## Powder X-ray diffraction of 3-(4-Morpholinyl)-1-(4-nitrophenyl)-5,6-dihydro-2 (1H)-pyridinone, $C_{15}H_{17}N_3O_4$

Qing Wang, Bin Tang, Pei Xiao Tang, Jia Wei He, and Hui Li

College of Chemical Engineering, Sichuan University, Chengdu 610065, China

(Received 11 June 2015; accepted 6 July 2015)

X-ray powder diffraction data for 3-(4-Morpholinyl)-1-(4-nitrophenyl)-5,6-dihydro-2(1H)-pyridinone,  $C_{15}H_{17}N_3O_4$ , are reported [a = 7.112(1) Å, b = 33.360(2) Å, c = 6.265(1) Å,  $\alpha = 90^\circ$ ,  $\beta = 94.037$  (1)°,  $\gamma = 90^\circ$ , V = 1483.08 Å<sup>3</sup>, Z = 4,  $\rho_{cal} = 1.358$  g cm<sup>-3</sup> and space group  $P2_1$ ]. All measured lines were indexed and are consistent with the  $P2_1$  space group. No detectable impurities were observed. © 2015 International Centre for Diffraction Data. [doi:10.1017/S0885715615000640]

Key words: 3-(4-Morpholinyl)-1-(4-nitrophenyl)-5, 6-dihydro-2(1H)-pyridinone, pharmaceutical intermediate, anticoagulant, apixaban

3-(4-Morpholinyl)-1-(4-nitrophenyl)-5,6-dihydro-2(1H)pyridinone is an intermediate in the synthesis of the anticoagulant, Apixaban (Jiang and Ji, 2013; Zikria and Ansell, 2009). The sample was prepared using 3,3-Dichloro-1-(4-nitrophenyl)-2-piperidinone and was recrystallized in ethanol and dried. The sample was then ground into powder (HPLC  $\ge 98\%$ ,  $\rho = 1.371 \text{ g cm}^{-3}$ ,  $T_{\text{melt}} = 166-168 \text{ °C}$ ) and mounted on a flat zero background plate. X-ray powder diffraction measurement was performed at room temperature using an X'Pert PRO diffractometer (PANalytical Co., Ltd., The Netherlands) with a PIXcel 1D detector and  $CuK\alpha$ radiation (generator setting: 40 kV and 40 mA). The diffraction data were collected over the angular range from  $4^{\circ}$  to  $50^{\circ}2\theta$  with a step size of  $0.013\,13^{\circ}2\theta$  and a counting time of 30 ms step<sup>-1</sup>. The software package Material Studio 8.0 (Accelrys Co., Ltd., CA, USA) was used to process the data in the Analytical & Testing Center (Sichuan University, China). The X-ray powder diffraction pattern was pre-treated by subtracting the background, smoothing, and stripping off the  $K\alpha_2$  component. Automatic indexing results were obtained by DICVOL91 method (Boultif and Louër, 1991). The following figures of merit were achieved:  $F_{20} = 36.0 \ (0.0103, 54) \ (Smith and Snyder, 1979) \ and \ M_{20}$ = 19.1 (de Wolff, 1968). The preliminary cell from indexing was refined using the Pawley method (Pawley, 1981). Pawley refinement results confirmed that the sample crystallizes in the monoclinic space group  $P2_1$  (4), with a = 7.112(1) Å, b = 33.360(2) Å, c = 6.265(1) Å,  $\alpha = 90^{\circ}$ ,  $\beta = 94.037$ (1)°,  $\gamma = 90^{\circ}$ ,  $V = 1483.08 \text{ Å}^3$ , Z = 4,  $\rho_{cal} = 1.358 \text{ g cm}^{-3}$ . Figure 1 shows the powder X-ray diffraction pattern of the compound.

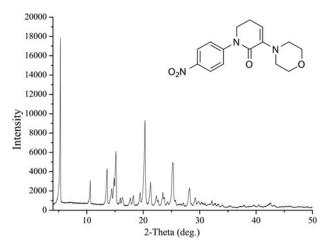


Figure 1. Powder x-ray diffraction pattern of 3-(4-morpholinyl)-1-(4-nitrophenyl)-5,6-dihydro-2(1H)-pyridinone.

## ACKNOWLEDGEMENTS

This work was supported by the Applied Basic Research Project of Sichuan Province (Grant no. 2014JY0042), the Testing Platform Construction of Technology Achievement Transform of Sichuan Province (Grant no. 13CGPT0049), and the National Development and Reform Commission and Education of China (Grant no. 2014BW011).

## SUPPLEMENTARY MATERIALS

For supplementary material for this article, please visit http://dx.doi.org/10.1017/S0885715615000640

Boultif, A. and Louër, D. (1991). "Indexing of powder diffraction patterns for low-symmetry lattices by the successive dichotomy method," J. Appl. Crystallogr. 24, 987–993.

- de Wolff, P. M. (1968). "A simplified criterion for the reliability of a powder pattern," J. Appl. Crystallogr. 1, 108–113.
- Jiang, J. and Ji, Y. (2013). "Alternate synthesis of apixaban (BMS-562247), an inhibitor of blood coagulation factor Xa," Synth. Commun. 43(1), 72–79.

<sup>&</sup>lt;sup>a)</sup>Author to whom correspondence should be addressed. Electronic mail: lihuilab@sina.com Data were submitted via Genie (http://www.icdd.com/ websubmission/launch.html), the ICDD Web Submission Page.

Pawley, G. S. (1981). "Unit-cell refinement from powder diffraction scans," J. Appl. Crystallogr. 14(6), 357–361.

Smith, G. S. and Snyder, R. L. (1979). "FN: a criterion for rating powder diffraction patterns and evaluating the reliability of powder indexing," J. Appl. Crystallogr. 12, 60–65. Zikria, J. and Ansell, J. (2009). "Oral anticoagulation with factor Xa and thrombin inhibitors: is there an alternative to warfarin?," Discov. Med. 8(43), 196–203.