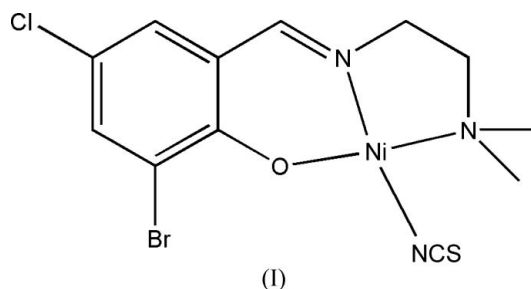


{2-Bromo-4-chloro-6-[2-(dimethylamino)-ethyliminomethyl]phenolato}thiocyanato-nickel(II)**Xiao-Fan Zhao**College of Chemistry & Chemical Engineering,
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Republic of ChinaCorrespondence e-mail:
xiaofan_zhao@126.com**Key indicators**Single-crystal X-ray study
 $T = 298$ K
Mean $\sigma(\text{C}-\text{C}) = 0.009$ Å
Disorder in main residue
 R factor = 0.043
 wR factor = 0.109
Data-to-parameter ratio = 14.1For details of how these key indicators were
automatically derived from the article, see
<http://journals.iucr.org/e>.

In the title mononuclear compound, $[\text{Ni}(\text{C}_{11}\text{H}_{13}\text{BrClN}_2\text{O})(\text{NCS})]$, the Ni^{II} atom is four-coordinated in a square-planar geometry by one O and two N atoms of the Schiff base ligand, and by one N atom of the thiocyanate anion. The molecule possesses crystallographic mirror symmetry.

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Transition metal complexes play an important role in both bioinorganic chemistry and coordination chemistry (Chun & Bernal, 2001; Jackson *et al.*, 2003; Nakashima *et al.*, 2003). Recently, the author has reported the crystal structure of a mononuclear Schiff base cobalt(III) complex (Zhao, 2006). As part of a further study of the crystal structures of such complexes, the title mononuclear nickel(II) complex, (I), is reported here.



The molecule of compound (I) possesses crystallographic mirror symmetry, with all atoms except atoms H8A, H8B (both attached to C8), C9, H9A, H9B and the methyl groups lying on the crystallographic mirror plane. The Ni^{II} atom is four-coordinated in a square-planar geometry by the Schiff base ligand and the thiocyanate ligand (Fig. 1). The Schiff base acts as a tridentate ligand through the phenolate O atom, the imine N atom and the amine N atom. The two *trans* angles subtended at the Ni^{II} atom deviate slightly from 180° (Table 1), and the other angles around the Ni^{II} atom deviate slightly from the ideal value (90°) for a square-planar coordination, ranging from $83.9(2)$ to $92.5(2)^\circ$. The Ni—O and Ni—N bond lengths are typical and comparable to those in similar Schiff base–nickel(II) complexes (Gomes *et al.*, 1999; Jircitano *et al.*, 1990; Kabak *et al.*, 1997).

Experimental

3-Bromo-5-chloro-2-hydroxybenzaldehyde (0.2 mmol, 47.0 mg), *N,N*-dimethylethane-1,2-diamine (0.2 mmol, 17.6 mg), ammonium thiocyanate (0.2 mmol, 15.2 mg) and nickel nitrate hexahydrate (0.2 mmol, 58.2 mg) were dissolved in methanol (30 ml) with stirring.

The mixture was stirred for 30 min at room temperature to give a red solution which was allowed to stand in air for one week, yielding red block-shaped crystals of (I).

Crystal data

[Ni(C₁₁H₁₃BrClN₂O)(NCS)]

M_r = 421.38

Orthorhombic, *Pnma*

a = 19.338 (1) Å

b = 6.994 (1) Å

c = 11.288 (2) Å

V = 1526.7 (4) Å³

Z = 4

D_x = 1.833 Mg m⁻³

Mo *K*α radiation

μ = 4.20 mm⁻¹

T = 298 (2) K

Block, red

0.17 × 0.15 × 0.12 mm

Data collection

Bruker SMART CCD area-detector diffractometer

ω scans

Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)

T_{min} = 0.536, *T_{max}* = 0.633

11929 measured reflections

1717 independent reflections

1427 reflections with *I* > 2σ(*I*)

R_{int} = 0.047

θ_{max} = 26.5°

Refinement

Refinement on *F*²

R[*F*² > 2σ(*F*²)] = 0.043

wR(*F*²) = 0.110

S = 1.05

1717 reflections

122 parameters

H-atom parameters constrained

w = 1/[σ²(*F_o*²) + (0.0414*P*)² + 3.2149*P*]

where *P* = (*F_o*² + 2*F_c*²)/3

(Δ/σ)_{max} = 0.001

Δρ_{max} = 0.82 e Å⁻³

Δρ_{min} = -1.00 e Å⁻³

Table 1

Selected geometric parameters (Å, °).

Ni1—O1	1.915 (5)	Ni1—N1	1.934 (6)
Ni1—N3	1.925 (6)	Ni1—N2	2.068 (6)
O1—Ni1—N3	91.9 (2)	O1—Ni1—N2	176.4 (2)
O1—Ni1—N1	92.5 (2)	N3—Ni1—N2	91.7 (2)
N3—Ni1—N1	175.6 (2)	N1—Ni1—N2	83.9 (2)

H atoms were placed in geometrically idealized positions and constrained to ride on their parent atoms, with C—H = 0.93 or 0.96 Å and *U*_{iso}(H) = 1.2*U*_{eq}(C) or 1.5*U*_{eq}(methyl C). C9, H9A and H9B are disordered about the mirror plane, with site occupancy factors of 0.5.

Data collection: *SMART* (Bruker, 2002); cell refinement: *SAIN*T (Bruker, 2002); data reduction: *SAIN*T; program(s) used to solve structure: *SHELXS97* (Sheldrick, 1997a); program(s) used to refine

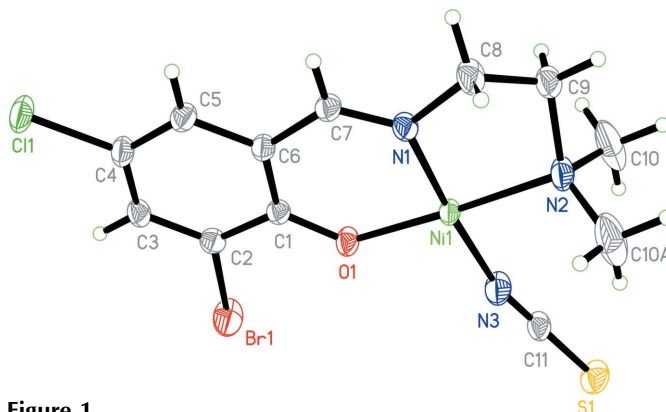


Figure 1

The molecular structure of (I), showing the atom-numbering scheme. Displacement ellipsoids are drawn at the 30% probability level. Only one disorder component is shown. [Symmetry code: (A) *x*, $\frac{1}{2}$ - *y*, *z*.]

structure: *SHELXL97* (Sheldrick, 1997a); molecular graphics: *SHELXTL* (Sheldrick, 1997b); software used to prepare material for publication: *SHELXL97*.

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