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The Structure of the [Platinum(ethylenediamine)(guanosine)₂]²⁺ Cation

Sir:

The surprisingly potent antitumor properties of cis- $Pt(NH_3)_2Cl_2$, $Pt(en)Cl_2$, and related compounds have been the subject of intensive investigation for the past few years.¹⁻³ Clinical trials using cis-Pt(NH₃)₂Cl₂ as a chemotherapeutic agent have been in progress in several hospitals since 1971.^{3,4} It is suspected that the activity of these compounds is related to their ability to interact with DNA,5-7 and several models involving the formation of platinum-bridged covalent cross-links have been proposed.⁶⁻¹⁰ These models generally involve a cis-Pt(amine)₂ fragment chelated by certain atoms from DNA, usually nitrogen or oxygen atoms from the purine or pyrimidine bases.⁷⁻¹⁰ Guanine in particular is thought to be especially vulnerable to attack, 9-12 since guanine derivatives are known to react more rapidly with platinum complexes than derivatives of the other nucleotide bases.¹² In this communication we wish to report the crystal structure of the [platinum(ethylenediamine)(guanosine)₂]²⁺ cation and suggest a possible model for Pt-DNA interaction.

 $[Pt(en)(Guo)_2]^{2+}$, prepared by treating $Pt(en)Cl_2$ with guanosine,¹³ was isolated as crystals of a mixed chlorideiodide salt¹⁴ by vapor diffusing¹⁶ the reaction mixture against 1-propanol. Crystals of $[Pt(en)(Guo)_2]Cl_{1.5}I_{0.5}$. $2H_2O$ are tetragonal (space group $I4_122$), with a = 17.557(4) Å, c = 23.883 (6) Å, and Z = 8. Two asymmetric sets of data were collected on a Nonius CAD-3 automated diffractometer using a $\theta/2\theta$ scan mode with Mo K α radiation



Figure 1. The geometry of the [Pt(en)(Guo)₂]²⁺ cation. A crystallographic twofold rotation axis bisects the N_7 -Pt- N_7 angle.



Figure 2. Part of the infinite spiral consisting of stacked [Pt(en)- $(Guo)_2]^{2+}$ units, showing the pairing of the guanine bases. The ribose groups have been omitted for clarity. A crystallographic 41 screw axis is approximately defined by the row of platinum atoms. The boxed area shows a Pt(en) fragment forming a bridge between two parallel guanine rings. Hydrogen bonds connect the NH2 groups of the ethylenediamine ligands to the carbonyl groups of the guanines. One of these is indicated (dotted line).

up to a 2θ limit of 50°. The data were corrected for absorption effects and merged to give 1146 nonzero reflections. The structure was solved by heavy atom methods and refined anisotropically to a final R factor of 5.9%.¹⁷

The square-planar geometry of the $[Pt(en)(Guo)_2]^{2+}$ cation is shown in Figure 1. A crystallographic twofold rotation axis passes through the platinum atom and the mid-point of the C-C bond of the ethylenediamine ligand. Each guanosine molecule coordinates in a monodentate fashion through the N7 position of the guanine base, making a Pt-N7 bond of 1.967 (15) Å and a N7-Pt-N7 angle of 87.0 (7)°. The planes of the guanine rings are tilted in such

a way that the normals to the planes intersect at an angle of 71°. Other distances and angles in the molecule are: Pt- $N(en) = 2.036 (17) \text{ Å}, N(en)-Pt-N(en) = 83.9 (9)^{\circ}, N_{7} Pt-N(en) = 94.6 (8)^{\circ}, 177.0 (8)^{\circ}, and N_7 - N_7 = 2.71 (2)$ Å. The ribose ring is in the C(3')-endo conformation and the torsion angles within the nucleoside are: $\chi = 224.0^{\circ}, \psi$ = 44.2°, $\psi' = 85.9°$, $\tau_0 = 358.7°$, $\tau_1 = 339.8°$, $\tau_2 = 35.9°$, $\tau_3 = 323.7°$, $\tau_4 = 22.9°$.¹⁸ The carbonyl oxygen at C₆ is hydrogen-bonded to an ethylenediamine nitrogen of a neighboring molecule at a distance of 2.84 Å. Additionally, each guanine ring is in perfectly parallel contact with a guanine ring on a neighboring cation at a distance of 3.31 Å. This effect, coupled with a crystallographic 4₁ screw axis which is approximately coincident with the row of platinum atoms, creates an infinite spiral of stacked [Pt(en)- $(Guo)_2$ ²⁺ units along the 4₁ axis (Figure 2).

Our result reinforces the belief that the preferred site for Pt complexation to guanosine is N_7 .²¹ This is significant since N7, an atom not involved in Watson-Crick base pairing, is exposed on the surface of the DNA molecule and would be a logical site for initial attack by an electrophilic species.

The orientation of the Pt(en) moiety with respect to a pair of parallel guanine rings from different cations suggests a possible model for initial attachment of a platinum-containing fragment to DNA. A closer examination of this region (boxed area in Figure 2) shows the Pt(en) group forming a bridge $[C_6=O...H-N(en)-Pt-N_7]$ across the pair of parallel guanine rings. The hydrogen bonding observed between the NH₂ group of the ethylenediamine ligand and the carbonyl group of the guanine is especially interesting in view of the fact that antitumor activity of cis- $Pt(amine)_2X_2$ complexes decreases markedly along the series $NH_3 \sim NH_2R > NHR_2 \gg NR_3$.^{2b}

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Evaluation of the Force Constant for the Carbon-Chlorine Stretching Vibration for tert-Butyl Chloride Using MINDO/3

Sir:

Recent work in these laboratories has illustrated the promise of kinetic isotope effects (KIE) as a chemical tool for studying reaction mechanisms and transition states.¹⁻⁵ In the course of such studies, vibrational frequencies, isotopic shifts, and vibrational force constants are necessary items for KIE calculations. The procedure involves the use of the Wilson FG method⁶ and the process is an iterative one, requiring experimental frequencies and an initial trial set of force constants. It would be desirable to have a theoretical means of obtaining these initial values. The purpose of this communication is to demonstrate that the MINDO/ 3 SCF MO method shows promise for producing such force constants for molecular vibrational modes of organic molecules.

The MINDO/3 SCF MO method developed by Dewar and co-workers has been described in detail recently in this journal.⁷⁻¹¹ It is a semiempirical SCF molecular orbital treatment which is quite successful in predicting such molecular properties as heats of formation, ionization energies, and dipole moments. The primary attribute of MINDO/3 of interest here is in optimizing molecular geometries of reactants, products, and transition states. The importance of this capability is evidenced by two studies employing an earlier version of MINDO (MINDO/2). In one study¹² where the molecular geometry was not optimized, the calculated stretching force constants agreed with experimental data to within a few tenths of a millidyne per angstrom for typical hydrocarbons but differed by more than 10% for all studied molecules containing heteroatoms. Much of the error could be attributed to the importance of the interactions of vibrational modes as the bond in question was