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Key indicators

Single-crystal X-ray study T = 293 KMean $\sigma(C-C) = 0.007 \text{ Å}$ Disorder in solvent or counterion R factor = 0.061 wR factor = 0.220 Data-to-parameter ratio = 10.9

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

N,N-Dimethyl-5-dimethylamino-3-(1-pyrrolyl)-2,4pentadieniminium tetrafluoroborate

In the crystal structure of the title salt, $C_{13}H_{20}N_3^+ \cdot BF_4^-$, the asymmetric unit has two independent cations and anions. The cations have similar markedly non-planar pentamethinecyanine moieties with a *trans-cis-cis-trans* conformation of the pentamethinecyanine NC₅N chain. The pyrrole substituents are rotated 21.14 (9) and 23.43 (19)° from planarity with the adjacent atoms in the pentamethinecyanine chains.

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Comment

The title salt, (1), containing the relatively electron-rich 1-pyrrolyl substituent was prepared in order to determine what effect the 1-pyrrolyl substitution would have on the conformation of the pentamethinecyanine moiety of the cation.



There are two independent cations and anions in the asymmetric unit of (1) with each of the anions disordered over two sites. The pentamethinecyanine moieties in the cations (Figs. 1 and 2) have similar, markedly non-planar, *trans-cis-cis-trans* conformations, with torsion angles Cn1-Cn2-Cn3-Cn4 and Cn2-Cn3-Cn4-Cn5 having values -22.8 (6), -30.7 (6), and -27.2 (6) and -26.4 (6)° for n = 1 and 2, respectively. It is interesting to compare these conformations with those found for the pentamethinecyanine moieties in 5-dimethylamino-N,N-dimethyl-3-(4-pyridyl)-2,4-pentadieniminium tetrafluorobate [(2); all-*trans*; Ferguson *et al.*, 2001] and in tris(2-dimethylaminoethenyl)methylium per-

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chlorate [(3); *trans-trans-cis-trans*; Dale *et al.*, 1988]. In (2), the relatively electron-poor 4-pyridyl substitutent is not conjugated with the pentamethinecyanine, and these two moieties form a dihedral angle of 60.58 (9)°. In (3), which may be regarded as a pentamethinecyanine carrying an electron-rich 2-dimethylaminoethenyl substituent at the central carbon, conjugative interaction results in a change in the conformation of the pentamethincyanine moiety in order to avoid a potentially severe steric interaction between C1-H (or C5-H) and C2-H of the 2-dimethylaminoethenyl substituent. In addition, the cation in (3), which has a threefold symmetry axis, is not planar but adopts a twisted-propeller shape.

In spite of their different conformations, bond lengths within the pentamethinecyanine moiety of (1) do not differ significantly from those reported for the all-trans pentamethinecyanine moieties in the simple parent salt (4) (Honda et al., 1986) and the 3-(4-pyridyl) derivative (2) (Ferguson et al., 2001). The bond lengths within the 1-pyrrolyl substituent of (1) are very similar to those reported (Ferguson & McAlees, 1990) for the 1-pyrrolyl fragment in the pyridinium salt (5), consistent with direct conjugation between the relatively electron-rich 1-pyrrolyl substituent and the electron-poor pentamethinecyanine and pyridinium moieties in (1) and (5), respectively. Such conjugation is also suggested by the relatively short bond between the pyrrole and pentamethinecyanine moieties [C13-N13 1.399 (5) Å and C23-N23 1.409 (5) Å]. With the pentamethinecyanine chain in the alltrans conformation, coplanarity with the pyrrole would result in severe steric interactions between the pyrrole α -H atoms and C1-H and C5-H. These and other interactions are avoided in (1) by adoption of the trans-cis-cis-trans conformation along with out-of-plane twisting as found for (3) (Dale et al., 1988).



Figure 1

View of one of the two independent cations of (1) with our numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.



Figure 2

Views of the second of the two independent cations of (1) with our numbering scheme. Displacement ellipsoids are drawn at the 30% probability level.

Experimental

A solution of dimethylamine (0.98 g, 22 mmol) in ethanol (7 ml) was added to a suspension of N-(2,4-dinitrophenyl)-4-(1-pyrrolyl)pyridinium chloride [(5); 3.47 g, 10.0 mmol; Ferguson & McAlees, 1990] in ethanol (20 ml) and the mixture was stirred at ambient temperature for 16 h. The resulting mixture was filtered, and the collected green-yellow powder (2,4-dinitroaniline) was washed with ethanol (3 \times 10 ml). The combined filtrate and washings were evaporated in vacuo to leave a clear orange gum which was taken up in water (50 ml) and filtered to remove yellow insoluble material. The aqueous filtrate was then washed with chloroform $(3 \times 25 \text{ ml})$ and treated with an aqueous solution (10 ml) of NaBF₄ (1.2 g, 11.0 mmol). The resulting mixture was extracted with dichloromethane (3 \times 30 ml), the combined extracts were dried (Na₂SO₄) and the solvent was evaporated to leave an orange-red gum. Crystallization from ethanol gave yellow hexagonal plates of (1) (2.32 g, 76% yield; m.p. 397.5-398.5 K).

Crystal data

 $\begin{array}{l} C_{13}H_{20}N_3^{+}\cdot BF_4^{-}\\ M_r = 305.13\\ Monoclinic, P_2/c\\ a = 14.882 \ (3) \ {\rm \AA}\\ b = 10.366 \ (2) \ {\rm \AA}\\ c = 24.064 \ (3) \ {\rm \AA}\\ \beta = 121.987 \ (11)^{\circ}\\ V = 3148.7 \ (10) \ {\rm \AA}^3\\ Z = 8 \end{array}$

 $D_x = 1.287 \text{ Mg m}^{-3}$ Mo K α radiation Cell parameters from 25 reflections $\theta = 10.4-12.8^{\circ}$ $\mu = 0.11 \text{ mm}^{-1}$ T = 293 (1) KPlate, yellow $0.43 \times 0.43 \times 0.30 \text{ mm}$ Data collection

Nonius CAD-4 diffractometer θ -2 θ scans 5728 measured reflections 4505 independent reflections 2271 reflections with $I > 2\sigma(I)$ $R_{int} = 0.036$ $\theta_{max} = 25.0^{\circ}$

Refinement

Refinement on F^2 $R[F^2 > 2\sigma(F^2)] = 0.061$ $wR(F^2) = 0.220$ S = 1.094505 reflections 414 parameters H-atom parameters constrained

Table 1

Selected geometric parameters (Å, °).

N11-C11	1.310 (5)	N21-C21	1.310 (5)
N11-C16	1.447 (5)	N21-C26	1.452 (6)
N11-C17	1.444 (6)	N21-C27	1.444 (6)
N12-C15	1.308 (5)	N22-C25	1.316 (5)
N12-C18	1.455 (5)	N22-C28	1.462 (5)
N12-C19	1.454 (6)	N22-C29	1.461 (5)
N13-C13	1.399 (5)	N23-C23	1.409 (5)
N13-C110	1.390 (5)	N23-C210	1.387 (5)
N13-C113	1.390 (5)	N23-C213	1.368 (5)
C11-C12	1.387 (6)	C21-C22	1.381 (5)
C12-C13	1.389 (6)	C22-C23	1.389 (5)
C13-C14	1.396 (5)	C23-C24	1.387 (6)
C14-C15	1.374 (5)	C24-C25	1.383 (5)
C110-C111	1.345 (6)	C210-C211	1.343 (6)
C111-C112	1.417 (7)	C211-C212	1.406 (7)
C112-C113	1.330 (6)	C212-C213	1.359 (6)
N11-C11-C12-C13	174.6 (4)	N21-C21-C22-C23	171.4 (4)
C11-C12-C13-C14	-22.8(6)	C21-C22-C23-C24	-27.2(6)
C12-C13-C14-C15	-30.7(6)	C22-C23-C24-C25	-26.4(6)
C13-C14-C15-N12	-179.3(4)	C23-C24-C25-N22	175.6 (4)

 $h = -17 \rightarrow 15$

3 standard reflections

frequency: 120 min

intensity decay: 1.0%

 $w = 1/[\sigma^2(F_o^2) + (0.1129P)^2]$

where $P = (F_o^2 + 2F_c^2)/3$

Extinction correction: SHELXL97

Extinction coefficient: 0.0060 (12)

+ 0.6033P]

 $(\Delta/\sigma)_{\rm max} < 0.001$

 $\Delta \rho_{\rm max} = 0.45 \text{ e } \text{\AA}^{-3}$

 $\Delta \rho_{\rm min} = -0.25 \text{ e } \text{\AA}^{-3}$

 $k = 0 \rightarrow 9$

 $l = 0 \rightarrow 28$

Compound (1) crystallized in the monoclinic system; space group $P2_1/c$ from the systematic absences. H atoms were treated as riding atoms with C-H 0.93 and 0.96 Å. Difference maps showed that methyl H atoms at C17, C19 and C29 were disordered and they were modelled using six H sites, each with occupancy 0.50, mutually offset by 60°. The H atoms of the other methyl groups were modelled with free rotation about the relevant C-C bond. It was also apparent that the F atoms of the BF₄⁻ ions were disordered over major and minor occupancy sites with common B atoms. The geometry of these tetrahedral BF₄⁻ anions was restrained to be near to tetrahedral by use of *DFIX* commands with an overall restrained B-F distance. The minor occupancy F atoms were allowed for with overall isotropic displacement parameters.

Data collection: *CAD*-4-*PC Software* (Nonius, 1992); cell refinement: *SET*4 and *CELDIM* (Nonius, 1992); data reduction: *HELENA* in *PLATON* (Spek, 2001); program(s) used to solve structure: *SHELXS*97 (Sheldrick, 1997); program(s) used to refine structure: *SHELXL*97 (Sheldrick, 1997); molecular graphics: *ORTEPII* (Johnson, 1976) and *PLATON*; software used to prepare material for publication: *SHELXL*97 and *WordPerfect* macro *PREP8* (Ferguson, 1998).

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