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Synthesis and Characterization of Fluorescent and Photoactivatable MIP-1 α Ligands and Interactions with Chemokine Receptors CCR1 and CCR5[§]

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Photoaffinity and fluorescent analogues of the 70-amino acid chemokine macrophage inflammatory protein-1 α (MIP-1 α) were designed, synthesized, characterized, and applied to probe MIP-1 α interactions with the chemokine receptors CCR1 and CCR5. The photoactivatable MIP-1 α ligand, BP-MIP-1 α , and the fluorescent ligand, Flu-MIP-1 α were prepared by selective chemical coupling of *p*-benzoylphenylthiocarbamyl or fluoresceinthiocarbamyl, respectively, at the N-terminus of MIP-1 α . Both ligands BP-MIP-1 α and Flu-MIP-1 α retained high binding affinity and agonist potency at CCR1 and CCR5. Photoaffinity labeling of CCR1 and CCR5 receptors stably expressed in CHO cells resulted in specific covalent attachment of [¹²⁵I]BP-MIP-1 α and production of protein complexes of 54 and 48 kDa, respectively, on SDS-PAGE. This represents the first photo-cross-linking between a chemokine and its receptor. Flu-MIP-1 α selectively labeled CCR1 or CCR5 receptors expressed in CHO cells and was used to characterize receptor binding domains. When bound to CCR1 or CCR5 receptors, the fluorescence signal of Flu-MIP-1 α was quenched by collision with iodide indicating that the N-terminal end of MIP-1 α is accessible to the solvent. These data strongly suggest that the N-terminal end of MIP-1 α interacts with domains of CCR1 or CCR5 receptors located at the extracellular surface. The photoactivatable BP-MIP-1 α described here should prove valuable for the identification of contact sites on receptors by photoaffinity labeling experiments.

Introduction

Chemokines represent a large family of 8–10-kDa proinflammatory peptides that play a role in the recruitment and activation of leukocytes at sites of inflammation.^{1,2} Chemokine actions are mediated by cell membrane receptors predicted to have seven transmembrane segments and coupled to G-proteins.³ Recently, some of these receptors have also been shown to act as co-receptors for HIV virus entry into cells.⁴ To date more than 40 chemokines are known and at least 15 receptors have been characterized. Structurally, chemokines can be classified in two subgroups based on the spacing of a motif containing cysteines near the N-terminus: the CXC chemokines in which two cysteine residues are spaced apart by one amino acid and the CC chemokines in which the two cysteines are adjacent. Most of the chemokine receptors are promiscuous, at least in vitro, in the sense that they recognize and transduce signals in response to more than one chemokine. For instance, the CCR1 receptor is a functional receptor for, at least, the chemokines RANTES, MIP-1 α , monocyte chemoat-

tractant peptide-2 and -3 (MCP-2, MCP-3), and MIP-5, although the sequence homology between these chemokines is low. Conversely, the chemokine MIP-1 α binds to and elicits a signaling response through, at least, the CCR1, CCR4, CCR5, and US28 receptors.^{2,3}

Little is known at the molecular level about chemokine ligand recognition by receptors. For most receptors studied so far, mutational analysis and chimeric receptor work suggests that chemokine binding determinants are broadly distributed over a large surface area including multiple domains on the extracellular face of the receptors. One exception is CCR2 for which the N-terminal extracellular extension is both necessary and sufficient for high-affinity binding of MCP-1.⁵ A two-site binding model has been proposed for the interaction of chemokines with their receptors.^{6,7} In this model, a first interaction with the extracellular domains of the receptor controls selectivity and a second interaction with the core of the receptor triggers the activation and the signaling response. Understanding the functional relationships between these two sites as well as the identification of residues involved in the binding and activation interactions will require more experimental work.

Direct covalent attachment of photolabile ligands is a method of choice for the determination of contact sites between hormones and their receptors. Among the different photochemically reactive functional groups, the benzophenone (BP) group offers the advantages of stability to ambient light, activation at a wavelength (350 nm) that does not damage proteins, and preferential insertion into C–H bonds even in the presence of

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[§] Abbreviations: HIV, human immunodeficiency virus; MIP-1 α , macrophage inflammatory protein-1 α ; RANTES, regulated on activation normal T-cell expressed and secreted; CCR1, CC chemokine receptor 1; CCR5, CC chemokine receptor 5; BP, 4-benzoylphenyl; Flu, 5-fluoresceinyl; CHO, Chinese hamster ovary; PBS, phosphate-buffered saline; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; amu, atomic mass unit; HA, hemagglutinin.

solvent water.⁸ Benzophenone probes have been used for photolabeling a variety of short to medium-size peptides binding to their cognate G-protein-coupled receptors such as substance P,^{9,10} angiotensin II,¹¹ parathyroid hormone,¹² secretin,¹³ or cholecystokinin.¹⁴ Fluorescent ligands have found many applications to study ligand–receptor interactions and for probing the microenvironment of binding pockets^{15–18} as well as for the detection of receptor expression in cells or tissues.¹⁹

Here we report the design, synthesis, and pharmacological characterization of the first photoactivatable and fluorescent analogues of the chemokine MIP-1 α with high affinity for the chemokine receptors CCR1 and CCR5. We show that these reagents can be used in photoaffinity cross-linking and fluorescence techniques to label the CCR1 and CCR5 receptors in transfected cell lines. Furthermore, these ligands should prove valuable to study the mode of recognition of the chemokine MIP-1 α by CCR1 and CCR5.

Experimental Section

Materials. Benzophenone 4-isothiocyanate was purchased from Molecular Probes (Eugene, OR) and fluoresceinyl isothiocyanate was purchased from Fluka (Buchs, Switzerland). [¹²⁵I]-MIP-1 α and [¹²⁵I]BP-MIP-1 α were prepared by Amersham (Buckinghamshire, U.K.) by oxidative ¹²⁵I₂ labeling. Wild-type MIP-1 α was from PeProTech (Rocky Hill, NJ).

General Methods. HPLC of proteins was effected on a Waters (Millipore Corp.) system, model 510 pumps linked to an automated gradient controller, model 481 UV/vis detector, using a Nucleosil 300-7 C8 column (Macherey-Nagel ET250/8/4); buffer A, 0.1% (w/v) TFA/water; buffer B, 0.09% TFA/80% acetonitrile/water; flow rate, 1.0 mL/min; gradient, 0–75% B at 1% B/min. Proteins were detected at 214 nm. Both BP-MIP-1 α and Flu-MIP-1 α were purified by HPLC, isolated as lyophilized CF₃CO₂H salts, and quantified by amino acid analysis. They were frequently checked by analytical HPLC.

Purified proteins and their derivatives were analyzed by Edman N-terminal sequencing on an Applied Biosystem, Inc. Procise 494. For compositional amino acid analysis, about 1 nmol of protein was hydrolyzed in gas phase under vacuum for 24 h in 6 M HCl containing 1 mg/mL phenol at 112 °C. The resultant amino acids were quantified using a Waters AccQTag chemistry and a Hewlett-Packard HP 1090 liquid chromatograph equipped with a HP 1046A fluorescence detector. Peptides were authenticated by MS (Micromass Q-TOF). Absorbance spectra were obtained using a Jasco V-550 UV/vis spectrophotometer using a bandwidth of 2 nm and a scanning speed of 100 nm/min. Data were collected every 0.5 nm (data pitch).

Synthesis of BP-MIP-1 α . The chemokine Δ Ala¹-MIP-1 α (A10T) was expressed in *E. coli* and purified as described.²⁰ To 1.0 mg of Δ Ala¹-MIP-1 α (A10T) (0.125 μ mol) in 100 μ L of water was added 100 μ L of sodium borate (50 mM, pH 8.0) at 4 °C. To this solution was added 14 μ L of a freshly prepared 10 mM solution of benzophenone 4-isothiocyanate in *N,N*-dimethylformamide. The mixture was stirred at room temperature for 3 h. The reaction was monitored by RP-HPLC using conditions described in General Methods. Portions of benzophenone 4-isothiocyanate (14 μ L) were added again at regular 2–3-h intervals until about 50% of the starting material had reacted. Attempts to push the reaction to completion resulted in the formation of di-BP and tri-BP MIP-1 α side products. Monolabeled BP-MIP-1 α was purified by HPLC as described in General Methods and lyophilized to a white powder. The yield was 30% (or 60% based on recovered starting material). Edman degradation for BP-MIP-1 α gave the sequence XLAADTPPT where X is a nonnatural amino acid, whereas analysis of Δ Ala¹-MIP-1 α (A10T) gave SLAADTPTT. Amino acid analysis agreed with the expected amino acid

composition. MS (electrospray ionization) for BP-MIP-1 α : calcd 7982.6 amu, found 7982.5 $\pm$ 0.4 amu.

Synthesis of Flu-MIP-1 α . To 50 μ L of a solution of Δ Ala¹-MIP-1 α (A10T) (1.25 mM) in water was added 100 μ L of sodium borate (50 mM, pH 8.0) at 4 °C. To this solution was added 3 μ L of a 25 mM solution of FITC in *N,N*-dimethylformamide. This represents a 1.2-fold molar excess of FITC over protein. The mixture was incubated at room temperature in the dark for 3 h with continuous stirring. Flu-MIP-1 α was purified directly from the reaction mixture by RP-HPLC under the conditions described in General Methods. Pooled fractions containing pure Flu-MIP-1 α were lyophilized. The yield was 62%. N-Terminal sequencing gave the expected sequence XLAADTPTA, where X means no amino acid detected in the first cycle consistent with the release of the FITC-Ser at the N-terminus. Amino acid analysis of Flu-MIP-1 α hydrolysate was in agreement with the expected theoretical composition. MS analysis: calcd 8131.45 amu, found 8131.53 $\pm$ 0.02 amu.

Cell Culture. Photoaffinity labeling and fluorescence experiments were performed with CHO cells stably transfected with the N-terminally HA-tagged human CCR1 receptor (CCR1/CHO cells) or the N-terminally HA-tagged human CCR5 receptor (CCR5/CHO cells) at expression levels of about 15 000 and 30 000 receptors/cell, respectively. The HA epitope (YPYDVPYASLRS) was added to the 5'-end of the receptor cDNA by PCR and the CHO cell lines were obtained as described.²³ The cells were cultured as monolayers in a humidified 5% CO₂ atmosphere at 37 °C, in Dulbecco's MEM/NUT mix F-12, supplemented with 10% fetal calf serum, 2% Pen-Strep, 2 mM L-glutamine. Cells were harvested at 80% confluence with PBS containing 1 mM EDTA and washed with ice-cold PBS. Membrane fractions were prepared as previously described.²¹

Radioligand Binding Assays. Competition binding experiments were carried out on CCR1/CHO cell membrane fractions using the scintillation proximity assay (SPA) methodology with [¹²⁵I]MIP-1 α as a tracer²¹ and CCR5/CHO whole cells with [¹²⁵I]-MIP-1 α as a tracer following published protocols.²⁰ All determinations were performed in triplicate, and the nonspecific binding was determined in the presence of 1 μ M MIP-1 α .

Ca²⁺ Mobilization Assays. Changes in intracellular Ca²⁺ concentration evoked by agonist stimulation were monitored spectrofluorometrically in CCR1/CHO and CCR5/CHO cells loaded with the Ca²⁺-sensitive dye Fura-2-AM. Cells grown in a 225-cm² flask were harvested at 80% confluence with PBS containing 1 mM EDTA, resuspended in 15 mL of cell culture medium and left for 1–2 h at 37 °C. Cells were spun down and the medium was removed. Cells were resuspended in 10 mL of Krebs-Ringer medium containing 10 μ g of bovine serum albumin and 20 μ L of a 1 mg/mL solution of fluorescent Ca²⁺-sensitive agent Fura-2-AM (Fluka) was added. The mixture was incubated at 37 °C in the dark for 30 min and then kept on ice until use. Cells were quickly spun down just before use and the medium was discarded. Cells were resuspended at 5 $\times$ 10⁶ cells/mL in Krebs-Ringer buffer at 37 °C. For the measurements, 0.5-mL aliquots were transferred to a quartz cuvette under constant stirring in a Jasco 777 spectrofluorometer. Irradiation was at 340 nm and fluorescence intensity was recorded at 505 nm (or 480 nm for assays with Flu-Mip-1 α). When the emission signal was stable, ligands were added in 5 μ L. Ca²⁺ concentration increase was expressed in percent (%) of maximal fluorescence intensity as determined by addition of Triton-X100. EC₅₀ for ligands were calculated by analysis of the data using software Grafit (Erithacus Software, Ltd.).

Photo-Cross-Linking Experiments. CHO cells stably transfected with either human CCR1, CCR5, or NK1 receptors were transferred to 24-well culture plates, 2 $\times$ 10⁵ cells/well, 24 h before the photo-cross-linking reaction. Cells were at about 80% confluency at the time of photo-cross-linking. Cells were washed in ice-cold buffer A (50 mM HEPES, 1 mM CaCl₂, 5 mM MgCl₂, 0.5% w/v bovine serum albumin, pH 7.2) and incubated with 1 nM [¹²⁵I]BP-MIP-1 α in buffer A for 4 h at 4 °C. Nonspecific cross-linking was determined in the presence

of 100 nM MIP-1 α . Cells were irradiated on ice for 20 min using an UV light at 365 nm (Philips HPR 125W lamp) at 6-cm distance. Cells were then washed twice in ice-cold PBS, scraped off and isolated by centrifugation (500*g*) for 15 min at 4 °C. The cell pellet was dissolved in 40 μ L of 10% SDS. After removal of cell debris by centrifugation, the extract was analyzed by denaturing PAGE under reducing conditions (4–20% Tris-glycine gel from Novex, 4 M urea, 2% β -mercaptoethanol). Cross-linking products were revealed by autoradiography.

Immunoprecipitation. CCR1/CHO cells labeled by photoaffinity with BP-MIP-1 α were incubated in buffer containing 300 mM NaCl, 2% w/v NP-40, 1% w/v deoxycholate, 0.2% w/v SDS in 100 mM Tris-HCl pH 7.5, on ice for 45 min. Cells were scraped off and centrifuged (100000*g*, 20 min, 4 °C) to remove the debris. Mouse monoclonal anti-HA antibody (12CA5 from Boehringer) and goat anti-mouse IgG magnetic beads (Dyna-beads) were added and the mixture was incubated with gentle stirring overnight at 4 °C. Beads were washed twice with PBS, 0.1% bovine serum albumin, and the complex was then released from the beads by incubating in 10% SDS for 30 min at 20 °C.

Alternatively, the CCR1–MIP-1 α complex was immunoprecipitated as described above using rabbit anti-MIP-1 α IgG (PeptoTech, Inc., Rocky Hill, NJ) and sheep anti-rabbit Dyna-beads.

Fluorescence Measurements. Binding of fluorescent Flu-MIP-1 α to the CCR1 or CCR5 receptors was measured on a suspension of cell membranes from stably transfected CHO cells expressing about 30 000 receptors/cell. Membrane suspensions at a total protein concentration of 0.7 mg/mL were incubated in 50 mM sodium Hepes, pH 7.2, 1 mM CaCl₂, 5 mM MgCl₂, 0.5% (w/v) BSA, 0.5 mM sodium azide with 50 nM Flu-MIP-1 α in the presence (nonspecific binding) or absence (total binding) of 17 μ M MIP-1 α at 20 °C for 1 h. As an additional control, membranes were also incubated in the absence of fluorescent ligand. Membranes were then washed twice with ice-cold PBS pH 7.2, resuspended in 500 μ L of ice-cold PBS and homogenized by sonication in an ice-cold bath for 3 min immediately prior to fluorescence measurements. Fluorescence spectra were recorded using a Jasco FP-750 spectrofluorometer with a 0.5 $\times$ 1.0-cm quartz cuvette. During the recording of emission spectra, membrane suspensions were continuously stirred at 20 °C. Fluorescence was corrected for light scattering and background fluorescence from control samples. The excitation wavelength was 475 nm and the emission was recorded from 500 to 600 nm. The spectra were generated using the following acquisition parameters: excitation and emission bandwidths, 5 and 10 nm, respectively; data pitch, 1 nm; scanning speed, 125 nm/min. The fluorescence measurements were made within the linear portion of the photon multiplier tube's response. Spectra were not corrected for lamp intensity or for PMT sensitivity.

For collisional quenching experiments, membrane suspensions were incubated as described above and increasing amounts of KI were added to a final concentration of 50 mM. Changes in fluorescence due to the addition of the quencher KI were corrected by subtracting the fluorescence measured in parallel control samples to account for autofluorescence quenching and dilution effects upon addition of the quencher. Typically, a complete quenching experiment was performed in about 6–8 min. Under these conditions the contribution of fluorescence signal from dissociated ligand was negligible. As a control for ligand dissociation, samples were incubated during the times typically allocated to quenching experiments, centrifuged and the measurement repeated. No changes in the fluorescence values are indicative of negligible dissociation of ligand. Also, we checked that the equilibrium binding of [¹²⁵I]-MIP-1 α to either CCR1 or CCR5 receptors was not affected by 50 mM KI. The quencher stock solution was 0.153 M KI containing 1 mM Na₂S₂O₃ to prevent formation of I₃[−]. The quenching of the fluorescence emission at 520 nm was calculated with the Stern–Volmer equation as described.¹⁵ The Stern–Volmer constant *K*_{sv} was determined from the slope of

Figure 1. Structure of the MIP-1 α derivatives BP-MIP-1 α and Flu-MIP-1 α .

the ratio of fluorescence intensities in the absence and in the presence of iodide, F_0/F , as a function of the iodide concentration.

Results

Synthesis of the Photoactivatable MIP-1 α Analogue BP-MIP- α . The chemokine Δ Ala¹-MIP-1 α (A10T), a variant of human MIP-1 α lacking the first amino acid Ala¹ and in which Ala¹⁰ was substituted with Thr for reasons not related to this study, was derivatized with the photoactivatable benzophenone group to yield the MIP-1 α analogue *N*- α -(4-benzoylphenyl)thiocarbamoyl- Δ Ala¹-MIP-1 α (A10T) (hereafter BP-MIP-1 α ; Figure 1) as described in the Experimental Section. Under these conditions, isothiocyanate electrophiles react preferentially with the low *pK*_a N-terminal amino group of peptides rather than with the high *pK*_a ϵ -amino groups on lysyl side chains.²² Comparative Edman degradation of BP-MIP-1 α and MIP-1 α and analysis of the phenylthiohydantoin derivatives at each cycle confirmed the preferential attachment of benzophenone at the N-terminal position of MIP-1 α . More than 95% of the benzophenone group was on the N-terminal amine.

Synthesis of the Fluorescent MIP-1 α Analogue Flu-MIP-1 α . Fluoresceinthiocarbamoyl- Δ Ala¹-MIP-1 α (A10T) (Flu-MIP-1 α ; Figure 1) was synthesized from fluorescein isothiocyanate and Δ Ala¹-MIP-1 α (A10T) as described in the Experimental Section. Flu-MIP-1 α was labeled at >95% on the N-terminal amine of the peptide as determined by Edman sequencing.

Pharmacological Characterization of BP-MIP-1 α and Flu-MIP-1 α . BP-MIP-1 α and Flu-MIP-1 α were assayed for chemokine receptor binding affinity by competitive displacement binding analysis with the [¹²⁵I]-labeled MIP-1 α ligand on membranes of CHO cells stably transfected with the CCR1 or CHO cells transfected with the CCR5 receptor tagged with a HA epitope at the extreme N-terminus (Figure 2A,B). It was shown previously that the tagged receptors retained high-affinity ligand binding and ligand specificity.²³ As shown in Table 1, BP-MIP-1 α conserved high-nanomolar affinity at CCR1 and was even binding tighter than MIP-1 α . BP-MIP-1 α bound to CCR5 with 20.3 nM affinity, comparable to wild-type MIP-1 α (8.9 nM) (Table 2). Flu-

Figure 2. Competition binding experiments in human CCR1 and CCR5 receptors: (A) CCR1 receptor; (B) CCR5 receptor. Binding experiments were performed on CHO cell membranes for CCR1 or whole cells for CCR5 as described in the Experimental Section. Competition binding of [125 I]MIP-1 α by Δ Ala¹-MIP-1 α (A10T) (●), BP-MIP-1 α (○), or Flu-MIP-1 α (×); average curves (means $\pm$ SE) of three independent experiments are shown.

Table 1. Pharmacological Constants for MIP-1 α Derivatives on CCR1^a

ligand	binding		Ca ²⁺ mobilization in CHO cells	
	K _i (nM)	change (-fold)	EC ₅₀ (nM)	change (-fold)
MIP-1 α	14.0 $\pm$ 1.1 (7)		1.23 $\pm$ 0.66 (2)	
Δ Ala ¹ -MIP-1 α (A10T)	8.50 $\pm$ 1.19 (5)	0.6	2.11 $\pm$ 0.53 (4)	1.7
BP-MIP-1 α	1.67 $\pm$ 0.45 (3)	0.1	1.00 $\pm$ 0.45 (3)	0.8
Flu-MIP-1 α	74.2 $\pm$ 12.4 (3)	5.3	75.0 $\pm$ 12.6 (3)	61

^a The number of independent experiments is shown in parentheses.

Table 2. Pharmacological Constants for MIP-1 α Derivatives on CCR5^a

ligand	binding		Ca ²⁺ mobilization in CHO cells	
	K _i (nM)	change (-fold)	EC ₅₀ (nM)	change (-fold)
MIP-1 α	8.9 $\pm$ 2.7 (4)		3.5 $\pm$ 0.9 (3)	
Δ Ala ¹ -MIP-1 α (A10T)	10.2 $\pm$ 1.9 (7)	1.1	59.3 $\pm$ 21.5 (4)	17
BP-MIP-1 α	20.3 $\pm$ 2.9 (4)	2.3	60.9 $\pm$ 31.4 (3)	17
Flu-MIP-1 α	28.1 $\pm$ 4.5 (3)	3.2	261 $\pm$ 24 (3)	83

^a The number of independent experiments is shown in parentheses.

MIP-1 α showed a 5-fold decrease in affinity at CCR1 and a 3-fold decrease at CCR5 (Tables 1 and 2). These results indicate that replacement of Ala¹ in MIP-1 α by the more bulky benzophenone group had no significant effect on the binding affinity. However, further increase of the molecular volume at the N-terminal position, as

Figure 3. Biological activity of MIP-1 α analogues. Stimulation of CCR1 (A) and CCR5 (B) in CHO cells assayed by mobilization of intracellular calcium in response to Δ Ala¹-MIP-1 α (A10T) (●), BP-MIP-1 α (○), or Flu-MIP-1 α (×). See the Experimental Section for details; average curves (means $\pm$ SE) of three independent experiments are shown.

in Flu-MIP-1 α , caused a slight decrease in receptor affinity. This can be best explained by the steric hindrance or creation of unfavorable electronic contacts between the fluorescein group in the receptor binding pocket.

The ability of N-terminally modified MIP-1 α analogues to activate either CCR1 or CCR5 receptors was assayed spectrophotometrically by measuring the agonist-evoked increase in intracellular concentration of calcium ions in CHO cells loaded with the dye Fura-2. Both BP-MIP-1 α and Flu-MIP-1 α conserved full agonist efficacy at the CCR1 receptor (Figure 3A). Though the potency of BP-MIP-1 α (EC₅₀ = 1 nM) and wild-type MIP-1 α (EC₅₀ = 1.2 nM) for CCR1 were virtually identical, Flu-MIP-1 α (EC₅₀ = 75 nM) showed a 60-fold decrease (Table 1). On CCR5, both N-terminally modified ligands were also full agonist albeit their potency was decreased 17-fold for BP-MIP-1 α and 83-fold for Flu-MIP-1 α , respectively, as compared to wild-type MIP-1 α . However this effect was much less pronounced when compared to the parent peptide Δ Ala-MIP-1 α (Figure 3B and Table 2). Similar variations in binding interactions and agonist potency at CCR1 and CCR5 receptors were recently reported for naturally occurring N-terminal isoforms of MIP-1 α .²⁶

Photoaffinity Labeling of CCR1 and CCR5. Irradiation at 365 nm of ¹²⁵I-labeled BP-MIP-1 α bound to CCR1/CHO or CCR5/CHO cells produced a cross-

Figure 4. Photoaffinity cross-linking of [^{125}I]BP-MIP-1 α to CCR1 (A) and CCR5 (B) in CHO cells. [^{125}I]BP-MIP-1 α (1 nM) was incubated with cells expressing CCR1 or CCR5 in the presence (+) or absence (-) of MIP-1 α (100 nM) or with cells expressing the human neurokinin NK1 receptor, then irradiated at 365 nm for 20 min and analyzed by sizing SDS-PAGE.

linked material of apparent molecular weight 54 or 48 kDa, respectively, on SDS-PAGE (Figure 4A,B). Several controls were carried out to ensure the specificity of photo-cross-linking. In the absence of light irradiation no product was observed. In the presence of a 100-fold excess of MIP-1 α competitor ligand, no cross-link material was visible. In addition [^{125}I]MIP-1 α lacking the benzophenone group was not complexed to the chemokine receptors upon light irradiation. The photo-cross-linking was specific for CCR1 or CCR5 and did not occur on CHO cells expressing the human neurokinin NK1 receptor. The photo-cross-linking of [^{125}I]BP-MIP-1 α was rapid, did not seem to progress further after 5 min, and was ligand concentration-dependent (Figure 5A–D). The apparent mass of 54 kDa is consistent with a 1:1 adduct between MIP-1 α (8 kDa) and CCR1. The calculated molecular weight of HA-CCR1 is 41 kDa, but the observed apparent migration on SDS polyacrylamide gel by Western blotting with the anti-HA monoclonal antibody 12CA5 is 46 kDa (data not shown). The 54-kDa band was also observed in immunoprecipitation experiments of the covalent photoproduct with either the monoclonal antibody 12CA5, directed against the HA-tag on CCR1, or with an anti-MIP-1 α antibody (Figure 6). In addition, chemical cross-linking of [^{125}I]MIP-1 α to CCR1 on CHO cells with *N*-ethyl-*N*-diethylaminocarbodiimide produced an adduct migrating at 54 kDa (data not shown). The apparent mass of the MIP-1 α -HA-CCR5 cross-linking complex was 48 kDa, a value in agreement with a 1:1 chemokine-receptor adduct.

Interaction of the Fluorescent Ligand Flu-MIP-1 α with CCR1 and CCR5 Receptors in Cells. Membrane fractions from CHO cells expressing the

Figure 5. Time and concentration dependence of photo-cross-linking of [^{125}I]BP-MIP-1 α to chemokine receptors: (A, C) CCR1; (B, D) CCR5. In panels A and B, the concentration of [^{125}I]BP-MIP-1 α was 0.5 nM and the times of irradiation were as indicated. In panels C and D, the time of irradiation was 20 min and the concentrations of [^{125}I]BP-MIP-1 α were as indicated.

CCR1 or CCR5 receptors were specifically labeled with the fluorescent chemokine Flu-MIP-1 α . Figure 7A shows the absorption and emission spectra. Figure 7B,C shows the total bound fluorescence (T), the nonspecific bound fluorescence (N), and the fluorescence from control membranes (C). For both receptors the nonspecific fluorescence signal N represents about 28% of the total signal, giving a ratio of signal-to-noise of about 3.5.

Figure 6. Immunoprecipitation of photoaffinity-labeled CCR1 receptor. Detection of the covalent complex between [125 I]BP-MIP1 α and HA-CCR1 by immunoprecipitation with the anti-HA monoclonal antibody 12CA5 (left lane) and with an anti MIP-1 α antibody (right lane).

Collisional quenching experiments were performed to probe the accessibility of the fluorescent group tagged at the N-terminus of the bound chemokine Flu-MIP-1 α in the ligand–receptor complexes with either CCR1 or CCR5. Time course measurements with excitation at 475 nm and emission at 522 nm were recorded using iodide as a quencher. Figure 8A,B shows representative traces of quenching analysis in triplicate performed in at least two independent experiments for each receptor. Quenching experiments with iodide gave similar Stern–Volmer constants (K_{sv}) for the fluorescent Flu-MIP-1 α when bound to CCR1 or CCR5 in CHO membranes compared to the ligand free in PBS solution. The averaged values of K_{sv} from three independent measurements were 11.0 ± 0.9 , 11.2 ± 1.5 , and 11.9 ± 1.9 M $^{-1}$ for Flu-MIP1 α free in solution, specifically bound to CCR1, and specifically bound to CCR5, respectively. In contrast, for Flu-MIP-1 α nonspecifically bound to CCR5/CHO membranes a K_{sv} value of 1.1 ± 1.0 M $^{-1}$ was obtained (data not shown). These results strongly suggest complete accessibility of the N-terminal site of the chemokine to the aqueous soluble quencher KI when the ligand is bound specifically to either CCR1 or CCR5 receptors. In contrast, the fluoresceinyl group is buried and not accessible to the aqueous phase quencher when the ligand interacts nonspecifically with the membranes.

Discussion

MIP-1 α is an important CC chemokine involved in many inflammatory processes. So far, there have been very few attempts to study the interactions between the prototypic MIP-1 α , or other chemokines in general, and their receptors at the molecular level. The little information available is inferred from site-directed mutagenesis or the construction of chimeric receptors. For either of these mutational mapping approaches, it is difficult to discriminate between direct effects on ligand binding affinity or allosteric effects on receptor folding. In this study, we have developed probes to map the binding site of the N-terminus of MIP-1 α to the CCR1 and CCR5 receptors. A photoactivatable analogue (BP-MIP-1 α) and a fluorescent analogue (Flu-MIP-1 α) were designed and prepared by replacing the Ala¹ residue of MIP-1 α with the benzophenyl and fluoresceinyl functional groups, respectively. Both analogues BP-MIP-1 α and Flu-MIP-1 α conserved high binding affinity and reasonable agonist potency in the calcium mobilization assay at CCR1 and CCR5, indicating that these modified chemokines bind to their cognate receptors in the same mode

Figure 7. Fluorescence spectra of Flu-MIP-1 α . Absorption and emission spectra of a 2.5 μ M solution of Flu-MIP-1 α in PBS are shown in panel A. Fluorescence of Flu-MIP-1 α bound to receptors in transfected CHO cell membranes is shown in panel B (CCR1) and panel C (CCR5) as total (trace T), control membranes alone (trace C), and nonspecific (trace N) emission spectra. See Experimental Section for details.

as the parent MIP-1 α . Similar observations were recently published for rhodamine-MIP-1 α in which the fluorophore was attached specifically to the terminal amino group of the peptide chain, albeit by a different procedure.²³ Also, natural isoforms of MIP-1 α in the N-terminal region show subtle variations in pharmacological properties at CCR1 and CCR5 that are best accounted for by local perturbation of ligand–receptor contacts rather than a totally different binding mode.²⁶

Photoaffinity cross-linking of BP-MIP-1 α to CCR1 stably expressed in CHO cells by irradiation at 365 nm cleanly produced a complex of apparent molecular mass 54 kDa consistent with a 1:1 MIP-1 α –CCR1 complex. Likewise, photo-cross-linking of BP-MIP-1 α to CCR5 gave a 1:1 adduct of apparent molecular mass 48 kDa.

Figure 8. Collisional quenching of the receptor-bound Flu-MIP-1 α fluorescence emission by water-soluble KI. Stern-Volmer plots of F_0/F versus quencher concentration for quenching of the fluoresceinyl fluorescence by water-soluble KI. The Stern-Volmer quenching constants (K_{sv}) (see text) were determined from the equation:²⁴ $F_0/F = 1 + K_{sv}[Q]$, where F_0/F is the ratio of fluorescence intensities in the absence and presence of quencher at a given concentration $[Q]$: (A) CCR1 receptors; (B) CCR5 receptors. A representative plot of three independent experiments is shown; ● = free in solution; ○ = specific bound; □ = nonspecific bound.

For both CCR1 and CCR5, photoaffinity labeling was dependent upon irradiation time and concentration of BP-MIP-1 α .

The observation of covalent attachment of bound BP-MIP-1 α indicates that the carbonyl oxygen of the benzophenone is within a distance 3.1 Å of the receptor proteins.⁸ Thus, the N-terminal position of bound MIP-1 α is in close contact to the CCR1 or CCR5 proteins. In the future, using proteolytic digestion and amino acid or MS analysis of the ligand-receptor complexes we hope to determine contact sites at the amino acid level.

Collisional quenching of fluorescence is a powerful technique that has been used to probe solvent accessibility of receptor-bound ligands and to characterize binding pockets.^{15,16} The fluorescence of Flu-MIP-1 α bound to either CCR1 or CCR5 was efficiently quenched by addition of increasing amounts of iodide to the solution, indicating that the N-terminal end of MIP-1 α was fully accessible to the solvent in the ligand-receptor complex. We infer from these data that the N-terminal position of bound MIP-1 α interacts with the extracellular domains and loops or the parts of the transmembrane helices that are at the water-membrane interface. From our results, it is unlikely that the N-terminal of MIP-1 α penetrates into the hydrophobic pocket formed by the seven transmembrane helices of the

receptors. We cannot totally exclude that increasing concentrations of KI cause conformational changes resulting in the N-terminus of Flu-MIP-1 α being exposed to the aqueous environment. However, this is very unlikely as we have shown that KI, at the concentrations used, does not interfere with binding of [¹²⁵I]MIP-1 α at equilibrium. In the future, fluorescence energy-transfer experiments may help define more accurately the position of bound MIP-1 α .²⁵

In conclusion, we have developed photoaffinity and fluorescent analogues of the chemokine MIP-1 α . In the absence of high-resolution structural data for the complexes between chemokines, or peptides in general, and their membrane receptors, the combination of biochemical analysis such as photoaffinity labeling and spectroscopic methods such as fluorescence provides valuable insights into ligand-receptor recognition patterns. The reagents described here permit to apply these methods to the important class of chemokine receptors. In particular, we showed that the N-terminal end of MIP-1 α is in the hydrophilic environment of the extracellular parts of CCR1 and CCR5. In the future, the unnatural amino acid benzoylphenylalanine could be iteratively incorporated into chemokines at different positions during chemical synthesis to create new probes for mapping contact sites between MIP-1 α or other chemokines and their cognate receptors by photoaffinity scanning. Eventually, approaches to understand the molecular basis of chemokine interactions with their receptors should help the design of chemokine antagonists to control inflammatory processes.

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