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Bioorganic & Medicinal Chemistry Letters

journal homepage: www.elsevier.com/locate/bmcl



Synthesis of novel 7-imino-2-thioxo-3,7-dihydro-2*H*-thiazolo [4,5-*d*] pyrimidine derivatives as adenosine A_{2A} receptor antagonists

Pratibha Mehta Luthra*, Chandra Bhushan Mishra, Pawan Kumar Jha, Sandeep Kumar Barodia

Medicinal Chemistry Division, Dr. B.R. Ambedkar Center for Biomedical Research, University of Delhi, Delhi 110 007, India

ARTICLE INFO

Article history: Received 28 August 2009 Revised 10 November 2009 Accepted 27 November 2009 Available online 4 December 2009

Keywords: Adenosine A_{2A}R Combinatorial chemistry GPCR Radioligand binding Parkinson's disease Thiazolopyrimidines

ABSTRACT

Novel bicyclic thiazolopyrimidine compounds (**15–26**) were synthesized to develop adenosine A_{2A} receptor ($A_{2A}R$) antagonist for the treatment of Parkinson's disease (PD). The binding affinity of the compounds (**15–26**) with $A_{2A}R$ was evaluated using radioligand binding assay on isolated membranes from stably transfected HEK293 cells. Selectivity of the compounds towards $A_{2A}R$ was assessed by comparing their binding affinities with A_1 receptors (A_1R). cAMP concentrations were measured from HEK293 cells treated with compounds (**15–26**) as compared to NECA ($A_{2A}R$ agonist). The compound (**16**) possessed strongest $A_{2A}R$ binding affinity (K_i value = 0.0038 nM) and selectivity (737-fold) versus A_1R . Decrease in $A_{2A}R$ -coupled release of endogenous cAMP from HEK293 cells treated with compounds (**15–26**) is evocative of their potential as $A_{2A}R$ antagonist.

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Parkinson's disease (PD) is a neurodegenerative disorder caused by the degeneration of dopaminergic neurons in nigrostriatal region of the brain and characterized by loss of motor coordination manifested as tremor and rigidity of the limbs and trunk.¹ Dopamine depletion due to degeneration of dopaminergic neurons can be supplemented by administration of the dopamine precursor L-DOPA, one of the major treatments for PD.² However, undesirable side effects such as dyskinesia (abnormal involuntary movements) induced by L-dopa therapy are one of the major challenges for the treatment of PD.³ Other available treatments (dopaminergic therapy) of PD include dopamine agonists and inhibitors of endogenous dopamine degradation by monoamine oxidase [MAO]B and catechol-O-methyltransferase[COMT]), and anticholinergics. As an alternative therapeutic approach, PD therapeutics involving A_{2A}Rs antagonists are gaining importance due to their capability to give partial symptomatic relief.^{4–6} Blockade of the A_{2A}Rs in striatopallidal neurons diminished postsynaptic effects of dopamine depletion, and in turn reduced the motor deficits of PD.⁷ A_{2A}R antagonists might partially improve not only the symptoms of PD but also its course, by lessening the prime neurodegeneration and reducing the maladaptive neuroplasticity³ A_{2A}R belongs to the family of seven trans-membrane G-protein coupled receptors (GPCRs) and is highly expressed in the nigrostriatum (basal ganglia) co-localized with dopamine D₂ receptors on striatopallidal

output neurons.⁸ A_{2A}R antagonist activated nigrostriatal pathway via D₂ receptor leading to antiparkinsonian effect. A_{2A}R and D₂R interact antagonistically in striatopallidal neurons. The effect of A_{2A}R activated adenylyl cyclase by persistent activation of D₂ receptor was antagonized at the level of the striatopallidal GABA neurons suggesting D₂ receptor mediated anti-parkinsonian role of A_{2A}R antagonists.⁹ Besides providing symptomatic relief in PD, A_{2A}R antagonists have also been shown to be neuroprotective in animal models of PD.¹⁰

Several xanthine and non-xanthine derivatives were synthesized in last decade in search of potent $A_{2A}R$ antagonist as novel antiparkinsonian agent (Fig. 1), however, not a single drug could be accomplished for the treatment of PD. Xanthine-based $A_{2A}R$ antagonists such as KW6002 showed efficacy in models of PD without inducing hyperactivity or inducing dyskinesia but failed in clinical trials.^{11–15} Non-xanthine compounds such as ZM241385 and SCH58261 possessed strong $A_{2A}R$ antagonist activity.^{16–19} However, suffered from several drawbacks including lower selectivity, poor solubility and poor pharmacokinetic profile.²⁰ The discovery of selective and potent $A_{2A}R$ antagonist still remains a challenge.

Thiazoles have emerged as an important class of compounds due to their anti-oxidant,²¹ anti-inflammatory,²² and neuroprotective effect.²³ A series of aryl/heteroaryl urea bearing thiazole compounds possessed selective cycline dependent kinase inhibiting activity.²⁴ Recently, thiazole derivatives with urea moiety have demonstrated A_{2A}R antagonist activity.²⁵ ECLiPSTM (Encoded

^{*} Corresponding author. Tel.: +91 11 27666272; fax: +91 11 27666248. E-mail addresses: pmluthra@acbr.du.ac.in, pmlsci@yahoo.com (P.M. Luthra).

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SCH 58261

Figure 1. Structures of A_{2A}R antagonists ZM241385, KW6002, and SCH58261. Binding affinities are as follows: (a) ZM241385 (*K*_i value; A_{2A}R = 0.8 nM, A₁R = 260 nM, and hA₁/hA₂ = 325)³³ (b) KW6002 (*K*_i value; A_{2A}R = 2.2 nM, A₁R = 150 nM, and hA₁/hA₂ = 68).³⁴ and (c) SCH58261 (*K*_i value; A_{2A}R = 1.23 nM, A₁R = 594.1 nM, and hA₁/hA₂ = 483).

combinatorial libraries on Polymeric Support) libraries showed aminothiazole core structure as active $A_{2A}R$ antagonists²⁶ Several pyrazolo-triazolo-pyrimidine derivatives were synthesized as potent A_{2A}R antagonist such as SCH58261.¹⁶ In our efforts to develop potent A_{2A}R antagonist, pyrazolo-triazolo-pyrimidine pharmacophore of known A_{2A}R antagonists (SCH58261) was replaced by thiazolopyrimidine pharmacophore, considered as thiazolo-analog of natural purine base along with open triazole ring to obtain 7imino-2-thioxo-3,7-dihydro-2H-thiazolo [4,5-d] pyrimidine pharmacophore with urea and furonamide moiety. Novel thiazolopyrimidine pharmacophore with urea moiety possessing aliphatic flexible groups, and aromatic planar structure was synthesized to give the compounds (15-21). Furan ring has been found essential for the activity of several non-xanthine derivatives to hold the molecule in active site cavity of A_{2A}R,²⁷ therefore, amino group of urea was replaced with furan ring to give furonamide series of the compounds (22-26). Since, non-xanthine compounds showed considerable selectivity for A2AR subtype, but suffered from low water solubility. We thought that thiazolopyrimidine ring possessing urea and furonamide derivatives would increase water solubility due to flexible polar interactions.

The binding affinity of the compounds (**15–26**) with human $A_{2A}R$ was evaluated in vitro using competitive binding with [³H]ZM241385. The $A_{2A}R$ selectivity of the compounds was determined using the A_1R selective antagonist [³H]DPCPX. $A_{2A}R$ -coupled cAMP concentrations were measured in HEK293 cells treated with compounds (**15–26**).²⁸

Synthesis of the compounds (**15–26**) was carried out according to scheme 1.²⁹ An equimolar mixture of isothiocyanate, malononitrile and sulfur powder in DMF was stirred in ice bath. After 15 min, triethylamine was added drop-wise to the mixture, and the reaction was continued for 4 h to give 4-amino-3-substituted-2-thioxo-2,3-dihydro-thiazole-5-carbonitrile derivatives (**1–7**). The carbonitrile compounds (**1–7**) were refluxed in toluene with triethylorthoformate (equimolar ratio) and *p*-toluene sulfonic acid (catalytic amount) for 6 h to yield imino-ether derivatives (**8–14**). The disappearance from IR spectra of an intense absorption band at 3230–3369 cm⁻¹ (free amino group), and the appearance of a singlet at 8.1–8.4 ppm (N=CH of methylene amine) in the 1H spectra validated the formation of Schiff's bases.

Mixture of imino-ether derivatives (8–14), semicarbazide hydrochloride/furoic acid hydrazide (equal mol) and TEA (catalyst) in ethanol was stirred at rt for 12 h to give the compounds (15–26).

The disappearance from IR spectra of an intense absorption band at 2205 cm⁻¹, and the appearance of a singlet at 8.1–8.8 ppm (N=CH of pyrimidine ring) in the 1H spectra confirmed the formation of bicyclic compounds. All the compounds have been fully characterized and their purity was checked by HPLC (Ref. 29 and Supplementary data).

The result of A_{2A}R binding assay are expressed as inhibition constants (K_i in nM). The A₁R/A_{2A}R describes their selectivity over A_1R^{30} In the set of thiazolopyrimidine urea derivatives (15–21, $R^2 = NH_2$), ethyl substitution (**15**, $R^1 = -C_2H_5$) exhibited significantly higher binding affinity with A_1R (0.00016 ± 0.007 nM) as compared to $A_{2A}R$ (0.09 ± 0.01 nM). Homologation of one carbon in compound 15 gave the propyl derivative of thiazolo pyrimidine urea (**16**, $\mathbb{R}^1 = -\mathbb{C}_3 \mathbb{H}_7$). In competitive radioligand binding assay, the displacement of [³H]ZM241385 with **16** was significantly increased (~25 times) as compared to 15, and with very high selectivity for A_{2A}R (737-fold selectivity over A₁R), and was better than the known antagonist SCH58261 ($K_i = 1.23 \pm 0.016$, $hA_1/hA_2 =$ 483).³¹ However 3-carbon chain with π -overlap in allyl derivative (**18**, $R^1 = -C_3H_5$) displayed good binding affinity ($K_1 = 0.092 \pm 0.01$) but reduced selectivity $(hA_1/hA_2 = 5)$. Further extending the alkyl chain to give butyl derivative of thiazolopyrimidine urea (17, $R^1 = -C_4H_9$) resulted in decreased selectivity over A₁R. Incorporation of aromatic ring (phenyl) substituent in thiazolopyrimidine urea (**19**, $R^1 = -C_6H_5$) gave enhanced binding affinity and selectivity. Moreover, *p*-iodophenyl substitution (**20**, $R^1 = -C_6H_4I$) on the pharmacophore improved both binding affinity and selectivity (144-fold). Insertion of one carbon homologation in planar aromatic ring of thiazolopyrimidine urea (**21**, $R^1 = -CH_2C_6H_5$) led decreased selectivity. We can conclude that both 19 and 20 possess promising activity, yet the compound (16) is the most active among all thiazolopyrimidine urea derivatives. The amino (NH₂) group of urea moiety of thiazolopyrimidine pharmacophore was replaced by furan ring to give another set of compounds (22-26, R^2 = -furyl). Overall substituent effects to binding affinity (propyl > butyl > allyl > aryl > ethyl) and selectivity (propyl > allyl > butyl > aryl > ethyl) profile of thiazolopyrimidine furonamide (22-26) decreased, however in the set of compound (22-26) propyl derivative (23, $R^1 = -C_3H_7$) showed maximum binding and selectivity to A2AR.

The finding clearly demonstrated that bicyclic thiazolopyrimidine urea derivatives (**15–21**) were more potent and selective than the corresponding bicyclic thiazolopyrimidine furonamide Download English Version:

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