

Chloroquine Derivatives. Conformation and Intermolecular Interactions

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Crystal structure of hydroxychloroquine (OHClQ) sulfate has been determined: $a=10.5437(3)\text{\AA}$, $b=8.8532(2)\text{\AA}$, $c=22.0923(8)\text{\AA}$, $\alpha=90^\circ$, $\beta=101.426(1)^\circ$, $\gamma=90^\circ$, $P2_1/c$, $Z=4$, in order to compare its conformation and intermolecular interactions to those in the crystalline chloroquine (ClQ) phosphate [1] and quinine salicylate (QSal) monohydrate [2].

Molecular conformations of OHClQ and ClQ are comparable in both salts; the differences between corresponding torsion angles are not greater than 10° . Each of the nitrogen atoms is a proton donor in the intermolecular hydrogen bonds with the oxygen atoms of sulfate or phosphate anions. While the parameters of the $N1-H1\cdots O$ and $N3-H3\cdots O$ bonds are similar, the distance $N\cdots O$ within the bond $N2-H2\cdots O$ is much shorter in the case of OHClQ. The -OH group of OHClQ forms an additional H-bond with the oxygen atom of SO_4^{2-} .

The comparison of the hydrogen bonds formed by OHClQ and ClQ with those of quininium anion in QSal shows that these antimalarial molecules may interact with their putative receptor in a similar way.

[1] Karle J.M., Karle I.L., *Acta Cryst.*, 1988, **C44**, 1605. [2] Oleksyn B.J., Serda P., *Acta Cryst.*, 1993, **B49**, 530.

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