amino acid sequence of SEQ ID NO:34.

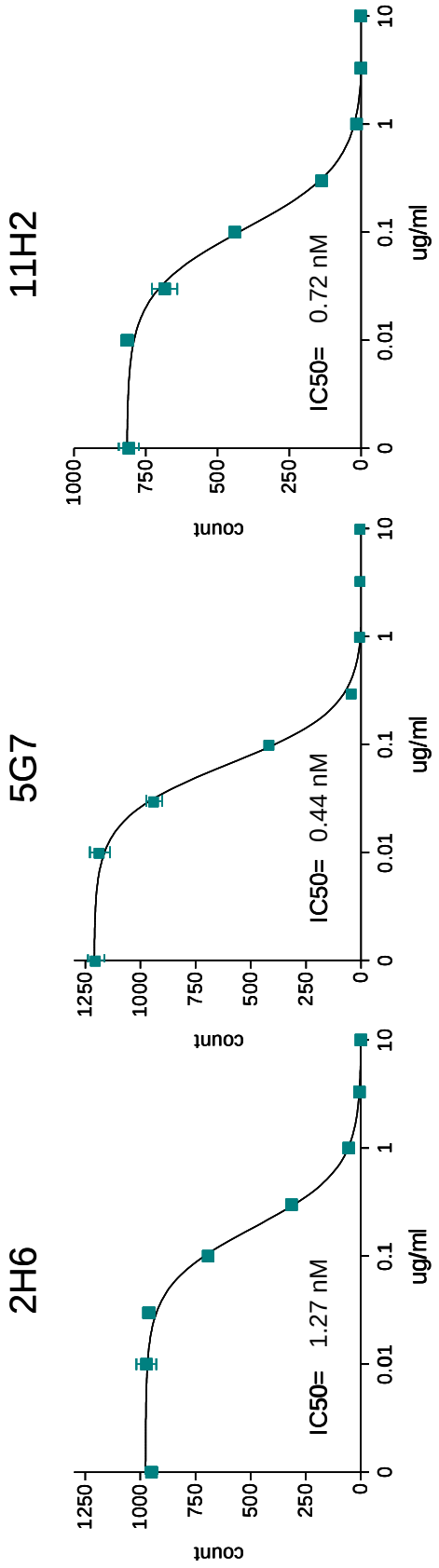
20. A pharmaceutical composition comprising the antibody according to claim 19 and a pharmaceutically acceptable carrier or additive.

21. The antibody according to claim 19, wherein the antibody is a humanized antibody.

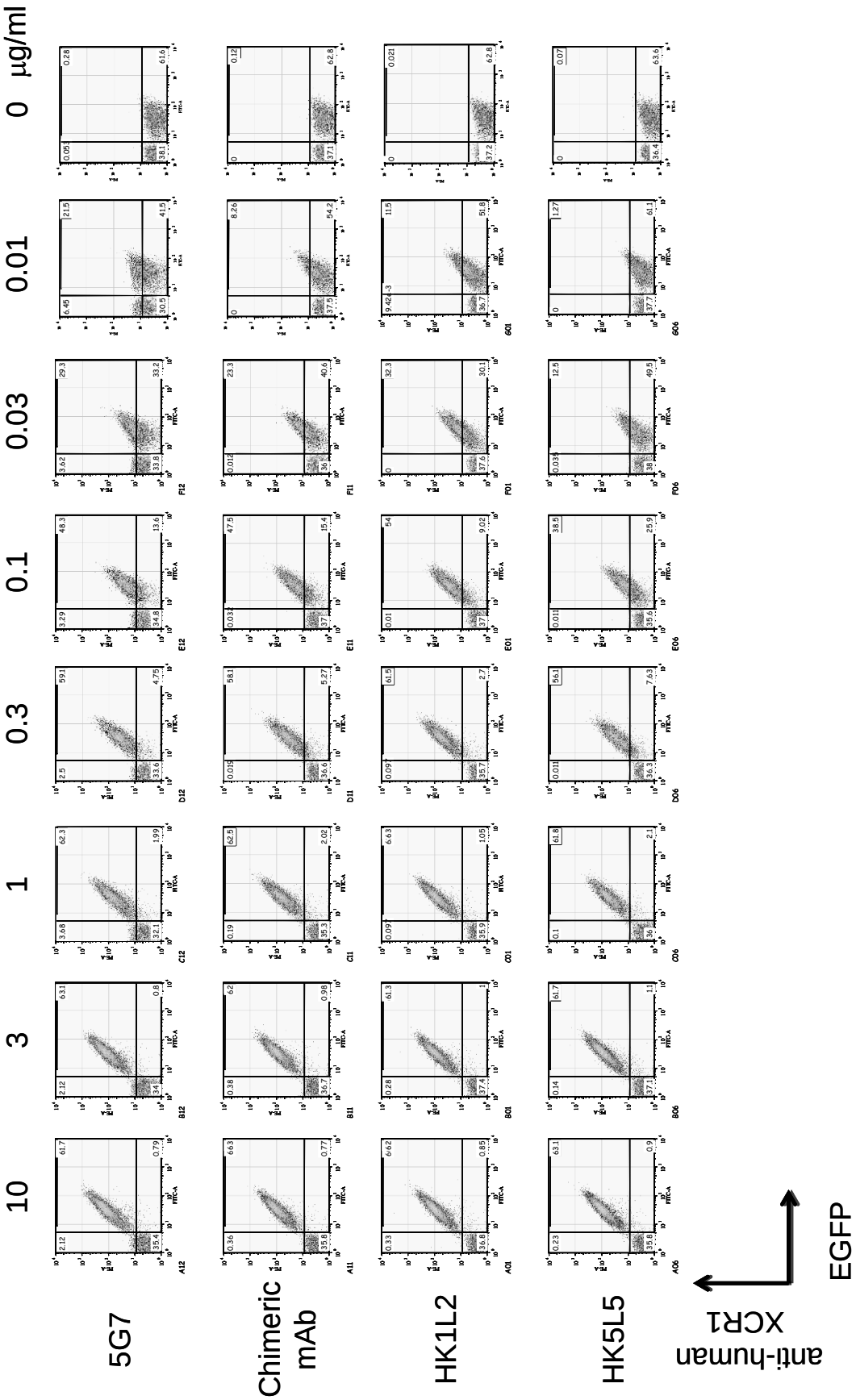
[Fig.1]



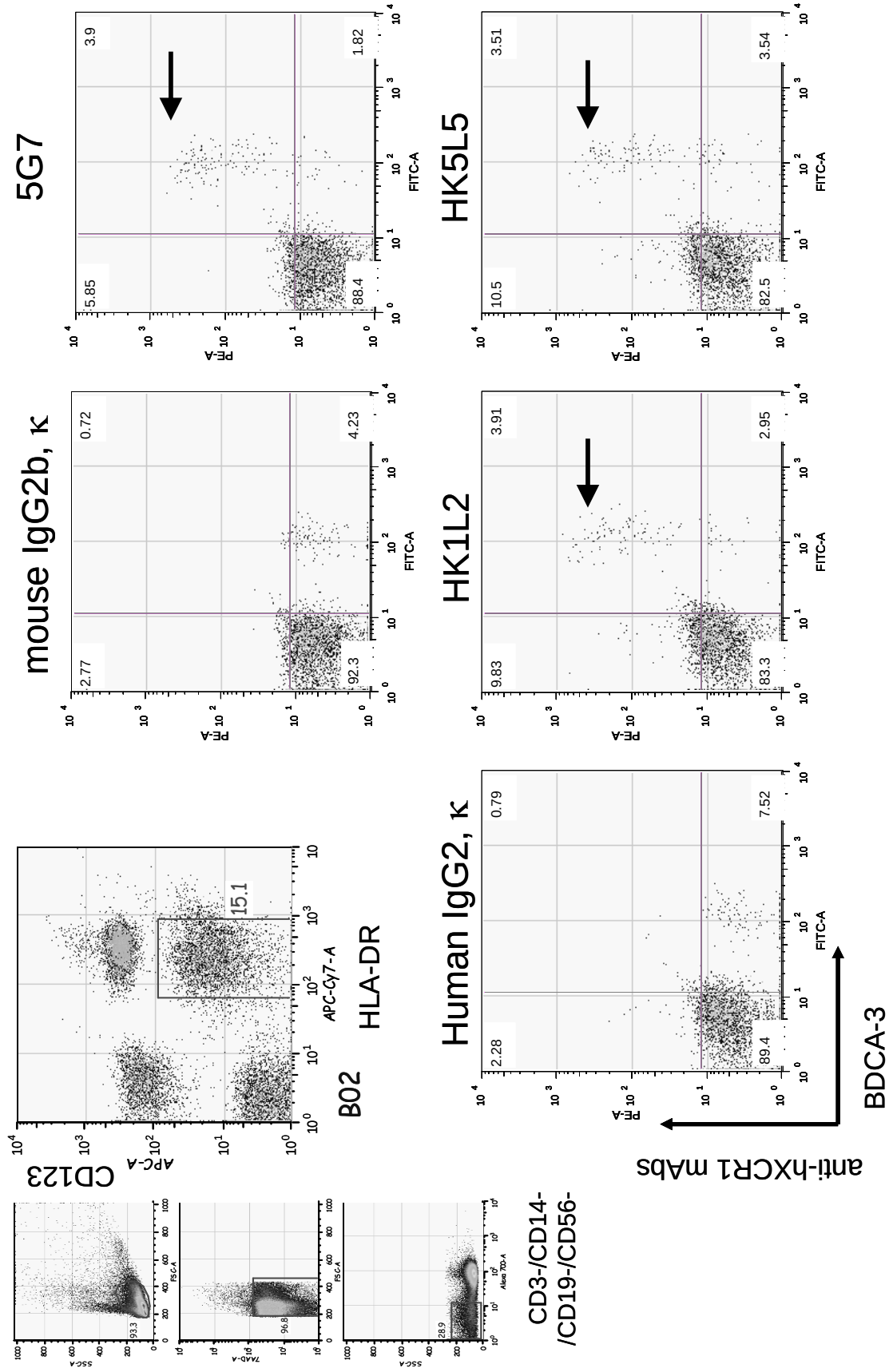
[Fig.2]


2H6
IC50= 1.27 nM
5G7
IC50= 0.44 nM
11H2
IC50= 0.72 nM
count
ug/ml
0
250
500
750
1000
1250
0.01
0.1
1
10

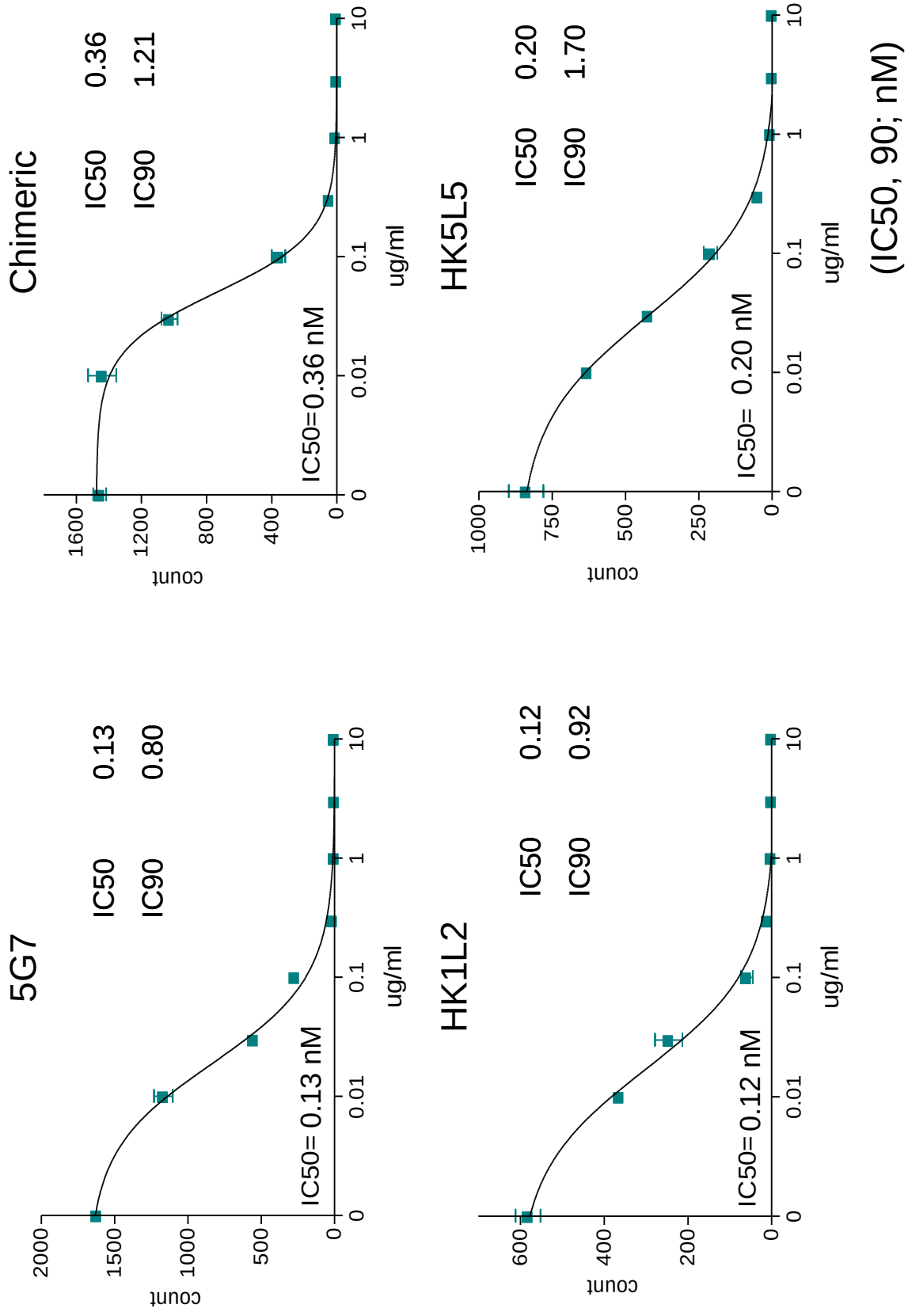
[Fig.3]


10
3
1
0.3
0.1
0.03
0.01
0 μg/ml
5G7
Chimeric mAb
HK1L2
HK5L5
anti-human XCR1
EGFP

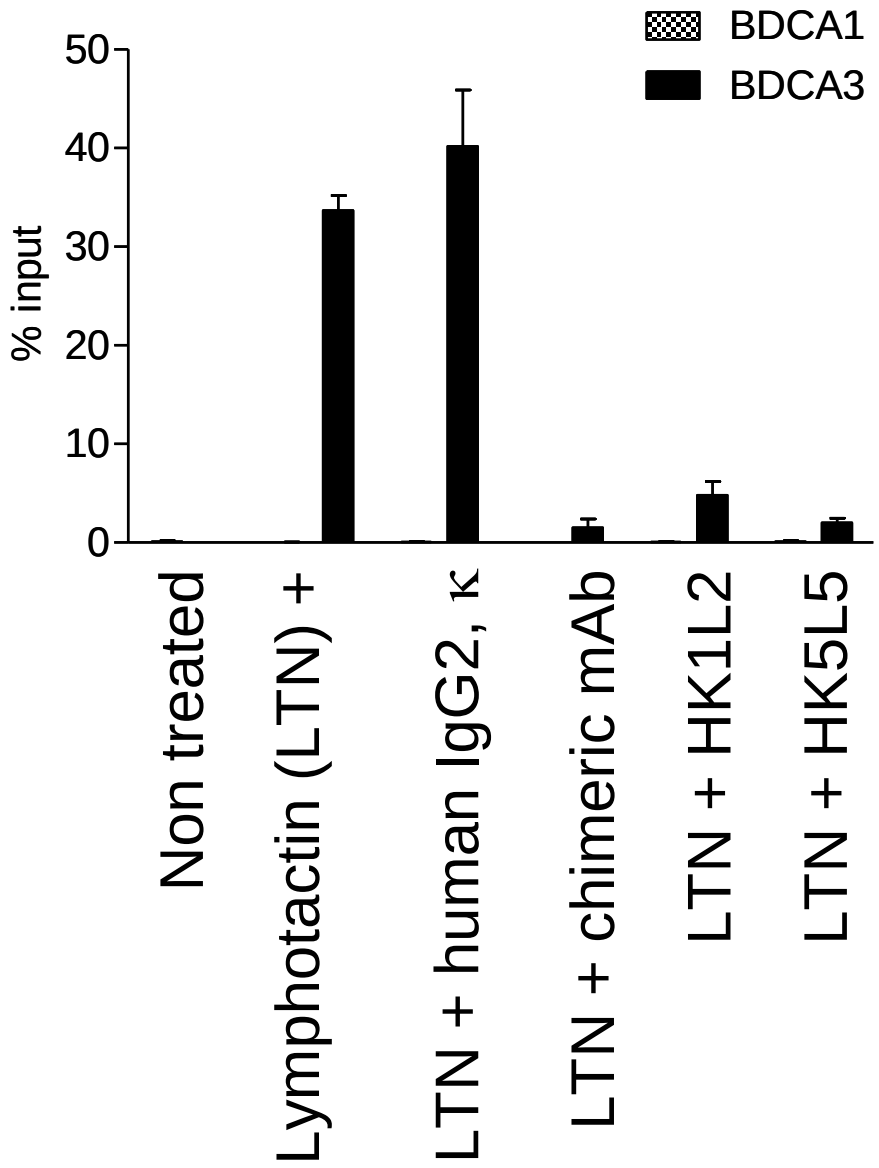
[Fig.4]


5G7
mouse IgG2b, κ
HK5L5
HK1L2
Human IgG2, κ
anti-hXCR1 mAbs
BDCA-3
CD123
HLA-DR
B02
CD3-/CD14-/CD19-/CD56-
FITC-A
PE-A
APC-A
APC-Cy7-A
SSC-A
FSC-A
7AAD-A
15.1
3.9
1.82
5.85
88.4
0.72
4.23
2.77
92.3
3.51
3.54
10.5
82.5
3.91
2.95
9.83
83.3
0.79
7.52
2.28
89.4
93.3
96.8
28.9

[Fig.5]


Chimeric
IC50 0.36
IC90 1.21
IC50=0.36 nM
5G7
IC50 0.13
IC90 0.80
IC50= 0.13 nM
HK5L5
IC50 0.20
IC90 1.70
IC50= 0.20 nM
HK1L2
IC50 0.12
IC90 0.92
IC50= 0.12 nM
count
ug/ml
(IC50, 90; nM)

[Fig.6]


BDCA1
BDCA3
% input
50
40
30
20
10
0
Non treated
Lymphotactin (LTN) +
LTN + human IgG2, κ
LTN + chimeric mAb
LTN + HK1L2
LTN + HK5L5

[Fig.7]

＜VH CDRs＞

CDR1

5G7 :SHN**L**H(SEQ ID No: 17)
2H6 :SHN**M**H(SEQ ID No: 29)
11H2:DYYVN(SEQ ID No: 41)
SHN**X**H(Seq ID No: 53)

CDR2

5G7 :AIYPG**N**GNT**A**YNQKFKG(SEQ ID No: 18)
2H6 :AIYPG**K**GNT**S**YNQKFKG(SEQ ID No: 30)
11H2:VSNPKNGDKSYNQKFKG(SEQ ID No: 42)
AIYPG**X**GNT**X**YNQKFKG(Seq ID No: 54)

CDR3

5G7 :WGSVVGDW•YFDV(SEQ ID No: 19)
2H6 :WGSVVGDW•YFDV(SEQ ID No: 31)
11H2:GLYYAGTYGYFDV(SEQ ID No: 42)
WGSVVGDW•YFDV(SEQ ID No: 55)

＜VL CDRs＞

CDR1

2H6 :RSS**QS**LVH**S**NGNTYLH(SEQ ID No: 20)
5G7 :RSS**LG**LVH**R**NGNTYLH(SEQ ID No: 32)
11H2:RASQDISNYLN (SEQ ID No: 44)
RSS**XX**LVH**X**NGNTYLH(SEQ ID No: 56)

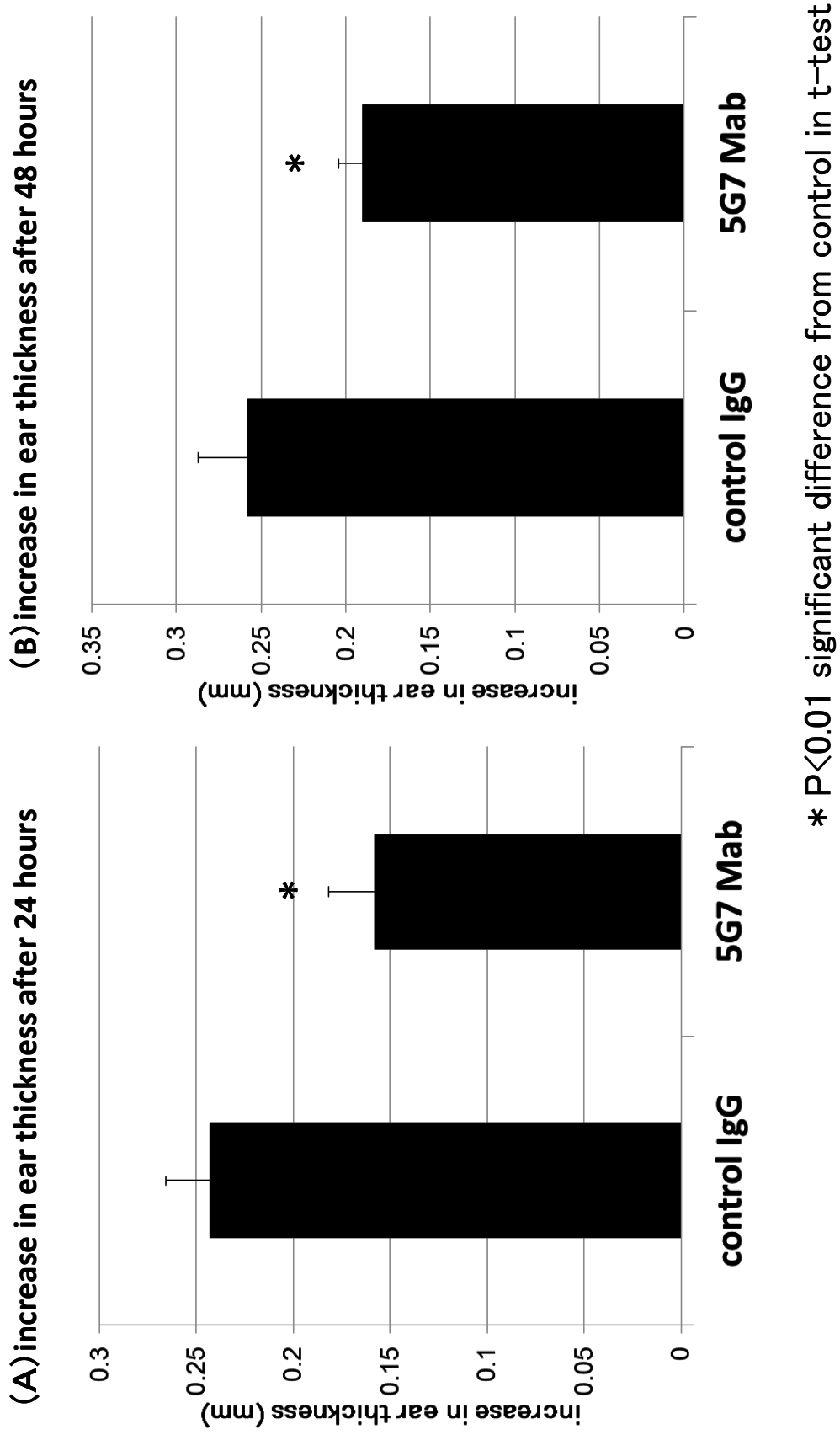
CDR2

5G7 :**K**VS**H**RFS(SEQ ID No: 21)
2H6 :**R**VS**N**RFS(SEQ ID No: 33)
11H2:YTSRLHS(SEQ ID No: 45)
XVS**X**RFS(SEQ ID No: 57)

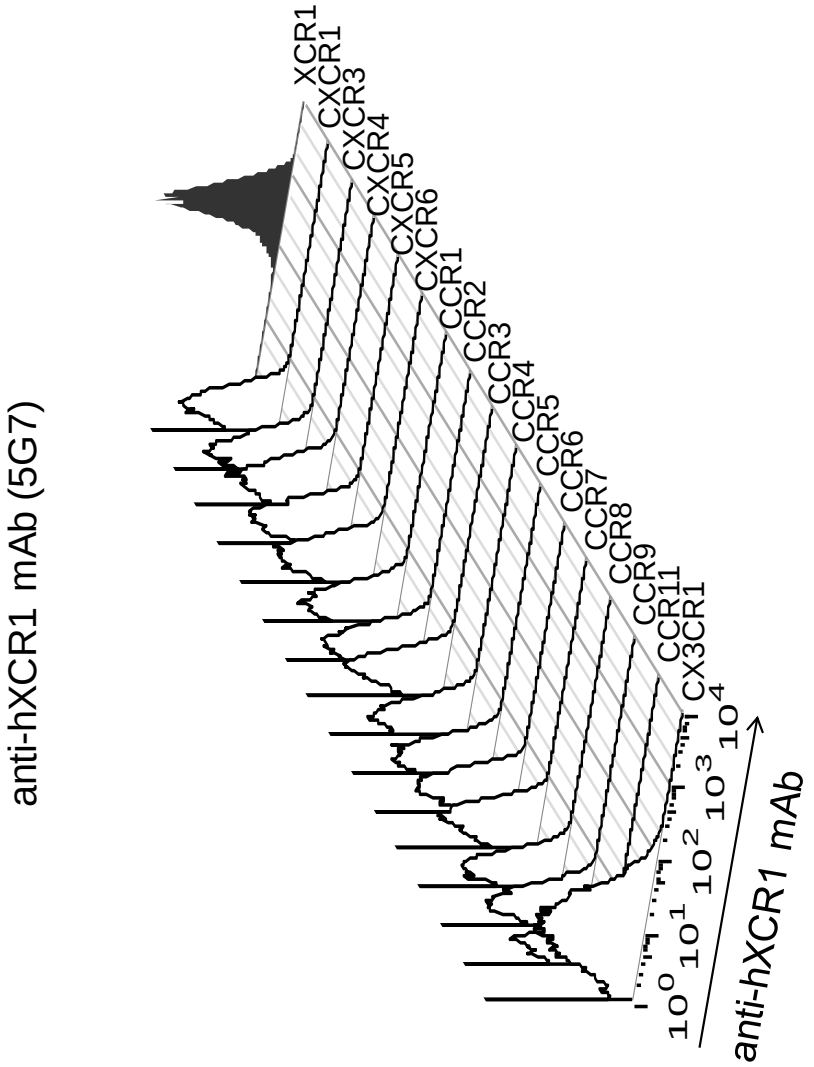
CDR3

5G7 :SQST**H**VPWT(SEQ ID No: 22)
2H6 :SQST**F**VPWT(SEQ ID No: 34)
11H2:QQGKTLPRT(SEQ ID No: 46)
SQST**X**VPWT(SEQ ID No: 58)

[Fig.8]


(A) increase in ear thickness after 24 hours
increase in ear thickness (mm)
0.3
0.25
0.2
0.15
0.1
0.05
0
*
control IgG
5G7 Mab
(B) increase in ear thickness after 48 hours
increase in ear thickness (mm)
0.35
0.3
0.25
0.2
0.15
0.1
0.05
0
*
control IgG
5G7 Mab
* P<0.01 significant difference from control in t-test

[Fig.9]

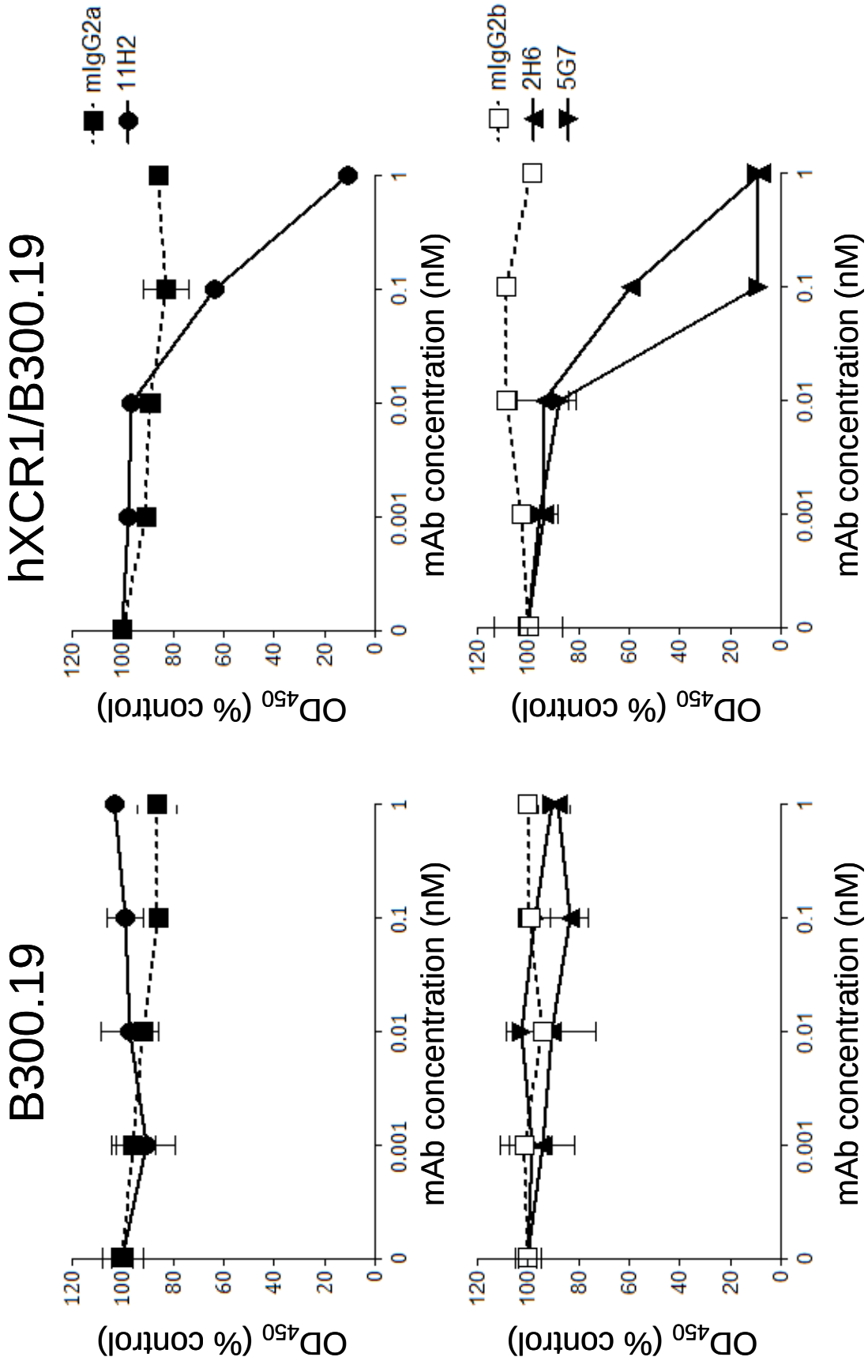

anti-hXCR1 mAb (5G7)
XCR1
CXCR1
CXCR3
CXCR4
CXCR5
CXCR6
CCR1
CCR2
CCR3
CCR4
CCR5
CCR6
CCR7
CCR8
CCR9
CCR11
CX3CR1
10^0 10^1 10^2 10^3 10^4
anti-hXCR1 mAb

[Fig.10]

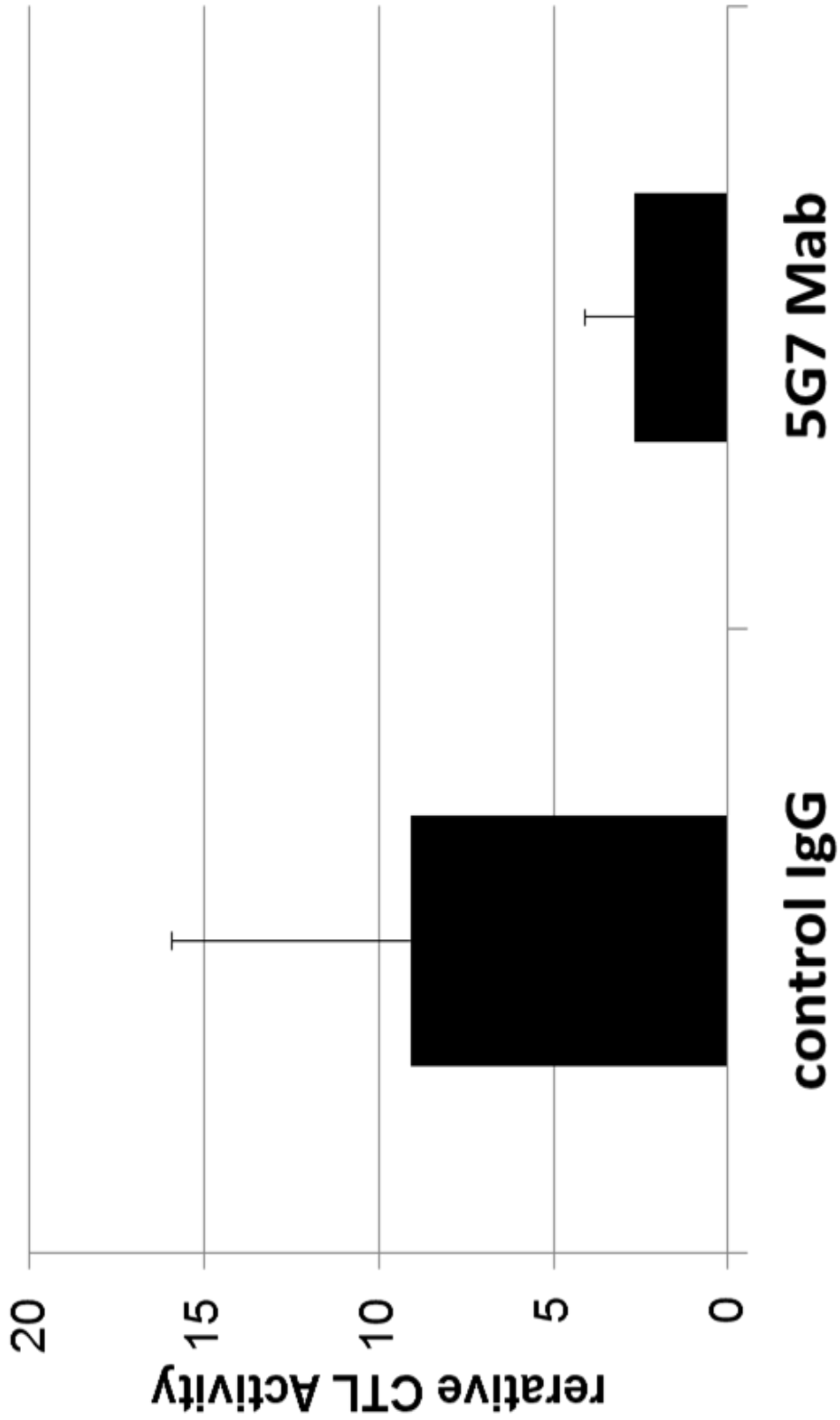
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<212> PRT
<213> Abramis brama

```
Met Glu Ser Ser Gly Asn Pro Glu Ser Thr Thr Phe Phe Tyr Tyr Asp
1               5                  10                  15
Leu Gln Ser Gln Pro Cys Glu Asn Gln Ala Trp Val Phe Ala Thr Leu
            20                  25                  30
Ala Thr Thr Val Leu Tyr Cys Leu Val Phe Leu Leu Ser Leu Val Gly
        35                  40                  45
Asn Ser Leu Val Leu Trp Val Leu Val Lys Tyr Glu Ser Leu Glu Ser
    50                  55                  60
Leu Thr Asn Ile Phe Ile Leu Asn Leu Cys Leu Ser Asp Leu Val Phe
65                  70                  75                  80
Ala Cys Leu Leu Pro Val Trp Ile Ser Pro Tyr His Trp Gly Trp Val
                85                  90                  95
Leu Gly Asp Phe Leu Cys Lys Leu Leu Asn Met Ile Phe Ser Ile Ser
            100                 105                 110
Leu Tyr Ser Ser Ile Phe Phe Leu Thr Ile Met Thr Ile His Arg Tyr
        115                 120                 125
Leu Ser Val Val Ser Pro Leu Ser Thr Leu Arg Val Pro Thr Leu Arg
    130                 135                 140
Cys Arg Val Leu Val Thr Met Ala Val Trp Val Ala Ser Ile Leu Ser
145                 150                 155                 160
Ser Ile Leu Asp Thr Ile Phe His Lys Val Leu Ser Ser Gly Cys Asp
                165                 170                 175
Tyr Ser Glu Leu Thr Trp Tyr Leu Thr Ser Val Tyr Gln His Asn Leu
            180                 185                 190
Phe Phe Leu Leu Ser Leu Gly Ile Ile Leu Phe Cys Tyr Val Glu Ile
        195                 200                 205
Leu Arg Thr Leu Phe Arg Ser Arg Ser Lys Arg Arg His Arg Thr Val
    210                 215                 220
Lys Leu Ile Phe Ala Ile Val Val Ala Tyr Phe Leu Ser Trp Gly Pro
225                 230                 235                 240
Tyr Asn Phe Thr Leu Phe Leu Gln Thr Leu Phe Arg Thr Gln Ile Ile
                245                 250                 255
Arg Ser Cys Glu Ala Lys Gln Gln Leu Glu Tyr Ala Leu Leu Ile Cys
            260                 265                 270
Arg Asn Leu Ala Phe Ser His Cys Cys Phe Asn Pro Val Leu Tyr Val
        275                 280                 285
Phe Val Gly Val Lys Phe Arg Thr His Leu Lys His Val Leu Arg Gln
    290                 295                 300
Phe Trp Phe Cys Arg Leu Gln Ala Pro Ser Pro Ala Ser Ile Pro His
305                 310                 315                 320
Ser Pro Gly Ala Phe Ala Tyr Glu Gly Ala Ser Phe Tyr
                325                 330
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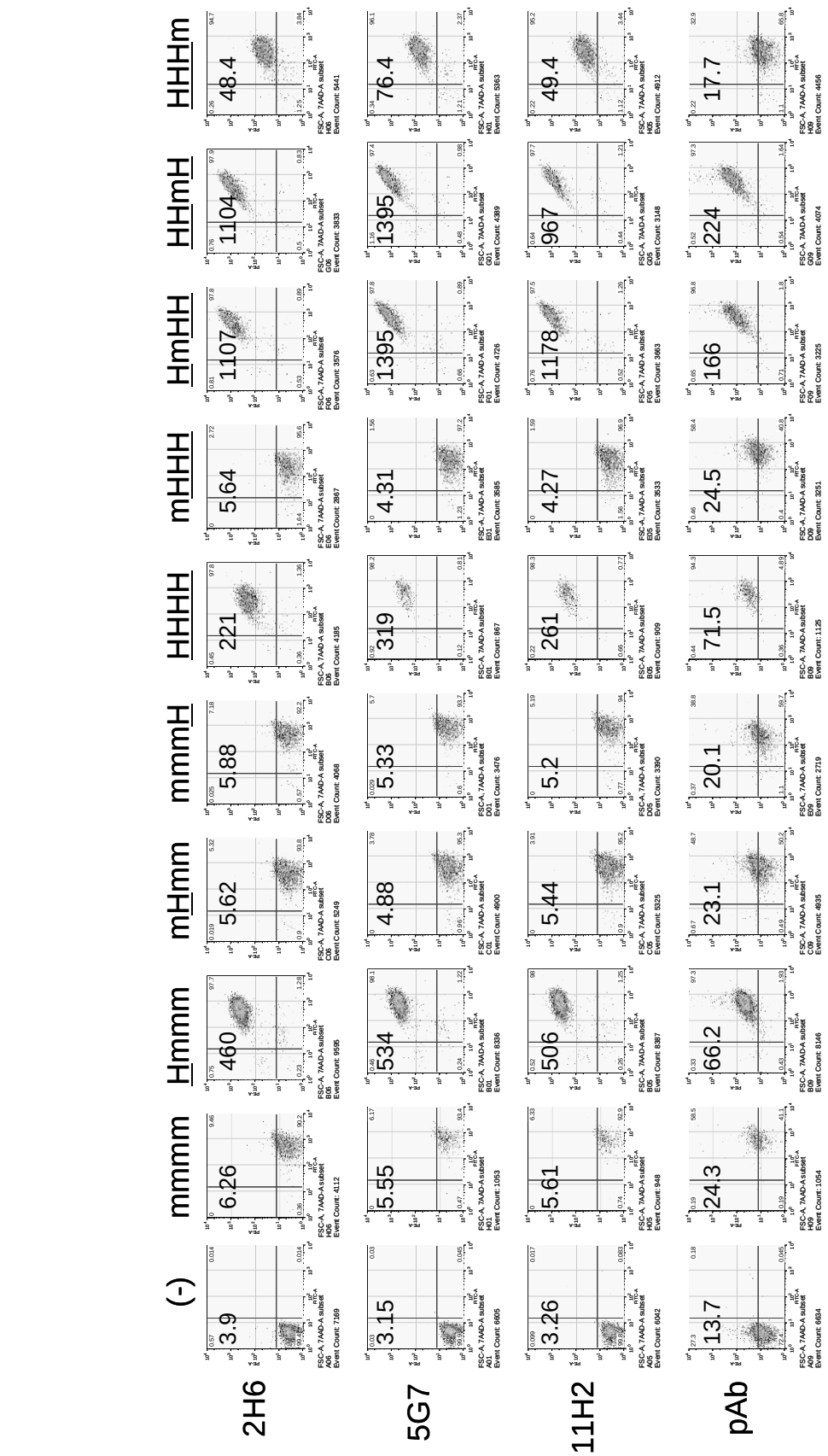
[Fig.11]


hXCR1/B300.19
B300.19
mIgG2a
11H2
mIgG2b
2H6
5G7
OD450 (% control)
mAb concentration (nM)
0
20
40
60
80
100
120
0
0.001
0.01
0.1
1

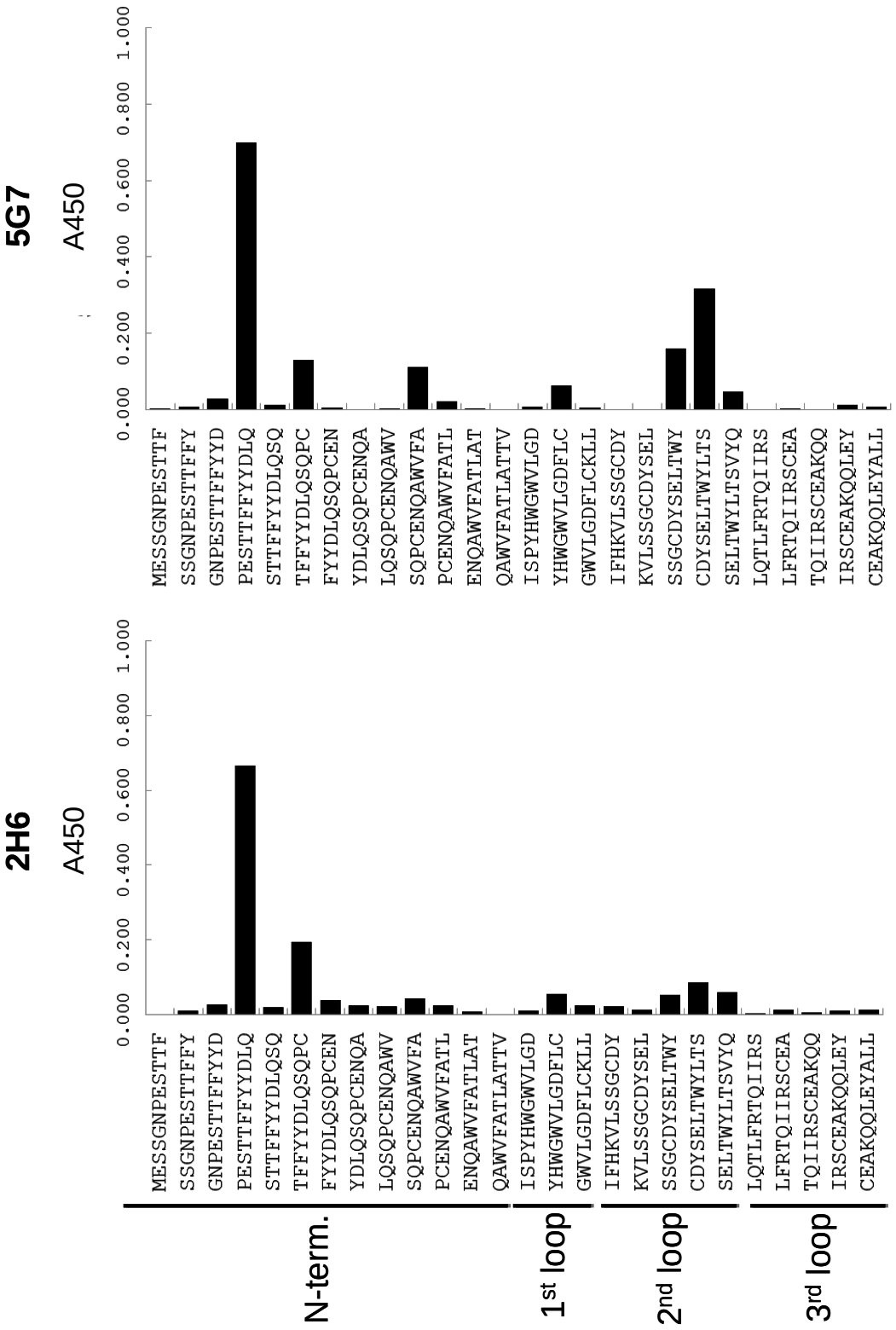
[Fig.12]



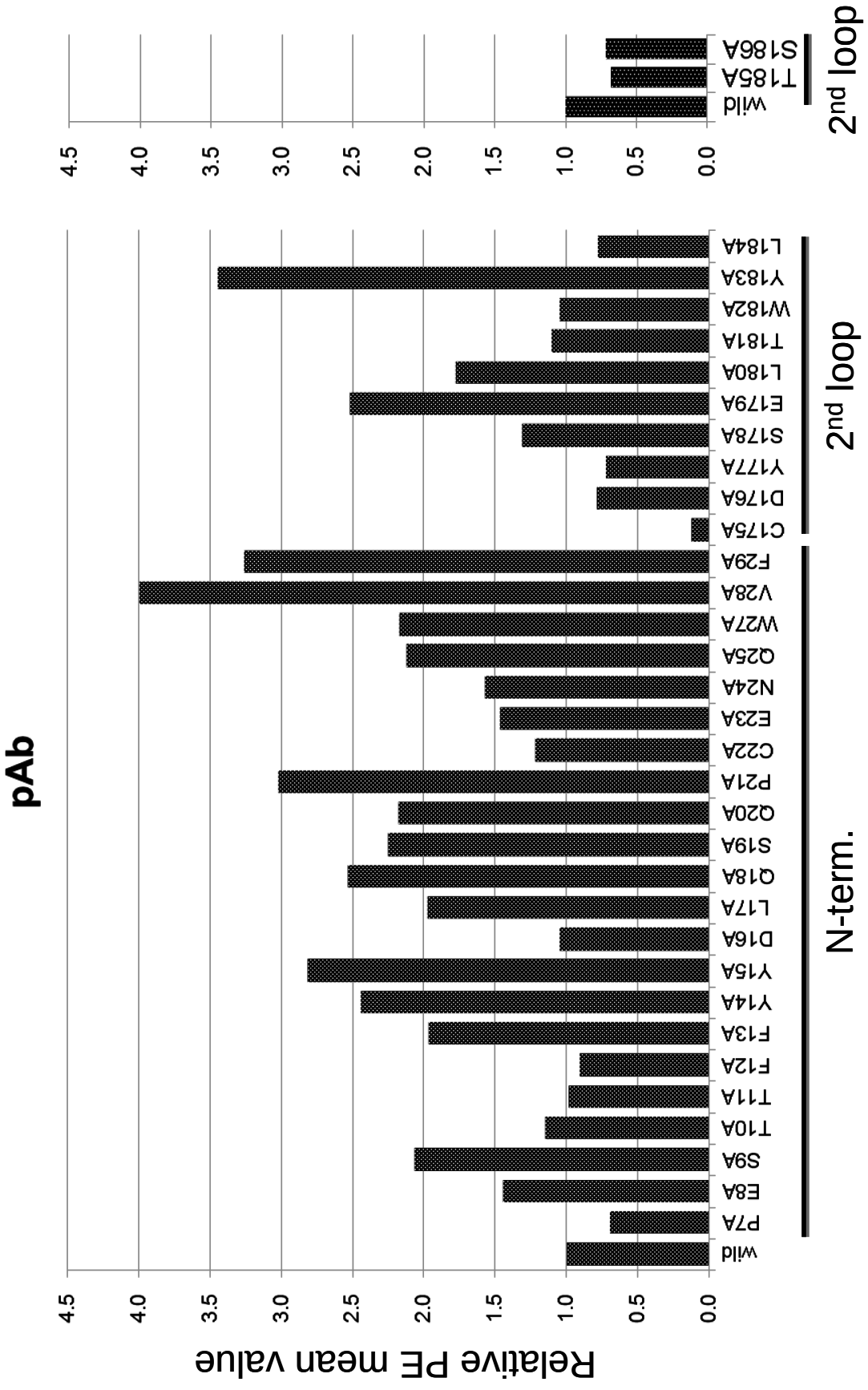
[Fig.13]


(-)
mmmm
Hmmm
mHmm
mmmH
HHHH
mHHH
HmHH
HHmH
HHHm
2H6
5G7
11H2
pAb
3.9
6.26
460
5.62
5.88
221
5.64
1107
1104
48.4
3.15
5.55
534
4.88
5.33
319
4.31
1395
1395
76.4
3.26
5.61
506
5.44
5.2
261
4.27
1178
967
49.4
13.7
24.3
66.2
23.1
20.1
71.5
24.5
166
224
17.7

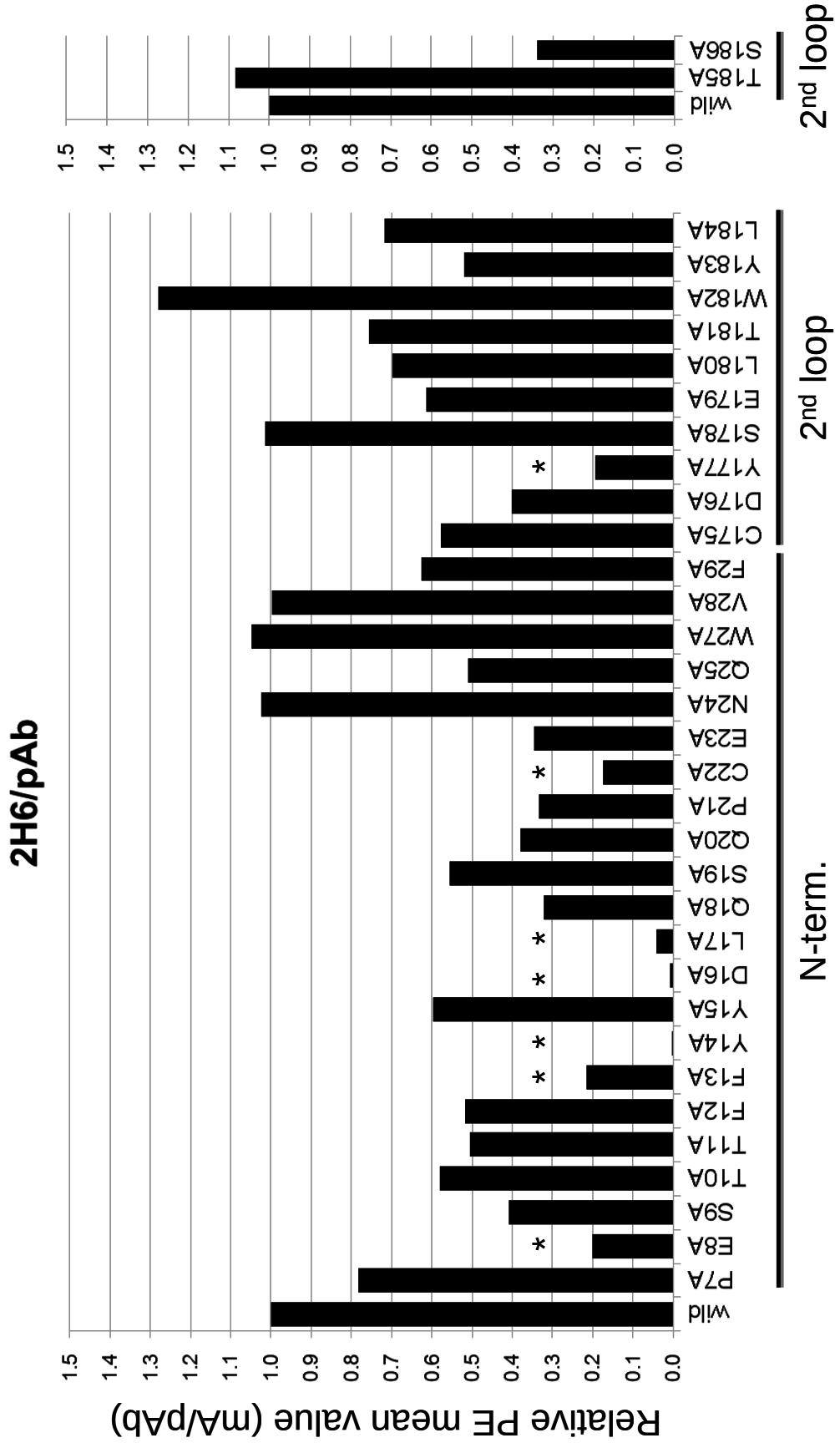
[Fig.14]


5G7
A450
0.000
0.200
0.400
0.600
0.800
1.000
MESSGNPESTTF
SSGNPESTTFFY
GNPESTTFFYYD
PESTTFFYYDLQ
STTFFYYDLQSQ
TFFYYDLQSQPC
FYYDLQSQPCEN
YDLQSQPCENQA
LQSQPCENQAWV
SQPCENQAWVFA
PCENQAWVFATL
ENQAWVFATLAT
QAWVFATLATTV
ISPYHWGWVLGD
YHWGWVLGDFLC
GWVLGDFLCKLL
IFHKVLSSGCDY
KVLSSGCDYSEL
SSGCDYSELTWY
CDYSELTWYLTS
SELTWYLTSVYQ
LQTLFRTQIIRS
LFRTQIIRSCEA
TQIIRSCEAKQQ
IRSCEAKQQLEY
CEAKQQLEYALL
2H6
A450
0.000
0.200
0.400
0.600
0.800
1.000
MESSGNPESTTF
SSGNPESTTFFY
GNPESTTFFYYD
PESTTFFYYDLQ
STTFFYYDLQSQ
TFFYYDLQSQPC
FYYDLQSQPCEN
YDLQSQPCENQA
LQSQPCENQAWV
SQPCENQAWVFA
PCENQAWVFATL
ENQAWVFATLAT
QAWVFATLATTV
ISPYHWGWVLGD
YHWGWVLGDFLC
GWVLGDFLCKLL
IFHKVLSSGCDY
KVLSSGCDYSEL
SSGCDYSELTWY
CDYSELTWYLTS
SELTWYLTSVYQ
LQTLFRTQIIRS
LFRTQIIRSCEA
TQIIRSCEAKQQ
IRSCEAKQQLEY
CEAKQQLEYALL
N-term.
1st loop
2nd loop
3rd loop

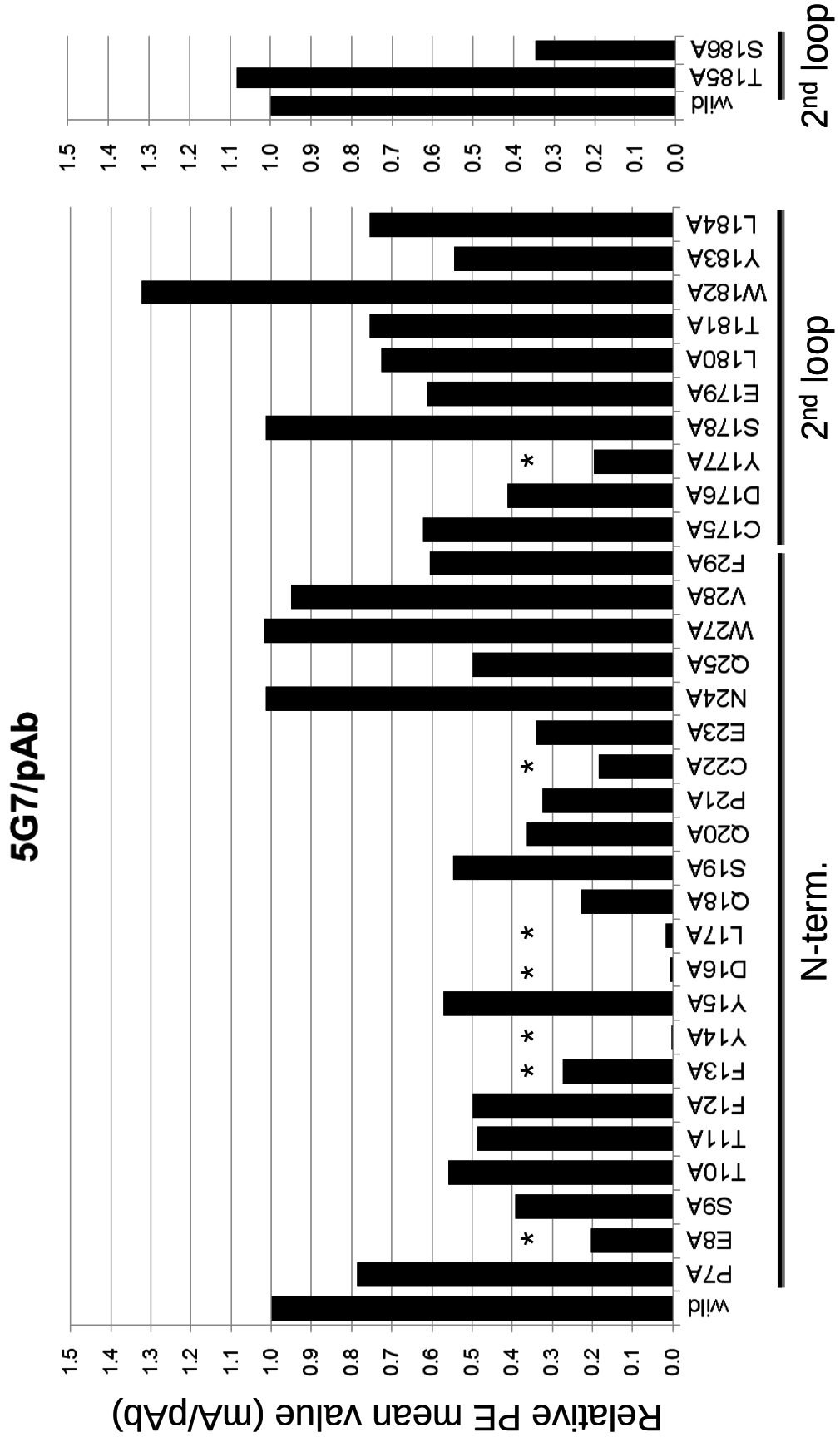
[Fig.15]


pAb
Relative PE mean value
4.5
4.0
3.5
3.0
2.5
2.0
1.5
1.0
0.5
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
N-term.
2nd loop
wild
T185A
S186A
2nd loop

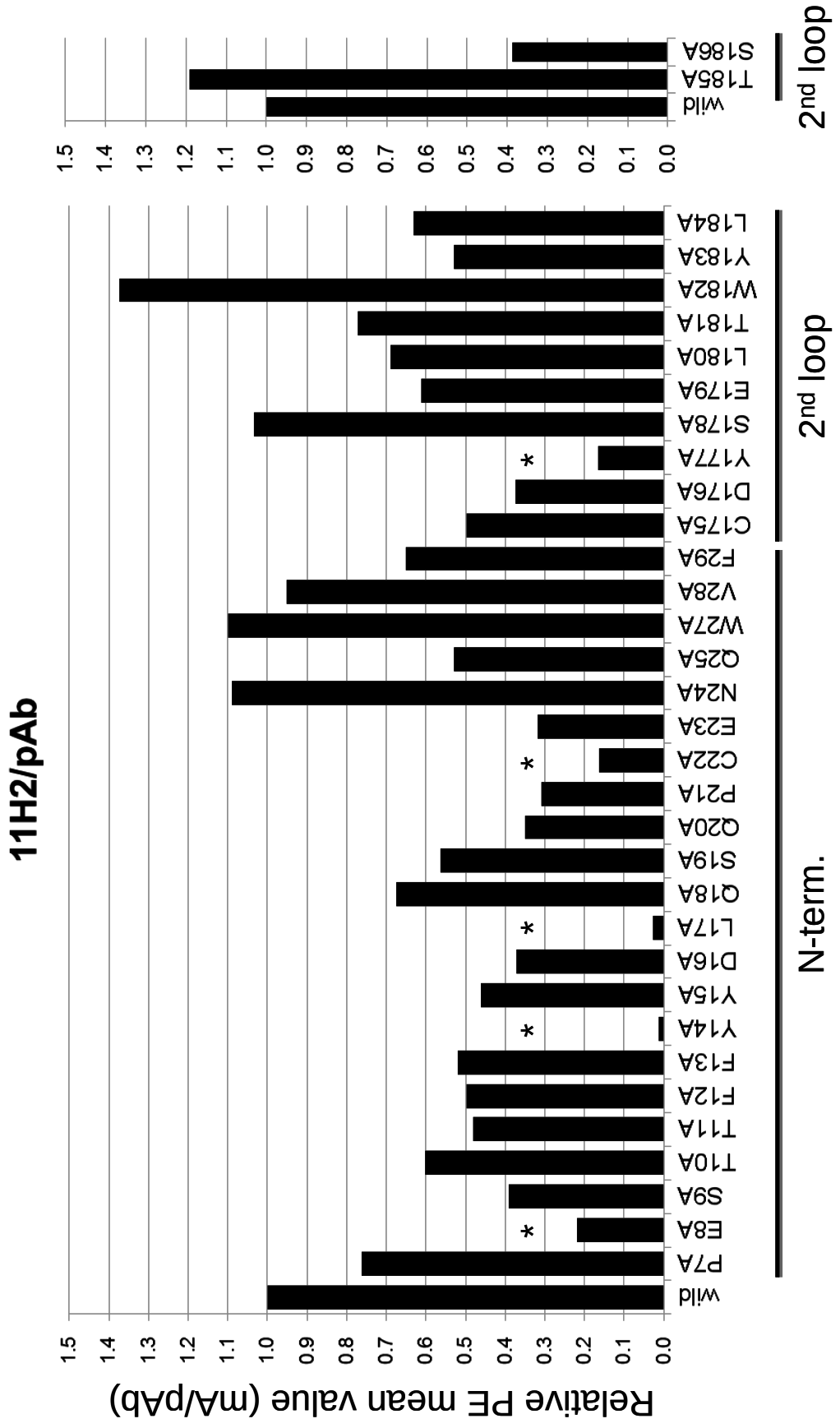
[Fig.16]


2H6/pAb
Relative PE mean value (mA/pAb)
1.5
1.4
1.3
1.2
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
N-term.
2nd loop
wild
T185A
S186A
2nd loop
*

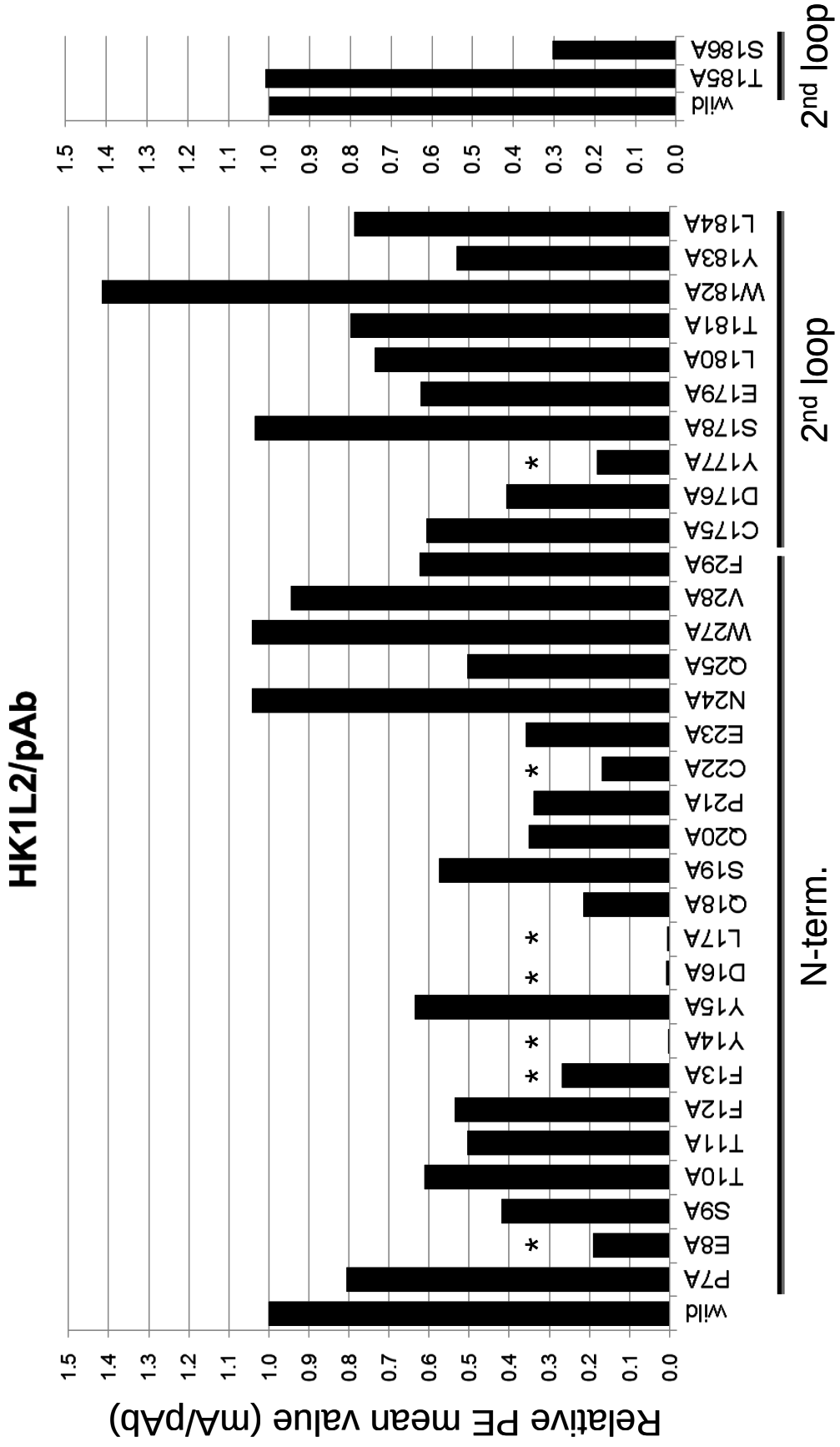
[Fig.17]


5G7/pAb
Relative PE mean value (mA/pAb)
1.5
1.4
1.3
1.2
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
S186A
N-term.
2nd loop
2nd loop
*

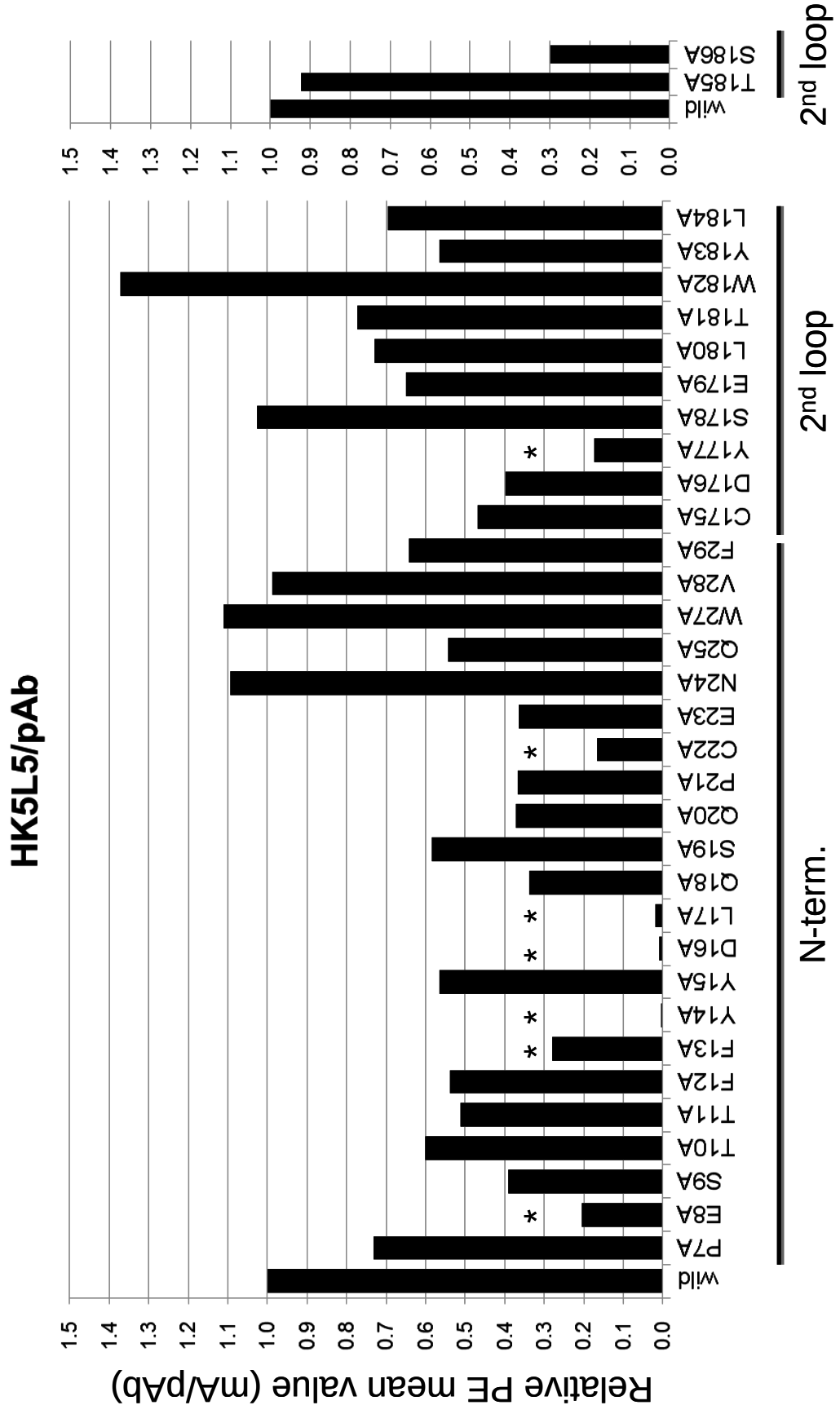
[Fig.18]


11H2/pAb
Relative PE mean value (mA/pAb)
1.5
1.4
1.3
1.2
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
N-term.
2nd loop
wild
T185A
S186A
2nd loop
*

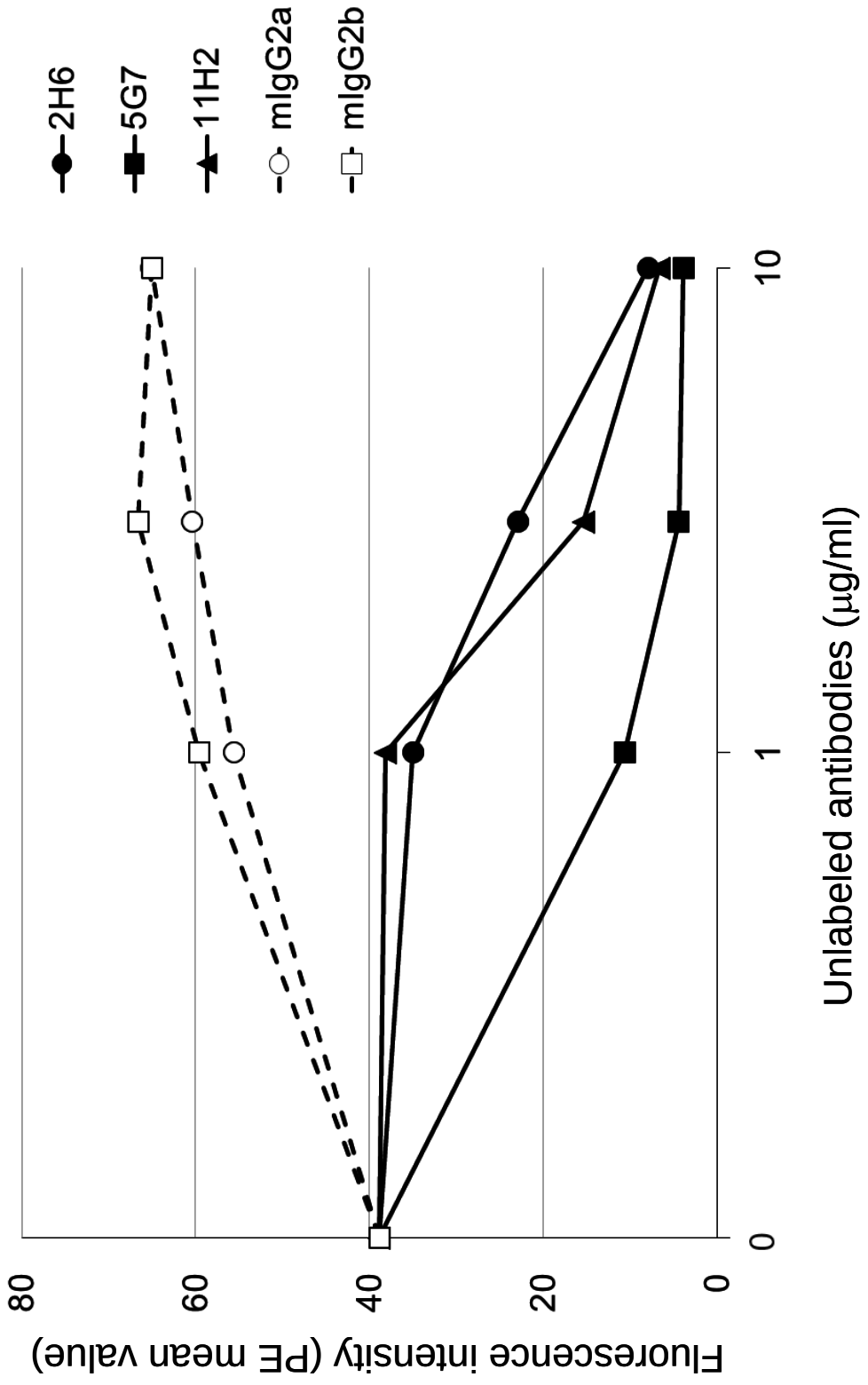
[Fig.19]


HK1L2/pAb
Relative PE mean value (mA/pAb)
1.5
1.4
1.3
1.2
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
N-term.
2nd loop
wild
T185A
S186A
2nd loop
*

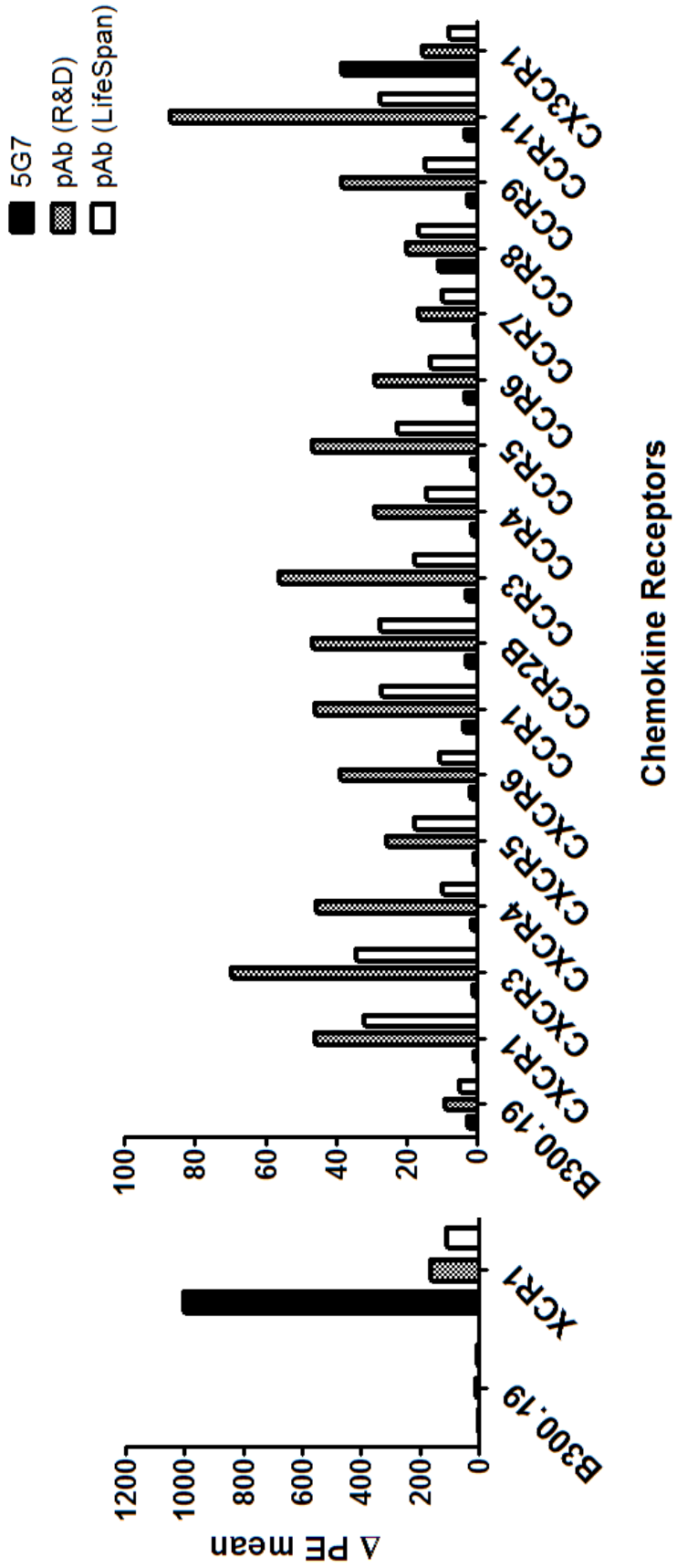
[Fig.20]


HK5L5/pAb
Relative PE mean value (mA/pAb)
1.5
1.4
1.3
1.2
1.1
1.0
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0.0
wild
P7A
E8A
S9A
T10A
T11A
F12A
F13A
Y14A
Y15A
D16A
L17A
Q18A
S19A
Q20A
P21A
C22A
E23A
N24A
Q25A
W27A
V28A
F29A
C175A
D176A
Y177A
S178A
E179A
L180A
T181A
W182A
Y183A
L184A
wild
T185A
S186A
N-term.
2nd loop
2nd loop
*

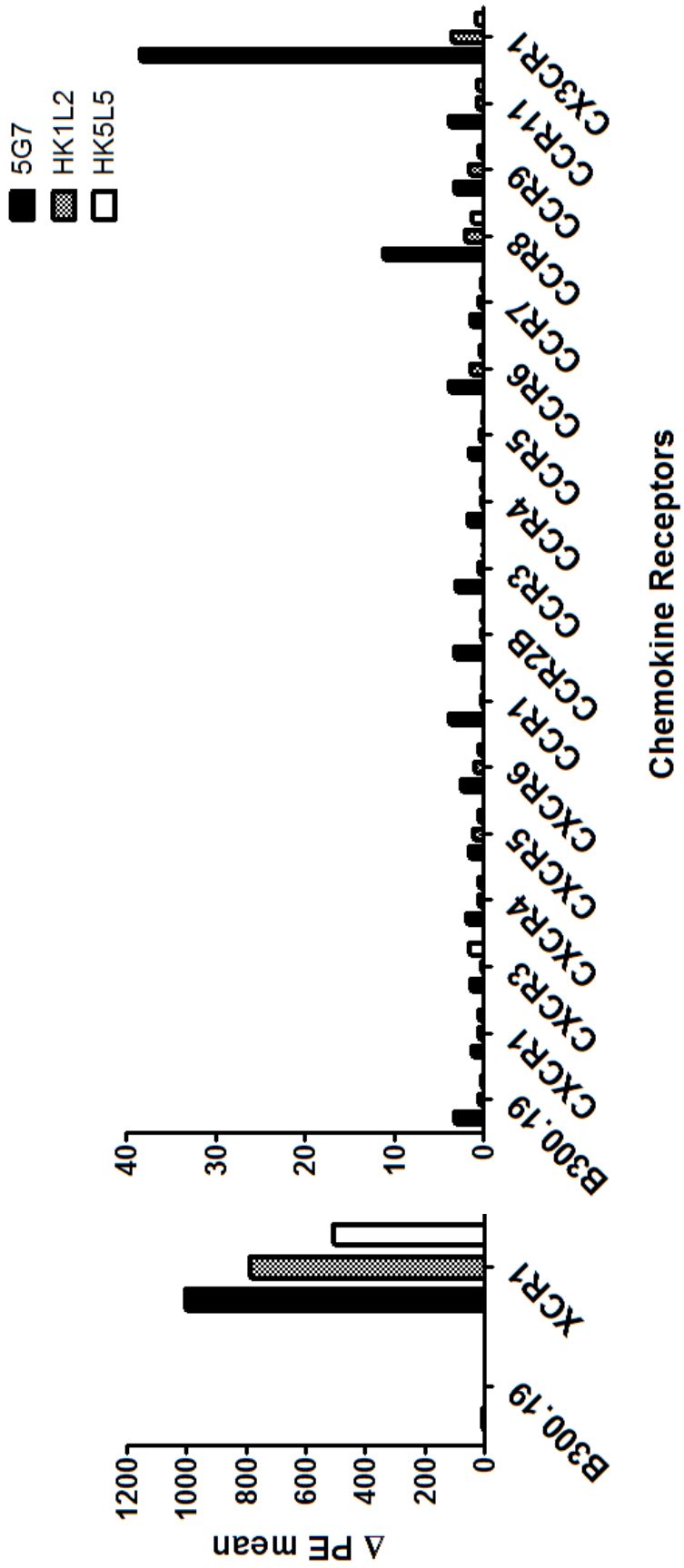
[Fig.21]


2H6
5G7
11H2
mIgG2a
mIgG2b
Fluorescence intensity (PE mean value)
0
20
40
60
80
0
1
10
Unlabeled antibodies (μg/ml)

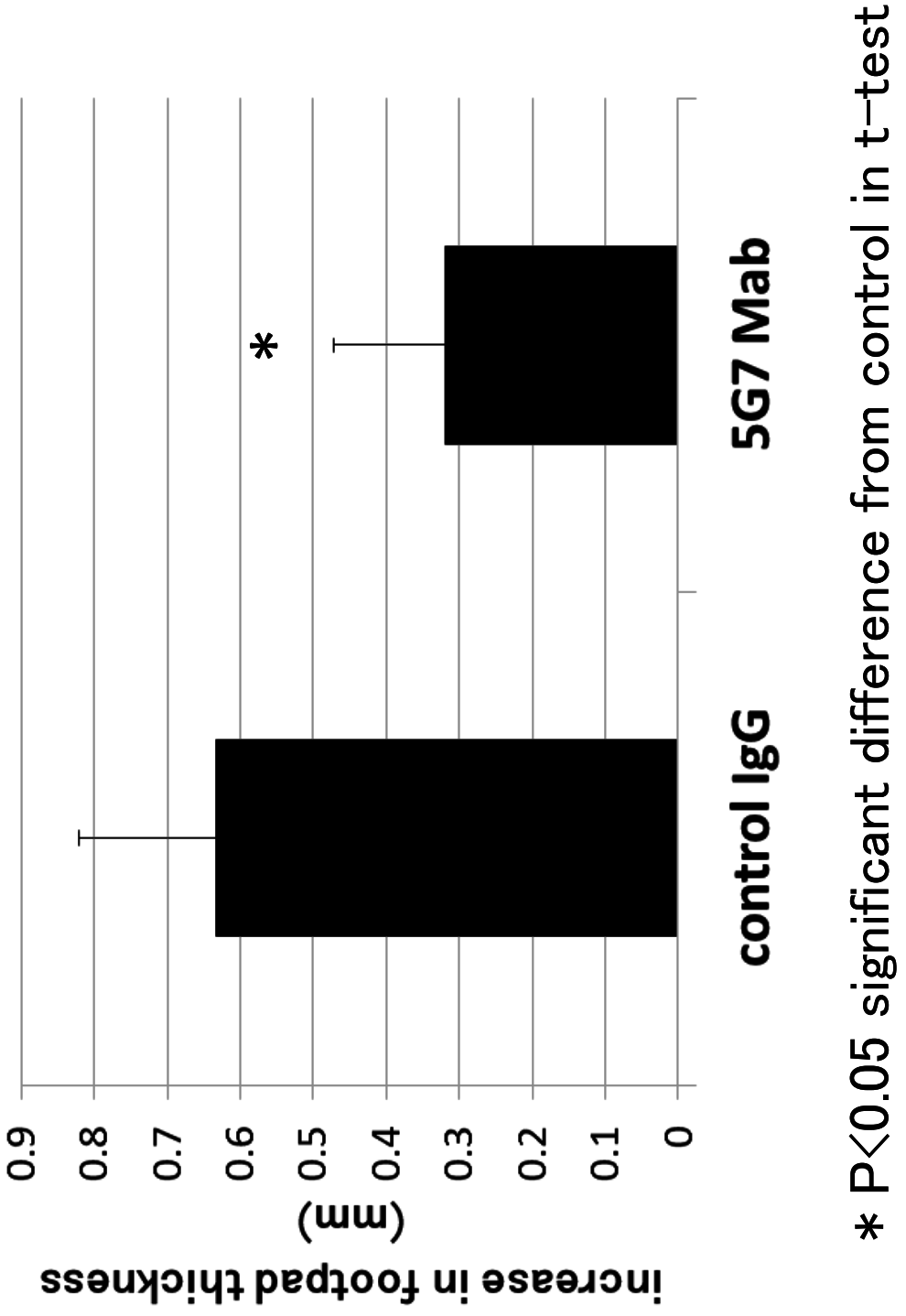
[Fig.22]


5G7
pAb (R&D)
pAb (LifeSpan)
Δ PE mean
1200
1000
800
600
400
200
0
B300.19
XCR1
100
80
60
40
20
0
B300.19
CXCR1
CXCR3
CXCR4
CXCR5
CXCR6
CCR1
CCR2B
CCR3
CCR4
CCR5
CCR6
CCR7
CCR8
CCR9
CCR11
CX3CR1
Chemokine Receptors

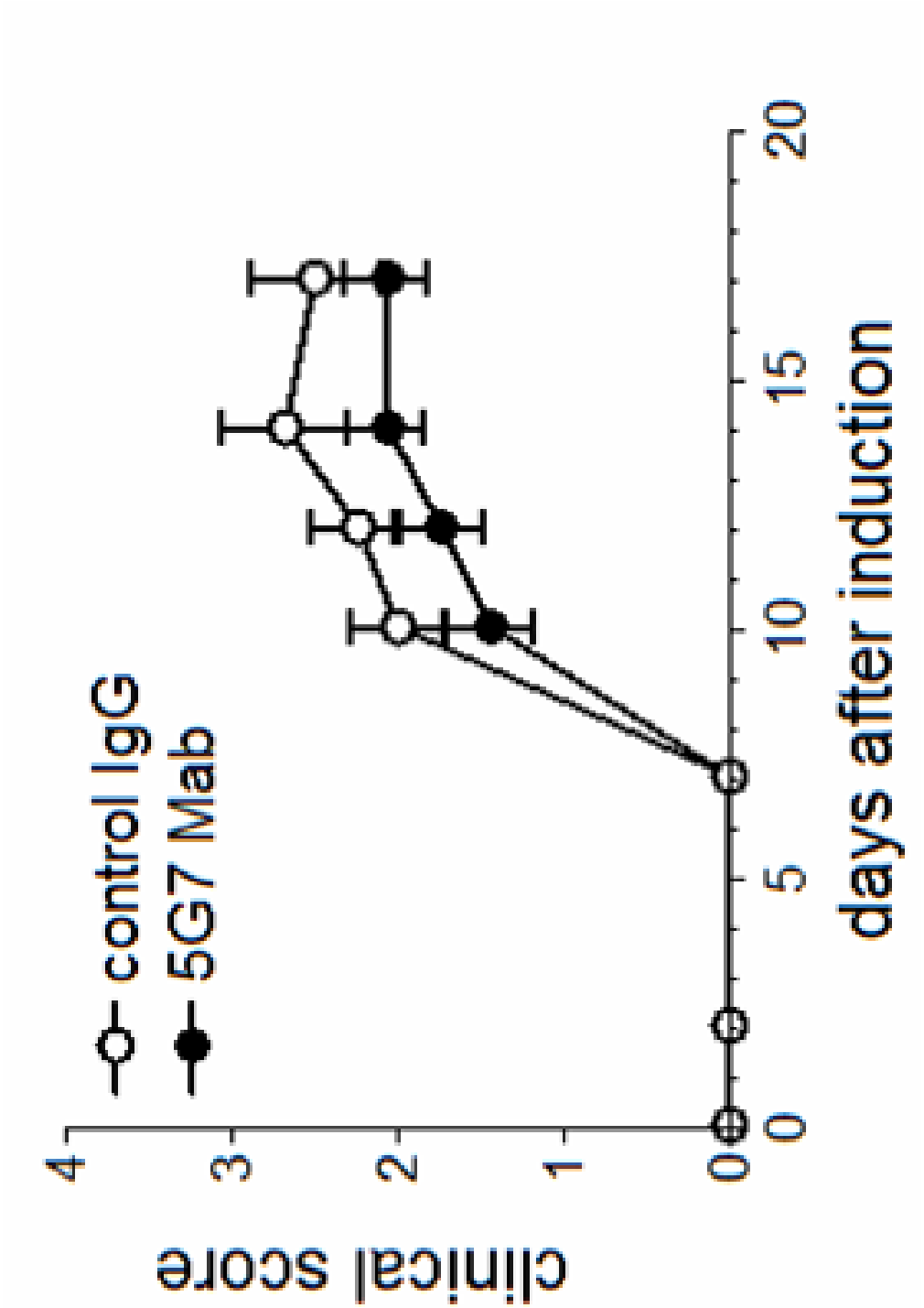
[Fig.23]


5G7
HK1L2
HK5L5
Δ PE mean
0
200
400
600
800
1000
1200
B300.19
XCR1
0
10
20
30
40
B300.19
CXCR1
CXCR3
CXCR4
CXCR5
CXCR6
CCR1
CCR2B
CCR3
CCR4
CCR5
CCR6
CCR7
CCR8
CCR9
CCR11
CX3CR1
Chemokine Receptors

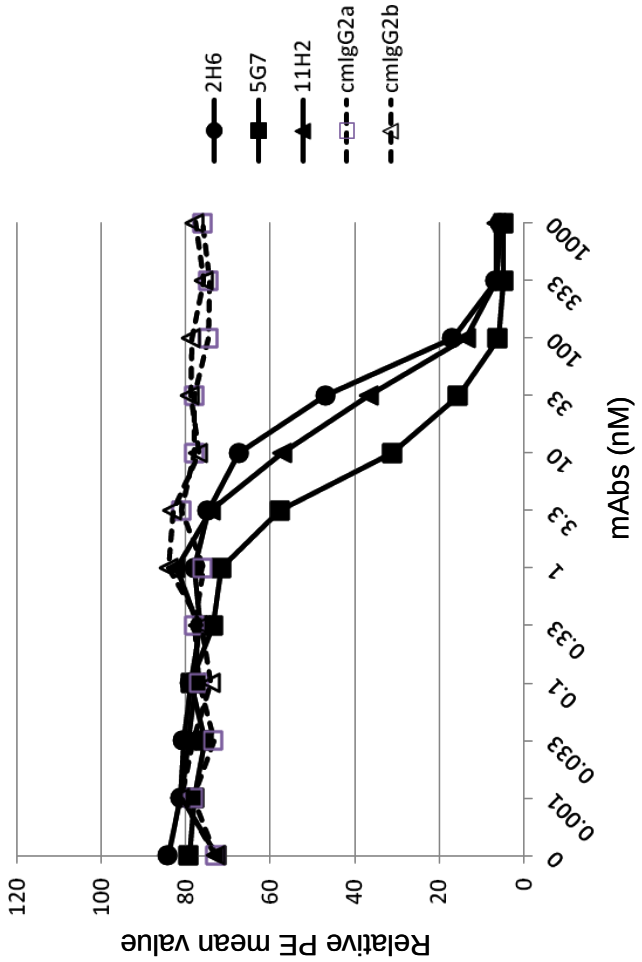
[Fig.24]

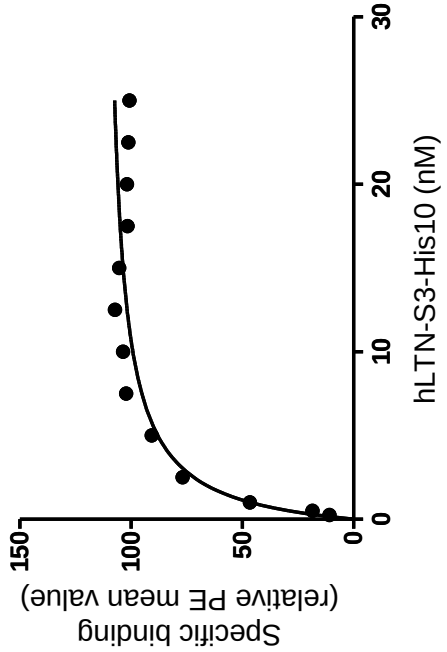

increase in footpad thickness
(mm)
0.9
0.8
0.7
0.6
0.5
0.4
0.3
0.2
0.1
0
*
control IgG
5G7 Mab
* P<0.05 significant difference from control in t-test

[Fig.25]


clinical score
days after induction
control IgG
5G7 Mab
0
1
2
3
4
0
5
10
15
20

[Fig.26]


2H6
5G7
11H2
cmIgG2a
cmIgG2b
Relative PE mean value
120
100
80
60
40
20
0
0
0.001
0.033
0.1
0.33
1
3.3
10
33
100
333
1000
mAbs (nM)


Specific binding
(relative PE mean value)
150
100
50
0
0
10
20
30
hLTN-S3-His10 (nM)