X-ray powder diffraction data for schisanhenol

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Experimental X-ray powder diffraction data, unit-cell parameters and space group for schisanhenol, $C_{23}H_{30}O_6$, are reported [a = 14.6157 Å, b = 12.8801 Å, c = 11.4907 Å, unit-cell volume V = 2163.14 Å³, Z = 4, and space group $P2_12_12_1$]. All of the measured lines were indexed and are consistent with the $P2_12_12_1$ space group. No detectable impurities were observed. © 2013 International Centre for Diffraction Data. [doi:10.1017/S0885715613000778]

Key words: X-ray powder diffraction, schisanhenol, Schisandra chinensis, Rietveld refinement

I. INTRODUCTION

Schisanhenol, systematic name (6*S*, 7*R*,)-2,3,10,11,12pentamethoxy-6,7-dimethyl-5,6,7,8-tetrahydrodibenzo[a,c] [8]annulen-1-ol, is an important lignan of *Schisandra chinensis* (Turcz.) Baill, which was first described by Liu *et al.* (1978). Schisanhenol (Figure 1) has strong biological activities and pharmacological properties, which was shown to be the most active one in inhibiting microsome lipid oxidation (Yu *et al.*, 2004). Schisanhenol absolutely inhibited the preoxidative damages of brain mitochondria and membrane of rats. The swelling and disintegration of brain mitochondria, as well as the reduction of brain fluidity were also restrained by schisanhenol (Xue *et al.*, 1992). In addition, schisanhenol exhibits various beneficial activities including antihepatitis, antitumor, and anti-HIV effects (Chen *et al.*, 2006).

The crystal structure of schisanhenol was solved by Xu *et al.* (1982), which was confirmed by single-crystal measurement.

II. EXPERIMENTAL

A. Sample preparation

The title compound was extracted, isolated, and purified from *S. chinensis*, and the natural sample was characterized by high-performance liquid chromatography (HPLC), as well as by UV, IR, and MS. Then, the pure schisanhenol (>98% purity) was recrystallized in petroleum ether–ethyl acetate (5:1, v/v).

B. Diffraction data collection and reduction

The diffraction pattern for the target compound was collected at room temperature using an X'Pert PRO diffractometer (PANalytical) with an X'celerator detector and Cu $K\alpha_1$ radiation ($\lambda = 1.54056$ Å, generator setting: 40 kV, 30 mA). The diffraction data were collected over the angular range from 5 to 50°2 θ with a step size of 0.01313°2 θ and a counting time of 10.16 ms step⁻¹. Data evaluation was



Figure 1. Structural formula of schisanhenol.



Figure 2. X-ray powder diffraction pattern of schisanhenol, using Cu $K\alpha_1$ radiation ($\lambda = 1.54056$ Å).

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TABLE I. Indexed X-ray powder diffraction data of schisanhenol, $C_{23}H_{30}O_6$. Only the peaks with I_{rel} of 1 or greater are reported [a = 14.6157 Å, b = 12.8801 Å, c = 11.4907 Å, unit-cell volume V = 2163.14 Å³, Z = 4, and space group P2₁2₁2₁]. All of the measured lines were indexed and are consistent with the $P2_12_12_1$ space group. The *d*-values were calculated using CuK α_1 radiation ($\lambda = 1.54056$ Å).

| $2\theta_{\rm obs}(^{\circ})$ | $d_{\rm obs}({\rm \AA})$ | $I_{\rm obs}$ | h | k | l | $2\theta_{\rm cal}(^\circ)$ | $d_{\rm cal}({\rm \AA})$ | $\Delta 2\theta$ |
|-------------------------------|--------------------------|---------------|---|---|---|-----------------------------|--------------------------|------------------|
| 9.1447 | 9.6625 | 4 | 1 | 1 | 0 | 9.1440 | 9.6633 | 0.0007 |
| 9.7487 | 9.0652 | 3 | 1 | 0 | 1 | 9.7832 | 9.0333 | -0.0345 |
| 10.3002 | 8.5811 | 1 | 0 | 1 | 1 | 10.3081 | 8.5745 | -0.0079 |
| 11.9415 | 7.4051 | 100 | 1 | 1 | 1 | 11.9567 | 7.3957 | -0.0152 |
| 12.0728 | 7.3248 | 89 | 2 | 0 | 0 | 12.1009 | 7.3079 | -0.0281 |
| 13.6747 | 6.4702 | 30 | 0 | 2 | 0 | 13.7388 | 6.4401 | -0.0641 |
| 13.8979 | 6.3667 | 20 | 2 | 1 | 0 | 13.9213 | 6.3561 | -0.0234 |
| 14.3312 | 6.1752 | 41 | 2 | 0 | 1 | 14.3518 | 6.1664 | -0.0206 |
| 15.0140 | 5.8959 | 2 | 1 | 2 | 0 | 15.0206 | 5.8933 | -0.0066 |
| 15.7624 | 5.6176 | 14 | 0 | 2 | 1 | 15.7615 | 5.6179 | 0.0009 |
| 15.8937 | 5.5715 | 33 | 2 | 1 | 1 | 15.9212 | 5.5619 | -0.0275 |
| 16.8785 | 5.2486 | 9 | 0 | 1 | 2 | 16.8835 | 5.247 | -0.0050 |
| 16.8916 | 5.2445 | 9 | 1 | 2 | 1 | 16.8936 | 5.2439 | -0.0020 |
| 17.9158 | 4.9469 | 56 | 1 | 1 | 2 | 17.9470 | 4.9384 | -0.0312 |
| 18.3228 | 4.8380 | 3 | 2 | 2 | 0 | 18.3471 | 4.8316 | -0.0243 |
| 19.4651 | 4.5565 | 3 | 3 | 1 | 0 | 19.4640 | 4.5568 | 0.0012 |
| 19.6358 | 4.5173 | 12 | 2 | 0 | 2 | 19.6389 | 4.5166 | -0.0031 |
| 19.7671 | 4.4876 | 27 | 3 | 0 | 1 | 19.7769 | 4.4854 | -0.0098 |
| 19.8722 | 4.4641 | 41 | 2 | 2 | 1 | 19.9182 | 4.4539 | -0.0460 |
| 20.6731 | 4.2929 | 11 | 0 | 3 | 0 | 20.6708 | 4.2934 | 0.0023 |
| 20.8044 | 4.2661 | 12 | 2 | 1 | 2 | 20.8238 | 4.2622 | -0.0194 |
| 20.9489 | 4.2370 | 4 | 3 | 1 | 1 | 20.9546 | 4.2359 | -0.0057 |
| 21.5397 | 4.1221 | 12 | 1 | 3 | 0 | 21.5547 | 4.1193 | -0.0150 |
| 21.5791 | 4.1147 | 9 | 1 | 2 | 2 | 21.5833 | 4.1139 | -0.0042 |
| 22.8659 | 3.8860 | 19 | 3 | 2 | 0 | 22.8693 | 3.8854 | -0.0034 |
| 22.9184 | 3.8772 | 11 | 1 | 3 | 1 | 22.9153 | 3.8/// | 0.0031 |
| 23.9294 | 3./156 | 25 | 3 | 0 | 2 | 23.9282 | 3./158 | 0.0013 |
| 23.9951 | 3.7050 | 33 21 | 1 | 0 | 3 | 23.9983 | 3.7051 | -0.0032 |
| 24.0213 | 3.7010 | 51 | 2 | 3 | 0 | 24.0200 | 3.7018 | 0.0013 |
| 24.2185 | 3.0/19 | 4 | 0 | 1 | 3 | 24.2220 | 3.0/13 | -0.0043 |
| 24.3303 | 3.0344 | 16 | 4 | 0 | 0 | 24.5597 | 3.0339 | -0.0032 |
| 24.0079 | 3.5740 | 10 | 3 | 1 | 2 | 24.9194 | 3.5702 | -0.0314 |
| 24.9799 | 3.5017 | 1 | 1 | 1 | 5 | 24.9609 | 3.5007 | -0.0071 |
| 25.2425 | 3.1842 | 1 | 4 | 0 | 1 | 25.2551 | 3.4821 | -0.0120 |
| 25.5445 | 3 4301 | 3 | 4 | 3 | 2 | 25.5004 | 3 4302 | -0.0139 |
| 25.8858 | 3 3967 | 4 | 2 | 0 | 2 | 25.8847 | 3 3025 | _0.0311 |
| 26.2141 | 3 3620 | 2 | 4 | 1 | 1 | 26.4946 | 3 3614 | -0.0048 |
| 26.5949 | 3 3490 | 2 | 1 | 3 | 2 | 26.4940 | 3 3478 | -0.0093 |
| 27.6716 | 3 2210 | 6 | 0 | 4 | 0 | 27 6807 | 3 22 | -0.0091 |
| 27.6847 | 3,2195 | 5 | 3 | 2 | 2 | 27.6939 | 3,2185 | -0.0092 |
| 27.7503 | 3,2121 | 1 | 1 | 2 | 3 | 27.7554 | 3,2115 | -0.0051 |
| 28.3675 | 3.1436 | 2 | 1 | 4 | 0 | 28.3581 | 3.1446 | 0.0093 |
| 28.7482 | 3.1028 | - 9 | 3 | 3 | 1 | 28.7606 | 3.1015 | -0.0124 |
| 29.0634 | 3.0699 | 5 | 4 | 2 | 1 | 29.1301 | 3.063 | -0.0667 |
| 29.4442 | 3.0310 | 3 | 1 | 4 | 1 | 29.4237 | 3.0331 | 0.0205 |
| 29.6411 | 3.0113 | 7 | 3 | 0 | 3 | 29.6436 | 3.0111 | -0.0025 |
| 31.0986 | 2.8735 | 2 | 0 | 0 | 4 | 31.1070 | 2.8727 | -0.0084 |
| 31.2824 | 2.8570 | 3 | 0 | 3 | 3 | 31.2688 | 2.8582 | 0.0136 |
| 31.3218 | 2.8535 | 2 | 2 | 4 | 1 | 31.3126 | 2.8543 | 0.0091 |
| 31.7157 | 2.8189 | 1 | 1 | 0 | 4 | 31.7185 | 2.8187 | -0.0028 |
| 31.8733 | 2.8054 | 2 | 1 | 3 | 3 | 31.8775 | 2.805 | -0.0042 |
| 32.1359 | 2.7830 | 5 | 4 | 3 | 0 | 32.1410 | 2.7826 | -0.0052 |
| 32.4247 | 2.7589 | 4 | 1 | 4 | 2 | 32.4295 | 2.7585 | -0.0048 |
| 32.4772 | 2.7546 | 2 | 1 | 1 | 4 | 32.4889 | 2.7536 | -0.0116 |
| 32.8055 | 2.7277 | 2 | 3 | 2 | 3 | 32.8060 | 2.7277 | -0.0005 |
| 34.1448 | 2.6237 | 1 | 0 | 2 | 4 | 34.1481 | 2.6235 | -0.0034 |
| 34.5650 | 2.5928 | 1 | 5 | 2 | 1 | 34.5610 | 2.5931 | 0.0040 |
| 36.9153 | 2.4329 | 1 | 3 | 4 | 2 | 36.9082 | 2.4334 | 0.0071 |
| 40.0140 | 2.2514 | 1 | 4 | 3 | 3 | 40.0176 | 2.2512 | -0.0035 |
| 41.2614 | 2.1862 | 1 | 5 | 2 | 3 | 41.2686 | 2.1858 | -0.0072 |
| 42.5876 | 2.1211 | 1 | 1 | 4 | 4 | 42.5919 | 2.1209 | -0.0043 |
| 44.0319 | 2.0548 | 4 | 7 | 0 | 1 | 44.0437 | 2.0543 | -0.0118 |
| 44.0975 | 2.0519 | 4 | 3 | 1 | 5 | 44.0956 | 2.052 | 0.0019 |
| 49.0477 | 1.8558 | 1 | 1 | 4 | 5 | 49.0554 | 1.8555 | -0.0078 |

performed using the software package Material Studio 4.2 (Accelrys Co., Ltd. USA).

Indexing was conducted using peak positions obtained from the powder diffraction profiles by X-cell method then the indexing result was refined with the type of Pawley (Pan *et al.*, 2012). MC/SA search algorithm in Powder Solve package (Engel *et al.*, 1999) was used to constantly adjust the conformation, position, and orientation of the trial model in the unit cell of schisanhenol, and then the result of Powder Solve was refined by Rietveld refinement techniques. In the Rietveld refinement (Young, 1993), variables defining the structural model and the powder diffraction profile were adjusted by least squares methods in order to obtain an optimal fit between the experimental pattern and the calculated pattern. After Rietveld refinement, the final R_{wp} was converged at 7.60%.

III. RESULTS

The experimental powder diffraction pattern is depicted in Figure 2. Indexing results show that schisanhenol is orthorhombic with space group $P2_12_12_1$ and unit-cell parameters: a = 14.6157 Å, b = 12.8801 Å, c = 11.4907 Å, unit-cell volume V = 2163.14 Å³, Z = 4, and space group $P2_12_12_1$ (Table I). All of the lines were indexed and are consistent with the $P2_12_12_1$ space group.

After Rietveld refinement, the structure of schisanhenol was solved. At the same time, single-crystal data were collected on an Oxford Diffraction Xcalibur Nova system with Mo $K\alpha$ radiation ($\lambda = 0.71073$ Å) at room temperature and θ from 3.05° to 28.77°, in that way to verify whether the powder data are accurate. A comparison of the unit-cell parameters from powder data and from single-crystal data reported by Xu *et al.* (1982) shows significant consistency, which indicates that X-ray powder diffraction as a means of crystal structure solution is credible.

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