

X-Ray Structure Analysis of Substituted 5-(2,3-Dihydro-7-benzofuryl)-1-methylpyrazolo[4,3-*d*]pyrimidin-7-one (Biagra) [1]

Wolfgang Voelter^{a,*}, Cäcilia Maichle-Mössmer^b, Monther A. Khanfar^b,
Mustafa M. El-Abadelah^{c,*}, and Salim S. Sabri^c

^a Abteilung für Physikalische Biochemie, Physiologisch-chemisches Institut der Universität Tübingen, Hoppe-Seyler-Straße 4, D-72076 Tübingen, Germany

^b Institut für Anorganische Chemie auf der Morgenstelle 18, Universität Tübingen, D-72076 Tübingen, Germany

^c Chemistry Department, Faculty of Science, University of Jordan, Amman-Jordan

* Reprint requests to Prof. Dr. Dr. h.c. W. Voelter.

E-mail: wolfgang.Voelter@uni-tuebingen.de or to Prof. Dr. M. El-Abadelah

Z. Naturforsch. **54b**, 1602–1605 (1999); received August 8, 1999

Dihydrobezofuryl-pyrazolo[4,3-*d*]pyrimidin-7-one, X-Ray Data

X-ray crystal structure data for the title compound dihydrobenzofuryl-pyrazolo[4,3-*d*]pyrimidin-7-one (**2**) reveal that the two bicyclic heteroaryl systems are virtually coplanar along their joint C(5)-C(11) axis with an interplanar angle of 1.13°. This planar conformation is held by an intramolecular hydrogen bond between the pyrimidinone N(6)-H and the O(17) lone pair of the dihydrobenzofuryl moiety. The spatial interatomic distance for O(17)-N(6), determined as 2.04 Å, favours this H-bridge.