

trans-Dichloridobis(4-nitroaniline- κN^1)-palladium(II)

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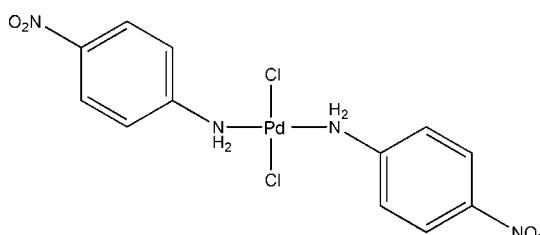
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Key indicators: single-crystal X-ray study; $T = 296\text{ K}$; mean $\sigma(\text{C}-\text{C}) = 0.008\text{ \AA}$; R factor = 0.021; wR factor = 0.077; data-to-parameter ratio = 10.3.

In the title compound, $[\text{PdCl}_2(\text{C}_6\text{H}_4\text{N}_2\text{O}_2)_2]$, the Pd^{II} atom is coordinated in a distorted square-planar geometry by two N atoms from two 4-nitroaniline ligands and two Cl atoms in a *trans* arrangement. Intermolecular N–H···Cl hydrogen bonds involving the amino groups and chloride anions lead to a chain along [100]. These chains are further self-assembled into a three-dimensional network through N–H···O and N–H···Cl hydrogen bonds.

Related literature

For background to the application of palladium compounds in catalysis, see: Hartley (1973); Padmanabhan *et al.* (1985). For related structures, see: Chen *et al.* (2002); Newkome *et al.* (1982).



Experimental

Crystal data

$[\text{PdCl}_2(\text{C}_6\text{H}_4\text{N}_2\text{O}_2)_2]$	$c = 11.0763\text{ (16) \AA}$
$M_r = 453.56$	$\beta = 99.828\text{ (2) }^\circ$
Monoclinic, Cc	$V = 1604.5\text{ (4) \AA}^3$
$a = 5.6014\text{ (8) \AA}$	$Z = 4$
$b = 26.246\text{ (4) \AA}$	Mo $K\alpha$ radiation

$\mu = 1.51\text{ mm}^{-1}$
 $T = 296\text{ K}$

$0.33 \times 0.30 \times 0.26\text{ mm}$

Data collection

Bruker APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.626$, $T_{\max} = 0.688$

4453 measured reflections
2135 independent reflections
2075 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.018$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.021$
 $wR(F^2) = 0.077$
 $S = 1.06$
2135 reflections
208 parameters
2 restraints

H-atom parameters constrained
 $\Delta\rho_{\max} = 0.50\text{ e \AA}^{-3}$
 $\Delta\rho_{\min} = -0.37\text{ e \AA}^{-3}$
Absolute structure: Flack (1983),
696 Friedel pairs
Flack parameter: 0.46 (4)

Table 1
Hydrogen-bond geometry (\AA , $^\circ$).

$D-\text{H}\cdots A$	$D-\text{H}$	$\text{H}\cdots A$	$D\cdots A$	$D-\text{H}\cdots A$
N1–H1A···Cl1 ⁱ	0.90	2.40	3.293 (5)	175
N1–H1B···Cl2 ⁱⁱ	0.90	2.71	3.474 (5)	143
N2–H2A···O3 ⁱⁱⁱ	0.90	2.52	3.287 (7)	143
N2–H2B···Cl2 ^{iv}	0.90	2.46	3.310 (5)	157

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

Data collection: *APEX2* (Bruker, 2007); cell refinement: *SAINT* (Bruker, 2007); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008); molecular graphics: *XP* in *SHELXTL* (Sheldrick, 2008) and *DIAMOND* (Brandenburg, 1999); 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: HY2592).

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supporting information

Acta Cryst. (2012). E68, m1351 [doi:10.1107/S1600536812042134]

***trans*-Dichloridobis(4-nitroaniline- κN^1)palladium(II)**

Tian-Jun Feng

S1. Comment

Palladium compounds have attracted much attention due to their applications in homogeneous and heterogeneous catalyses (Padmanabhan *et al.*, 1985). Some dramatic results in the homogeneous catalysis of the reactions of organic compounds, particularly the successful commercial exploitation of the Wacker one stage process for the homogeneous catalytic oxidation of ethylene to acetaldehyde in the presence of palladium(II) chloride (Hartley, 1973), have contributed to this interest. In this paper, we report the crystal structure of the title compound, a new palladium(II) complex obtained by the reaction of 4-nitroaniline with palladium chloride in ethanol solution.

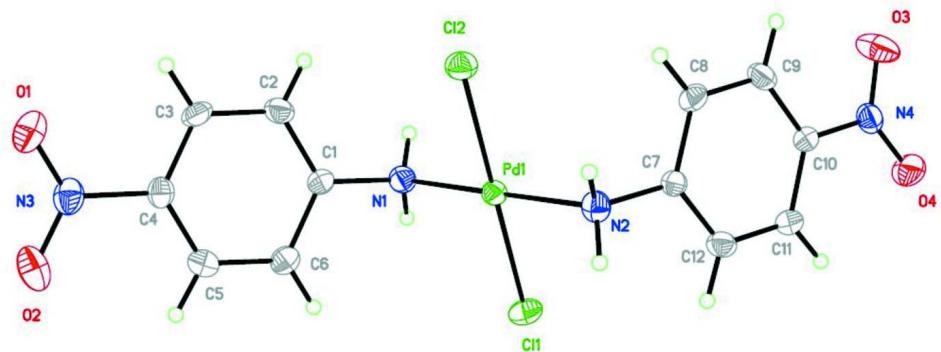
As illustrated in Fig. 1, the Pd^{II} atom exhibits a distorted square-planar coordination geometry, defined by two N atoms from two 4-nitroaniline ligands and two chloride atoms. The molecule adopts the *trans* configuration. The bond distances of Pd—N [2.061 (2) Å] and Pd—Cl [2.302 (3) Å] are comparable with the values found in related complexes (Chen *et al.*, 2002; Newkome *et al.*, 1982). The dihedral angle between the aromatic ring plane and the square plane around Pd1 is 70.54 (2)°. Intermolecular N—H···Cl hydrogen bonds involving the amino groups and chlorine anions (Table 1) lead to a chain along [100] (Fig. 2). These chains are further self-assembled into a three-dimensional network through N—H···O and N—H···Cl hydrogen bonds (Fig. 3).

S2. Experimental

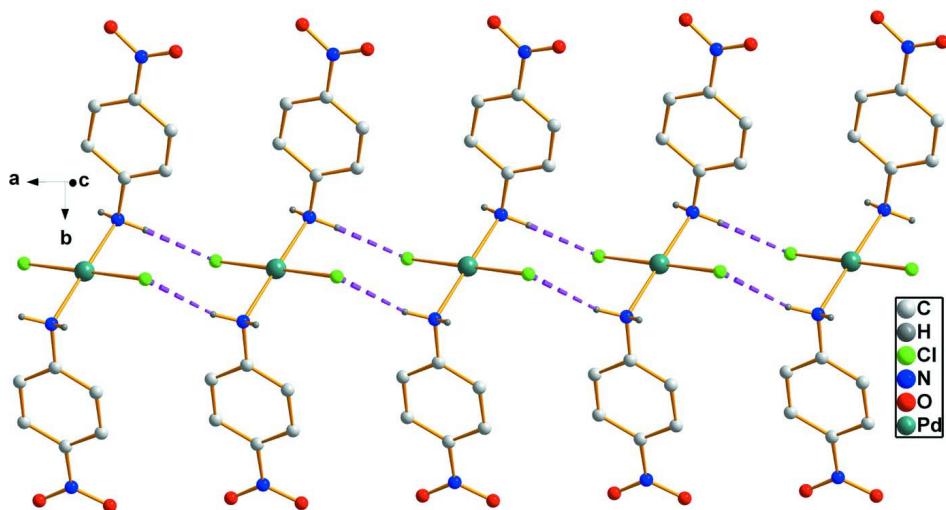
A mixture of palladium chloride (0.1 mmol, 0.018 g) and 4-nitroaniline (0.2 mmol, 0.027 g) in 10 ml of anhydrous ethanol was sealed in an autoclave equipped with a Teflon liner (23 ml) and then heated at 353 K for 2 days. Yellow crystals were obtained by slow evaporation of the solvent at room temperature (yield: 49% based on 4-nitroaniline).

S3. Refinement

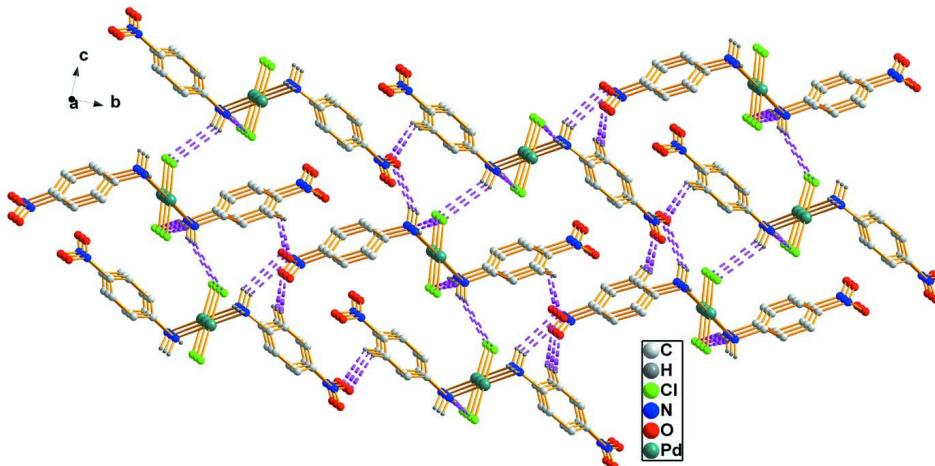
All H atoms were positioned geometrically and refined using a riding model, with C—H = 0.93 and N—H = 0.90 Å and with $U_{\text{iso}}(\text{H}) = 1.2U_{\text{eq}}(\text{C}, \text{N})$. The highest peak is located 0.99 Å from Pd1 and the deepest hole is located 1.24 Å from H5.

**Figure 1**

The molecular structure of the title compound, showing 30% probability displacement ellipsoids.

**Figure 2**

View of the chain structure in the title compound. Hydrogen bonds are shown as pink dashed lines. H atoms not involved in hydrogen bonds have been omitted for clarity.

**Figure 3**

Crystal packing of the title compound. Hydrogen bonds are shown as pink dashed lines. H atoms not involved in hydrogen bonds have been omitted for clarity.

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Crystal data



$M_r = 453.56$

Monoclinic, Cc

Hall symbol: C -2yc

$a = 5.6014 (8)$ Å

$b = 26.246 (4)$ Å

$c = 11.0763 (16)$ Å

$\beta = 99.828 (2)^\circ$

$V = 1604.5 (4)$ Å³

$Z = 4$

$F(000) = 896$

$D_x = 1.878 \text{ Mg m}^{-3}$

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

Cell parameters from 5300 reflections

$\theta = 1.3\text{--}28.0^\circ$

$\mu = 1.51 \text{ mm}^{-1}$

$T = 296$ K

Block, yellow

$0.33 \times 0.30 \times 0.26$ mm

Data collection

Bruker APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

φ and ω scans

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

$T_{\min} = 0.626$, $T_{\max} = 0.688$

4453 measured reflections

2135 independent reflections

2075 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.018$

$\theta_{\max} = 25.2^\circ$, $\theta_{\min} = 2.4^\circ$

$h = -6 \rightarrow 6$

$k = -28 \rightarrow 31$

$l = -13 \rightarrow 10$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.021$

$wR(F^2) = 0.077$

$S = 1.06$

2135 reflections

208 parameters

2 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

$w = 1/[\sigma^2(F_o^2) + (0.059P)^2 + 0.1195P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.002$

$\Delta\rho_{\max} = 0.50 \text{ e } \text{\AA}^{-3}$

$\Delta\rho_{\min} = -0.37 \text{ e } \text{\AA}^{-3}$

Absolute structure: Flack (1983), 696 Friedel pairs

Absolute structure parameter: 0.46 (4)

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>	$U_{\text{iso}}^*/U_{\text{eq}}$
C1	0.7813 (8)	0.46008 (19)	0.2257 (5)	0.0324 (10)
C2	0.6105 (10)	0.4507 (2)	0.2984 (6)	0.0405 (13)
H2	0.4896	0.4745	0.3037	0.049*
C3	0.6200 (9)	0.4053 (2)	0.3641 (5)	0.0406 (12)
H3	0.5098	0.3985	0.4161	0.049*
C4	0.7981 (9)	0.37081 (19)	0.3496 (5)	0.0358 (11)
C5	0.9644 (10)	0.3795 (2)	0.2764 (5)	0.0403 (12)
H5	1.0798	0.3547	0.2692	0.048*
C6	0.9651 (10)	0.4245 (2)	0.2123 (5)	0.0380 (12)
H6	1.0801	0.4311	0.1629	0.046*
C7	1.1775 (9)	0.66743 (19)	0.2754 (5)	0.0313 (10)
C8	0.9834 (10)	0.6949 (2)	0.3044 (5)	0.0371 (11)
H8	0.8961	0.6828	0.3628	0.045*
C9	0.9216 (10)	0.7412 (2)	0.2439 (5)	0.0380 (12)
H9	0.7917	0.7605	0.2606	0.046*
C10	1.0586 (8)	0.75750 (18)	0.1588 (4)	0.0307 (10)
C11	1.2582 (9)	0.73067 (19)	0.1320 (5)	0.0358 (11)
H11	1.3503	0.7432	0.0762	0.043*
C12	1.3145 (9)	0.6845 (2)	0.1918 (5)	0.0366 (11)
H12	1.4445	0.6653	0.1753	0.044*
Cl1	1.2293 (4)	0.55635 (8)	0.0903 (2)	0.0436 (6)
Cl2	0.7793 (4)	0.56838 (7)	0.4000 (2)	0.0408 (5)
N1	0.7776 (8)	0.50756 (15)	0.1568 (4)	0.0356 (9)
H1A	0.6253	0.5198	0.1427	0.043*
H1B	0.8208	0.5010	0.0837	0.043*
N2	1.2299 (8)	0.61827 (16)	0.3332 (4)	0.0365 (9)
H2A	1.2109	0.6202	0.4122	0.044*
H2B	1.3851	0.6099	0.3316	0.044*
N3	0.8016 (11)	0.3224 (2)	0.4165 (6)	0.0463 (15)
N4	0.9877 (9)	0.80407 (15)	0.0883 (5)	0.0404 (10)
O1	0.6631 (11)	0.3158 (2)	0.4860 (6)	0.0769 (16)
O2	0.9473 (9)	0.28974 (16)	0.3938 (5)	0.0609 (12)
O3	0.7966 (11)	0.8248 (2)	0.1012 (5)	0.0674 (18)

O4	1.1159 (9)	0.82003 (16)	0.0195 (5)	0.0601 (12)
Pd1	1.00552 (7)	0.562350 (12)	0.24606 (5)	0.02914 (13)

Atomic displacement parameters (\AA^2)

	U^{11}	U^{22}	U^{33}	U^{12}	U^{13}	U^{23}
C1	0.027 (2)	0.035 (3)	0.035 (3)	-0.0012 (19)	0.005 (2)	0.000 (2)
C2	0.029 (3)	0.037 (3)	0.056 (4)	0.001 (2)	0.010 (3)	-0.008 (3)
C3	0.034 (3)	0.045 (3)	0.047 (3)	-0.006 (2)	0.020 (2)	-0.003 (3)
C4	0.038 (3)	0.032 (3)	0.036 (3)	-0.006 (2)	0.004 (2)	-0.002 (2)
C5	0.039 (3)	0.041 (3)	0.042 (3)	0.006 (2)	0.010 (2)	-0.003 (2)
C6	0.038 (3)	0.044 (3)	0.033 (3)	-0.003 (2)	0.011 (2)	0.002 (2)
C7	0.029 (2)	0.029 (2)	0.036 (2)	-0.0016 (19)	0.005 (2)	-0.005 (2)
C8	0.040 (3)	0.039 (3)	0.036 (3)	-0.005 (2)	0.016 (2)	0.004 (2)
C9	0.035 (2)	0.033 (3)	0.046 (3)	0.0080 (19)	0.007 (3)	-0.005 (2)
C10	0.028 (2)	0.031 (2)	0.033 (3)	-0.0011 (18)	0.006 (2)	0.000 (2)
C11	0.033 (2)	0.038 (3)	0.039 (3)	0.000 (2)	0.013 (2)	-0.001 (2)
C12	0.029 (2)	0.040 (3)	0.041 (3)	0.004 (2)	0.007 (2)	-0.002 (2)
Cl1	0.0384 (10)	0.0581 (11)	0.0374 (11)	0.0009 (7)	0.0154 (8)	0.0016 (7)
Cl2	0.0420 (10)	0.0447 (9)	0.0380 (11)	0.0022 (6)	0.0133 (8)	-0.0019 (7)
N1	0.033 (2)	0.035 (2)	0.038 (2)	-0.0002 (17)	0.0059 (18)	-0.0011 (18)
N2	0.036 (2)	0.036 (2)	0.036 (2)	0.0006 (17)	0.0009 (18)	0.0037 (18)
N3	0.059 (3)	0.043 (3)	0.037 (3)	-0.003 (3)	0.008 (3)	0.004 (2)
N4	0.045 (2)	0.031 (2)	0.045 (3)	0.0020 (19)	0.007 (2)	0.001 (2)
O1	0.084 (4)	0.069 (3)	0.087 (4)	-0.002 (3)	0.040 (3)	0.028 (3)
O2	0.076 (3)	0.044 (2)	0.056 (3)	0.008 (2)	-0.005 (2)	0.006 (2)
O3	0.086 (4)	0.068 (4)	0.057 (3)	0.042 (3)	0.036 (3)	0.024 (3)
O4	0.064 (3)	0.053 (3)	0.071 (3)	0.004 (2)	0.031 (2)	0.023 (2)
Pd1	0.02850 (17)	0.02877 (19)	0.03039 (19)	0.00214 (16)	0.00572 (12)	0.00207 (17)

Geometric parameters (\AA , ^\circ)

C1—C2	1.374 (8)	C9—H9	0.9300
C1—C6	1.416 (8)	C10—C11	1.395 (7)
C1—N1	1.459 (7)	C10—N4	1.468 (6)
C2—C3	1.394 (9)	C11—C12	1.390 (7)
C2—H2	0.9300	C11—H11	0.9300
C3—C4	1.376 (8)	C12—H12	0.9300
C3—H3	0.9300	Cl1—Pd1	2.305 (2)
C4—C5	1.354 (8)	Cl2—Pd1	2.297 (2)
C4—N3	1.470 (8)	N1—Pd1	2.060 (4)
C5—C6	1.379 (8)	N1—H1A	0.9000
C5—H5	0.9300	N1—H1B	0.9000
C6—H6	0.9300	N2—Pd1	2.062 (4)
C7—C12	1.375 (8)	N2—H2A	0.9000
C7—C8	1.387 (7)	N2—H2B	0.9000
C7—N2	1.448 (7)	N3—O1	1.196 (9)
C8—C9	1.403 (8)	N3—O2	1.238 (8)

C8—H8	0.9300	N4—O4	1.207 (7)
C9—C10	1.381 (8)	N4—O3	1.231 (7)
C2—C1—C6	122.3 (5)	C12—C11—C10	117.6 (5)
C2—C1—N1	120.6 (5)	C12—C11—H11	121.2
C6—C1—N1	117.0 (5)	C10—C11—H11	121.2
C1—C2—C3	119.4 (5)	C7—C12—C11	119.9 (5)
C1—C2—H2	120.3	C7—C12—H12	120.1
C3—C2—H2	120.3	C11—C12—H12	120.1
C4—C3—C2	117.7 (5)	C1—N1—Pd1	113.1 (3)
C4—C3—H3	121.1	C1—N1—H1A	109.0
C2—C3—H3	121.1	Pd1—N1—H1A	109.0
C5—C4—C3	123.1 (5)	C1—N1—H1B	109.0
C5—C4—N3	119.7 (5)	Pd1—N1—H1B	109.0
C3—C4—N3	117.2 (5)	H1A—N1—H1B	107.8
C4—C5—C6	120.9 (5)	C7—N2—Pd1	111.5 (3)
C4—C5—H5	119.5	C7—N2—H2A	109.3
C6—C5—H5	119.5	Pd1—N2—H2A	109.3
C5—C6—C1	116.5 (5)	C7—N2—H2B	109.3
C5—C6—H6	121.8	Pd1—N2—H2B	109.3
C1—C6—H6	121.8	H2A—N2—H2B	108.0
C12—C7—C8	122.4 (5)	O1—N3—O2	123.6 (6)
C12—C7—N2	119.7 (5)	O1—N3—C4	119.7 (6)
C8—C7—N2	117.9 (5)	O2—N3—C4	116.6 (6)
C7—C8—C9	118.6 (5)	O4—N4—O3	122.9 (5)
C7—C8—H8	120.7	O4—N4—C10	119.2 (5)
C9—C8—H8	120.7	O3—N4—C10	117.9 (5)
C10—C9—C8	118.3 (5)	N1—Pd1—N2	178.86 (17)
C10—C9—H9	120.8	N1—Pd1—Cl2	91.68 (14)
C8—C9—H9	120.8	N2—Pd1—Cl2	88.43 (15)
C9—C10—C11	123.2 (5)	N1—Pd1—Cl1	87.97 (15)
C9—C10—N4	119.3 (4)	N2—Pd1—Cl1	91.91 (15)
C11—C10—N4	117.5 (4)	Cl2—Pd1—Cl1	179.48 (11)

Hydrogen-bond geometry (Å, °)

D—H···A	D—H	H···A	D···A	D—H···A
N1—H1A···Cl1 ⁱ	0.90	2.40	3.293 (5)	175
N1—H1B···Cl2 ⁱⁱ	0.90	2.71	3.474 (5)	143
N2—H2A···O3 ⁱⁱⁱ	0.90	2.52	3.287 (7)	143
N2—H2B···Cl2 ^{iv}	0.90	2.46	3.310 (5)	157

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