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catena-Poly[[dichloridocobalt(II)]- μ -4,4'-bis(benzimidazol-1-yl)biphenyl]

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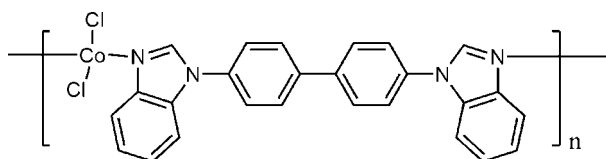
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Key indicators: single-crystal X-ray study; $T = 293$ K; mean $\sigma(\text{C}-\text{C}) = 0.004$ Å; R factor = 0.053; wR factor = 0.124; data-to-parameter ratio = 18.0.

In the title compound, $[\text{CoCl}_2(\text{C}_{26}\text{H}_{18}\text{N}_4)]_n$, the Co^{II} atom (site symmetry 2) is tetrahedrally coordinated by two chloride ions and two N atoms from 4,4'-bis(benzimidazol-1-yl)biphenyl ligands: the complete ligand is generated by crystallographic twofold symmetry. The dihedral angle between the benzene rings is $34.67(8)^\circ$ and the angle between the benzene ring and the adjacent benzimidazole ring system is $43.26(10)^\circ$. The bridging ligand links the Co^{II} atoms into chains propagating in $[\bar{1}01]$.

Related literature

For background to benzimidazole-based ligands in crystal engineering, see: Jin *et al.* (2006); Li *et al.* (2009); Su *et al.* (2003).



Experimental

Crystal data

 $[\text{CoCl}_2(\text{C}_{26}\text{H}_{18}\text{N}_4)]$ $M_r = 516.27$

Monoclinic, $C2/c$
 $a = 12.878(3)$ Å
 $b = 15.181(3)$ Å
 $c = 11.136(2)$ Å
 $\beta = 91.37(3)^\circ$
 $V = 2176.5(8)$ Å³

$Z = 4$
 Mo $K\alpha$ radiation
 $\mu = 1.06$ mm⁻¹
 $T = 293$ K
 $0.25 \times 0.20 \times 0.15$ mm

Data collection

Rigaku Mercury CCD diffractometer
 Absorption correction: multi-scan (*CrystalClear*; Rigaku/MS, 2005)
 $T_{\text{min}} = 0.776$, $T_{\text{max}} = 0.853$

13945 measured reflections
 2696 independent reflections
 2361 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.053$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.124$
 $S = 1.12$
 2696 reflections

150 parameters
 H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.48$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.71$ e Å⁻³

Table 1

Selected bond lengths (Å).

Co1—N1	2.022 (2)	Co1—Cl1	2.2491 (8)
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Data collection: *CrystalClear* (Rigaku/MS, 2005); cell refinement: *CrystalClear*; data reduction: *CrystalClear*; 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: HB5820).

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catena-Poly[[dichloridocobalt(II)]- μ -4,4'-bis(benzimidazol-1-yl)biphenyl]

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S1. Comment

Benzimidazole has been well used in crystal engineering, and a large number of benzimidazole-containing flexible ligands have been extensively studied (Su *et al.*,2003; Jin *et al.*,2006). However, to our knowledge, the research on benzimidazole ligands bearing rigid spacers is still less developed (Li *et al.*,2009).

Single-crystal X-ray diffraction analysis reveals that the title compound (I) crystallizes in the monoclinic space group $C2/c$. The geometry of the Co^{II} ion is surrounded by two benzimidazole rings of distinct L ligands and two chlorine anions, which illustrates a slightly distorted tetrahedral coordination environment (Fig. 1). Notably, as shown in Fig. 2, the four-coordinated Co^{II} center is bridged by the linear ligand L to form an infinite one-dimensional architecture. The dihedral angle between the biphenyl rings is $34.67(8)^\circ$.

S2. Experimental

A mixture of CH_3OH and CHCl_3 (1:1, 8 ml), as a buffer layer, was carefully layered over a solution of 4,4'-Bis(benzimidazol-1-yl)terphenyl (L, 0.06 mmol) in CHCl_3 (6 ml). Then a solution of CoCl_2 (0.06 mmol) in CH_3OH (6 ml) was layered over the buffer layer, and the resultant reaction was left to stand at room temperature. After *ca* three weeks, blue block single crystals appeared at the boundary. Yield: ~40% (based on L).

S3. Refinement

C-bound H atoms were positioned geometrically and refined in the riding-model approximation, with $\text{C-H} = 0.93\text{\AA}$ and $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}$.

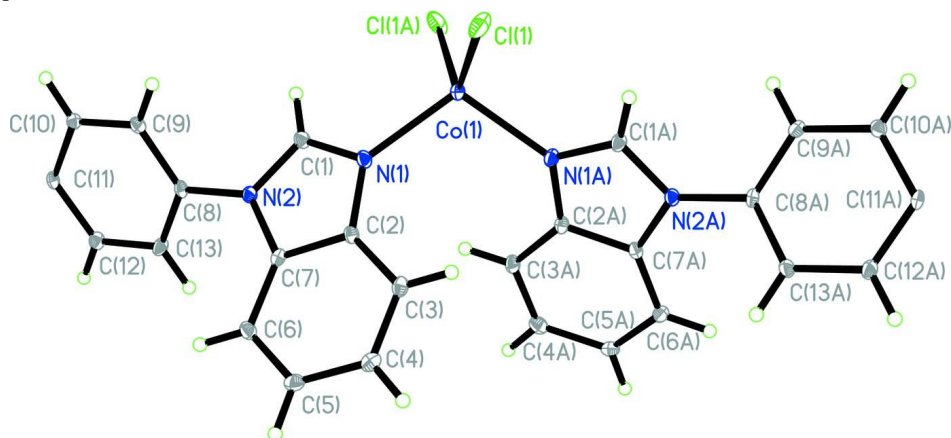
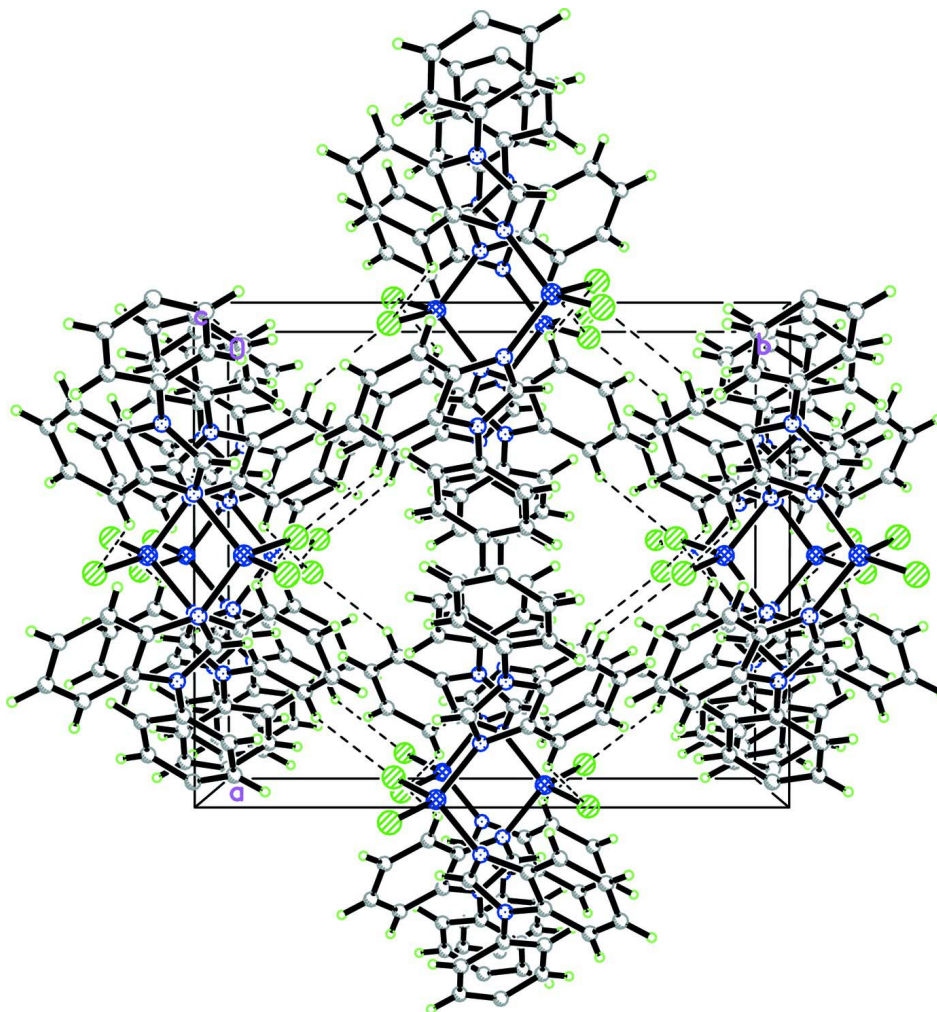


Figure 1

The molecular structure of (I). Displacement ellipsoids are drawn at the 30% probability level and H atoms are shown as small spheres of arbitrary radius. Atoms with suffix A are generated by $(-x, y, 3/2-z)$.

**Figure 2**

The crystal packing for (I).

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Crystal data

[CoCl₂(C₂₆H₁₈N₄)]

$M_r = 516.27$

Monoclinic, C2/c

Hall symbol: -C 2yc

$a = 12.878 (3) \text{ \AA}$

$b = 15.181 (3) \text{ \AA}$

$c = 11.136 (2) \text{ \AA}$

$\beta = 91.37 (3)^\circ$

$V = 2176.5 (8) \text{ \AA}^3$

$Z = 4$

$F(000) = 1052$

$D_x = 1.576 \text{ Mg m}^{-3}$

Mo $K\alpha$ radiation, $\lambda = 0.71073 \text{ \AA}$

Cell parameters from 3011 reflections

$\theta = 2.1\text{--}28.3^\circ$

$\mu = 1.06 \text{ mm}^{-1}$

$T = 293 \text{ K}$

Block, blue

$0.25 \times 0.20 \times 0.15 \text{ mm}$

Data collection

Rigaku Mercury CCD

diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

Detector resolution: 9 pixels mm^{-1}

ω scans

Absorption correction: multi-scan
 (*CrystalClear*; Rigaku/MS, 2005)
 $T_{\min} = 0.776$, $T_{\max} = 0.853$
 13945 measured reflections
 2696 independent reflections
 2361 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.053$
 $\theta_{\max} = 28.3^\circ$, $\theta_{\min} = 2.1^\circ$
 $h = -17 \rightarrow 17$
 $k = -20 \rightarrow 20$
 $l = -14 \rightarrow 14$

Refinement

Refinement on F^2
 Least-squares matrix: full
 $R[F^2 > 2\sigma(F^2)] = 0.053$
 $wR(F^2) = 0.124$
 $S = 1.12$
 2696 reflections
 150 parameters
 0 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
 $w = 1/[\sigma^2(F_o^2) + (0.0551P)^2 + 4.8457P]$
 where $P = (F_o^2 + 2F_c^2)/3$
 $(\Delta/\sigma)_{\max} < 0.001$
 $\Delta\rho_{\max} = 0.48 \text{ e } \text{\AA}^{-3}$
 $\Delta\rho_{\min} = -0.71 \text{ e } \text{\AA}^{-3}$

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)

	<i>x</i>	<i>y</i>	<i>z</i>	$U_{\text{iso}}^*/U_{\text{eq}}$
Co1	0.0000	0.40291 (3)	0.7500	0.01602 (16)
Cl1	0.03587 (6)	0.32578 (5)	0.91836 (7)	0.0311 (2)
N2	0.24698 (16)	0.52363 (14)	0.58018 (19)	0.0139 (4)
N1	0.11781 (16)	0.48022 (14)	0.69588 (19)	0.0151 (4)
C8	0.31898 (18)	0.52081 (17)	0.4838 (2)	0.0134 (5)
C11	0.46103 (18)	0.51614 (17)	0.2986 (2)	0.0140 (5)
C3	0.1150 (2)	0.61615 (17)	0.8273 (2)	0.0162 (5)
H3	0.0601	0.5985	0.8743	0.019*
C12	0.4046 (2)	0.59193 (17)	0.3225 (2)	0.0169 (5)
H12	0.4150	0.6419	0.2761	0.020*
C7	0.23231 (19)	0.59156 (16)	0.6623 (2)	0.0139 (5)
C13	0.3332 (2)	0.59502 (17)	0.4139 (2)	0.0172 (5)
H13	0.2955	0.6461	0.4278	0.021*
C1	0.17714 (19)	0.45999 (17)	0.6047 (2)	0.0153 (5)
H1	0.1715	0.4075	0.5620	0.018*
C10	0.44322 (18)	0.44110 (17)	0.3680 (2)	0.0145 (5)
H10	0.4781	0.3890	0.3514	0.017*
C9	0.37368 (19)	0.44371 (17)	0.4617 (2)	0.0151 (5)
H9	0.3638	0.3942	0.5093	0.018*

C2	0.15099 (19)	0.56315 (17)	0.7347 (2)	0.0138 (5)
C6	0.2823 (2)	0.67174 (17)	0.6813 (2)	0.0173 (5)
H6	0.3365	0.6901	0.6336	0.021*
C4	0.1643 (2)	0.69603 (18)	0.8462 (2)	0.0193 (5)
H4	0.1423	0.7328	0.9073	0.023*
C5	0.2473 (2)	0.72257 (17)	0.7745 (2)	0.0185 (5)
H5	0.2795	0.7762	0.7905	0.022*

Atomic displacement parameters (Å²)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
Co1	0.0152 (3)	0.0153 (3)	0.0178 (3)	0.000	0.00656 (19)	0.000
Cl1	0.0347 (4)	0.0324 (4)	0.0267 (4)	0.0187 (3)	0.0143 (3)	0.0116 (3)
N2	0.0139 (10)	0.0151 (10)	0.0127 (10)	-0.0009 (8)	0.0033 (8)	-0.0021 (8)
N1	0.0146 (10)	0.0170 (10)	0.0138 (10)	-0.0027 (8)	0.0039 (8)	-0.0005 (8)
C8	0.0113 (10)	0.0178 (12)	0.0111 (11)	-0.0016 (9)	0.0032 (9)	0.0000 (9)
C11	0.0112 (11)	0.0184 (12)	0.0126 (12)	-0.0020 (9)	0.0041 (9)	0.0012 (9)
C3	0.0151 (11)	0.0207 (13)	0.0130 (12)	0.0016 (9)	0.0029 (10)	0.0012 (10)
C12	0.0169 (12)	0.0189 (12)	0.0152 (12)	0.0011 (9)	0.0045 (10)	0.0044 (10)
C7	0.0121 (11)	0.0163 (12)	0.0135 (12)	0.0027 (9)	0.0018 (9)	-0.0005 (9)
C13	0.0180 (12)	0.0177 (12)	0.0161 (12)	0.0046 (10)	0.0063 (10)	0.0030 (10)
C1	0.0168 (12)	0.0170 (12)	0.0124 (12)	-0.0024 (9)	0.0036 (10)	-0.0005 (9)
C10	0.0117 (11)	0.0144 (11)	0.0175 (13)	-0.0001 (9)	0.0002 (9)	-0.0015 (10)
C9	0.0158 (11)	0.0151 (12)	0.0144 (12)	-0.0029 (9)	0.0022 (9)	0.0018 (9)
C2	0.0129 (11)	0.0169 (12)	0.0116 (12)	-0.0007 (9)	0.0017 (9)	0.0006 (9)
C6	0.0153 (12)	0.0162 (12)	0.0205 (13)	-0.0013 (9)	0.0023 (10)	0.0036 (10)
C4	0.0216 (13)	0.0185 (13)	0.0177 (13)	0.0036 (10)	0.0009 (10)	-0.0023 (10)
C5	0.0214 (12)	0.0129 (11)	0.0214 (14)	0.0015 (10)	0.0011 (11)	0.0018 (10)

Geometric parameters (Å, °)

Co1—N1	2.022 (2)	C3—H3	0.9300
Co1—N1 ⁱ	2.022 (2)	C12—C13	1.388 (4)
Co1—Cl1 ⁱ	2.2491 (8)	C12—H12	0.9300
Co1—Cl1	2.2491 (8)	C7—C6	1.391 (3)
N2—C1	1.352 (3)	C7—C2	1.404 (3)
N2—C7	1.394 (3)	C13—H13	0.9300
N2—C8	1.436 (3)	C1—H1	0.9300
N1—C1	1.321 (3)	C10—C9	1.392 (4)
N1—C2	1.395 (3)	C10—H10	0.9300
C8—C13	1.384 (3)	C9—H9	0.9300
C8—C9	1.391 (3)	C6—C5	1.378 (4)
C11—C12	1.390 (4)	C6—H6	0.9300
C11—C10	1.399 (3)	C4—C5	1.408 (4)
C11—C11 ⁱⁱ	1.494 (5)	C4—H4	0.9300
C3—C4	1.383 (4)	C5—H5	0.9300
C3—C2	1.396 (4)		

N1—Co1—N1 ⁱ	109.02 (13)	N2—C7—C2	105.3 (2)
N1—Co1—C11 ⁱ	101.21 (7)	C8—C13—C12	118.9 (2)
N1 ⁱ —Co1—C11 ⁱ	114.22 (7)	C8—C13—H13	120.5
N1—Co1—C11	114.22 (7)	C12—C13—H13	120.5
N1 ⁱ —Co1—C11	101.21 (7)	N1—C1—N2	112.9 (2)
C11 ⁱ —Co1—C11	117.25 (5)	N1—C1—H1	123.6
C1—N2—C7	107.1 (2)	N2—C1—H1	123.6
C1—N2—C8	125.1 (2)	C9—C10—C11	120.5 (2)
C7—N2—C8	127.8 (2)	C9—C10—H10	119.7
C1—N1—C2	105.6 (2)	C11—C10—H10	119.7
C1—N1—Co1	123.10 (18)	C8—C9—C10	119.6 (2)
C2—N1—Co1	131.16 (17)	C8—C9—H9	120.2
C13—C8—C9	120.7 (2)	C10—C9—H9	120.2
C13—C8—N2	119.5 (2)	N1—C2—C3	130.1 (2)
C9—C8—N2	119.7 (2)	N1—C2—C7	109.1 (2)
C12—C11—C10	118.4 (2)	C3—C2—C7	120.8 (2)
C12—C11—C11 ⁱⁱ	120.15 (16)	C5—C6—C7	116.5 (2)
C10—C11—C11 ⁱⁱ	121.49 (16)	C5—C6—H6	121.8
C4—C3—C2	117.3 (2)	C7—C6—H6	121.8
C4—C3—H3	121.3	C3—C4—C5	121.1 (2)
C2—C3—H3	121.3	C3—C4—H4	119.5
C13—C12—C11	121.8 (2)	C5—C4—H4	119.5
C13—C12—H12	119.1	C6—C5—C4	122.2 (3)
C11—C12—H12	119.1	C6—C5—H5	118.9
C6—C7—N2	132.6 (2)	C4—C5—H5	118.9
C6—C7—C2	122.0 (2)		
N1 ⁱ —Co1—N1—C1	-143.7 (2)	C8—N2—C1—N1	-177.9 (2)
C11 ⁱ —Co1—N1—C1	-23.0 (2)	C12—C11—C10—C9	-2.5 (4)
C11—Co1—N1—C1	104.0 (2)	C11 ⁱⁱ —C11—C10—C9	177.1 (3)
N1 ⁱ —Co1—N1—C2	31.89 (19)	C13—C8—C9—C10	-0.2 (4)
C11 ⁱ —Co1—N1—C2	152.6 (2)	N2—C8—C9—C10	179.8 (2)
C11—Co1—N1—C2	-80.5 (2)	C11—C10—C9—C8	2.2 (4)
C1—N2—C8—C13	135.7 (3)	C1—N1—C2—C3	179.4 (3)
C7—N2—C8—C13	-41.8 (4)	Co1—N1—C2—C3	3.3 (4)
C1—N2—C8—C9	-44.3 (4)	C1—N1—C2—C7	0.3 (3)
C7—N2—C8—C9	138.2 (3)	Co1—N1—C2—C7	-175.87 (17)
C10—C11—C12—C13	1.0 (4)	C4—C3—C2—N1	179.3 (2)
C11 ⁱⁱ —C11—C12—C13	-178.6 (3)	C4—C3—C2—C7	-1.6 (4)
C1—N2—C7—C6	178.6 (3)	C6—C7—C2—N1	-178.9 (2)
C8—N2—C7—C6	-3.6 (4)	N2—C7—C2—N1	-0.3 (3)
C1—N2—C7—C2	0.1 (3)	C6—C7—C2—C3	1.8 (4)
C8—N2—C7—C2	178.0 (2)	N2—C7—C2—C3	-179.5 (2)
C9—C8—C13—C12	-1.3 (4)	N2—C7—C6—C5	-178.7 (3)
N2—C8—C13—C12	178.7 (2)	C2—C7—C6—C5	-0.5 (4)
C11—C12—C13—C8	0.8 (4)	C2—C3—C4—C5	0.2 (4)
C2—N1—C1—N2	-0.2 (3)	C7—C6—C5—C4	-0.9 (4)

Co1—N1—C1—N2	176.35 (16)	C3—C4—C5—C6	1.1 (4)
C7—N2—C1—N1	0.0 (3)		

Symmetry codes: (i) $-x, y, -z+3/2$; (ii) $-x+1, y, -z+1/2$.