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# *catena*-Poly[($\mu$-2-amino-1,3,4-thiadiazole-$\kappa^2N^3$:$N^4$)di-$\mu$-chlorido-cadmium]

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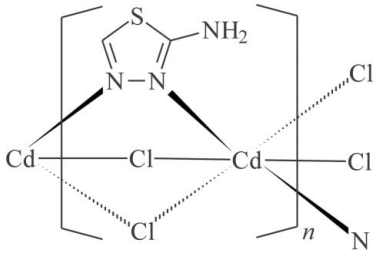
Key indicators: single-crystal X-ray study; $T$ = 297 K; mean $\sigma$(N–C) = 0.004 Å; $R$ factor = 0.021; $wR$ factor = 0.056; data-to-parameter ratio = 16.6.

In the title coordination polymer, $[CdCl_2(C_2H_3N_3S)]_n$, the $Cd^{II}$ cation is coordinated by four $Cl^-$ anions and two N atoms from two *trans* 2-amino-1,3,4-thiadiazole (*L*) ligands in a distorted octahedral geometry. The *L* ligand and $Cl^-$ anions bridge adjacent Cd cations, forming a polymeric chain along the *b* axis; the separation between adjacent Cd cations is 3.619 (1) Å. In the crystal, the polymeric chains are interlinking through N—H···Cl hydrogen bonds between the *L* ligands and $Cl^-$ anions.

## Related literature

For background to coordination polymers, see: Kitagawa *et al.* (2004); Chiang *et al.* (2008); Yeh *et al.* (2008, 2009); Hsu *et al.* (2009). For related Cd coordination polymers, see: Suen & Wang (2007*a*,*b*).




## Experimental

### Crystal data

$[CdCl_2(C_2H_3N_3S)]$
$M_r$ = 284.43
Monoclinic, $P2_1/n$
$a$ = 7.7264 (6) Å
$b$ = 7.2227 (6) Å
$c$ = 12.7608 (11) Å
$\beta$ = 95.489 (2)°
$V$ = 708.86 (10) Å$^3$
$Z$ = 4
Mo $K\alpha$ radiation
$\mu$ = 4.04 mm$^{-1}$
$T$ = 297 K
0.48 × 0.46 × 0.34 mm

### Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2000)
$T_{min}$ = 0.170, $T_{max}$ = 0.341
3718 measured reflections
1381 independent reflections
1354 reflections with $I > 2\sigma(I)$
$R_{int}$ = 0.019

### Refinement

$R[F^2 > 2\sigma(F^2)]$ = 0.021
$wR(F^2)$ = 0.056
$S$ = 1.16
1381 reflections
83 parameters
1 restraint
H-atom parameters constrained
$\Delta\rho_{max}$ = 0.61 e Å$^{-3}$
$\Delta\rho_{min}$ = −0.76 e Å$^{-3}$

**Table 1**
Selected bond lengths (Å).

| | | | |
|---|---|---|---|
| Cd—N1 | 2.361 (2) | Cd—Cl1$^{ii}$ | 2.6697 (7) |
| Cd—N2$^{i}$ | 2.341 (2) | Cd—Cl2 | 2.6583 (7) |
| Cd—Cl1 | 2.6262 (7) | Cd—Cl2$^{ii}$ | 2.6222 (7) |

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

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

| $D$—H···$A$ | $D$—H | H···$A$ | $D$···$A$ | $D$—H···$A$ |
|---|---|---|---|---|
| N3—H3$A$···Cl2$^{iii}$ | 0.86 | 2.60 | 3.390 (3) | 154 |
| N3—H3$B$···Cl2$^{iv}$ | 0.86 | 2.77 | 3.216 (3) | 114 |

Symmetry codes: (iii) $x, y+1, z$; (iv) $-x+2, -y+1, -z+2$.

Data collection: *APEX2* (Bruker, 2010); cell refinement: *SAINT* (Bruker, 2010); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: DAIMOND (Brandenburg, 2010); software used to prepare material for publication: *SHELXL97*.

We are grateful to the National Science Council of the Republic of China and the Nanya Institute of Technology for support.

Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: XU5265).

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# supporting information

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## *catena*-Poly[($\mu$-2-amino-1,3,4-thiadiazole-$\kappa^2N^3$:$N^4$)di-$\mu$-chlorido-cadmium]

**Maw-Cherng Suen, Chun-Wei Yeh and Chi-Hsiung Jou**

### S1. Comment

The synthesis of metal coordination polymers has been a subject of intense research due to their interesting structural chemistry and potential applications in gas storage, separation, catalysis, magnetism, luminescence, and drug delivery (Kitagawa *et al.*, 2004). Roles of anion, solvent and ligand comformations in self-assembly of coordination complexes containing polydentate nitrogen ligands are very intersting (Chiang *et al.*, 2008; Yeh *et al.*, 2008; Hsu *et al.*, 2009; Yeh *et al.*, 2009). Tha Cd(II) complexes containing polydentate ligands showing various type frameworks are also reported (Suen & Wang, 2007*a*,*b*). The $Cd^{2+}$ cations are six-coordinate, which are coordinated with four Cl atoms and two N atoms from two *L* ligands (Fig. 1). The Cd···Cd distance separated by the bridging *L* ligands and Cl atoms is 10.257 (1) and 3.619 (1) Å. The one-dimensional polymeric chains are interlinking through N—H···Cl hydrogen bonds between the *L* ligands and Cl anions in the crystal structure (Fig. 2, Tab.1).

### S2. Experimental

An aqueous solution (5.0 ml) of cadmium chloride (1.0 mmol) was layered carefully over a methanolic solution (5.0 ml) of 2-amino-1,3,4-thiadiazole (1.0 mmol) in a tube. Colourless crystals were obtained after several weeks. These were washed with methanol and collected in 68.7% yield.

### S3. Refinement

H atoms were contrained to ideal geometries with C—H = 0.93 and N—H = 0.86 Å and $U_{iso}$(H) = 1.2$U_{eq}$(C,N).

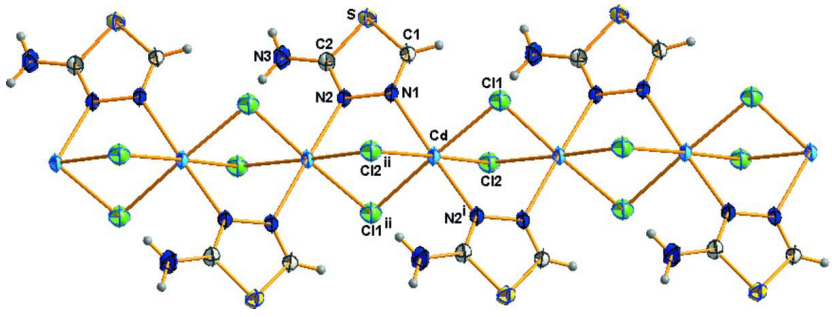


**Figure 1**

A portion of the one-dimensional chain. Ellipsoids are drawn at 30% probability level, and H atoms of spheres of arbitrary radius. Symmetry codes: (i) -*x* + 3/2, *y* - 1/2, -*z* + 3/2; (ii) -*x* + 3/2, *y* + 1/2, -*z* + 3/2.

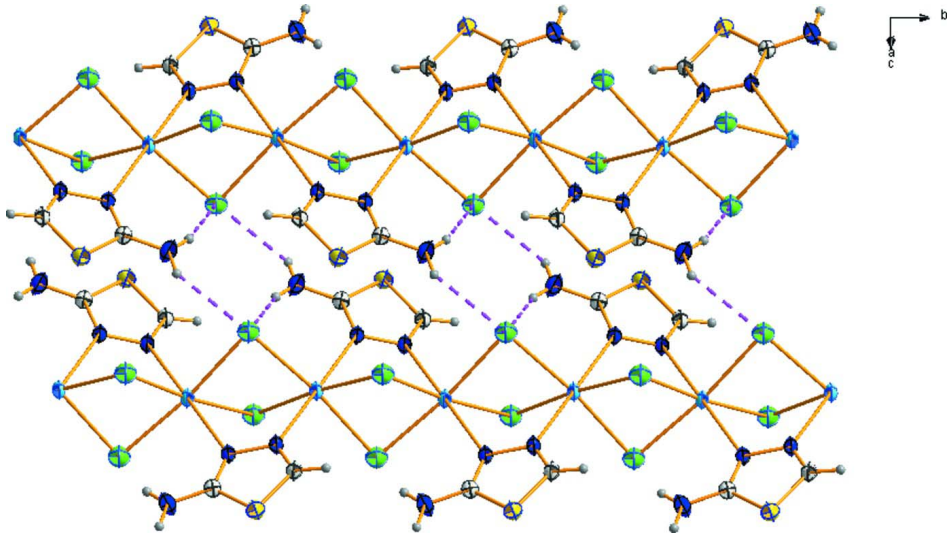


**Figure 2**

The packing diagram shows the N—H···Cl hydrogen bonds among the one-dimensional Chains.

**catena-Poly[(μ-2-amino-1,3,4-thiadiazole- κ²N³:N⁴)di-μ-chlorido-cadmium]**

*Crystal data*

$[CdCl_2(C_2H_3N_3S)]$
$M_r = 284.43$
Monoclinic, $P2_1/n$
Hall symbol: -P 2yn
$a = 7.7264$ (6) Å
$b = 7.2227$ (6) Å
$c = 12.7608$ (11) Å
$\beta = 95.489$ (2)°
$V = 708.86$ (10) Å$^3$
$Z = 4$
$F(000) = 536$
$D_x = 2.665$ Mg m$^{-3}$
Mo $K\alpha$ radiation, $\lambda = 0.71073$ Å
Cell parameters from 3467 reflections
$\theta = 2.7$–$26.0°$
$\mu = 4.04$ mm$^{-1}$
$T = 297$ K
Parallelepiped, colourless
0.48 × 0.46 × 0.34 mm

*Data collection*

Bruker APEXII CCD diffractometer
Radiation source: fine-focus sealed tube
Graphite monochromator
$\varphi$ and $\omega$ scans
Absorption correction: multi-scan (*SADABS*; Bruker, 2000)
$T_{min} = 0.170$, $T_{max} = 0.341$
3718 measured reflections
1381 independent reflections
1354 reflections with $I > 2\sigma(I)$
$R_{int} = 0.019$
$\theta_{max} = 26.0°$, $\theta_{min} = 3.0°$
$h = -9 \rightarrow 9$
$k = -8 \rightarrow 7$
$l = -15 \rightarrow 10$

*Refinement*

Refinement on $F^2$
Least-squares matrix: full
$R[F^2 > 2\sigma(F^2)] = 0.021$
$wR(F^2) = 0.056$
$S = 1.16$
1381 reflections
83 parameters
1 restraint
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
$w = 1/[\sigma^2(F_o^2) + (0.0313P)^2 + 0.7123P]$ where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{max}$ = 0.001
$\Delta\rho_{max}$ = 0.61 e Å$^{-3}$
$\Delta\rho_{min}$ = −0.76 e Å$^{-3}$

Extinction correction: *SHELXL97* (Sheldrick, 2008), $Fc^{*}=kFc[1+0.001xFc^{2}\lambda^{3}/\sin(2\theta)]^{-1/4}$
Extinction coefficient: 0.0097 (6)

*Special details*

**Experimental**. 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.

**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 (Å$^2$)*

| | *x* | *y* | *z* | $U_{iso}$*/$U_{eq}$ |
|---|---|---|---|---|
| Cd | 0.73571 (3) | 0.17322 (3) | 0.745512 (14) | 0.02441 (12) |
| Cl1 | 0.51717 (10) | −0.08486 (9) | 0.79344 (6) | 0.02966 (18) |
| Cl2 | 0.97270 (10) | −0.06290 (9) | 0.83088 (6) | 0.03076 (18) |
| S | 0.70167 (11) | 0.40964 (10) | 1.09911 (6) | 0.03228 (19) |
| N1 | 0.7182 (3) | 0.3292 (3) | 0.90672 (19) | 0.0253 (5) |
| N2 | 0.7623 (3) | 0.5146 (3) | 0.91436 (18) | 0.0245 (5) |
| N3 | 0.7988 (4) | 0.7503 (4) | 1.0408 (2) | 0.0412 (7) |
| H3A | 0.8242 | 0.8302 | 0.9947 | 0.049* |
| H3B | 0.7969 | 0.7827 | 1.1056 | 0.049* |
| C1 | 0.6831 (4) | 0.2586 (4) | 0.9949 (2) | 0.0281 (6) |
| H1A | 0.6504 | 0.1356 | 1.0016 | 0.034* |
| C2 | 0.7623 (4) | 0.5757 (4) | 1.0118 (2) | 0.0260 (6) |

*Atomic displacement parameters (Å$^2$)*

| | $U^{11}$ | $U^{22}$ | $U^{33}$ | $U^{12}$ | $U^{13}$ | $U^{23}$ |
|---|---|---|---|---|---|---|
| Cd | 0.03477 (16) | 0.01300 (14) | 0.02532 (16) | 0.00063 (7) | 0.00221 (9) | −0.00127 (6) |
| Cl1 | 0.0319 (4) | 0.0198 (3) | 0.0383 (4) | −0.0011 (3) | 0.0083 (3) | 0.0025 (3) |
| Cl2 | 0.0327 (4) | 0.0180 (3) | 0.0395 (4) | −0.0019 (3) | −0.0070 (3) | 0.0008 (3) |
| S | 0.0457 (5) | 0.0284 (4) | 0.0229 (4) | −0.0039 (3) | 0.0040 (3) | 0.0012 (3) |
| N1 | 0.0344 (13) | 0.0146 (11) | 0.0270 (12) | 0.0000 (9) | 0.0037 (10) | −0.0012 (9) |
| N2 | 0.0324 (12) | 0.0156 (11) | 0.0255 (11) | −0.0012 (9) | 0.0025 (9) | −0.0007 (9) |
| N3 | 0.0633 (19) | 0.0271 (14) | 0.0331 (14) | −0.0125 (13) | 0.0037 (13) | −0.0069 (11) |
| C1 | 0.0370 (16) | 0.0189 (14) | 0.0282 (14) | −0.0014 (12) | 0.0027 (11) | 0.0014 (11) |
| C2 | 0.0298 (15) | 0.0218 (14) | 0.0258 (14) | 0.0001 (11) | −0.0006 (11) | −0.0006 (10) |

*Geometric parameters (Å, º)*

| | | | |
|---|---|---|---|
| Cd—N1 | 2.361 (2) | N1—C1 | 1.287 (4) |
| Cd—N2$^{i}$ | 2.341 (2) | N1—N2 | 1.383 (3) |
| Cd—Cl1 | 2.6262 (7) | N2—C2 | 1.320 (4) |
| Cd—Cl1$^{ii}$ | 2.6697 (7) | N2—Cd$^{ii}$ | 2.341 (2) |
| Cd—Cl2 | 2.6583 (7) | N3—C2 | 1.337 (4) |
| Cd—Cl2$^{ii}$ | 2.6222 (7) | N3—H3A | 0.8600 |
| S—C1 | 1.715 (3) | N3—H3B | 0.8600 |
| S—C2 | 1.731 (3) | C1—H1A | 0.9300 |
| | | | |
| N2$^{i}$—Cd—N1 | 177.00 (9) | C1—S—C2 | 87.11 (14) |
| N2$^{i}$—Cd—Cl2$^{ii}$ | 94.97 (6) | C1—N1—N2 | 113.1 (2) |
| N1—Cd—Cl2$^{ii}$ | 83.84 (6) | C1—N1—Cd | 127.38 (19) |
| N2$^{i}$—Cd—Cl1 | 85.04 (6) | N2—N1—Cd | 119.28 (16) |
| N1—Cd—Cl1 | 92.52 (6) | C2—N2—N1 | 111.6 (2) |
| Cl2$^{ii}$—Cd—Cl1 | 102.52 (2) | C2—N2—Cd$^{ii}$ | 131.18 (19) |
| N2$^{i}$—Cd—Cl2 | 88.94 (6) | N1—N2—Cd$^{ii}$ | 115.76 (16) |
| N1—Cd—Cl2 | 92.51 (6) | C2—N3—H3A | 120.0 |
| Cl2$^{ii}$—Cd—Cl2 | 173.27 (2) | C2—N3—H3B | 120.0 |
| Cl1—Cd—Cl2 | 83.24 (2) | H3A—N3—H3B | 120.0 |
| N2$^{i}$—Cd—Cl1$^{ii}$ | 95.37 (6) | N1—C1—S | 114.6 (2) |
| N1—Cd—Cl1$^{ii}$ | 87.23 (6) | N1—C1—H1A | 122.7 |
| Cl2$^{ii}$—Cd—Cl1$^{ii}$ | 83.09 (2) | S—C1—H1A | 122.7 |
| Cl1—Cd—Cl1$^{ii}$ | 174.324 (18) | N2—C2—N3 | 123.8 (3) |
| Cl2—Cd—Cl1$^{ii}$ | 91.11 (2) | N2—C2—S | 113.5 (2) |
| Cd—Cl1—Cd$^{i}$ | 86.22 (2) | N3—C2—S | 122.6 (2) |
| Cd$^{i}$—Cl2—Cd | 86.53 (2) | | |

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

*Hydrogen-bond geometry (Å, º)*

| *D*—H···*A* | *D*—H | H···*A* | *D*···*A* | *D*—H···*A* |
|---|---|---|---|---|
| N3—H3*A*···Cl2$^{iii}$ | 0.86 | 2.60 | 3.390 (3) | 154 |
| N3—H3*B*···Cl2$^{iv}$ | 0.86 | 2.77 | 3.216 (3) | 114 |

Symmetry codes: (iii) $x, y+1, z$; (iv) $-x+2, -y+1, -z+2$.