

angles in the heterocyclic ring.

Keywords: organophosphorus compounds, dioxaphosphepine rings, anticancer agents

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Crystal structure and quantum mechanical calculations of an oxime compound, $C_9H_{11}N_3O_2$

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Oxime and dioxime derivatives are very important compounds in the chemical industry and medicine [1]. The oxime (-C=N-OH) moiety is potentially ambidentate, with possibilities of coordination through the N and/or O atoms. It is a functional group that has not been extensively explored in crystal engineering. In the solid state, oximes are usually associated via O-H...N hydrogen bonds of length 2.8 Å. Oxime groups possess stronger hydrogen-bonding capabilities than alcohols, phenols and carboxylic acids [2]. The hydrogen-bond systems in the crystals of oximes have been analysed and a correlation between a pattern of hydrogen bonding and N-O bond lengths has been suggested [3]. The configurational and/or conformational isomers of glyoxime derivatives (dioximes) have also been analysed [4]. The experimental geometry of $C_9H_{11}N_3O_2$ obtained from single-crystal X-ray diffraction was compared with those obtained from quantum-mechanical calculations. It is found that the bond lengths and angles obtained from two methods are in good agreement with each other.

[1] Sevagapandian, S., Rajagopal, G., Nehru, K. & Athappan, P. (2000). *Transition Met. Chem.* **25**, 388-393. [2] Marsman, A. W., Leussing, E. D., Zwikker, J. W. & Jenneskens, L. W. (1999). *Chem. Mater.* **11**, 1484-1491. [3] Bertolasi, V., Gilli, G. & Veronese, A. C. (1982). *Acta Cryst.* **B38**, 502-511. [4] Chertanova, L., Pascard, C. & Sheremetev, A. (1994). *Acta Cryst.* **B50**, 708-716.

Keywords: oxime, X-ray diffraction, quantum mechanical methods

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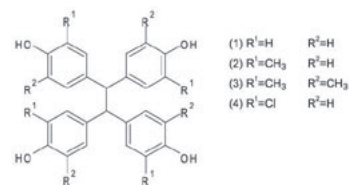
Crystal structure analysis of inclusion crystals with tetrapodal host molecules

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Inclusion crystals have unique characteristics such as guest solubility or temperature dependent guest releasing, which is relevant to the structure of host framework enclosing guest molecules. Hence it is of great importance to investigate and develop host frameworks to design new functional materials. 1,1,2,2-tetrakis (4-hydroxyphenyl) ethane, TEP (1), has four hydroxyl phenyl groups in different directions to build different types of hydrogen bonding host frameworks corresponding to guest molecule types. In our previous study, three types of host frameworks accommodating different kind

of guest molecules via hydrogen bonds were analyzed to show TEP has well-qualified inclusion ability as an inclusion host molecule. By introducing one or two substituents next to hydroxyl group, (2)-(4), hydrogen bonding direction and pattern should be restricted and such TEP derivatives are expected to show different hydrogen bonding pattern which leads new host frameworks. In this study, inclusion crystals of TEP derivative are synthesized and their crystal structures are analyzed to compare the host framework structures.



Keywords: inclusion compounds, molecular interactions, crystal engineering

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A detailed study of the helical conformation in 2-aminothiazole derivatives

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Aminothiazoles have many applications such as antiviral, antimicrobial, bactericidal and fungicidal agents. Aminothiazoles metal complexes display anti cancer activity.¹ The crystal structures of 2-Aminothiazole N-Boc protected Leucine and Valine belongs to the space group $P 2_12_12_1$ and $P 4_12_12$ with cell dimensions $a=10.5247(14)\text{Å}$, $b=11.2407(15)\text{Å}$, $c=30.3258(3)\text{Å}$ and $a=9.2614(4)\text{Å}$, $b=9.2614(4)\text{Å}$, $c=36.6412(3)\text{Å}$ respectively. The molecules are packed via intermolecular N-H...O and N-H...N hydrogen bonds giving rise to the formation of helix like structure. The helical nature and the tilt associated with the title compounds compares well with those of the B-DNA. The pitch per one complete turn of the helix is 30.30Å and 36.64Å while the pitch in B-DNA is reported to be 34Å and this aspect might have consequences related to the activity profiles of this compound. The idea is to exploit the hydrogen bonding potential in the title compound to get the supramolecular assembly into a helix and examine its features with respect to DNA.

1. Lynch, D. E. *et al*, *Acta Cryst.*, 1999, **B55**, 758

Keywords: supramolecular assemblies, DNA, hydrogen bonding recognition



Valine Thiazole