

# CCR6

---

**Joshua Marion Farber\***

Laboratory of Clinical Investigation, National Institute of Allergy and Infectious Diseases, NIH,  
10 Center Drive, Room 11N-228, MSC 1888, Bethesda, MD 20892-1888, USA

\*corresponding author tel: 301-402-4910, fax: 301-402-0627, e-mail: joshua\_farber@nih.gov

DOI: 10.1006/rwcy.2000.22006.

## SUMMARY

CCR6 is a seven transmembrane domain G protein-coupled receptor for MIP-3 $\alpha$ , a CC chemokine expressed by a variety of cell types, including epithelial cells, in response to inflammatory stimuli. CCR6 is expressed on memory T cells, B cells, and CD34<sup>+</sup> bone marrow progenitor-derived dendritic cells. CCR6 is active on resting memory cells and can mediate both chemotaxis and adhesion of these cells to ICAM-1. It is presumed that CCR6 is involved in the recruitment of memory T cells at the initiation of a response at inflammatory sites and/or in reactive lymphoid tissue. Data also suggest that CCR6 on immature dendritic cells may be important for the recruitment of these cells to inflammatory sites. Since the MIP-3 $\alpha$  gene is also expressed in dendritic cells, CCR6 may mediate aggregation of dendritic cells, memory T cells, and B cells.

## BACKGROUND

### Discovery

CCR6 was discovered by several groups as an orphan receptor as part of screening for new G protein-coupled receptors or specifically for new chemokine receptors. The first sequence entered in the database was a genomic sequence, GPR-CY4, and the first published sequences were a genomic sequence designated CKR-L3 (Zaballos *et al.*, 1996) and cDNA and genomic sequences for a gene designated *STRL22* (Liao *et al.*, 1997c). The protein encoded by this gene was described as a receptor for the chemokine LARC/MIP-3 $\alpha$ /Exodus/CK- $\beta$ 4 by several groups (Baba *et al.*, 1997; Liao *et al.*, 1997a; Power *et al.*, 1997; Greaves *et al.*, 1997) and was thereby named CCR6.

### Alternative names

None currently in use. Old names include GPR-CY4, CKR-L3, STRL22, DCCR2, BN-1, and DRY6.

### Structure

Like other chemokine receptors, CCR6 is a seven transmembrane domain G protein-coupled receptor. CCR6 is predicted to be 374 amino acids in length.

### Main activities and pathophysiological roles

CCR6 is a receptor for the chemokine LARC/MIP-3 $\alpha$ /Exodus/CK- $\beta$ 4 (Baba *et al.*, 1997; Greaves *et al.*, 1997; Liao *et al.*, 1997a; Power *et al.*, 1997), a chemokine that, based on the expression of mRNA, is produced by activated macrophages, endothelial cells, and dendritic cells, and at some mucosal sites. CCR6 is expressed and functional on CD34<sup>+</sup> bone marrow progenitor cell-derived dendritic cells (Greaves *et al.*, 1997; Power *et al.*, 1997; Liao *et al.*, 1999), particularly on 'immature' dendritic cells early in their differentiation *in vitro* (Dieu *et al.*, 1998), as well as on CD4<sup>+</sup> and CD8<sup>+</sup> T cells (Baba *et al.*, 1997; Greaves *et al.*, 1997; Power *et al.*, 1997; Liao *et al.*, 1999). In the case of T cells, CCR6 is expressed exclusively on those of the memory phenotype (Liao *et al.*, 1999). CCR6 is also expressed on B cells (Zaballos *et al.*, 1996; Baba *et al.*, 1997; Liao *et al.*, 1997b, 1999; Varona *et al.*, 1998), and MIP-3 $\alpha$  has been reported to be active on B cells from Peyer's patches (Tanaka *et al.*, 1999). It is presumed that CCR6 is involved in the trafficking of dendritic cells and T cells, and perhaps B cells, particularly at mucosal sites.

**GENE**

**Accession numbers**

Human gene: U45985, Z79784, U68031  
 Human cDNAs: U68030, U68032  
 Mouse gene: AJ222714  
 Mouse cDNA (KY411): AB009369

The human CCR6 gene is on chromosome 6q27, a site without other known chemokine receptors (Liao *et al.*, 1997c).

**Sequence**

See **Figure 1**.

**PROTEIN**

**Accession numbers**

Human: AAC51124  
 Mouse: BAA23776

**Sequence**

See **Figure 2**.

**Description of protein**

No information is available other than what can be predicted from the primary structure. CCR6 is

**Figure 1** Nucleotide sequences for human and mouse CCR4.

**Human gene:**

```

AACCTGTAGTGCATTTTGCCTTCTTTCTTCTTAGAGTCACCTCTACTTTCTGCTACCGCTGCCTGTGAGCTGAAGGGGCTGAACCATACACTCCCTT
TTTCTACAACCAGCTTGCAATTTTTCTGCCACAATGAGCGGGTAAGATTTTTATTTTTGGCAAGGGGTATAATTTGGGTTCACTGTGGCTACTTG
AACACTACACTGCAGCTAACTCTATCTTTGTTTCTTTCCAGGAATCAATGAATTTCCAGCGATGTTTTCGACTCCAGTGAAGATTATTTGTGTCAG
TCAATACTTTCATATTACTCAGTTGATTCTGAGATGTTACTGTGCTCCTTGCCAGGAGGTCAGGCAGTTCTCCAGGCTATTTGTACCGATTGCCTACTC
CTTGATCTGTGCTTTGGCTCCTGGGGAATATCTGGTGGTGATCACCTTGGCTTTTATAAGAAGGCCAGGCTATGACAGACGCTATCTCTTTG
AACATGGCCATTGCGACATCCTCTTTGTTCTTACTCTCCCATCTGGGCAGTGAGTCATGCCACTGGTGCCTGGGTTTTTCAGCAATGCCACGTGCA
AGTTGCTAAAAGGCATCTATGCCATCACTTTAACTGCGGGATGCTGCTCTGACTTGCATTAGCATGGACCGGTACATCGCCATTGTACAGGCGAC
TAAGTCATTCGGCTCCGATCCAGAACACTACCGCGCAGCAAAATCATCTGCCTGTTGTGTGGGGGCTGTGTCAGTCATCATCTCCAGCTCAACTTTT
GTCTTCAACCAAAAATACAAACCCAAAGGCAGCGATGTCTGTGAACCCAAAGTACCAGACTGTCTCGGAGCCCATCAGGTGGAAGCTGTGATGTTGG
GGCTTGAGCTACTCTTTGGTTTCTTTATCCCTTTGATGTTTCATGATATTTTGTACACGTTTCATGTCAAACCTTGGTGCAAGCTCAGAATTTCTAA
AAGGCACAAAGCCATCCGTGTAATCATAGCTGTGGTGTGTTTCTGGCTTGTGAGATTCCATACATGCTGCTTGTGACGGCTGCAAAAT
TTGGGTAAAATGAACCGATCCTGCCAGAGCGAAAAGCTAAATGGCTATACGAAAAGTGCACAGAAAGTCTGGCTTCTGCACTGCTGCCTGAACC
CTGTGCTCTACGCTTTTATTGGGCAGAAAGTTCAGAAACTACTTTCTGAAGACTTGAAGGACCTGTGGTGTGTGAGAAGGAAGTACAAGTCCCTCAGG
CTTCTCCTGTGCCGGGAGTACTCAGAAAACATTTCTCGGCAGACCAGTGAGACCCGAGATAACGACAATGCGTCCTTCACTATGTGATAGAAAA
GCTGAGTCTCCCTAAGGCATGTGTGAAACATACTCATAGATGTTATGCAAAAAAAGTCTATGGCCAGGTATGCATGGAAAATGTGGGAATTAAGCA
AAATCAAGCAAGCCTCTCCTGCGGGACTTAACTGTCTCATGGCTGTGTGATCTCTTCCAGGTGGGGTGGTCTCTGATAGGTAGCATTTCAGC
ACTTTGCAAGGAATGTTTTGTAGCTCTAGGGTATATATCCGCCTGGCATTTCACAAAACAGCCTTTGGGAAATGCTGAATTAAGTGAATTTGTGAC
AAATGTAACATTTTCAGAAATATTCATGAAGCGGTACAGATCACAGTGTCTTTGGTTACAGCACAAAATGATGGCAGTGGTTTGA AAAACTAAA
ACAGAAAAAAAATGGAAGCCACACATCACTCATTTTAGGCAATGTTTAAACATTTTATCTATCAGAATGTTTATTGTTGCTGGTTATAAGCAG
CAGGATTGCGCGGCTAGTGTTCCTCTCATTTCCCTTTGATACAGTCAACAAGCCTGACCCTGTAAAATGGAGGTGGAAGACAAGTCAAGTCAAGTGTTC
ACAACCTGGAAGTGCTTCGGAAGAAGGGGACAATGGCAGAACAGGTGTTGGTGACAATTGTCAACCAATTGGATAAAGCAGCTCAGGTTGTAGTGGG
CCATTAGGAAACTGTCGGTTTGCCTTTGATTTCCCTGGGAGCTGTCTCTGTGCTGAGTGTCTCTGTCTAAACGTCCTATTAAGCTGAGAGTGTATG
AAGACAGGATCTAGAATAATCTTGTCTCACAGCTGTGCTCTGAGTGCCTAGCGGAGTTCAGCAGAAACAAAATGGACTCAAGAGAGATTTGATTAATGA
ATCGTAATGAAGTTGGGGTTTATGTACAGTTTAAAATGTTAGATGTTTTTAAATTTTTTAAATAAATGGAATACTTTTTTTTTTTTTTAAAGAAAGC
AACTTTACTGAGACAATGTAGAAAAGAGTTTGTTCCTTTTAAATGTGGTTGAAGAGCAATGTGTGGCTGAAGACTTTTTGTTATGAGGAGCTGC
AGATTAGCTAGGGGACAGCTGGAATTTATGCTGGCTTCTGATAAATTTTAAAGGGTCTGAAATTTGTGATGGAATCAGATTTTAAACAGCTCTCTT
CAATGACATAGAAAAGTTCATGGAACCTCATGTTTTTAAAGGGCTATGTAATATATGAACATTAGAAAATAGCAACTTGTGTTACAAAATACAAAC
ACATGTTAGGAAGTACTGTTCATGGCTAGGCATGGTGGCTCACACCTGTAATCCAGCATTTTGGGAAGCTAAGATGGGTGGATCATTGAGGTCA
GGAGTTTGAAGACCAGCCTGGCCAACATGGCGAAAACCCCTCTCTACTAAAATAACAAAATTTGCCAGGCGTGGTGGCGGGTGCCTGTAATCCCAGCT
ACTTGGGAGGCTGAGGCAAGAGAATCGCTTGAACCCAGGAGGCAGAGGTTGCAGTGAGCCGAGATCGTGCCATTGCACTCCAGCCTGGGTGACAAAAG
CGAGACTCCATCTCAAAAAAAAAAAAAAAAAAAAAAAAAAAGGAAAGAACTGTATGTAACATAACCGACATGTTTAAACCTGACAATGGTGTATTATGAAA
CTTTATATTGTTCTTGTAAAGCTTTAACTATATCTCTCTTTAAAATGCAAAAATAATGTCTTAAAGATTCAAAGTCTGTATTTTTAAAGCATGGCTTTGG
CTTTGCAAAAATAAAAATGTGTTTTGTACATGAAGTAGGAATCGTATTTTCAAGTTCAGATTGAGGGGCCACTGTTTGGAGAGGATGGTA
TTCAGGCTTCTCATGTCTTCAAATCTGTAGCGTTTACTCTAGAAAATCAAAGCAAAGGAGTGGTTACCCAGACACTTCTTTTGGTGTGATCAAT
GCGCTGATGTGATCTATGAAGATGATTCATGCTTGA AAACTAGCACAGAAACATCTTGCTTATTTGCCAAAGCTGGGAGATGAGCTTCTCTGCATAA
TTTTAAATGTTTCAATAAATGAAGCTGACTTATTTAAGCAATAACCTTTTTAAACATTTTAACTAAGTAAATGATAAAAATGTTTCCAAAATATACCAT
ACTTTATTTCTTCTTAAATGTAGTACATTAAGTTACATATTTTCTTCTGTCTTGGGCATCAAAAACAGGTGCCATGGTAACCTGACACTCTCAGG
AGACATTAAGATAGAAGGGGCTGTCTTTCAGTGGTTCCCAATTGATTTCTCCCATATCTTTTGTCTCTCAGGCTCTGGCCGTCTCTCTGAGCCTTA
ACTGTGT
    
```

Figure 1 (Continued)

## Human cDNA:

CAAACGTTCCCAAATCTTCCAGTCCGGCTTGACAGACTCCTTGCTCCAGGAGATAACCAGAAGCTGCATCTTATTGACAGATGGTCATCACATTG  
 GTGAGCTGGAGTCATCAGATTGTGGGCCCCGAGTGAAGGAGTGGATCAGAGCACTGCCTGAGAGTACCTCTACTTTCTGCTACCGCT  
 GCCTGTGAGCTGAAGGGGCTGAACCATACTCCTTTTCTACAACCAGCTTGCAATTTTTCTGCCACAATGAGCGGGGAATCAATGAATTTGAG  
 GATGTTTTGACTCCAGTGAAGATTATTTGTGTGTCAGTCAACTTCTATATTACTCAGTTGATCTGAGATGTACTGTGCTCCTTGACGAGGTC  
 GCAGTTCTCCAGGCTATTTGTACCGATTGCCTACTCCTTGATCTGTGCTTTGGCCTCCTGGGAATATTCTGGTGGTATCACCTTTGCTTTTFA  
 TAAGAAGGCCAGGTCTATGACAGACGCTATCTCTTGAACATGGCCATTGCAGACATCCTCTTTGTTCTTACTCTCCCATCTGGGACGTGAGTCAT  
 GCCACTGGTGCCTGGGTTTTCAGCAATGCCAGTGCAGATTGCTAAAAGGCATCTATGCCATCAACTTTAACTGCGGGATGCTGCTCCTGACTTGCA  
 TTAGCATGGACCGGTACATCGCCATTGTACAGGCGACTAAGTCATTCCGGCTCCGATCCAGAACAACCTACCGCGCACGAAAATCATCTGCCTTGTGT  
 GTGGGGCTGTGAGTCATCATCTCCAGCTCACTTTTGTCTTCAACCAAAAATCAACACCAAGGACGAGTGTCTGTGAACCAAGTACCAGACT  
 GTCTCGAGCCCATCAGTGGAAAGCTGTGATGTGGGGCTTGAGCTACTCTTTGGTTTCTTATCCCTTTGATGTCATGATATTTTGTACACGT  
 TCATTGTCAAAAACCTTGGTGCAGCTCAGAAATCTAAAAGGCACAAAGCCATCCGTGTAATCATAGCTGTGGTCTTGTGTTTCTGGCTTGTGAGAT  
 TCCTCATAACATGGTCTGCTTGTGACGGCTGCAAATTTGGGTAAAATGAACCGATCTGCCAGAGCGAAAAGCTAATGGCTATACGAAAACGTG  
 ACAGAAGTCTGGCTTTCTGCACTGCTGCCTGAACCTGTGCTCTACGCTTTTATGGGCAGAAGTTCAGAAACTACTTTCTGAAGATCTGAAGG  
 ACCTGTGGTGTGTGAGAAGGAAGTACAAGTCTCAGGCTTCTCCTGTGCGGGAGGTAAGTACTCAGAAAACATTTCTCGGCAGACAGTGTGAGCCG  
 TAACGACAATCGCTCGTCTTCACTATGTGATAGAAAGCTGAGTCTCCCTAAGGCATGTGTGAAACATCTCATAGATGTTATGCAAAAAAAGTCT  
 ATGGCCAGGTATGCATGAAGAGCTCAGGTTGTAGTGGGCTTAGGCAAAATCAAGCAAGCTCTCTCTGCGGGACTTAACGTGCTCATGGCTGTGATCTCTC  
 AGGGTGGGGTGGTCTCTGATAGGTAGCATTTTCCAGCACTTTGCAAGGAATGTTTTGTAGCTCTAGGGTATATATCCGCTGGCATTTCACAAAACA  
 GCCTTTGGGAAATGCTGAATTAAGTGAATTTGTGACAAATGTAACATTTTTCAGAAATATTCATGAAGCGGTACAGATCACAGTGTCTTTTGGTT  
 ACAGCACAAAATGATGCAAGTGGTTTGAAGAACTAAAACAGAAAAAATGGAAGCAACACATCACTCATTTTAGGCAATGTTTAAACATTTT  
 ATCTATCAGAATGTTTATGTTGCTGGTTATAAGCAGCAGGATGGCCGGCTAGTGTTCCTCTCATTTCCCTTTGATACAGTCAACAAGCCTGACC  
 CTGTAAAATGGAGTGGAAAGACAAGCTCAAGTGTTCACAACCTGGAAGTCTTCCGGAAGAAGGGACAATGGCAGAACAGGTGTTGGTGACAAT  
 GTCAACCAATTGGATAAAGCAGCTCAGGTTGTAGTGGGCTTAGGAAACTGTCGGTTTGC'TTGA'TTCCCTGGGAGCTGTTCTCTGTCTGTGAGTGT  
 CTCTTGTCTAAACGTCATTAAGCTGAGAGTGTATGAAGACAGGATCTAGAATAATCTTGCTCACAGCTGTGCTCTGAGTGCCTAGCGGAGTTCCA  
 GCAACAAAATGGACTCAAGAGAGATTTGATTAATGAATCGTAATGAAGTGGGGTTTATTGTACAGTTAAAATGTTAGATGTTTTAATTTTTTA  
 AATAAATGGAATACTTTTTTTTTTTTTTAAAGAAAGCACTTACTGAGACAATGTAGAAGAAGTTTGTTCGGTTCTTTAATGTGGTTGAAGAGC  
 AATGTGTGGCTGAAGACTTTTGTATGAGGAGCTGCAGATTAGCTAGGGGACAGCTGGAATATGCTGGCTCTGATAATTTTAAAGGGGTCTG  
 AAATTTGTGATGGAATCAGATTTTACAGCTCTCTCAATGACATGAAAGTTCATGGAACATCATGTTTAAAGGGCTATGTAATATATGAACAT  
 TAGAAAAATGCAACTTGTGTTTACAAAAATACAAACACATGTTAGGAAGTACTGTAGGCTAGGCTGAGTGCCTACAGCTGTAATCCAGCAT  
 TTTGGGAAGCTAAGATGGGTGGATCACTTGAGGTGAGGAGTTGAGACCAGCTGGCAACATGGCGAAACCCCTCTACTAAAAATACAAAAAT  
 TGCCAGGCTGGTGGCGGGTGCCTGTAATCCAGCTACTTGGGAGGCTGAGGCAAGAGAATCGCTTGAACCCAGGAGGAGGTTGCAGTGTGAGCC  
 AGATCGTGCCATGCACTCCAGCCTGGGTGACAGAGCGAGACTCCTACTCAAAAAAAAAAAAAAAAAA

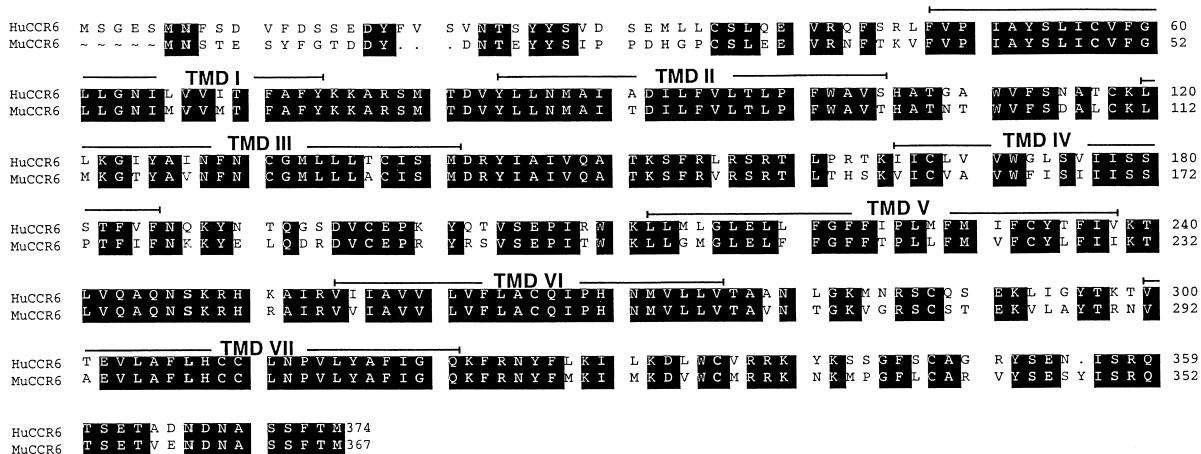
## Mouse gene:

TCTGCTCTCCCAACATCTGCACTAGTGTGAGAGTGTGGTTGAACTGCCCACTTCCCTTCTACACCAGATCTGGCTCTCCCATCCACATAGAGAACCAC  
 GCCTGCCTGGGGTGTGAAATCTACTTTATCTTGGCAGGGACTCTGGCATGGCTAGGTGTGGTTGCTTGAATCACACTGTACAGATTCTATTTTCAT  
 TATCATTCAGGAATGAATCCACAGAGTCTACTTTGGAACGGATGATTATGACAACACAGAGTATTATTTCTATTCCTCCAGACCATGGGCCATGCT  
 CCTAGAAAGAGTTCAGAACTCACCAGGATTTGTGCCAATGCTACTCCTTAATATGTGCTTTGGCCTCCTGGGCAACATTATGGTGGTGAT  
 GACCTTTGCCTTCTACAAGAAAGCCAGATCCATGACTGACGTCTACCTGTTGAACATGGCCATCACAGACATACTTTGTCTCACCTACCCTACCGTT  
 TGGGAGTTACTCATGCCACCAACACTTGGGTTTTCAGCGATGCATGTGTAACATGATGAAAGGCACATATGCGGTCACTTTAACTGTGGGATGC  
 TGCTCCTGGCCTGTATCAGCATGGACCGGTACATTTGCCATCTGTCAGGCAACCAAACTTTCCGGGTACGCTCCAGAACACTGACGCACAGTAAGT  
 CATCTGTGTGGCAGTGTGGTTTCACTCTCCATCATCTCAAGCCCTACATTTATCTTCAACAAGAAATACGAGCTGCAGGATCGTGATGTCTGTGAG  
 CCACGGTACAGGTCTGTCTCAGAGCCATCAGTGGAAAGTCTGGGTATGGGACTGGAGTGTCTTTGGGTTCTTACCCTTTGCTGTTTATGG  
 GTTCTGCTATCTGTTTATTAACAAGACCTTGGTGCAGGCCAGAACTCCAAGAGGCACAGAGCCTCCGAGTCTGTGATCGCTGTGGTTCTCGTGT  
 CCTGGCTTGTGATCCTCACAACATGGTCTCTCTGCTGACTGCGGTCAACCGGGCAAAGTGGGCGGAGCTGCAGCACCGGAGAAAGTCTCTGCC  
 TACACGGAACAGTGGCCGAGTCTGGCTTTCTGCAATGCTGCTCAACCCGTTGTATGCGTTTATTGGACAGAAATCAGAAACTACTTCA  
 TGAAGATCATGAAGGATGTGTGGTGTATGAGAAGGAAGAATAAGATGCTGCTGCTTCTGTGCCGGGTTACTCGGAAAGCTACATCTCCAGGCA  
 GACCAGTGTGAGCCGTCGAAAATGATAATGCATCGTCTTTACCATGTAACACGAGAGCACAAAGCAACATTGCCCAAAAGCCTTGGTGAACCTTGC  
 TATTACATATGAAAAAAGCCATGCCAAATATGTACAGTAACTATGAAATTCAGCAAAGACTTCTGCAAGTTTCAAGAACAGCCATGAG  
 GTGGCACTATCAGCCAAATCTTCCAGGTTGTGGTTGACAAGAAACATTGAGCTCCTCCAGGTTTGGTTTCAAAAATGAGATGGGAAATGCC  
 AGATTACTGGGTTTGTGTTAATGAACATAAACATATCCAGAAAGCTTTCATGAAGGGGTTTACAGAACTAGTTGACCCCTAACCCCCATG  
 CCACAAAACAAGGATGTTACCTTGA

## Mouse cDNA:

GGCTGCGAGAAGACGACAGAAGGGGAGCACTGCTGGTTGTGTCTGTCAACAGAATAGTCCACATCTTAGACTGGAGCCTGGATAAACCACTGA  
 GGCAGGAGTACCTGGCCAGTCTACTTTGGAGCTCAGCATTTTCTGGGGAATGAATTCACAGAGTCTACTTTGGAACGGATGATTATGACAACACA  
 GAGTATTATTCTATTCCTCCAGACCATGGGCCATGCTCCCTAGAAGAGTTCAGAACTTCACCAAGGATTTGTGCCAATGCTACTCCTTAATAT  
 GTGTCTTTGGCCTCCTGGGCAACATTATGGTGGTGTGACCTTTGCCTTCTACAAGAAAGCCAGATCCATGACTGACGTCTACCTGTTGAACATGCC  
 CATCACAGACATACTTGTCTCACCTACCCTACCGTTCTGGGAGTTACTCTATGCAACCAACACTTGGGTTTTCAGCGATGCCTGTGTAACATGATG  
 AAAGGCACATATGCGGTCAACTTTAACTGTGGGATGCTGCTCTGCTGATGCTGATGAGTGTGATGCGTTTATTGGACAGAAATCAGAAACTCTT  
 TCCGGGTACGCTCCAGAACACTGACGCACAGTAAGTCACTCTGTGTGCGAGTGTGGTTTCACTCCATCATCATCTCCAGCTCATCTTATCTTCAA  
 CAAGAAATACGAGCTGCAGGATCGTGATGTCTGTGAGCCACGGTACAGTCTGTCTCAGAGCCATCAGTGGAAAGTGTGGTATGGGACTGGAG  
 CTGTTCTTTGGGTTCTTACCCTTTGCTGTTTATGGTGTCTGCTATCTGTTTATTAACAAGCCTTGGTGCAGGCCAGAATCCAAGAGGCACA  
 GAGCCATCCGAGTCTGTGATCGCTGTGGTTCTCGTGTCTGCTGCTGATCCCTCACAACATGGTCTCTCTGACTGCGGTCAACACGGGCAA  
 AGTGGGCGGAGCTGCAGCACCGGAGAAAGTCTCGCTACACCAGGAACGTGGCCGAGGTCCTGGCTTTCTGCAATGCTGCTCAACCCGTTGT  
 TATGCGTTTATTGGACAGAAATCAGAACTACTTCAATGAAGATCATGAAGGATGTGTGGTGTATGAGAAGGAAGAATAAGATGCTGGCTTCTCT  
 GTGCGGGTTTACTCCGAAAGCTACATCTCCAGGACAGCAAGTGCAGCAAGCTGCAAAAATGATAATGCATCGTCTTTACCATGTAACACGAGAGC  
 AAAGCAACATTGCCCAAAAGCCTTGGTGAACCTTGTCTATT

**Figure 2** Comparison between human and mouse CCR6 amino acid sequences. Numbers at the right indicate the positions of the residues at the end of each line. Solid backgrounds indicate identities between the two proteins. Dots mark gaps introduced to create an optimal alignment. Tildes mark positions without corresponding residues. Putative transmembrane domains are indicated. The alignment was created using the PileUp and PrettyBox programs of the Wisconsin Sequence Analysis Package, Genetics Computer Group, Madison, WI.



predicted to contain 374 amino acids, with features that are typical for a member of the chemokine receptor subfamily of the seven transmembrane domain G protein-coupled receptor superfamily. These include an acidic N-terminal domain, a small and basic third intracellular loop, cysteine residues in the N-terminal domain (C36) and the third extracellular loop (C288), and a conserved DRY motif following the third transmembrane domain.

## Relevant homologies and species differences

While obviously related to other chemokine receptors by sequence, CCR6 is not closely related to other receptors, showing greatest similarity to CCR7 (39% identity). Overall, human CCR6 is identical at 74% of its residues, as compared with mouse protein. As shown in Figure 2, mismatched residues are found disproportionately in the N-terminal region.

## Affinity for ligand(s)

$K_d$  for MIP-3 $\alpha$  on CCR6-transfected cells is 0.1–12 nM (Baba *et al.*, 1997; Greaves *et al.*, 1997; Power *et al.*, 1997) and on lymphocytes is 0.4 nM (Hieshima *et al.*, 1997).

## Cell types and tissues expressing the receptor

CCR6 mRNA is expressed in lymphoid tissue including spleen, lymph node, thymus, appendix, and peripheral blood lymphocytes (Zaballos *et al.*, 1996; Liao *et al.*, 1997c; Varona *et al.*, 1998) and mRNA and protein are found in/on CD4<sup>+</sup> CD8<sup>+</sup> T cells, dendritic cells derived from CD34<sup>+</sup> bone marrow progenitors, and B cells (Zaballos *et al.*, 1996; Baba *et al.*, 1997; Greaves *et al.*, 1997; Liao *et al.*, 1997b, 1999; Power *et al.*, 1997; Varona *et al.*, 1998). On T cells, CCR6 is limited to those with a memory phenotype (Liao *et al.*, 1999). CCR6 is found not only on freshly isolated peripheral blood T cells, but also on tumor-infiltrating lymphocytes that have been repeatedly activated *in vitro* (Liao *et al.*, 1997a,b,c).

## Regulation of receptor expression

Some investigators have reported induction of CCR6 mRNA on T cells after treatment with IL-2 (Baba *et al.*, 1997), although others have reported down-regulation of mRNA after T cell activation with anti-CD3 and PMA (Greaves *et al.*, 1997). Activation of T cells *in vitro* with anti-CD3 or IL-2 did not upregulate CCR6 surface expression (Liao *et al.*, 1999) or responses to LARC/MIP-3 $\alpha$ . Maturation of CD34<sup>+</sup> progenitor-derived dendritic cells *in vitro* was

associated with downregulation of CCR6 mRNA and responses to LARC/MIP-3 $\alpha$  (Dieu *et al.*, 1998).

## SIGNAL TRANSDUCTION

### Cytoplasmic signaling cascades

Calcium signaling was blocked or diminished in tumor-infiltrating lymphocytes and in transfected cells by the addition of pertussis toxin, implicating G $\alpha_i$  proteins (Liao *et al.*, 1997a; Power *et al.*, 1997), and in transfected cells by inhibiting phospholipase C and depleting stores of intracellular calcium (Power *et al.*, 1997).

## BIOLOGICAL CONSEQUENCES OF ACTIVATING OR INHIBITING RECEPTOR AND PATHOPHYSIOLOGY

### Unique biological effects of activating the receptors

CCR6 is unusual in that it is a receptor for an inflammation-induced chemokine that functions well on freshly isolated lymphocytes, i.e. on nonactivated cells, producing chemotaxis (Hieshima *et al.*, 1997), calcium flux (Liao *et al.*, 1997a), and rapid adherence to ICAM-1-coated glass under conditions of flow (Campbell *et al.*, 1998). These activities were found on both CD4 $^+$  and CD8 $^+$  T cells, and were limited to T cells with a memory phenotype (Campbell *et al.*, 1998; Liao *et al.*, 1999). Consequently, CCR6 is the only chemokine receptor that functions well in the above assays specifically on nonactivated memory T cells. CCR6 is also unusual in being expressed and functional on a subset of immature dendritic cells and then being downregulated as the dendritic cells mature (Dieu *et al.*, 1998).

## References

Baba, M., Imai, T., Nishimura, M., Kakizaki, M., Takagi, S., Hieshima, K., Nomiyama, H., and Yoshie, O. (1997). Identification of CCR6, the specific receptor for a novel lymphocyte-directed CC chemokine LARC. *J. Biol. Chem.* **272**, 14893–14898.

Campbell, J. J., Hedrick, J., Zlotnik, A., Siani, M. A., Thompson, D. A., and Butcher, E. C. (1998). Chemokines and the arrest of lymphocytes rolling under flow conditions. *Science* **279**, 381–384.

Dieu, M. C., Vanbervliet, B., Vicari, A., Bridon, J. M., Oldham, E., Ait-Yahia, S., Briere, F., Zlotnik, A., Lebecque, S., and Caux, C. (1998). Selective recruitment of immature and mature dendritic cells by distinct chemokines expressed in different anatomic sites. *J. Exp. Med.* **188**, 373–386.

Greaves, D. R., Wang, W., Dairaghi, D. J., Dieu, M. C., Saint-Vis, B., Franz-Bacon, K., Rossi, D., Caux, C., McClanahan, T., Gordon, S., Zlotnik, A., and Schall, T. J. (1997). CCR6, a CC chemokine receptor that interacts with macrophage inflammatory protein 3 $\alpha$  and is highly expressed in human dendritic cells. *J. Exp. Med.* **186**, 837–844.

Hieshima, K., Imai, T., Opdenakker, G., Van Damme, J., Kusuda, J., Tei, H., Sakaki, Y., Takatsuki, K., Miura, R., Yoshie, O., and Nomiyama, H. (1997). Molecular cloning of a novel human CC chemokine liver and activation-regulated chemokine (LARC) expressed in liver. Chemotactic activity for lymphocytes and gene localization on chromosome 2. *J. Biol. Chem.* **272**, 5846–5853.

Liao, F., Alderson, R., Su, J., Ullrich, S. J., Kreider, B. L., and Farber, J. M. (1997a). STRL22 is a receptor for the CC chemokine MIP-3 $\alpha$ . *Biochem. Biophys. Res. Commun.* **236**, 212–217.

Liao, F., Alkhatib, G., Peden, K. W. C., Sharma, G., Berger, E. A., and Farber, J. M. (1997b). STRL33, a novel chemokine receptor-like protein, functions as a fusion cofactor for both macrophage-tropic and T cell line-tropic HIV-1. *J. Exp. Med.* **185**, 2015–2023.

Liao, F., Lee, H.-H., and Farber, J. M. (1997c). Cloning of STRL22, a new human gene encoding a G protein-coupled receptor related to chemokine receptors and located on chromosome 6q27. *Genomics* **40**, 175–180.

Liao, F., Rabin, R. L., Smith, C. S., Sharma, G., Nutman, T. B., and Farber, J. M. (1999). CC-chemokine receptor 6 is expressed on diverse memory subsets of T cells and determines responsiveness to macrophage inflammatory protein 3  $\alpha$ . *J. Immunol.* **162**, 186–194.

Power, C. A., Church, D. J., Meyer, A., Alouani, S., Proudfoot, A. E. I., Clark-Lewis, I., Sozzani, S., Mantovani, A., and Wells, T. N. C. (1997). Cloning and characterization of a specific receptor for the novel CC chemokine MIP-3 $\alpha$  from lung dendritic cells. *J. Exp. Med.* **186**, 825–835.

Tanaka, Y., Imai, T., Baba, M., Ishikawa, I., Uehira, M., Nomiyama, H., and Yoshie, O. (1999). Selective expression of liver and activation-regulated chemokine (LARC) in intestinal epithelium in mice and humans. *Eur. J. Immunol.* **29**, 633–642.

Varona, R., Zaballos, A., Gutierrez, J., Martin, P., Roncal, F., Albar, J. P., Ardavin, C., and Marquez, G. (1998). Molecular cloning, functional characterization and mRNA expression analysis of the murine chemokine receptor CCR6 and its specific ligand MIP-3 $\alpha$ . *FEBS Lett.* **440**, 188–194.

Zaballos, A., Varona, R., Gutierrez, J., Lind, P., and Marquez, G. (1996). Molecular cloning and RNA expression of two new human chemokine receptor-like genes. *Biochem. Biophys. Res. Commun.* **227**, 846–853.

## LICENSED PRODUCTS

Anti-human CCR6 mouse IgG<sub>2B</sub> monoclonal antibody, clone 53103.111 from R&D Systems for flow cytometry.

