

SYNTHESIS OF A PYRIMIDINE BY ELIMINATION OF NITROGEN FROM A TRIAZOLO[4,5-*D*]PYRIMIDINE

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Summary. Treatment of 3-phenyl-1,2,3-triazolo[4,5-*d*]pyrimidine-5,7-dithione with butyl lithium and an alkylating agent afforded a bis-alkylated pyrimidine, incorporating one butyl side chain.

We have recently proposed a model of the adenosine A1 and A2 receptors that defines the receptors as having three binding domains; a hydrophobic binding domain, an aromatic binding domain and a ribose binding domain.¹ The aromatic binding domain is described by a conserved 6 membered ring. We now report the synthesis of a six membered ring heterocyclic compound, 4-(*n*-butylthio)-6-(phenylamino)-2-propionamidylthiopyrimidine by elimination of nitrogen from a triazolopyrimidine. This appears to be the first monocyclic heterocyclic compound to exhibit binding to the adenosine receptor.

Treatment of 3-phenyl-1,2,3-triazolo[4,5-*d*]pyrimidine-5,7-dithione (**1**)² in THF at -70° with *n*-butyllithium (3.0 equiv, hexane, 5 min) followed by 2-bromopropionamide (2.0 equiv in THF, over 20 min) and stirring for a further 45 min at -70° gave complete reaction (tlc analysis:- hexane - ethyl acetate 1:1). Quenching with water, extractive workup, silica gel flash chromatography (ethyl acetate - hexane 1:1) and recrystallization from ethyl acetate afforded **2** in 30% yield.³ The product contained only one amide side chain, one methyl doublet at δ 1.56 in the ¹H NMR and a carbonyl and methine signals at δ 175.3 and 41.6 respectively in the ¹³C NMR. The presence of a butyl side chain was apparent from both ¹H and ¹³C NMR which showed signals for three methylenes and one methyl, the chain was apparently attached to a heteroatom as methylene signals occurred at δ 3.06 in the ¹H and δ 31.3 in the ¹³C NMR. Mass spectral analysis showed a base peak at 362 which together with the spectroscopic data indicated a formula of C₁₇H₂₂N₄OS₂. The loss of nitrogen from the precursor triazolopyrimidine was confirmed by a X-ray crystallographic analysis (Figure 1). Crystal/refinement data: -C₁₇H₂₂N₄OS₂, *M* = 362.6, Triclinic, space group *P* $\bar{1}$, *a* = 11.184(5), *b* = 10.066(11), *c* = 9.509(5)Å, α = 86.55(7), β = 68.61(3), γ = 68.40(7)°, *V* = 923 Å³. *D*_c (*Z* = 2) = 1.30g cm⁻³. *F*(000) = 384. Monochromatic MoK α radiation, λ = 0.71069Å, closely twinned crystal *ca* 0.2mm, μ_{Mo} = 2.5cm⁻¹, no absorption or extinction corrections. $2\theta_{\text{max}}$ = 45°, *N*_{indep} = 2414, *N*_{obs} = 1567; *R*_F = 0.064, *R*_W = 0.069 (statistical weights). Anisotropic thermal parameter refinement for non-hydrogen atoms; (*x*,*y*,*z*,*U*_{iso})_H constrained at estimated values in full matrix least squares refinement.⁴ The refinement model is consistent with the spectroscopic evidence.

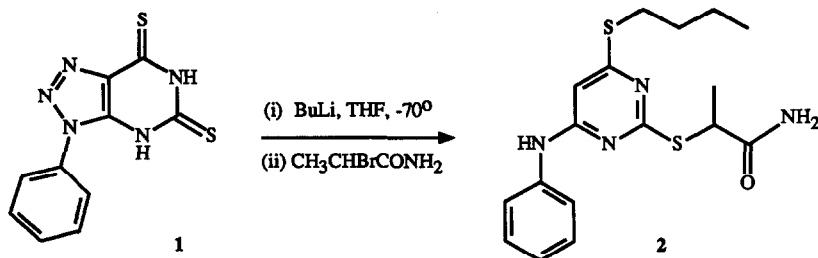


Table 1. Non-hydrogen atom coordinates

Atom	x	y	z
S(1)	0.1628(2)	0.8670(2)	0.3302(2)
S(2)	0.3758(2)	0.5368(2)	-0.1736(6)
N(1)	0.3912(6)	0.9056(6)	-0.0072(7)
C(2)	0.2973(7)	1.0026(7)	0.1089(7)
O(2)	0.3259(5)	1.0975(4)	0.1521(5)
C(3)	0.1609(7)	0.9870(6)	0.1822(7)
C(31)	0.0413(8)	1.1288(8)	0.2451(9)
C(4)	0.2216(6)	0.6959(6)	0.2344(7)
N(5)	0.2716(5)	0.6831(5)	0.0847(6)
C(6)	0.3112(7)	0.5464(6)	0.0231(7)
C(61)	0.4265(7)	0.3498(6)	-0.2275(8)
C(62)	0.4927(7)	0.3281(6)	-0.3987(7)
C(63)	0.5284(7)	0.1726(7)	-0.4535(8)
C(64)	0.5927(9)	0.1487(8)	-0.6252(9)
C(7)	0.2956(7)	0.4391(6)	0.1095(7)
C(8)	0.2433(6)	0.4666(6)	0.2675(7)
N(9)	0.2062(5)	0.5989(5)	0.3320(6)
N(10)	0.2298(5)	0.3591(5)	0.3559(6)
C(11)	0.1759(6)	0.3551(6)	0.5111(7)
C(12)	0.1234(7)	0.4727(7)	0.6184(8)
C(13)	0.0727(8)	0.4538(7)	0.7679(8)
C(14)	0.0673(8)	0.3267(8)	0.8204(8)
C(15)	0.1188(8)	0.2117(7)	0.7150(8)
C(16)	0.1701(7)	0.2271(7)	0.5659(8)

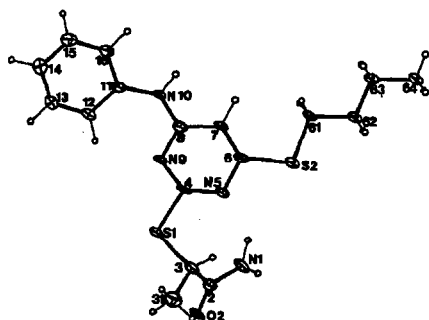


Figure 1 Projection of a single molecule normal to the central ring plane; 20% thermal ellipsoids are shown for the non-hydrogen atoms. Hydrogen atoms have arbitrary radii of 0.1 Å.

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References and Notes

- R. J. Quinn, M. J. Dooley, A. Escher, F. A. Harden and H. Jayasuriya, *Nucleosides and Nucleotides*, in press
- prepared following the method of A. Albert, C. I. Lin and I. Perkins, *J. Chem. Soc.*, 210 (1977).
- Spectral data for 2: m.p.: 142.7- 143.1°C; $^1\text{H-NMR}$ (250MHz, CDCl_3): δ 0.91 (t, J 7.5 Hz, 3H, CH_3), 1.42 (sextet, 2H, CH_2); 1.56 (d, J 7.5Hz, 3H, CH_3); 1.61 (quintet, 2H, CH_2); 3.06 (ABX₂ system J_{AB} 13.1 Hz, J_{AX} 7.2Hz, J_{BX} 7.2Hz, δ_A 3.1, δ_B 3.01, 2H); 4.37 (q, J 7.5Hz, 1H, C-H); 5.75 (br s, 1H, amide N-H); 6.31 (s, 1H, C-H); 7.06 (br s, 1H, amide N-H); 7.17-7.41 (m, 5H, aromatic H); 7.52 (br s, 1H, N-H); $^{13}\text{C NMR}$ (CDCl_3): δ 13.49 (CH_3), 16.34 (CH_3), 21.81 (CH_2), 29.29 (CH_2), 31.29 (CH_2), 41.55 (C-H), 95.66 (C-H), 123.01 (C², C⁶), 125.51 (C⁴), 129.63 (C³, C⁵), 137.67 (C¹), 159.62 (C), 169.32 (C), 170.00 (C), 175.27 (C=O); I.R. 3340 (N-H broad), 3180, 3230 (N-H), 2970, 2960, 2940, 1680, 1620, 1550, 1540 cm^{-1} ; UV (methanol): $\lambda_{\text{max}} = 252, 305 \text{ nm}$ ($\epsilon = 26780, 16380$), base shift: $\lambda_{\text{max}} 252, 306 \text{ nm}$; acid shift: $\lambda_{\text{max}} 248, 311 \text{ nm}$.
- Tables of structure factor amplitudes, thermal and hydrogen parameters and molecular geometries have been deposited at the Cambridge Crystallographic Centre.