

The common C-terminal sequences of substance P and neurokinin A contact the same region of the NK-1 receptor

Andrew A. Bremer, Susan E. Leeman, Norman D. Boyd*

Department of Pharmacology and Experimental Therapeutics, Boston University School of Medicine, L-611, 80 E. Concord Street, Boston, MA 02118, USA

Received 18 August 2000; revised 10 October 2000; accepted 20 October 2000

First published online 16 November 2000

Edited by Richard Cogdell

Abstract Although neurokinin A (NKA), a tachykinin peptide with sequence homology to substance P (SP), is a weak competitor of radiolabeled SP binding to the NK-1 receptor (NK-1R), more recent direct binding studies using radiolabeled NKA have demonstrated an unexpected high-affinity interaction with this receptor. To document the site of interaction between NKA and the NK-1R, we have used a photoreactive analogue of NKA containing *p*-benzoyl-L-phenylalanine (Bpa) substituted in position 7 of the peptide. Peptide mapping studies of the receptor photolabeled by ¹²⁵I-iodohistidyl¹-Bpa⁷NKA have established that the site of photoinsertion is located within a segment of the receptor extending from residues 178 to 190 (VVCMIIEW-PEHPNR). We have previously shown that ¹²⁵I-BH-Bpa⁸SP, a photoreactive analogue of SP, covalently attaches to M¹⁸¹ within this same receptor sequence. Importantly, both of these peptides (¹²⁵I-iodohistidyl¹-Bpa⁷NKA and ¹²⁵I-BH-Bpa⁸SP) have the photoreactive amino acid in an equivalent position within the conserved tachykinin carboxyl-terminal tail. In this report, we also show that site-directed mutagenesis of M¹⁸¹ to A¹⁸¹ in the NK-1R results in a complete loss of photolabeling of both peptides to this receptor site, indicating that the equivalent position of SP and NKA, when bound to the NK-1R, contact the same residue. © 2000 Federation of European Biochemical Societies. Published by Elsevier Science B.V. All rights reserved.

Key words: Substance P; Neurokinin A; NK-1 receptor; Photoaffinity labeling

1. Introduction

The tachykinins comprise a family of bioactive peptides which are structurally characterized by the conserved C-terminal sequence of -Phe-Xaa-Gly-Leu-Met-NH₂ [1–5]¹. Two

mammalian tachykinins are substance P (SP) (Arg-Pro-Lys-Pro-Gln-Gln-Phe-Phe-Gly-Leu-Met-NH₂) and neurokinin A (NKA) (His-Lys-Thr-Asp-Ser-Phe-Val-Gly-Leu-Met-NH₂) [4–6]. The mRNAs encoding both SP and NKA arise from tissue-specific differential RNA processing of the preprotachykinin-A gene [7–9]; post-translational modification by endoproteases and C-terminus amidation of protein precursors then yield the mature bioactive peptides.

The NK-1 receptor (NK-1R), the well-documented site of action for SP, is an integral membrane protein belonging to the rhodopsin-type family of G-protein coupled receptors [10–12]. Competition binding assays using radiolabeled SP and the NK-1R have shown that (i) SP binds with high affinity to the NK-1R and that (ii) NKA is a poor competitor for radiolabeled SP binding [13]. Since NKA displaces SP binding to the NK-1R only at high concentrations, the notion that NKA might functionally interact with the NK-1R was not immediately entertained. Recently, however, functional assays and direct binding studies using radiolabeled NKA have demonstrated that NKA can interact with the NK-1R with high affinity to elicit biological responses [14,15], despite its relative inability to compete for receptor binding with SP. The most straightforward explanation for this phenomenon is that the two peptides interact on the receptor at distinct sites. An alternative explanation, however, is the existence of multiple ligand-specific receptor conformations. We have addressed these issues directly by comparing the site of photoincorporation of a photoreactive analogue of NKA, ¹²⁵I-iodohistidyl¹-(*p*-benzoyl-L-phenylalanine)⁷NKA (¹²⁵I-Bpa⁷NKA), with that for the corresponding photoreactive analogue of SP, ¹²⁵I-(*N*-succinimidyl-3[4-hydroxyphenyl]propionate)³-Bpa⁸SP (¹²⁵I-BH-Bpa⁸SP).

This study presents the first direct biochemical evidence for an NKA/NK-1R interaction, and moreover shows that the structurally conserved C-terminal sequences of both SP and NKA interact with a common region of the NK-1R.

2. Materials and methods

2.1. Materials

Purified SP and NKA were purchased from Sigma. Bpa⁸SP was synthesized as described previously [16], and Bpa⁷NKA was obtained from Jeff Kelly (Scripps Institute, San Diego, CA, USA). ¹²⁵I-Bolton Hunter (BH) reagent and ¹²⁵I (each with specific activities of 2200 Ci/mmol) were obtained from NEN Life Sciences. SP and Bpa⁸SP were radioiodinated by coupling the Lys³ residue ε-NH₂ group to the ¹²⁵I-BH reagent as described previously [16], generating ¹²⁵I-BH-SP and ¹²⁵I-BH-Bpa⁸SP, respectively. NKA and Bpa⁷NKA were radioiodinated by coupling the His¹ residue to the solid phase oxidant 1,3,4,6-tetrachloro-3α,6α-diphenylglycoluril (ODO-GEN[®]) iodination

Abbreviations: Bpa, *p*-benzoyl-L-phenylalanine; BH, Bolton Hunter (*N*-succinimidyl-3[4-hydroxyphenyl]propionate); CHO, Chinese hamster ovary; DTT, D,L-dithiothreitol; E2, NK-1R second extracellular loop; ¹²⁵I-Bpa⁷NKA, ¹²⁵I-iodohistidyl¹-Bpa⁷NKA; ¹²⁵I-BH-Bpa⁸SP, ¹²⁵I-Bolton-Hunter³-Bpa⁸SP; MALDI, matrix-assisted laser desorption/ionization; NKA, neurokinin A; NK-1R, NK-1 receptor; rNK-1R, rat NK-1 receptor; SDS-PAGE, sodium dodecyl sulfate-polyacrylamide gel electrophoresis; SP, substance P

¹ Three letter amino acid abbreviations denote amino acids of the corresponding ligand.

Table 1
Average IC₅₀ values for rNK-1R ligands

Competing ligand	¹²⁵ I-BH-SP binding IC ₅₀ (nM ± S.E.M.)	¹²⁵ I-NKA binding IC ₅₀ (nM ± S.E.M.)
SP	0.9 ± 0.3	0.09 ± 0.02
Bpa ⁸ SP	1 ± 0.3	0.1 ± 0.03
NKA	46 ± 4	5 ± 2
Bpa ⁷ NKA	48 ± 4	6 ± 2

reagent, Pierce) in the presence of ¹²⁵I, generating ¹²⁵I-iodohistidyl¹-NKA (¹²⁵I-NKA) and ¹²⁵I-Bpa⁷NKA, respectively.

2.2. Cell culture and site-directed mutagenesis

Chinese hamster ovary (CHO) cells stably transfected with the cDNA encoding a geneticin resistance gene and (i) the full-length rat NK-1R (rNK-1R) [17], (ii) a site-directed rNK-1R mutant in which M¹⁷⁴ was substituted by alanine² or (iii) a site-directed rNK-1R mutant in which M¹⁸¹ was substituted by alanine were kindly provided by Dr. J.E. Krause (Neurogen, Brandford, CT, USA). All transfected CHO cells were maintained as monolayer cultures in α -minimal essential medium (Gibco BRL Life Technologies) supplemented with 10% (v/v) Cool Cal[®] 2 (Sigma) and 1 mg/ml geneticin (G-418 sulfate) (Gibco BRL Life Technologies), as described previously [18,19]. Cells were grown in an atmosphere of 95% air and 5% CO₂ at 37°C, and were harvested for experiments using phosphate-buffered saline based enzyme free dissociation buffer (Specialty Media).

2.3. Equilibrium displacement competition assay

Transfected CHO cells were harvested as described above and resuspended in ice-cold KRH buffer (20 mM HEPES, 1 mM CaCl₂, 2.2 mM MgCl₂, 5 mM KCl, 120 mM NaCl, pH 7.4) supplemented with 6 mg/ml glucose and 0.6 mg/ml bovine serum albumin (BSA). Cells were incubated for 2 h at 4°C with the radiolabeled ligands ¹²⁵I-BH-SP or ¹²⁵I-NKA, and binding was measured either alone or in the presence of increasing concentrations of unlabeled inhibitor. In all experiments, non-specific ¹²⁵I-BH-SP or ¹²⁵I-NKA binding was defined as binding in the presence of 1 μ M unlabeled peptide. To separate bound ligand from free ligand, cells were filtered after incubation through Whatman GF/C filter paper (soaked > 2 h in 0.1% polyethylenimine) and washed three times in ice-cold KRH buffer (pH 7.4) with a Brandel Harvester apparatus. Bound radioactivity on the filters was quantified with γ -emission spectrometry. Competition assays were performed in triplicate and repeated at least three times.

2.4. Photoaffinity labeling of transfected CHO cells and identification of photolabeled receptors

Stably transfected CHO cells were photolabeled with ¹²⁵I-BH-Bpa⁸SP or ¹²⁵I-Bpa⁷NKA using the procedure described previously [18,19]. Transfected cells were harvested, pelleted and resuspended in ice-cold KRH buffer (pH 7.4) supplemented with 6 mg/ml glucose, 0.6 mg/ml BSA, 3 μ g/ml chymostatin, 5 μ g/ml leupeptin and 30 μ g/ml bacitracin. The cell resuspensions were then incubated in the dark with ¹²⁵I-BH-Bpa⁸SP (added to a final concentration of 0.5–1.0 nM) or ¹²⁵I-Bpa⁷NKA (added to a final concentration of 3.0–3.5 nM) for 2 h at 4°C with gentle agitation. Competing peptides or non-peptide antagonists were added at the concentrations indicated. Following incubation, the mixtures were diluted 1:1 with ice-cold KRH buffer (pH 7.4) and irradiated at 365 nm by exposure to a 100 W long-wave UV lamp for 15 min at a distance of 6 cm. Cell membranes were then prepared as described previously [18,19]. ¹²⁵I-BH-Bpa⁸SP- or ¹²⁵I-Bpa⁷NKA-labeled membranes were next solubilized in sample buffer (0.125 M Tris, 2% sodium dodecyl sulfate (SDS), 10% glycerol, 0.01% bromophenol blue, pH 6.8), heated at 55°C for 10 min, and then subjected to SDS–polyacrylamide gel electrophoresis (PAGE) as described by Laemmli [20].

The percent photoincorporation of ¹²⁵I-BH-Bpa⁸SP or ¹²⁵I-Bpa⁷NKA into the wildtype (WT), M174A and M181A rNK-1Rs

was determined as described previously [16] by comparing the amount of photoligand incorporated into the receptors after UV exposure (as determined by SDS–PAGE resolution, autoradiography and γ -emission spectrometric analysis) to the amount of photoligand specifically bound to the receptors before UV exposure (as determined by equilibrium filtration binding assay).

2.5. Tryptic digestion of photolabeled receptors

¹²⁵I-BH-Bpa⁸SP- or ¹²⁵I-Bpa⁷NKA-labeled membranes were resuspended in 0.1% SDS, 50 mM Tris, 1 mM CaCl₂, pH 8.0, and digested at room temperature for 2 h with 0.02 mg/ml L-1-tosylamide-2-phenylethylchloromethyl ketone-treated bovine trypsin (Sigma). *N* α -p-tosyl-L-lysine chloromethyl ketone was added to the reaction mixtures in a 1:1000 dilution following incubation to terminate enzymatic activity. Sample buffer (\pm 30 mM D,L-dithiothreitol, DTT) was then added to the samples. The mixtures (\pm 30 mM DTT) were allowed to incubate for 1 h at room temperature before being heated for 10 min at 55°C, as described above. Tryptic cleavage fragments were separated and analyzed using the tricine-gel system of SDS–PAGE [21].

3. Results

3.1. Affinity of photoreactive tachykinin analogues for the NK-1R as measured by homologous and heterologous competition binding assays

To gain information on the bimolecular complexes formed between SP and the NK-1R as well as NKA and the NK-1R, a photoreactive analogue of each peptide was synthesized (Bpa⁸SP and Bpa⁷NKA, respectively) (Fig. 1). Importantly, the introduction of the photoreactive Bpa residue into either the eighth position of SP (normally a Phe) or the seventh position of NKA (normally a Val) does not significantly alter the receptor binding characteristics of these tachykinin analogues, making them useful for direct receptor photolabeling experiments. Both Bpa⁸SP and Bpa⁷NKA also elicit a calcium response from rNK-1R-expressing CHO cells (data not shown), consistent with their being functional agonists at this receptor. Therefore, since Bpa⁸SP and Bpa⁷NKA bind the NK-1R in a manner indistinguishable from that of SP and NKA, respectively, the introduction of the Bpa residue into the photoreactive tachykinin analogues does not significantly alter the functional properties of the ligands.

3.2. ¹²⁵I-Bpa⁷NKA specifically photolabels the NK-1R

Our laboratory has successfully used Bpa⁸SP as an efficient photoreactive ligand for the rNK-1R [18,22]; moreover, a radiolabeled derivative of Bpa⁸SP, ¹²⁵I-BH-Bpa⁸SP, has been used to identify M¹⁸¹ as a site of covalent attachment of SP to the rNK-1R [19]. In this report, we show that the

Bpa⁸SP: Arg - Pro - Lys - Pro - Gln - Gln - Phe - **Bpa⁸** - Gly - Leu - Met - NH₂

Bpa⁷NKA: His - Lys - Thr - Asp - Ser - Phe - **Bpa⁷** - Gly - Leu - Met - NH₂

Fig. 1. Amino acid sequences of Bpa⁸SP and Bpa⁷NKA. Note that the Bpa residue of both Bpa⁸SP and Bpa⁷NKA is located in an equivalent position within the conserved tachykinin C-terminal tail.

² Single letter amino acid abbreviations denote amino acids of the NK-1 receptor.

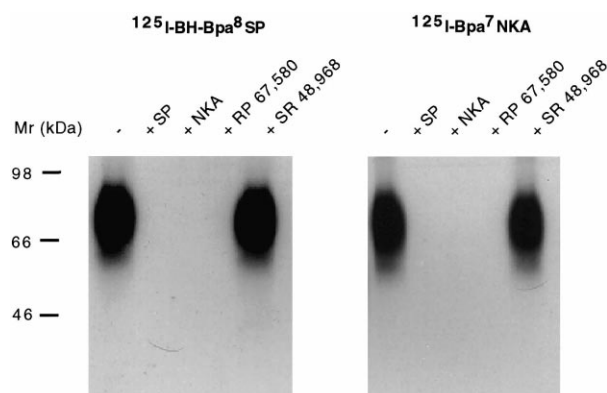


Fig. 2. Specific photolabeling of the rNK-1R with ^{125}I -BH-Bpa 8 SP and ^{125}I -Bpa 7 NKA. Both ^{125}I -BH-Bpa 8 SP and ^{125}I -Bpa 7 NKA specifically photolabel the same protein which migrates diffusely at $M_r \sim 80$ kDa on SDS-PAGE due to the heterogeneity of receptor glycosylation [37].

tachykinin derivative Bpa 7 NKA is also a high-affinity ligand for the rNK-1R (Table 1), and that its radiolabeled derivative, ^{125}I -Bpa 7 NKA, specifically photolabels the rNK-1R as well (Fig. 2). Pharmacological evidence of the specificity of the ^{125}I -BH-Bpa 8 SP-rNK-1 and ^{125}I -Bpa 7 NKA-rNK-1R interaction is provided by results showing that the addition of 1 μM SP, NKA or RP 67 580 (a rNK-1R non-peptide antagonist [23]) prevents rNK-1R labeling by both ^{125}I -BH-Bpa 8 SP and ^{125}I -Bpa 7 NKA, whereas the addition of 1 μM of SR 48 968 (a rNK-2R non-peptide antagonist [24]) has no effect on receptor photolabeling.

3.3. ^{125}I -Bpa 7 NKA photolabels the same residue on the NK-1R as ^{125}I -BH-Bpa 8 SP

Since the seventh amino acid of NKA is in the equivalent position as the eighth amino acid of SP (Fig. 1), the use of Bpa 7 NKA as a photoligand is strategic in that it allows direct comparison of ^{125}I -Bpa 7 NKA-rNK-1R peptide mapping data

with the published results of extensive peptide mapping studies performed on the ^{125}I -BH-Bpa 8 SP-labeled rNK-1R [18,19,22]. In the absence of DTT, trypsinization of the ^{125}I -BH-Bpa 8 SP-labeled rNK-1R results in the generation of two receptor fragments linked by a disulfide bond between C 105 and C 180 , with a combined $M_r \sim 8$ kDa. Reduction with DTT yields a smaller fragment ($M_r \sim 3$ kDa) that contains residues 178–190 of the NK-1R second extracellular (E2) loop. Large-scale isolation of this particular ~ 3 kDa rNK-1R fragment and characterization by matrix-assisted laser desorption/ionization (MALDI) mass spectrometry has identified the residue within this sequence that serves as the site of covalent attachment of ^{125}I -BH-Bpa 8 SP to the rNK-1R as M 181 [19].

Peptide mapping studies performed on the ^{125}I -Bpa 7 NKA-labeled rNK-1R show a remarkable similarity to those discussed above for the ^{125}I -BH-Bpa 8 SP-labeled rNK-1R, i.e. the generation of a fragment at $M_r \sim 8$ kDa following trypsinization, which is reduced in size to $M_r \sim 3$ kDa following reduction with DTT (Fig. 3). These data suggest that the site of covalent attachment of ^{125}I -Bpa 7 NKA to the rNK-1R is also to a residue in the $M_r \sim 3$ kDa fragment of the E2 loop extending from residues 178–190.

To test whether ^{125}I -Bpa 7 NKA specifically photolabels rNK-1R residue M 181 , photolabeling experiments using mutant rNK-1Rs were performed (Fig. 4). Two specific mutant rNK-1Rs, in which an alanine residue is substituted for either M 174 (M174A) or M 181 (M181A), have been characterized (Table 2) and are both functional. These two mutant receptors were constructed on the basis of M 181 serving as the site of photoinsertion of ^{125}I -BH-Bpa 8 SP to the rNK-1R [19], and M 174 serving as the site of photoinsertion of another photo-reactive SP analogue designed in our laboratory, ^{125}I -BH-Bpa 4 SP, to the rNK-1R [25]. In this report, we show that when alanine is substituted for M 181 , although ^{125}I -BH-Bpa 8 SP binding is not significantly affected (IC_{50} values of 0.77 ± 0.12 nM for WT versus 0.82 ± 0.16 nM for M181A), the efficiency of photolabeling is significantly reduced (relative

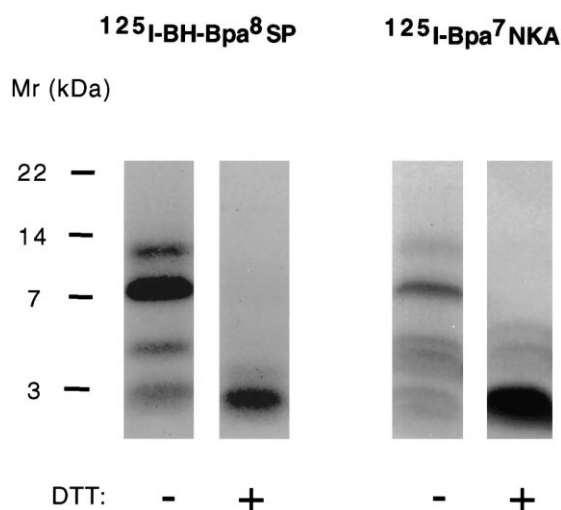


Fig. 3. Tryptic digestion pattern of ^{125}I -BH-Bpa 8 SP- and ^{125}I -Bpa 7 NKA-photolabeled rNK-1Rs. Tryptic digestion (0.02 mg/ml) of ^{125}I -BH-Bpa 8 SP-labeled rNK-1R generates a major fragment at $M_r \sim 8$ kDa in the absence of 30 mM DTT (–), which is reduced in size to $M_r \sim 3$ kDa in the presence of 30 mM DTT (+). An identical tryptic digest pattern \pm DTT is produced with ^{125}I -Bpa 7 NKA-labeled rNK-1R.

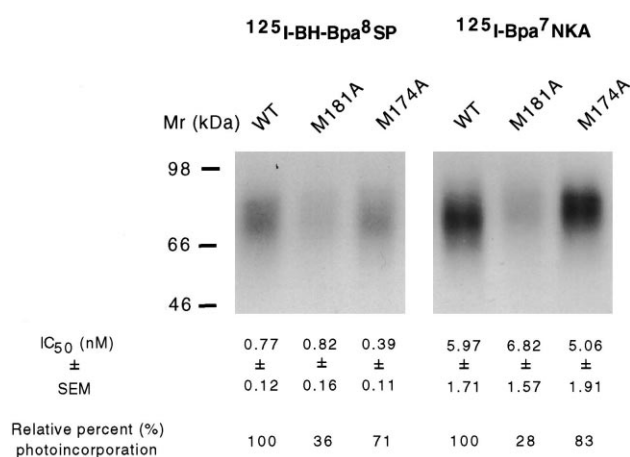


Fig. 4. Photoincorporation of ^{125}I -BH-Bpa 8 SP and ^{125}I -Bpa 7 NKA into WT and mutant rNK-1Rs. The photoincorporation of ^{125}I -BH-Bpa 8 SP into the WT rNK-1R and M174A mutant rNK-1R is comparable; however, its photoincorporation into the M181A mutant rNK-1R is markedly reduced. A similar pattern of receptor photoincorporation efficiencies is reported when ^{125}I -Bpa 7 NKA is the photoligand. Importantly, the M181A and M174A mutant rNK-1Rs display only slight differences in their binding affinity for the photoligands as compared to WT rNK-1R; IC_{50} values are shown.

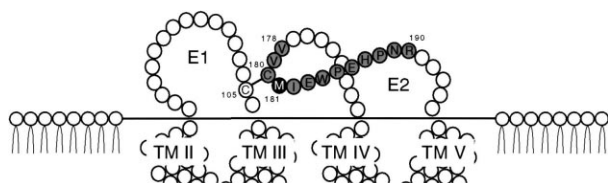


Fig. 5. Schematic diagram of the first (E1) and second (E2) extracellular loops of the rNK-1R. The residues shown in gray correspond to the $M_r \sim 3$ kDa trypsin-generated rNK-1R fragment (residues 178–190). A disulfide bond exists between C¹⁰⁵ and C¹⁸⁰, linking E1 and E2 near the membrane surface. The E2 loop residue Met¹⁸¹, the direct site of ¹²⁵I-BH-Bpa⁸SP and ¹²⁵I-Bpa⁷NKA photoinsertion to the rNK-1R, is shown in black. TM = transmembrane domain.

percent photoincorporation of 100% for WT versus 36% for M181A). However, when alanine is substituted for M¹⁷⁴, which is not the precise site of photoincorporation of ¹²⁵I-BH-Bpa⁸SP, there is only a modest effect on binding affinity (IC_{50} values of 0.77 ± 0.12 nM for WT versus 0.39 ± 0.11 nM for M174A) and only a small decrease in photolabeling efficiency (relative percent photoincorporation of 100% for WT versus 71% for M174A).

These two mutant receptors were also photolabeled with ¹²⁵I-Bpa⁷NKA (Fig. 4), and the binding and photolabeling results compared with those obtained with ¹²⁵I-BH-Bpa⁸SP. In parallel with the results seen with ¹²⁵I-BH-Bpa⁸SP, (i) a marked decrease in the efficiency of photoincorporation of ¹²⁵I-Bpa⁷NKA into the M181A mutant was observed (relative percent photoincorporation of 100% for WT versus 28% for M181A) and (ii) a comparable efficiency of photoincorporation of ¹²⁵I-Bpa⁷NKA into the M174A mutant was observed (relative percent photoincorporation of 100% for WT versus 83% for M174A). Again, paralleling the results seen with ¹²⁵I-BH-Bpa⁸SP, the substitution of alanine for either M¹⁸¹ or M¹⁷⁴ had no marked effect on ¹²⁵I-Bpa⁷NKA binding affinity (Fig. 4). Taken together, these results provide strong evidence that the site of ¹²⁵I-Bpa⁷NKA covalent attachment to the rNK-1R is the same as that of ¹²⁵I-BH-Bpa⁸SP (i.e. E2 loop residue M¹⁸¹) (Fig. 5). Interestingly, the mutant receptors in which alanine was substituted for M¹⁷⁴ and M¹⁸¹ display only slight differences in their binding affinity for Bpa⁸SP and Bpa⁷NKA, indicating that neither M¹⁷⁴ nor M¹⁸¹ per se contribute significantly to the total binding energy despite their location at the ligand–receptor interface. These findings suggest that several ligand–receptor contacts exist in the high-affinity binding complex, and are consistent with the view that multiple peptide–receptor interactions participate in peptide recognition and receptor activation.

4. Discussion

In this study, we (i) confirm published reports establishing that NKA binds to the NK-1R with high affinity [14,15] and

(ii) present the first data directly identifying a site of interaction between NKA and the NK-1R at the bimolecular interface. First, we demonstrate that ¹²⁵I-Bpa⁷NKA is an efficient and specific photoligand for the rNK-1R. Second, we restrict the site of photoinsertion of ¹²⁵I-Bpa⁷NKA to an amino acid of the rNK-1R between residues 178 and 190 on the E2 loop. As previously reported, this same restricted segment of the rNK-1R contains M¹⁸¹, the site of ¹²⁵I-BH-Bpa⁸SP photoinsertion as documented by MALDI mass spectrometry [19]. (iii) We show that a comparable loss of photoinsertion efficiency results when either ¹²⁵I-BH-Bpa⁸SP or ¹²⁵I-Bpa⁷NKA is used to photolabel a mutant rNK-1R in which the side-chain of M¹⁸¹ has been removed. Taken together, these results indicate that in the high-affinity bound state, the equivalent positions within the common C-terminal sequences of SP and NKA are in close proximity to M¹⁸¹ on the E2 loop of the rNK-1R.

In our laboratory, we have demonstrated that when the side-chain of an identified contact residue of a photoligand on a receptor is removed, a marked loss of photoincorporation results in the absence of a significant affect on peptide binding [26]. Pertinent to this investigation, we show that when M¹⁸¹ of the rNK-1R is mutated to an alanine, the photolabeling efficiency of ¹²⁵I-BH-Bpa⁸SP is dramatically reduced. Since we had restricted the site of ¹²⁵I-Bpa⁷NKA photoinsertion to a region of the rNK-1R containing the site of ¹²⁵I-BH-Bpa⁸SP covalent attachment (i.e. M¹⁸¹), we found it plausible to speculate that both photoligands might have the same contact residue, particularly since the photoreactive amino acid of each ligand is in an equivalent position within the conserved tachykinin C-terminal tail. Paralleling the results obtained when ¹²⁵I-BH-Bpa⁸SP is used to photolabel the M181A mutant receptor, we in fact demonstrate a significant loss of photoincorporation of ¹²⁵I-Bpa⁷NKA into the M181A mutant receptor, validating its contact site to the rNK-1R as M¹⁸¹. Thus, the strategy of using specifically mutated rNK-1Rs and photoaffinity labeling experiments to characterize the site of ¹²⁵I-Bpa⁷NKA covalent attachment to the rNK-1R was both novel and successful, and eliminated the necessity of purifying the site of peptide–receptor interaction for analysis by mass spectrometry.

It is important to note that the observation of two structurally related peptides displaying differences in their binding affinity for a given receptor target depending on the assay used (i.e. homologous versus heterologous competition assays) is not limited to SP and NKA interacting with the NK-1R, but rather is emerging as a more general phenomenon for other structurally related peptide families [27]. For example, although vasoactive intestinal peptide (VIP) competes for the binding of the structurally similar peptide pituitary adenylate cyclase activating polypeptide (PACAP) to a receptor target that has been designated the PACAP receptor (PACAP-R) with low affinity, in direct binding experiments radiolabeled VIP is found to bind the PACAP-R with high affinity [28].

Table 2
Binding affinities of SP and NKA to WT, M174A and M181A rNK-1Rs

Receptor	¹²⁵ I-BH-SP binding versus unlabeled SP IC_{50} (nM \pm S.E.M.)	¹²⁵ I-NKA binding versus unlabeled NKA IC_{50} (nM \pm S.E.M.)
WT	0.9 ± 0.3	5.0 ± 2.0
M174A	1.4 ± 0.4	5.3 ± 2.1
M181A	1.5 ± 0.6	6.1 ± 2.4

The most straightforward explanation for the observation that two ligands can interact at the same receptor with high affinity but not readily compete is that they each interact with the receptor at distinct sites [27]. To address the issue as to whether SP and NKA interact with the NK-1R at the same or different sites, Wijkhuysen et al. recently followed a molecular recognition theory to identify regions of the NK-1R whose hydropathic profiles were opposed to that of the common tachykinin C-terminal sequence, and then performed binding experiments on receptors mutated in those particular regions [29]. They subsequently reported the identification of an NK-1R domain located at the distal end of the E2 loop that interacts with the C-terminus of NKA but not of SP, suggesting the existence of distinct binding sites on the receptor for the C-terminal sequences of these two peptides. In the current study, however, we have taken a direct approach to study the tachykinin/NK-1R complex by comparing the site of ^{125}I -Bpa⁷NKA photoinsertion to the rNK-1R with that of the well-studied corresponding photoreactive analogue of SP, ^{125}I -BH-Bpa⁸SP. By reporting that both ^{125}I -Bpa⁷NKA and ^{125}I -BH-Bpa⁸SP photolabel the same residue on the rNK-1R, we provide evidence against the existence of a distinct, NKA-preferring binding site, and conclude that the equivalent positions of SP and NKA within the conserved tachykinin C-terminal tail are in close proximity to M¹⁸¹ in the peptide/rNK-1R complex.

An alternative and somewhat more complex possibility to account for the apparent discrepancy in homologous versus heterologous competition assay results is that the NK-1R exists in multiple peptide-selective conformations that not only display differences in their ligand specificities [27], but also potentially in their G-protein and effector molecule interactions and evoked physiological responses [30]. We favor this multiple conformation theory. Particularly since SP and NKA arise from the same gene and are often co-expressed and co-released in regions of the central nervous system and periphery without detectable NK-2 receptors [31–33], that both peptides may stimulate peptide-specific effector responses through functional interaction with the NK-1R may have important physiological consequences.

Using the direct and efficient methods of photochemistry, combined with specific receptor residue site-directed mutagenesis, we have demonstrated that the common C-terminal sequences of SP and NKA contact the same region of the NK-1R. We furthermore predict that the divergent N-terminal regions of these two peptides will interact with different regions of the NK-1R. Extensive biochemical analysis of the high-affinity SP/rNK-1R complex has already been performed [18,19,22,34–37]. These studies directly document the importance of both the extracellular N-terminus and the E2 loop of the rNK-1R for peptide recognition, and support the viewpoint that several ligand–receptor interactions participate in binding and activation [38–40]. Just as multiple contacts exist between SP and the NK-1R in the bound state, so too will multiple interactions undoubtedly occur between NKA and the NK-1R. By directly evaluating and comparing the tachykinin/rNK-1R bimolecular interface using multiple photoreactive SP and NKA derivatives, we can use a photoaffinity-scanning approach to further investigate the existence of these non-distinct peptide-preferring receptor conformations, and determine which residue–residue contacts participate in the

molecular processes of peptide recognition and receptor activation.

Acknowledgements: This work is sponsored by the Public Health Service grant from the National Institutes of Health: NS 3134.

References

- [1] Erspamer, V. (1971) *Annu. Rev. Pharmacol.* 11, 327–350.
- [2] Erspamer, V. (1981) *Trends Neurosci.* 4, 267–269.
- [3] Pernow, B. (1983) *Pharmacol. Rev.* 35, 85–141.
- [4] Nakanishi, S. (1987) *Physiol. Rev.* 67, 1117–1142.
- [5] Maggio, J.E. (1988) *Annu. Rev. Neurosci.* 11, 13–28.
- [6] Nakanishi, S. (1986) *Trends Neurosci.* 9, 41–44.
- [7] Nawa, H., Kotani, H. and Nakanishi, S. (1984) *Nature* 312, 729–734.
- [8] Kawaguchi, Y., Hoshimaru, M., Nawa, H. and Nakanishi, S. (1986) *Biochem. Biophys. Res. Commun.* 139, 1040–1046.
- [9] Krause, J.E., Chirgwin, J.M., Carter, M.S., Xu, Z.S. and Hershey, A.D. (1987) *Proc. Natl. Acad. Sci. USA* 84, 881–885.
- [10] Yokota, Y., Sasai, Y., Tanaka, K., Fujiwara, T., Tsuchida, K., Shigemoto, R., Kakizuka, A., Ohkubo, H. and Nakanishi, S. (1989) *J. Biol. Chem.* 264, 17649–17652.
- [11] Hershey, A.D. and Krause, J.E. (1990) *Science* 247, 958–961.
- [12] Takeda, Y., Chou, K.B., Takeda, J., Sachais, B.S. and Krause, J.E. (1991) *Biochem. Biophys. Res. Commun.* 179, 1232–1240.
- [13] Ingi, T., Kitajima, Y., Minamitake, Y. and Nakanishi, S. (1991) *J. Pharmacol. Exp. Ther.* 259, 968–975.
- [14] Hastrup, H. and Schwartz, T.W. (1996) *FEBS Lett.* 399, 264–266.
- [15] Sagan, S., Beaujouan, J.-C., Torrens, Y., Saffroy, M., Chassaing, G., Glowinski, J. and Lavielle, S. (1997) *Mol. Pharmacol.* 52, 120–127.
- [16] Boyd, N.D., White, C.F., Cerpa, R., Kaiser, E.T. and Leeman, S.E. (1991) *Biochemistry* 30, 336–342.
- [17] Takeda, Y., Blount, P., Sachais, B.S., Hershey, A.D., Raddatz, R. and Krause, J.E. (1992) *J. Neurochem.* 59, 740–745.
- [18] Boyd, N.D., Kage, R., Dumas, J.J., Krause, J.E. and Leeman, S.E. (1996) *Proc. Natl. Acad. Sci. USA* 93, 433–437.
- [19] Kage, R., Leeman, S.E., Krause, J.E., Costello, C.E. and Boyd, N.D. (1996) *J. Biol. Chem.* 271 (42), 25797–25800.
- [20] Laemmli, U.K. (1970) *Nature (Lond.)* 227, 680–685.
- [21] Schagger, H. and Jagow, G.V. (1987) *Anal. Biochem.* 166, 368–379.
- [22] Boyd, N.D. and Leeman, S.E. (1995) *Ann. N. Y. Acad. Sci.* 757, 405–409.
- [23] Garret, C., Carruette, A., Fardin, V., Moussaoui, S., Peyronel, J.-F., Blanchard, J.-C. and Laduron, P.M. (1991) *Proc. Natl. Acad. Sci. USA* 88, 10208–10212.
- [24] Emonds-Alt, X., Vilain, P., Goulauic, P., Proietto, V., Broeck, D.V., Advenier, C., Naline, E., Neliat, G., Fur, G.L. and Breliere, J.C. (1992) *Life Sci.* 50, 101–106.
- [25] Li, H. (1998) Ph.D. thesis, Boston University School of Medicine.
- [26] Macdonald, D.M. (1998) Ph.D. thesis, Boston University School of Medicine.
- [27] Maggi, C.A. and Schwartz, T.W. (1997) *Trends Pharmacol. Sci.* 18, 351–355.
- [28] Hashimoto, H., Ogawa, N., Hagihara, N., Yamamoto, K., Im-anishi, K., Nogi, H., Nishino, A., Fujita, T., Matsuda, T., Nagata, S. and Baba, A. (1997) *Mol. Pharmacol.* 52, 128–135.
- [29] Wijkhuysen, A., Sagot, M.-A., Frobert, Y., Creminon, C., Grassi, J., Boquet, D. and Couraud, J.-Y. (1999) *FEBS Lett.* 447, 155–159.
- [30] Sagan, S., Chassaing, G., Pradier, L. and Lavielle, S. (1996) *J. Pharmacol. Exp. Ther.* 276, 1039–1048.
- [31] Sasai, Y. and Nakanishi, S. (1989) *Biochem. Biophys. Res. Commun.* 165, 695–702.
- [32] Tsuchida, K., Shigemoto, R., Yokota, Y. and Nakanishi, S. (1990) *Eur. J. Biochem.* 193, 751–757.
- [33] Otsuka, M. and Yoshioka, K. (1993) *Physiol. Rev.* 73, 229–308.
- [34] Boyd, N.D., Macdonald, S.G., Kage, R., Lubner-Narod, J. and Leeman, S.E. (1991) *Ann. N. Y. Acad. Sci.* 632, 79–93.

- [35] Kage, R., Leeman, S.E. and Boyd, N.D. (1993) *J. Neurochem.* 60, 347–351.
- [36] Boyd, N.D., Kage, R.K. and Leeman, S.E. (1994) in: *The Tachykinin Receptors* (Buck, S.H., Ed.), pp. 219–236, Humana Press, NJ.
- [37] Kage, R., Hershey, A.D., Krause, J.E., Boyd, N.D. and Leeman, S.E. (1995) *J. Neurochem.* 64, 316–321.
- [38] Fong, T.M., Huang, R.-R.C. and Strader, C.D. (1992) *J. Biol. Chem.* 267, 25664–25667.
- [39] Yokota, Y., Akazawa, C., Ohkubo, H. and Nakanishi, S. (1992) *EMBO J.* 11, 3585–3591.
- [40] Gether, U., Johansen, T.E. and Schwartz, T.W. (1993) *J. Biol. Chem.* 268, 7893–7898.