

Use of receptor chimeras to identify small molecules with high affinity for the dynorphin A binding domain of the κ opioid receptor

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Abstract—A series of 2-substituted sulfamoyl arylacetamides of general structure **2** were prepared as potent κ opioid receptor agonists and the affinities of these compounds for opioid and chimeric receptors were compared with those of dynorphin A. Compounds **2e** and **2i** were identified as non-peptide small molecules that bound to chimeras 3 and 4 with high affinities similar to dynorphin A, resulting in K_i values of 1.5 and 1.2 nM and 1.3 and 2.2 nM, respectively.

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The identification of the mu (μ), delta (δ), and kappa (κ) subtypes of the opioid receptor led to the suggestions that agonists selective for receptor subtypes might be effective analgesics with fewer serious side effects.¹ Even though the arylacetamide series of κ opioid receptor agonists lack μ opioid receptor-mediated side effects, the utility of these agonists as antinociceptive agents is limited due to side effects such as dysphoria, diuresis, and psychotomimesis.^{2–4} In clinical trials, the naturally occurring peptide κ opioid receptor agonist, dynorphin A, mediates analgesia without dysphoria, diuresis, and psychosis, indicating that the antinociceptive effects of κ opioid receptor agonists could be dissociated from their side effects.^{5,6} A metabolically stable analog of dynorphin A, E2078, is an effective analgesic in post-surgical patients at doses that produce no side effects.⁷ This exemplifies that there are opportunities for identifying metabolically stable small peptides or small molecule κ opioid receptor agonists as effective analgesics that lack the side effect profile of the arylacetamides.

This distinction in the side effect profiles of arylacetamides and dynorphin A could in part be related to the different binding regions for the κ opioid receptor^{8,9} which were observed through the use of chimeric receptors composed of sequences derived from κ and μ opioid

receptors. These different modes of binding have led to a hypothesis that different domain selectivity of agonists that bind to the κ receptor might be related to different patterns of side effects.⁹ Therefore, in an effort to discover small molecule κ opioid receptor agonists that have a therapeutic profile similar to that of dynorphin A and related compounds, we have recently described¹⁰ the design and construction of two μ/κ chimeric receptors composed primarily of amino acid residues derived from the μ opioid receptor for the screening of compounds.

The chimeric receptors used in this study include one of the chimeric receptors used in the earlier study (designated chimera 3)¹⁰ and another chimeric receptor (designated chimera 4). These receptors are depicted in Figure 1 in which filled circles represent amino acids derived from the κ opioid receptor and open circles represent amino acids derived from the μ opioid receptor. For chimera 3, the 25 amino acids of the putative second extracellular loop of the μ opioid receptor were replaced with the 28 amino acids (8 identical) of the putative second extracellular loop of the κ opioid receptor. The chimera 4 construct was made using a synthetic oligonucleotide corresponding to the Bcl1–Styl region (343 bp) of the human κ opioid receptor in which amino acid numbers 86–178 were replaced with the corresponding amino acids of the human μ opioid receptor. This construct is a human κ opioid receptor where the first and second intracellular loops, the first extracellular loop, and the second and third transmembrane regions were replaced

Keywords: κ Opioid receptor agonists; Dynorphin A; Chimeric receptors.

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