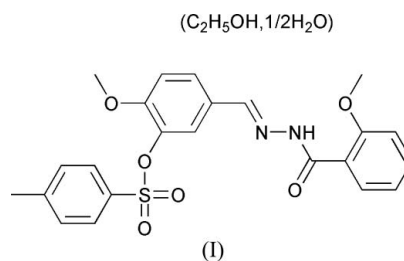


**(E)-2-Methoxy-N'-[4-methoxy-3-(4-methylphenyl-sulfonyloxy)benzylidene]benzohydrazide ethanol solvate hemihydrate****Xin Chen\* and Ming Yu**College of Sciences, Tianjin University of  
Science and Technology, Tianjin 300222,  
People's Republic of ChinaCorrespondence e-mail:  
chen\_xin9999@163.com**Key indicators**Single-crystal X-ray study  
 $T = 294$  K  
Mean  $\sigma(\text{C}-\text{C}) = 0.004$  Å  
Disorder in solvent or counterion  
R factor = 0.060  
wR factor = 0.189  
Data-to-parameter ratio = 13.7For details of how these key indicators were  
automatically derived from the article, see  
<http://journals.iucr.org/e>.

In the title compound,  $\text{C}_{23}\text{H}_{22}\text{N}_2\text{O}_6\text{S}\cdot\text{C}_2\text{H}_6\text{O}\cdot 0.5\text{H}_2\text{O}$ , the isovanillin group makes dihedral angles of  $50.57$  (8) and  $8.89$  (11) $^\circ$ , respectively, with the methyl- and methoxy-substituted benzene rings. Intramolecular  $\text{N}-\text{H}\cdots\text{O}$  hydrogen bonds help to stabilize the molecular conformation, while intermolecular  $\text{O}-\text{H}\cdots\text{O}$  and  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bonds link adjacent molecules, forming an infinite network.

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Metal complexes based on Schiff bases have attracted much attention because of their biological activity (Kahwa *et al.*, 1986; Klayman *et al.*, 1979). Many Schiff base derivatives have been synthesized and employed to develop protein and enzyme mimics (Santos *et al.*, 2001), such as models to mimic hydrolase in the hydrolysis of *p*-nitrophenyl picolinate (Li *et al.*, 2005). Structural information is useful when investigating the coordination properties of Schiff bases functioning as ligands. We report here the synthesis and molecular structure of the title Schiff base compound, (I) (Fig. 1).



In (I), bond lengths and angles are within normal ranges (Allen *et al.*, 1987). The isovanillin group (C8–C13/C15/O3/O5) is nearly planar, with an r.m.s. deviation for fitted atoms of  $0.0378$  Å. This plane makes dihedral angles of  $50.57$  (8) and  $8.89$  (11) $^\circ$ , respectively, with the C1–C6 and C16–C21 benzene rings. The dihedral angle between these two benzene rings is  $53.47$  (10) $^\circ$ .

An intramolecular hydrogen bond links the NH group to O6, thereby influencing the molecular conformation. There are two  $\text{O}-\text{H}\cdots\text{O}$  intermolecular hydrogen bonds and one weak non-classical intermolecular  $\text{C}-\text{H}\cdots\text{O}$  hydrogen bond linking the main molecule and the solvent molecules (Table 1 and Fig. 2). These hydrogen bonds link molecules into an infinite network.

**Experimental**

An anhydrous ethanol solution (50 ml) of 5-formyl-2-methoxyphenyl 4-methylbenzenesulfonate (3.06 g, 10 mmol) was added to an anhy-

drous ethanol solution (50 ml) of 2-methoxybenzohydrazide (1.66 g, 10 mmol) and the mixture stirred at 350 K for 3 h under N<sub>2</sub>, giving a white precipitate. The product was isolated, recrystallized from ethanol, and then dried in a vacuum to give pure compound (I) in 81% yield. Colourless single crystals of (I) suitable for X-ray analysis were obtained by slow evaporation of a 95% ethanol–water solution.

Crystal data

C<sub>23</sub>H<sub>22</sub>N<sub>2</sub>O<sub>6</sub>S·C<sub>2</sub>H<sub>6</sub>O·0.5H<sub>2</sub>O  
*M<sub>r</sub>* = 509.57  
 Triclinic, *P* $\bar{1}$   
*a* = 7.9106 (16) Å  
*b* = 8.6824 (17) Å  
*c* = 19.266 (4) Å  
 $\alpha$  = 87.61 (3)°  
 $\beta$  = 80.12 (3)°  
 $\gamma$  = 81.39 (3)°  
*V* = 1288.8 (5) Å<sup>3</sup>  
*Z* = 2  
*D<sub>x</sub>* = 1.313 Mg m<sup>-3</sup>  
 Mo *K*α radiation  
 $\mu$  = 0.17 mm<sup>-1</sup>  
*T* = 294 (2) K  
 Block, colourless  
 0.14 × 0.12 × 0.10 mm

Data collection

Bruker SMART APEX CCD area-detector diffractometer  
 $\varphi$  and  $\omega$  scans  
 Absorption correction: multi-scan (SADABS; Sheldrick, 1996)  
*T<sub>min</sub>* = 0.954, *T<sub>max</sub>* = 0.983  
 7839 measured reflections  
 4492 independent reflections  
 3294 reflections with *I* > 2σ(*I*)  
*R<sub>int</sub>* = 0.027  
 $\theta_{max}$  = 25.0°

Refinement

Refinement on *F*<sup>2</sup>  
*R*[*F*<sup>2</sup> > 2σ(*F*<sup>2</sup>)] = 0.060  
*wR*(*F*<sup>2</sup>) = 0.189  
*S* = 1.07  
 4492 reflections  
 329 parameters  
 H-atom parameters constrained  
 $w = 1/[\sigma^2(F_o^2) + (0.1233P)^2 + 0.0157P]$   
 where  $P = (F_o^2 + 2F_c^2)/3$   
 $(\Delta/\sigma)_{max} = 0.020$   
 $\Delta\rho_{max} = 0.40 \text{ e \AA}^{-3}$   
 $\Delta\rho_{min} = -0.30 \text{ e \AA}^{-3}$

Table 1

Hydrogen-bond geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
N2—H2...O6	0.86	1.97	2.641 (3)	134
O8—H8B...O4 <sup>i</sup>	0.87	2.37	2.938 (11)	124
O7—H7...O4 <sup>i</sup>	0.82	2.39	2.814 (4)	113
C15—H15...O8 <sup>ii</sup>	0.93	2.49	3.414 (8)	172
C14—H14C...O1 <sup>iii</sup>	0.96	2.42	3.223 (4)	141

Symmetry codes: (i) *x* + 1, *y* − 1, *z*; (ii) *x* − 1, *y*, *z*; (iii) *x*, *y* − 1, *z*.

The H atoms were included in calculated positions and refined using a riding-model approximation. Constrained C—H and N—H bond lengths and isotropic U parameters: 0.93 Å and *U<sub>iso</sub>*(H) = 1.2*U<sub>eq</sub>*(C) for *Csp*<sup>2</sup>, 0.97 Å and *U<sub>iso</sub>*(H) = 1.2*U<sub>eq</sub>*(C) for methylene, 0.96 Å and *U<sub>iso</sub>*(H) = 1.5*U<sub>eq</sub>*(C) for methyl, 0.85 Å and *U<sub>iso</sub>*(H) = 1.2*U<sub>eq</sub>*(O) for water, 0.82 Å and *U<sub>iso</sub>*(H) = 1.5*U<sub>eq</sub>*(O) for hydroxy, and 0.86 Å and *U<sub>iso</sub>*(H) = 1.2*U<sub>eq</sub>*(N) for imino H atoms. The water molecule (O8, H8A and H8B) was refined using a disorder model, with occupancy factors constrained to be 0.50.

Data collection: SMART (Bruker, 1999); cell refinement: SAINT (Bruker, 1999); data reduction: SAINT; program(s) used to solve structure: SHELXS97 (Sheldrick, 1997a); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997a); molecular graphics: SHELXTL (Sheldrick, 1997b); software used to prepare material for publication: SHELXTL.

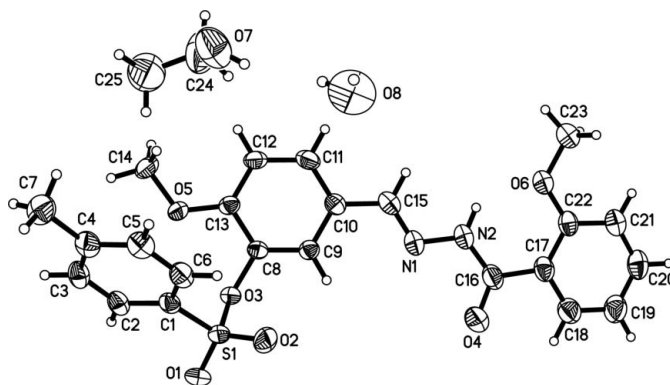


Figure 1

The asymmetric unit of (I), with displacement ellipsoids for non-H atoms drawn at the 30% probability level.

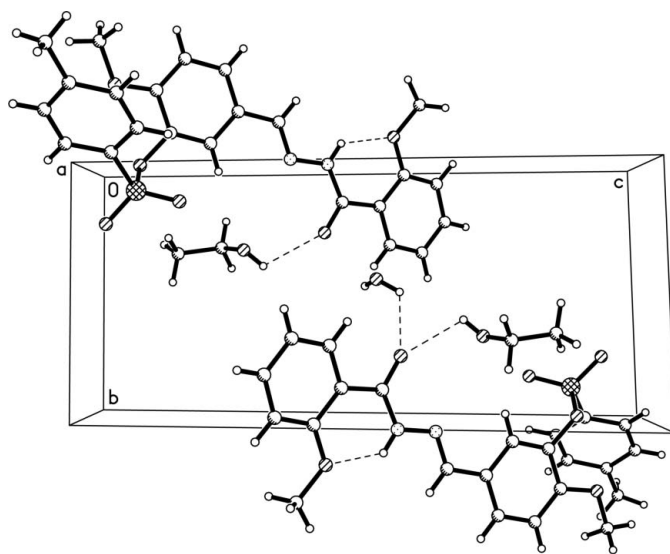


Figure 2

A packing diagram for (I), with hydrogen bonds shown as dashed lines.

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