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Key indicators

Single-crystal X-ray study T = 150 KMean σ (C–C) = 0.003 Å R factor = 0.046 wR factor = 0.131 Data-to-parameter ratio = 9.1

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

4-{[3,5-Bis(methoxycarbonyl)-4,5-dihydro-1*H*-pyrazol-5-yl]methylamino}pyridine 1-oxide monohydrate

The title compound, $C_{13}H_{16}N_4O_5 \cdot H_2O$, was isolated following the cycloaddition reaction of 4-azidopyridine 1-oxide with an excess of methyl acrylate. The compound arises from a reaction sequence whereby the expected triazoline product undergoes rearrangement and a second cycloaddition reaction to yield the title pyrazoline product without the presence of a strong base which is usually required for such transformations.

Comment

The title compound, (I), is formed by the thermal reaction of 4-azidopyridine 1-oxide with methyl acrylate. 1,3-Dipolar cycloaddition reactions of aryl azides, particularly phenyl azide and its derivatives, with alkenes and alkynes have been well studied. In general terms, the expected outcome of such a reaction is a triazole (in the case of alkynes) or triazoline (in the case of alkenes) compound (Patai, 1971; Sustmann & Trill, 1972). However, under the reaction conditions employed, the triazoline compound generated from this reaction undergoes rearrangement to generate an diazoalkane intermediate, which in turn undergoes a second 1,3-cycloaddition reaction with methyl acrylate to generate the pyrazoline product. The X-ray crystal structure indicates that water co-crystallizes with the title compound, (I), in association with the *N*-oxide compound.



The tautomer shown in the *ORTEP* (Johnson, 1976) representation of the structure (Fig. 1) is consistent with the NMR data obtained for this compound. Peaks observed at 8.88 and 6.73 p.p.m. in d_6 -dimethyl sulfoxide (DMSO) are not observed in d_4 -methanol, indicating that these resonances correspond to labile N—H protons (at the N1 position of the pyrazoline ring, and N position of the aminopyridine 1-oxide fragment, respectively, confirmed by two-dimensional NMR experiments). Such reactions have been observed in substituted phenyl azides, but these have required the presence of a moderately strong base, such as 1,4-diazabicyclo[2.2.2]octane (DABCO), to catalyse the triazoline–diazoalkane rearrangement (Yadav *et al.*, 2002). In this particular case, either water itself is sufficiently basic to catalyse the reaction, or no such base-catalysis is required.

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Experimental

A solution of 4-azidopyridine 1-oxide in methyl acrylate and acetonitrile (approximate 1:1 ratio) was deoxygenated by a thin stream of bubbling argon and heated at 333 K overnight. Slow evaporation of the solvent yielded crystalline chunks of the title compound which were isolated from the supernatant and characterized.

Crystal data

 $\begin{array}{l} C_{13}H_{16}N_4O_5 \cdot H_2O\\ M_r = 326.31\\ \text{Triclinic, } P\overline{1}\\ a = 6.0678 \ (7) \ \text{\AA}\\ b = 8.7476 \ (9) \ \text{\AA}\\ c = 15.0449 \ (18) \ \text{\AA}\\ \alpha = 104.435 \ (6)^{\circ}\\ \beta = 94.885 \ (6)^{\circ}\\ \gamma = 103.676 \ (6)^{\circ} \end{array}$

Data collection

Nonius KappaCCD diffractometer ω scans Absorption correction: multi-scan

(Otwinowski & Minor, 1997) $T_{\min} = 0.936, T_{\max} = 0.969$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.046$ $wR(F^2) = 0.131$ S = 1.082559 reflections 280 parameters V = 742.57 (15) Å³ Z = 2 D_x = 1.459 Mg m⁻³ Mo - Kα radiation μ = 0.11 mm⁻¹ T = 150 K Chunk, colourless 0.40 × 0.35 × 0.29 mm

8766 measured reflections 2559 independent reflections 1906 reflections with $I > 2.0\sigma(I)$ $R_{\text{int}} = 0.029$ $\theta_{\text{max}} = 27.8^{\circ}$

All H-atom parameters refined $w = 1/[\sigma^2(F_o^2) + (0.0806P)^2]$ where $P = (F_o^2 + 2F_c^2)/3$ $(\Delta/\sigma)_{max} < 0.001$ $\Delta\rho_{max} = 0.47$ e Å⁻³ $\Delta\rho_{min} = -0.24$ e Å⁻³

Coordinates and isotropic displacement parameters for all H atoms were refined [C-H = 0.86 (5)-1.02 (2) Å].

Data collection: *COLLECT* (Nonius, 1998); cell refinement: *DENZO/SCALEPACK* (Otwinowski & Minor, 1997); data reduction: *DENZO/SCALEPACK*; program(s) used to solve structure: *SIR2002* (Burla *et al.*, 2003); program(s) used to refine structure: *SHELXL97* (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON* (Spek, 2003); software used to prepare material for publication: *SHELXL97* and local programs.



Figure 1

View of (I) showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 50% probability level and H atoms are shown as small spheres of arbitrary radii.

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