

trans-Bis[4-amino-N-(pyrimidin-2-yl)- κ N]benzenesulfonamido- κ N]bis(N,N-dimethylformamide- κ O)cobalt(II)

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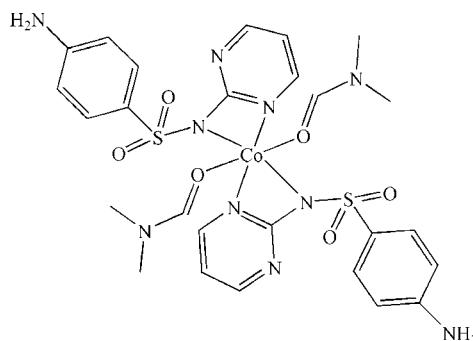
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(C-C) = 0.004$ Å; disorder in main residue; R factor = 0.040; wR factor = 0.095; data-to-parameter ratio = 15.7.

The title complex, $[Co(C_{10}H_9N_4O_2S)_2(C_3H_7NO)_2]$, lies across an inversion center. The Co^{II} atom is coordinated in a slightly distorted octahedral geometry by four N atoms from two bidentate 4-amino-N-(pyrimidin-2-yl)benzenesulfonamide (sulfadiazine) anions and two O atoms from two dimethylformamide (DMF) ligands. The dihedral angle between the benzene and pyrimidine rings is 82.37 (13)°. A three-dimensional network is generated by N–H···O hydrogen bonds between the amino groups and of the sulfonamide O atoms of neighbouring molecules. The DMF ligand is disordered over two sets of sites in a 0.559 (4):0.441 (4) ratio.

Related literature

For background to sulfonamides, see: Connor (1998). For background to metal complexes of sulfadiazine, see: Wang *et al.* (2009, 2010); Ajibade *et al.* (2006); Hossain *et al.* (2011); Tommasino *et al.* (2011); Ghosh *et al.* (2011).



Experimental

Crystal data

$[Co(C_{10}H_9N_4O_2S)_2(C_3H_7NO)_2]$	$V = 1614.67$ (16) Å ³
$M_r = 703.67$	$Z = 2$
Monoclinic, $P2_1/c$	Mo $K\alpha$ radiation
$a = 8.9008$ (6) Å	$\mu = 0.72$ mm ⁻¹
$b = 11.2078$ (6) Å	$T = 298$ K
$c = 16.5565$ (9) Å	$0.28 \times 0.25 \times 0.20$ mm
$\beta = 102.147$ (6)°	

Data collection

Bruker SMART CCD diffractometer	13220 measured reflections
Absorption correction: multi-scan (<i>SADABS</i> ; Bruker, 2001)	3495 independent reflections
$T_{min} = 0.815$, $T_{max} = 0.869$	2989 reflections with $I > 2\sigma(I)$
	$R_{int} = 0.028$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.040$	36 restraints
$wR(F^2) = 0.095$	H-atom parameters constrained
$S = 0.99$	$\Delta\rho_{\text{max}} = 0.33$ e Å ⁻³
3495 reflections	$\Delta\rho_{\text{min}} = -0.35$ e Å ⁻³
222 parameters	

Table 1
Selected bond lengths (Å).

Co1—O1	2.065 (19)	Co1—N2	2.1460 (18)
Co1—N1	2.121 (2)		

Table 2
Hydrogen-bond geometry (Å, °).

D—H···A	D—H	H···A	D···A	D—H···A
N4—H4B···O3 ⁱ	0.86	2.31	3.112 (3)	155
N4—H4C···O3 ⁱⁱ	0.86	2.27	2.951 (3)	136

Symmetry codes: (i) $x, -y + \frac{3}{2}, z + \frac{1}{2}$; (ii) $-x + 1, y + \frac{1}{2}, -z + \frac{1}{2}$.

Data collection: *SMART* (Bruker, 2001); cell refinement: *SAINT* (Bruker, 2001); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *SHELXTL* (Sheldrick, 2008); software used to prepare material for publication: *SHELXTL*.

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: WM2693).

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supporting information

Acta Cryst. (2012). E68, m1398–m1399 [doi:10.1107/S160053681204336X]

trans-Bis[4-amino-N-(pyrimidin-2-yl-κN)benzenesulfonamido-κN]bis(N,N-dimethylformamide-κO)cobalt(II)

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S1. Comment

Sulfonamides are among the most widely used antibacterial agents (Connor, 1998), because of their low cost, low toxicity, and excellent activity against bacterial diseases. Sulfadiazine, or 4-amino-*N*-pyrimidin-2-yl-benzenesulfonamide, is a sulfanilamide antibiotic and its metal complexes have been studied previously (Wang *et al.* 2009, 2010; Ajibade *et al.*, 2006; Hossain *et al.*, 2011; Tommasino *et al.*, 2011; Ghosh *et al.*, 2011). The crystal structure of the title compound, $[\text{Co}(\text{C}_{10}\text{H}_9\text{N}_4\text{O}_2\text{S})_2(\text{C}_3\text{H}_7\text{NO})_2]$, a cobalt sulfadiazine complex with additional dimethylformamide ligands, (I), is presented herein.

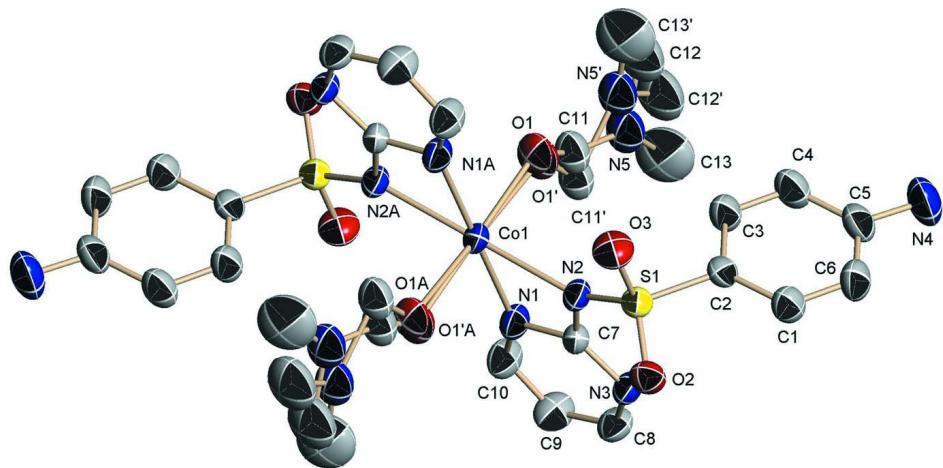
The molecular structure of (I) is shown in Fig. 1. The Co^{II} ion lies on a centre of inversion and is coordinated by four N atoms from two symmetry-related 4-amino-*N*-pyrimidin-2-yl-benzenesulfonamide anions and two O atoms from two dimethylformamide ligands, forming a slightly distorted octahedral N_4O_2 geometry (Table 1). The dihedral angle between the phenyl and pyrimidine rings is $82.37\ (13)\ ^\circ$. A three-dimensional hydrogen-bonded network is generated by N—H \cdots O interactions (Table 2; Fig.2).

S2. Experimental

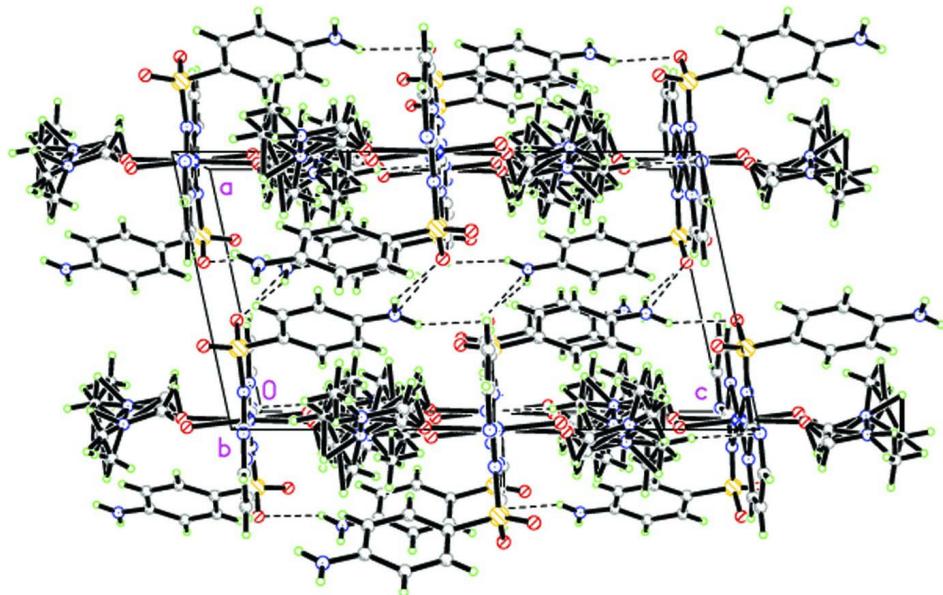
Complex (I) was prepared from a mixture of sulfadiazine (1 mmol, 0.250 g), $\text{Co}(\text{NO}_3)_2 \cdot 6\text{H}_2\text{O}$ (0.5 mmol, 0.145 g), triethylamine (0.5 ml) and *N,N*-dimethylformamide (8 ml) sealed in a 15 ml teflon-lined stainless steel bomb, and kept at 373 K for 96 h under autogenous pressure. After the reaction was slowly cooled to room temperature, red block-like crystals were obtained (yield: 72% based on cobalt). Anal./calc. for $\text{C}_{26}\text{H}_{32}\text{N}_{10}\text{CoO}_6\text{S}_2$ (%): C 44.38; H 4.58; N 19.90. Found(%): C 44.34; H 4.61; N 19.95.

S3. Refinement

H atoms were positioned geometrically and refined with a riding model, with distances 0.86 Å (N—H), 0.96 Å (CH_3) or 0.93 Å (aromatic ring), and with $U_{\text{iso}}(\text{H}) = 1.2 U_{\text{eq}}$ (aromatic ring, N—H) or $U_{\text{iso}}(\text{H}) = 1.5 U_{\text{eq}}(\text{CH}_3)$. The DMF ligand is positionally disordered over two sets of sites in a 0.559 (4):0.441 (4) ratio.

**Figure 1**

The molecular structure of (I), showing 30% probability displacement ellipsoids. [Symmetry code (*a*): $-x, 1 - y, -z$.] Both parts of the disordered DMF ligand are shown.

**Figure 2**

Packing diagram of (I) viewed down [010]. Dashed lines denote N—H···O hydrogen bonds.

trans-Bis[4-amino-*N*-(pyrimidin-2-yl- κ *N*)benzenesulfonamido- κ *N*]bis(*N,N*- dimethylformamide- κ *O*)cobalt(II)

Crystal data



$M_r = 703.67$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 8.9008$ (6) Å

$b = 11.2078$ (6) Å

$c = 16.5565$ (9) Å

$\beta = 102.147$ (6)°

$V = 1614.67$ (16) Å³

$Z = 2$

$F(000) = 730$

$D_x = 1.447$ Mg m⁻³

Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å

Cell parameters from 4388 reflections

$\theta = 3.0\text{--}28.7^\circ$

$\mu = 0.72$ mm⁻¹

$T = 298\text{ K}$
Block, red

$0.28 \times 0.25 \times 0.20\text{ mm}$

Data collection

Bruker SMART CCD
diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
Detector resolution: 0 pixels mm^{-1}
 ϕ and ω scans
Absorption correction: multi-scan
(*SADABS*; Bruker, 2001)
 $T_{\min} = 0.815$, $T_{\max} = 0.869$

13220 measured reflections
3495 independent reflections
2989 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.028$
 $\theta_{\max} = 27.0^\circ$, $\theta_{\min} = 3.0^\circ$
 $h = -11 \rightarrow 11$
 $k = -14 \rightarrow 13$
 $l = -21 \rightarrow 19$

Refinement

Refinement on F^2
Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.040$
 $wR(F^2) = 0.095$
 $S = 0.99$
3495 reflections
222 parameters
36 restraints
Primary atom site location: structure-invariant
direct methods
Secondary atom site location: difference Fourier
map

Hydrogen site location: inferred from
neighbouring sites
H-atom parameters constrained
 $w = 1/[\sigma^2(F_o^2) + (0.0311P)^2 + 1.5416P]$
where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.33\text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.35\text{ e } \text{\AA}^{-3}$
Extinction correction: *SHELXL97* (Sheldrick,
2008), $F_c^* = kFc[1 + 0.001xFc^2\lambda^3/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0012 (1)

Special details

Geometry. All e.s.d.'s (except the e.s.d. in the dihedral angle between two l.s. planes) are estimated using the full covariance matrix. The cell e.s.d.'s are taken into account individually in the estimation of e.s.d.'s in distances, angles and torsion angles; correlations between e.s.d.'s in cell parameters are only used when they are defined by crystal symmetry. An approximate (isotropic) treatment of cell e.s.d.'s is used for estimating e.s.d.'s involving l.s. planes.

Refinement. Refinement of F^2 against ALL reflections. The weighted R -factor wR and goodness of fit S are based on F^2 , conventional R -factors R are based on F , with F set to zero for negative F^2 . The threshold expression of $F^2 > \sigma(F^2)$ is used only for calculating R -factors(gt) etc. and is not relevant to the choice of reflections for refinement. R -factors based on F^2 are statistically about twice as large as those based on F , and R -factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters (\AA^2)

	x	y	z	$U_{\text{iso}}^* / U_{\text{eq}}$	Occ. (<1)
C1	0.4095 (3)	0.9002 (2)	0.13724 (16)	0.0476 (6)	
H1A	0.4447	0.9344	0.0935	0.057*	
C2	0.3251 (2)	0.7957 (2)	0.12498 (14)	0.0363 (5)	
C3	0.2761 (3)	0.7454 (3)	0.19139 (15)	0.0507 (6)	
H3A	0.2212	0.6742	0.1843	0.061*	
C4	0.3079 (3)	0.7994 (3)	0.26800 (16)	0.0572 (7)	
H4A	0.2744	0.7640	0.3119	0.069*	
C5	0.3892 (3)	0.9061 (3)	0.28047 (16)	0.0508 (7)	
C6	0.4421 (3)	0.9543 (3)	0.21390 (17)	0.0568 (7)	
H6A	0.5002	1.0240	0.2213	0.068*	
C7	-0.0153 (2)	0.73277 (19)	0.00525 (13)	0.0319 (5)	
C8	-0.1683 (3)	0.8945 (2)	-0.00174 (16)	0.0438 (6)	

H8A	-0.1808	0.9767	0.0005	0.053*	
C9	-0.2982 (3)	0.8242 (2)	-0.01745 (18)	0.0513 (7)	
H9A	-0.3961	0.8573	-0.0260	0.062*	
C10	-0.2766 (3)	0.7031 (2)	-0.01992 (17)	0.0497 (6)	
H10A	-0.3616	0.6528	-0.0293	0.060*	
Co1	0.0000	0.5000	0.0000	0.03466 (14)	
N1	-0.1368 (2)	0.65637 (17)	-0.00916 (13)	0.0402 (5)	
N2	0.1174 (2)	0.66834 (16)	0.01562 (12)	0.0350 (4)	
N3	-0.0252 (2)	0.85105 (17)	0.01044 (12)	0.0372 (4)	
N4	0.4124 (3)	0.9634 (3)	0.35511 (15)	0.0724 (8)	
H4B	0.3765	0.9335	0.3950	0.087*	
H4C	0.4629	1.0294	0.3621	0.087*	
O2	0.29575 (19)	0.81638 (15)	-0.03312 (10)	0.0437 (4)	
O3	0.38724 (19)	0.62601 (16)	0.02969 (12)	0.0519 (5)	
S1	0.28414 (6)	0.72636 (5)	0.02742 (3)	0.03470 (15)	
O1	0.0274 (19)	0.4809 (16)	0.1263 (12)	0.053 (2)	0.559 (4)
C11	-0.0332 (9)	0.5365 (7)	0.1739 (6)	0.0563 (17)	0.559 (4)
H11A	-0.1225	0.5630	0.1388	0.084*	0.559 (4)
C13	-0.1386 (10)	0.6711 (8)	0.2810 (5)	0.107 (2)	0.559 (4)
H13A	-0.1134	0.6764	0.3402	0.160*	0.559 (4)
H13B	-0.1218	0.7471	0.2577	0.160*	0.559 (4)
H13C	-0.2446	0.6489	0.2632	0.160*	0.559 (4)
C12	0.0603 (12)	0.5055 (8)	0.3034 (5)	0.098 (2)	0.559 (4)
H12A	0.1286	0.4397	0.3204	0.147*	0.559 (4)
H12B	0.1008	0.5758	0.3334	0.147*	0.559 (4)
H12C	-0.0388	0.4876	0.3145	0.147*	0.559 (4)
N5	-0.0499 (7)	0.5829 (5)	0.2463 (3)	0.0665 (13)	0.559 (4)
O1'	-0.009 (3)	0.501 (2)	0.1318 (16)	0.053 (2)	0.441 (4)
C11'	-0.0812 (12)	0.5748 (10)	0.1647 (7)	0.0563 (17)	0.441 (4)
H11B	-0.1748	0.6128	0.1375	0.084*	0.441 (4)
C13'	0.1448 (14)	0.4546 (11)	0.2886 (6)	0.107 (2)	0.441 (4)
H13D	0.1783	0.4035	0.2493	0.160*	0.441 (4)
H13E	0.2274	0.5065	0.3134	0.160*	0.441 (4)
H13F	0.1146	0.4068	0.3306	0.160*	0.441 (4)
C12'	-0.0588 (13)	0.5888 (10)	0.3139 (5)	0.098 (2)	0.441 (4)
H12D	-0.0379	0.5933	0.3731	0.147*	0.441 (4)
H12E	-0.0382	0.6648	0.2916	0.147*	0.441 (4)
H12F	-0.1647	0.5681	0.2937	0.147*	0.441 (4)
N5'	0.0174 (10)	0.5244 (7)	0.2573 (5)	0.0665 (13)	0.441 (4)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.0493 (14)	0.0501 (15)	0.0422 (13)	-0.0135 (12)	0.0066 (11)	0.0020 (12)
C2	0.0331 (11)	0.0383 (12)	0.0342 (11)	-0.0024 (9)	-0.0004 (9)	0.0011 (10)
C3	0.0565 (15)	0.0515 (16)	0.0409 (13)	-0.0211 (13)	0.0031 (11)	0.0010 (12)
C4	0.0653 (17)	0.0698 (19)	0.0340 (13)	-0.0220 (15)	0.0047 (12)	0.0016 (13)
C5	0.0473 (14)	0.0605 (17)	0.0386 (13)	-0.0069 (12)	-0.0049 (11)	-0.0041 (12)

C6	0.0605 (17)	0.0523 (16)	0.0527 (16)	-0.0222 (14)	0.0006 (13)	-0.0068 (13)
C7	0.0351 (11)	0.0280 (11)	0.0313 (11)	-0.0031 (9)	0.0038 (9)	-0.0032 (9)
C8	0.0448 (13)	0.0333 (12)	0.0533 (15)	0.0049 (10)	0.0103 (11)	-0.0007 (11)
C9	0.0366 (12)	0.0485 (15)	0.0697 (18)	0.0047 (11)	0.0130 (12)	0.0027 (13)
C10	0.0359 (13)	0.0469 (15)	0.0657 (17)	-0.0097 (11)	0.0095 (12)	-0.0025 (13)
Co1	0.0448 (3)	0.0246 (2)	0.0328 (2)	-0.00372 (17)	0.00405 (18)	-0.00398 (17)
N1	0.0384 (10)	0.0322 (10)	0.0489 (12)	-0.0077 (8)	0.0066 (9)	-0.0043 (9)
N2	0.0353 (9)	0.0254 (9)	0.0416 (10)	-0.0006 (7)	0.0021 (8)	-0.0041 (8)
N3	0.0375 (10)	0.0269 (9)	0.0457 (11)	-0.0010 (8)	0.0056 (8)	-0.0031 (8)
N4	0.0836 (18)	0.0854 (19)	0.0428 (13)	-0.0255 (16)	0.0012 (13)	-0.0148 (13)
O2	0.0474 (9)	0.0473 (10)	0.0369 (9)	-0.0027 (8)	0.0099 (7)	0.0049 (8)
O3	0.0441 (9)	0.0492 (11)	0.0617 (11)	0.0162 (8)	0.0092 (8)	-0.0004 (9)
S1	0.0324 (3)	0.0339 (3)	0.0365 (3)	0.0030 (2)	0.0044 (2)	0.0001 (2)
O1	0.078 (7)	0.052 (5)	0.033 (2)	-0.005 (4)	0.020 (4)	-0.008 (3)
C11	0.062 (4)	0.067 (4)	0.041 (2)	-0.019 (3)	0.014 (3)	-0.008 (3)
C13	0.140 (5)	0.126 (5)	0.068 (3)	0.001 (4)	0.051 (3)	0.000 (3)
C12	0.128 (5)	0.118 (5)	0.049 (3)	-0.017 (4)	0.020 (3)	-0.012 (3)
N5	0.089 (3)	0.070 (3)	0.0436 (18)	-0.019 (2)	0.021 (2)	-0.007 (2)
O1'	0.078 (7)	0.052 (5)	0.033 (2)	-0.005 (4)	0.020 (4)	-0.008 (3)
C11'	0.062 (4)	0.067 (4)	0.041 (2)	-0.019 (3)	0.014 (3)	-0.008 (3)
C13'	0.140 (5)	0.126 (5)	0.068 (3)	0.001 (4)	0.051 (3)	0.000 (3)
C12'	0.128 (5)	0.118 (5)	0.049 (3)	-0.017 (4)	0.020 (3)	-0.012 (3)
N5'	0.089 (3)	0.070 (3)	0.0436 (18)	-0.019 (2)	0.021 (2)	-0.007 (2)

Geometric parameters (\AA , $^\circ$)

C1—C6	1.381 (4)	Co1—O1'	2.20 (2)
C1—C2	1.383 (3)	N2—S1	1.5941 (18)
C1—H1A	0.9300	N4—H4B	0.8600
C2—C3	1.385 (3)	N4—H4C	0.8600
C2—S1	1.760 (2)	O2—S1	1.4412 (17)
C3—C4	1.380 (4)	O3—S1	1.4470 (17)
C3—H3A	0.9300	O1—C11	1.216 (13)
C4—C5	1.390 (4)	C11—N5	1.342 (10)
C4—H4A	0.9300	C11—H11A	0.9300
C5—N4	1.370 (3)	C13—N5	1.457 (9)
C5—C6	1.396 (4)	C13—H13A	0.9602
C6—H6A	0.9300	C13—H13B	0.9597
C7—N3	1.333 (3)	C13—H13C	0.9600
C7—N1	1.360 (3)	C13—H12E	0.8765
C7—N2	1.364 (3)	C12—N5	1.489 (11)
C8—N3	1.339 (3)	C12—H12A	0.9599
C8—C9	1.378 (3)	C12—H12B	0.9600
C8—H8A	0.9300	C12—H12C	0.9601
C9—C10	1.373 (4)	O1'—C11'	1.244 (17)
C9—H9A	0.9300	C11'—N5'	1.694 (14)
C10—N1	1.327 (3)	C11'—H11B	0.9593
C10—H10A	0.9300	C13'—N5'	1.386 (13)

Co1—O1 ⁱ	2.065 (19)	C13'—H13D	0.9600
Co1—O1	2.065 (19)	C13'—H13E	0.9600
Co1—N1	2.121 (2)	C13'—H13F	0.9600
Co1—N1 ⁱ	2.121 (2)	C12'—N5'	1.458 (13)
Co1—N2	2.1460 (18)	C12'—H12D	0.9597
Co1—N2 ⁱ	2.1460 (18)	C12'—H12E	0.9601
Co1—O1 ⁱⁱ	2.20 (2)	C12'—H12F	0.9601
C6—C1—C2	120.6 (2)	N5—C13—H13B	107.4
C6—C1—H1A	119.7	H13A—C13—H13B	109.5
C2—C1—H1A	119.7	N5—C13—H13C	106.4
C1—C2—C3	118.7 (2)	H13A—C13—H13C	109.4
C1—C2—S1	120.66 (19)	H13B—C13—H13C	109.5
C3—C2—S1	120.62 (18)	N5—C13—H12E	53.8
C4—C3—C2	120.9 (2)	H13A—C13—H12E	77.9
C4—C3—H3A	119.6	H13B—C13—H12E	85.1
C2—C3—H3A	119.6	H13C—C13—H12E	159.2
C3—C4—C5	120.9 (3)	N5—C13—H12F	63.8
C3—C4—H4A	119.6	H13A—C13—H12F	83.9
C5—C4—H4A	119.6	H13B—C13—H12F	166.5
N4—C5—C4	120.8 (3)	H13C—C13—H12F	65.9
N4—C5—C6	121.2 (3)	H12E—C13—H12F	96.5
C4—C5—C6	117.9 (2)	N5—C12—H12A	157.8
C1—C6—C5	121.0 (2)	N5—C12—H12B	88.6
C1—C6—H6A	119.5	H12A—C12—H12B	109.5
C5—C6—H6A	119.5	N5—C12—H12C	74.6
N3—C7—N1	125.2 (2)	H12A—C12—H12C	109.5
N3—C7—N2	125.86 (19)	H12B—C12—H12C	109.5
N1—C7—N2	108.91 (19)	N5—C12—H13E	125.9
N3—C8—C9	123.7 (2)	H12A—C12—H13E	53.6
N3—C8—H8A	118.2	H12B—C12—H13E	70.9
C9—C8—H8A	118.2	H12C—C12—H13E	159.1
C10—C9—C8	117.0 (2)	N5—C12—H13F	153.6
C10—C9—H9A	121.5	H12B—C12—H13F	117.7
C8—C9—H9A	121.5	H12C—C12—H13F	92.7
N1—C10—C9	121.3 (2)	H13E—C12—H13F	70.1
N1—C10—H10A	119.3	C11—N5—C13	141.8 (7)
C9—C10—H10A	119.3	C11—N5—C12	99.2 (6)
O1 ⁱ —Co1—O1	180.0 (10)	C13—N5—C12	119.0 (6)
O1 ⁱ —Co1—N1	84.2 (3)	C11—N5—H12E	149.4
O1—Co1—N1	95.8 (3)	C12—N5—H12E	95.2
O1 ⁱ —Co1—N1 ⁱ	95.8 (3)	C11—N5—H12F	132.3
O1—Co1—N1 ⁱ	84.2 (3)	C13—N5—H12F	49.4
N1—Co1—N1 ⁱ	180.00 (11)	C12—N5—H12F	92.0
O1 ⁱ —Co1—N2	89.0 (6)	H12E—N5—H12F	73.5
O1—Co1—N2	91.0 (6)	C11'—O1'—Co1	124.5 (17)
N1—Co1—N2	62.59 (7)	O1'—C11'—N5'	87.5 (14)
N1 ⁱ —Co1—N2	117.41 (7)	O1'—C11'—H11B	124.6

O1 ⁱ —Co1—N2 ⁱ	91.0 (6)	N5'—C11'—H11B	143.7
O1—Co1—N2 ⁱ	89.0 (6)	N5'—C13'—H12A	99.0
N1—Co1—N2 ⁱ	117.41 (7)	N5'—C13'—H13D	115.4
N1 ⁱ —Co1—N2 ⁱ	62.59 (7)	H12A—C13'—H13D	126.7
N2—Co1—N2 ⁱ	180.0	N5'—C13'—H13E	108.1
O1 ⁱ —Co1—O1 ⁱ	10.6 (8)	H12A—C13'—H13E	95.5
O1—Co1—O1 ⁱ	169.4 (8)	H13D—C13'—H13E	109.5
N1—Co1—O1 ⁱ	94.1 (5)	N5'—C13'—H13F	104.7
N1 ⁱ —Co1—O1 ⁱ	85.9 (5)	H13D—C13'—H13F	109.5
N2—Co1—O1 ⁱ	90.2 (7)	H13E—C13'—H13F	109.5
N2 ⁱ —Co1—O1 ⁱ	89.8 (7)	N5'—C12'—H13A	154.8
O1 ⁱ —Co1—O1'	169.4 (8)	N5'—C12'—H12B	60.6
O1—Co1—O1'	10.6 (8)	H13A—C12'—H12B	117.7
N1—Co1—O1'	85.9 (5)	N5'—C12'—H12C	54.9
N1 ⁱ —Co1—O1'	94.1 (5)	H13A—C12'—H12C	150.1
N2—Co1—O1'	89.8 (7)	H12B—C12'—H12C	75.3
N2 ⁱ —Co1—O1'	90.2 (7)	N5'—C12'—H12D	131.7
O1 ⁱ —Co1—O1'	180.000 (3)	H13A—C12'—H12D	66.3
C10—N1—C7	117.6 (2)	H12B—C12'—H12D	78.5
C10—N1—Co1	147.50 (17)	H12C—C12'—H12D	92.6
C7—N1—Co1	94.85 (14)	N5'—C12'—H12E	92.3
C7—N2—S1	123.96 (15)	H13A—C12'—H12E	62.9
C7—N2—Co1	93.64 (13)	H12B—C12'—H12E	85.1
S1—N2—Co1	142.24 (11)	H12C—C12'—H12E	146.9
C7—N3—C8	115.21 (19)	H12D—C12'—H12E	109.5
C5—N4—H4B	120.0	N5'—C12'—H12F	102.2
C5—N4—H4C	120.0	H13A—C12'—H12F	83.3
H4B—N4—H4C	120.0	H12B—C12'—H12F	158.7
O2—S1—O3	115.11 (11)	H12C—C12'—H12F	84.4
O2—S1—N2	113.67 (10)	H12D—C12'—H12F	109.5
O3—S1—N2	104.78 (10)	H12E—C12'—H12F	109.5
O2—S1—C2	107.31 (11)	C13'—N5'—C12'	119.6 (8)
O3—S1—C2	108.17 (11)	C13'—N5'—C11'	138.9 (8)
N2—S1—C2	107.48 (10)	C12'—N5'—C11'	101.2 (8)
C11—O1—Co1	129.0 (12)	C13'—N5'—H12B	70.6
O1—C11—N5	157.5 (12)	C12'—N5'—H12B	57.6
O1—C11—H11A	101.2	C11'—N5'—H12B	137.0
N5—C11—H11A	101.2	C13'—N5'—H12C	87.5
O1—C11—H11B	118.0	C12'—N5'—H12C	49.6
N5—C11—H11B	84.2	C11'—N5'—H12C	126.1
N5—C13—H13A	114.5	H12B—N5'—H12C	71.3

Symmetry code: (i) $-x, -y+1, -z$.

Hydrogen-bond geometry (\AA , $^\circ$)

$D\text{—H}\cdots A$	$D\text{—H}$	$H\cdots A$	$D\cdots A$	$D\text{—H}\cdots A$
N4—H4B ⁱⁱ —O3 ⁱⁱ	0.86	2.31	3.112 (3)	155

N4—H4C···O3 ⁱⁱⁱ	0.86	2.27	2.951 (3)	136
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Symmetry codes: (ii) $x, -y+3/2, z+1/2$; (iii) $-x+1, y+1/2, -z+1/2$.