MS21-P08

Crystal and electronic structure of Ruthenium(II) complex

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The crystal structure of Ruthenium(II) complex, $C_{24}H_{26}Cl-N_2O_4Ru\cdot PF_6$, has been determined and analyzed in terms of connectivity and packing patterns. The compound crystal-lizes in the monoclinic crystal system in the space group $P2_1/c$. with one cation and one anion in the asymmetric unit [1].

Biological effects of ruthenium complexes and their potential use as metal-based drugs, especially as anticancer agents are one of the leading area. Excluding widespread use of platinum-based chemotherapeutics, ruthenium-based complexes are among the most promising alternatives.

Data collection was performed on a Stoe STADIVARI diffractometer with a Dectris Pilatus 300K detector and with an Incoatec IµS Ag microfocus source (Ag-K α , λ = 0.56083 Å) at 100 K using a nitrogen gas open-flow cooler Cobra from Oxford Cryosystems. Data reduction was processed using X-Area [2]. For numerical absorption corrections a crystal-shape model with 10 faces was employed. An average redundancy of 18.28 gives Rint of 3.18%. A Hirshfeld surface analysis was carried out and two-dimensional (2D) fingerprint plots [3] were generated to visualize the intermolecular interactions and to provide quantitative data for their relative contributions. Direction cosines were applied for the anisotropic secondary extinction correction. The results of multipole refinement were performed on F² using XD suite of programs. After the multipole refinement a topological analysis of the charge density was done. In addition, these results were compared with the electrostatic potential from Hirshfeld surfaces.

Acknowledgement

This work has been carried out with the support of the Ministry of Education, Science, Research and Sport of the Slovak Republic within the Research and Development Operational Programme for the project "University Science Park of STU Bratislava", ITMS 26240220084, co-funded by the European Regional Development Fund and also by support of the Research and Development Agency under the contract No. APVV-15-0079 and Scientific Grant Agency of the Slovak Republic VEGA (Project Nos. 1/0871/16 and 1/0598/14).

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Keywords: ruthenium(II) complex, electrostatic potential, charge density

MS22- Molecular structure and chemical properties: chemistry meets charge density

Chairs: Dr. Vladimir Stilinovic, Dr. Jacob Overgaard

MS22-P01

HAR and TAAM eefinements of model crystal structures using CuKα and MoKα X-ray diffraction data

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Commonly, the Independent Atom Model (IAM) of electron density is used in the case of routine X-ray data analysis. However, this model does not give a quantitative description of electron density distribution because atoms are assumed to be neutral and spherical. A far better model that allows for modelling of deformation of spherical charge density was introduced by Hansen and Coppens¹ and is called a pseudoatom model of electron density. Application of this kind of model requires an excellent quality crystals and high resolution XRD data. Quite often, this is difficult to be fulfilled. Therefore, new methods have been developed that enable reconstruction of electron density i.e. Hirshfeld Atom Refinement (HAR)² or Transferable Aspherical Atom Model (TAAM)³.

Validation of application of HAR and TAAM methods for low-resolution data will be presented. For three compounds following refinements of CuK α and MoK α data will be presented: IAM, TAAM (isotropic H atoms), TAAM (using Shade), HAR (isotropic H atoms), HAR (anisotropic H atoms), HAR (using Shade). Additionally, the multipole model and higher order of TAAM and HAR refinements will be presented for MoK α X-ray diffraction data. Analysis of geometry, ADPs (Fig. 1), fractal dimension plots and residual density maps will be shown in comparison with neutron diffraction data.

Fig.1 PEANUT representations of the difference between ADPs obtained from neutron data refinement and HAR, TAAM refinements for MoKa and CuKa data. Root mean square difference scale of 2 was used.

