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Key indicators

Single-crystal X-ray study T = 293 K Mean σ (C–C) = 0.004 Å R factor = 0.031 wR factor = 0.089 Data-to-parameter ratio = 8.4

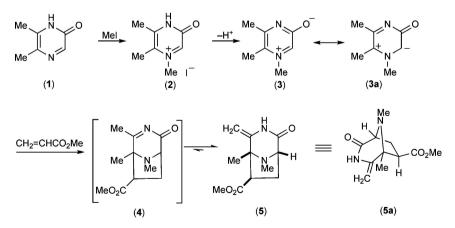
For details of how these key indicators were automatically derived from the article, see http://journals.iucr.org/e.

The dipolar cycloaddition of methyl acrylate to 1,5,6-trimethyl-3-oxidopyrazinium

5,6-Dimethylpyrazin-2-one reacts with iodomethane to give a quaternary salt, deprotonation of which liberates a 3-oxidopyrazinium which undergoes a 1,3-dipolar cycloaddition with methyl acrylate to form methyl 5,8-dimethyl-4-methyl-ene-2-oxo-3,8-diazabicyclo[3.2.1]octane-6-*exo*-carboxylate, $C_{11}H_{16}N_2O_3$, as the major product.

Comment

We have been investigating the 1,3-dipolar cycloaddition chemistry of 3-oxidopyraziniums (Kiss et al., 1987; Allway et al., 1990; Yates et al., 1995). These reactions efficiently produce bridged bicyclic systems, viz. 3,8-diazabicyclo[3.2.1]octanes, which comprise key structural components of such biologically active natural products as anticancer quinocarcin (Takahashi & Tomita 1983; Tomita et al., 1983; Hirayama & Shirahata, 1983) and antibiotic lemonomycin (He et al., 2000). Our studies were initially inspired by the series of benchmark papers by Katritzky and co-workers [for reviews, see Dennis et al. (1976) and Katritzky & Dennis (1989)] on the cycloadditions of 3-oxidopyridiniums. In neither Katritzky's extensive studies nor our own on 3-oxidopyraziniums had the possible influence of a substituent on the 1,3-dipole at one (or both) of the future ring-junction positions been assessed. This report describes our first study to remedy this omission, in which the reactivity of 1,5,6-trimethyl-3-oxidopyrazinium, (3) (see scheme; synthesis of 1,5,6-trimethyl-3-oxidopyrazinium and its reaction with methyl acrylate), was assessed.



5,6-Dimethylpyrazin-2-one, (1) (Jones, 1949; Karmas & Spoerri, 1952), was reacted with iodomethane to produce the methiodide (2), treatment of which with triethylamine allowed the generation of the zwitterion (3), *in situ* and in the presence of methyl acrylate. The reactivity and regioselectivity of such 3-oxidodiaziniums is easily understood in terms of a resonance contributor [(3a) in this case]. The immediate products of the

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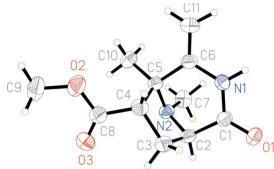


Figure 1

Plot of (4), with displacement ellipsoids drawn at the 50% probability level.

cycloadditions [(4) in this case] are not isolated, but tautomerize to the enamide structure [(5) in this case], (5a) showing better the bicyclic nature of the product. A mixture of two isomeric products was formed from which the major isomer was isolated, crystalline, allowing an X-ray analysis to show that it was the exo-ester (4) (Fig. 1). Thus, the additional oxidopyrazinium-6-methyl, appearing at the ring junction (C-5) in the cycloadduct, did not affect the efficiency or stereoselectivity of the cycloaddition, compared with the comparable reaction of 1,5-dimethyl-3-oxidopyrazinium which also gave a 6-exo-ester as the major product (Yates et al., 1995).

Experimental

5,6-Dimethylpyrazin-2-one (Jones, 1949; Karmas & Spoerri, 1952) (800 mg, 6.5 mmol) and iodomethane (2 ml, 32.5 mmol, 5 equivalents) were heated under reflux in MeCN (150 ml) under nitrogen for 24 h. The solvent was evaporated under vacuum and the residue extracted with CH₂Cl₂. Insoluble material was removed by filtration and the solution evaporated, leaving 3,4-dihydro-1,5,6-trimethyl-3oxopyrazinium iodide as a dark-brown crystalline solid (1.12 g, 66%; m.p. >523 K); ¹H NMR (D₂O, 300 MHz, δ , p.p.m.)): 8.20 (1H, s, C2-H), 4.15 (3H, s, NMe), 2.50 and 2.45 (2 \times s, 2 \times 3H, 2 \times CMe). Analysis found: C 32.31, H 4.02, N 10.39%; C₇H₁₁N₂O requires: C 31.60, H 4.17, N 10.53%

A solution of 3,4-dihydro-1,5,6-trimethyl-3-oxopyrazinium iodide (1.5 g, 5.6 mmol), Et₃N (1.6 ml, 11.2 mmol, 2 equivalents), and methyl acrylate (1.52 ml, 16.8 mmol, 3 equivalents) in dry MeCN (100 ml) was heated under reflux for 2 h. Solvents were removed from the resulting orange solution under vacuum, H2O (30 ml) was added and the product extracted into CH_2Cl_2 (3 × 30 ml). The combined dried extract was evaporated leaving a brown oil (0.88 g, 70%) from which, by careful chromatography over silica, eluting with n-hexane-EtOAc (1:1), the major (thin-layer chromatography) adduct was obtained as colourless plates (330 mg, 26%; m.p. 383-388 K).

Crystal data

$C_{11}H_{16}N_2O_3$	$D_x = 1.330 \text{ Mg m}^{-3}$
$M_r = 224.26$	Mo $K\alpha$ radiation
Monoclinic, $P2_1/c$	Cell parameters from 17414
a = 9.818 (10) Å	reflections
b = 7.89(2) Å	$\theta = 2.1 - 25.0^{\circ}$
c = 14.80 (3) Å	$\mu = 0.10 \text{ mm}^{-1}$
$\beta = 102.23 \ (8)^{\circ}$ V = 1120 (4) Å ³	T = 293 (2) K
$V = 1120 (4) \text{ Å}^3$	Plate, colourless
Z = 4	$0.3 \times 0.2 \times 0.1 \text{ mm}$

Data collection

A 1 1 K F K

Rigaku R-AXIS II diffractometer φ scans Absorption correction: none 17414 measured reflections 1767 independent reflections 1516 reflections with $I > 2\sigma(I)$	$\begin{aligned} R_{\text{int}} &= 0.037\\ \theta_{\text{max}} &= 25.0^{\circ}\\ h &= 0 \rightarrow 11\\ k &= 0 \rightarrow 9\\ l &= -17 \rightarrow 16 \end{aligned}$
Refinement	

Refinement on F^2	$w = 1/[\sigma^2(F_o^2) + (0.0555P)^2]$
$R[F^2 > 2\sigma(F^2)] = 0.031$	+ 0.1608P]
$wR(F^2) = 0.089$	where $P = (F_0^2 + 2F_c^2)/3$
S = 1.03	$(\Delta/\sigma)_{\rm max} = 0.003$
1767 reflections	$\Delta \rho_{\rm max} = 0.16 \text{ e } \text{\AA}^{-3}$
210 parameters	$\Delta \rho_{\rm min} = -0.13 \ {\rm e} \ {\rm \AA}^{-3}$
All H-atom parameters refined	Extinction correction: SHELXL97
	(Shaldmink 1007)

(Sheldlick, 1997)			
Extinction	coefficient: 0.037 (6)		

Table 1			
Hydrogen-bond	geometry	(Å,	°).

$D - H \cdots A$	D-H	$H \cdots A$	$D \cdots A$	$D - \mathbf{H} \cdots A$
$N1 - H1 \cdots O1^i$	0.913 (18)	2.082 (19)	2.980 (6)	167.6 (15)
Symmetry code: (i) $-x + 1$, $y - \frac{1}{2}$, $-z + \frac{3}{2}$.				

H atoms were found by difference Fourier methods and refined isotropically, with refined C-H distances in the range 0.932 (15)-1.041 (19) Å and an N-H distance of 0.913 (18) Å.

Data collection: MSC Diffractometer Control Software (Molecular Structure Corporation, 1992); cell refinement: DENZO (Otwinowski & Minor, 1987); data reduction: DENZO; program(s) used to solve structure: SAPI91 (Fan, 1991); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Bruker, 2001); software used to prepare material for publication: TEXSAN (Molecular Structure Corporation, 1995).

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