Crystal Structure of a Polymorph of Carnidazole from Synchrotron X-ray Powder Diffraction

Hector Novoa de Armas^a, Maurice O. Peeters ^a, Dirk J.A. De Ridder^b, Henk Schenk^b, Norbert Blaton^a, ^aKatholieke Universiteit Leuven, Laboratorium voor Analytische Chemie en Medicinale Fysicochemie, Faculteit Farmaceutische Wetenschappen, Van Evenstraat 4, 3000 Leuven, Belgium. ^bUniversieit van Amsterdam, Laboratorium voor Kristallografie, IMC, Nieuwe Achtergracht 166, 1018 WV Amsterdam, The Netherlands. E-mail: hector.novoa@pharm.kuleuven.ac.be

The crystal structure of a polymorphic form of carnidazole, a nitroheterocyclic compound active against both anaerobic protozoa and bacteria, has been determined using synchrotron powder diffraction. The simulated annealing approach, as implemented in the program FOX [1], was used to obtain the initial model. The model was refined with the Rietveld refinement program Fullprof [2]. This monoclinic polymorph crystallizes in $P2_1/n$ space group with a=13.907(3), b=8.091(2), c=10.643(2), $\beta=110.831(5)$, Z=4. The imidazole ring is planar. The molecules are held in the crystal forming two infinite zig-zag chains along [010] via hydrogen bonds of the type N-H...N. A structural comparison with the previously reported polymorph and the monohydrate forms of this drug is presented.

[1] Favre-Nicolin V., Cerny R., J. Appl. Cryst., 2002, 35, 734. [2] Rodriguez-Carvajal J., 2001, FullProf, version 1.9c, LLB, CEA/Saclay, France.

Keywords: carnidazole, powder diffraction, polymorphism