

Design and synthesis of melatonin receptors agonists and antagonists

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Abstract

We review our work towards the design and synthesis of high-affinity melatonin (*N*-acetyl-5-methoxytryptamine) agonist and antagonist compounds. High affinity melatonergic agonists were obtained by shifting the melatonin side chain from C₃ to N₁ of the indole ring system. Conversely, by moving the side chain from C₃ to C₂ it was possible to obtain melatonin antagonist compounds, albeit of moderate affinity. © 2000 Elsevier Science S.A. All rights reserved.

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1. Introduction

Melatonin (*N*-acetyl-5-methoxytryptamine, MLT (**1**) Fig. 1) is the principal hormone of the vertebrate pineal gland by which it is secreted during darkness [1]. Much has been claimed for its physiological role and therapeutic effects [2–10], but the situation has yet to be clarified [10].

Two distinct types of MLT receptors have been identified: one of high affinity (ML₁) and one of low affinity (ML₂) [11]. ML₁ (or Mel₁) receptors have been identified in and cloned from several species including humans [12–17] and are further classified into three subtypes Mel_{1A}, Mel_{1B} and Mel_{1C}; of these three receptor subtypes (G-protein coupled receptors) only Mel_{1A} and Mel_{1B} are present in mammals. Mel_{1A} and Mel_{1B} are now designated as mt₁ and MT₂ [18].

2. Discussion of the project and results

Our research focused initially on the design of several indole melatonin agonists to contribute to the knowledge of the mode of interaction of MLT with its

receptors [19–23]. A number of structure–affinity relationships were identified, and recently we and other authors proposed molecular models of the putative melatonin binding site [21,22,24–28]. These groups produced several high-affinity non-selective melatonin agonists and it is now possible to reliably design this type of compounds. The situation is quite different in the case of the antagonists, as very few high-affinity competitive antagonists have been reported to date [12,29–36] and the lack of selective agonists and antagonists endowed with high affinity slows progress in the field. The structure of MLT is reported in Fig. 1.

In agreement with and in addition to previous findings the following results were obtained:

- Insertion of a suitable substituent (Br, Ph, COOR) at C₂ enhances affinity via interaction with complementary sites of the receptor [19,20,37].
- The substitution at C₄ with Br or at C₇ with OMe decreased the affinity.
- The size of the acyl group is important for the binding of the side chain to the receptor and in some cases (cycloalkyl groups) it decreases the intrinsic activity of the corresponding compounds [19–21,23,37,38].
- A 3D-QSAR model describing quantitative–affinity relationships within different classes of melatonergic ligands was derived by applying the CoMFA methodology; this model offers the possibility of

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predicting affinities of designed compounds for the target MLT receptors [21].

- By using conformationally restricted MLT congeners (Fig. 1) it is seen that one of the bioactive conformations of the 3-ethylamido side chain of MLT is in a folded *gauche/anti* oriented form and is almost orthogonal to the plane of the indole [22].
- The 5-OMe group can be replaced by suitable substituents (i.e. 5-Br or 5-Cl) [23].
- The translocation of the ethylamido side chain from C₃ to N₁ resulted in high affinity MLT agonists that obey the same SARs ascertained for the natural ligand [37].
- The translocation of the melatonin side chain from C₃ to C₂ yields melatonin antagonists; the length of the amidoethyl side chain can be shortened or lengthened by one C unit [38].
- Relative to the 2-alkylamido compounds, suitable combinations of the OMe group (C₄- or C₆-) of the acyl residue and of the N-1 indole substituent enhance the binding affinity of the ligands for the melatonin receptors and, based on preliminary data, the MT₂ selectivity [37].
- The effects on binding affinity and efficacy deriving from variation of the acyl group are similar to those observed in the natural series of melatonergic agonists [37,38].
- In the case of the 2-alkylamido compounds, substitution with 3-Br does not lead to an increase of affinity [38].

2.1. Notes on the *in vitro* testing methodologies

The evaluation of our compounds was performed according to the following models used in the following sequences.

2.1.1. Affinity

The affinity of several compounds prepared in this study was determined using 2-[¹²⁵I]iodomelatonin (100 pM) as a radioligand in competition binding analyses on cloned human mt₁ receptor subtype stably expressed in NIH3T3 mouse fibroblast cells [39]. The affinity of some of these compounds was evaluated prior to the cloning of human melatonin receptors, and it was done on quail optic tecta membranes [40].

2.1.2. Intrinsic activity

In vitro functional data were obtained by using the following assays: (a) effects on forskolin-stimulated cAMP accumulation; (b) effects of co-incubation with GTPγS on the IC₅₀ values (GTPγS index) [37]; and (c) [³⁵S]GTPγS radioligand binding assay in N11-13M mouse fibroblast cells expressing human mt₁ melatonin receptor [38,39].

The relative intrinsic activity values were obtained by dividing the maximal G-protein activation of a test compound by that of MLT. Thus, the intrinsic activity of MLT or of full agonists is always equal or very close to 1, between 0 and 1 for partial agonists, and below 0 for inverse agonists.

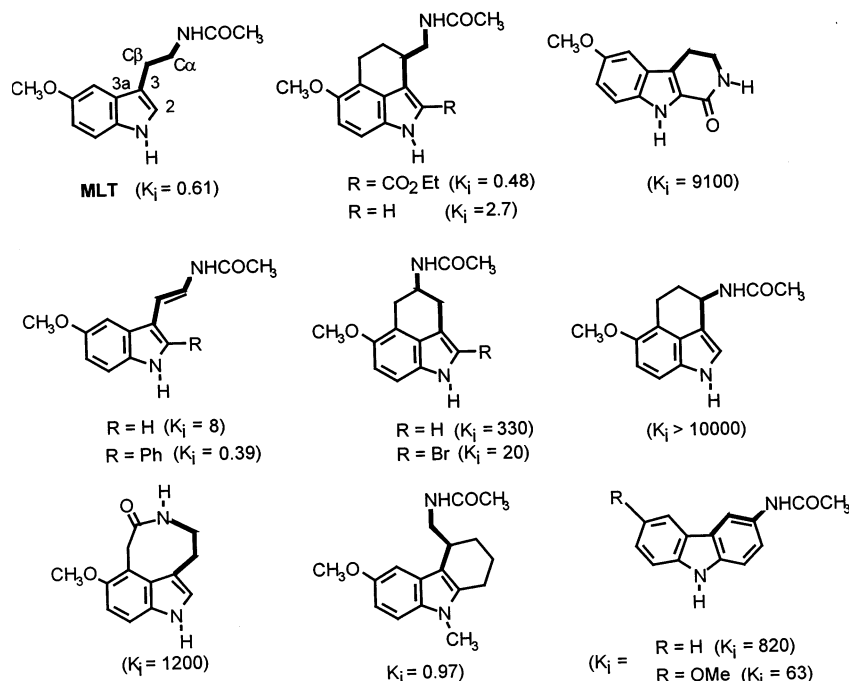


Fig. 1. Melatonin and conformationally restricted melatonin analogs (K_i = nM).

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