

Chlorido(4,4',4''-tri-tert-butyl-2,2':6',2''-terpyridine)platinum(II) chloride toluene monosolvate

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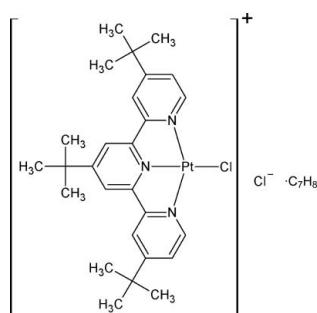
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Key indicators: single-crystal X-ray study; $T = 100$ K; mean $\sigma(\text{C}-\text{C}) = 0.007$ Å; disorder in solvent or counterion; R factor = 0.032; wR factor = 0.108; data-to-parameter ratio = 18.6.

In the title compound, $[\text{PtCl}(\text{C}_{27}\text{H}_{35}\text{N}_3)]\text{Cl} \cdot \text{C}_7\text{H}_8$, the Pt^{II} atom is coordinated in a pseudo-square-planar fashion by the N atoms of a 4,4',4''-tri-tert-butyl-2,2':6',2''-terpyridine (tbtrpy) ligand and a Cl atom. The Pt–N distance of the N atom on the central pyridine is 1.941 (4) Å, while the peripheral N atoms have Pt–N distances of 2.015 (4) and 2.013 (4) Å. The Pt–Cl bond distance is 2.3070 (10) Å. The cations pack as dimers in a head-to-tail orientation with an intermolecular Pt···Pt distance of 3.2774 (3) Å and Pt···N distances of 3.599 (4), 3.791 (4) and 4.115 (4) Å. The solvent molecule is disordered and occupies two positions with a ratio of 0.553 (6):0.447 (6).

Related literature

For crystal structures of the title cation, $[(\text{tbtrpy})\text{PtCl}]^+$, see: Batrice *et al.* (2010); Lai *et al.* (1999). For head-to-tail packing of related terpyridine complexes with close Pt···Pt distances, see: Bailey *et al.* (1995); Sengul (2004). For the synthesis of $[(\text{tbtrpy})\text{PtCl}]\text{Cl}$, see: Howe-Grant & Lippard (1980).



Experimental

Crystal data

$[\text{PtCl}(\text{C}_{27}\text{H}_{35}\text{N}_3)]\text{Cl} \cdot \text{C}_7\text{H}_8$
 $M_r = 759.70$
Monoclinic, $P2_1/c$
 $a = 9.4418$ (3) Å
 $b = 20.0002$ (7) Å
 $c = 17.2321$ (6) Å
 $\beta = 91.948$ (1)°

$V = 3252.19$ (19) Å³
 $Z = 4$
Mo $K\alpha$ radiation
 $\mu = 4.51$ mm⁻¹
 $T = 100$ K
 $0.26 \times 0.21 \times 0.09$ mm

Data collection

Bruker SMART APEXII CCD diffractometer
Absorption correction: multi-scan (*SADABS*; Bruker, 2007)
 $T_{\min} = 0.387$, $T_{\max} = 0.677$

37919 measured reflections
6647 independent reflections
6232 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.030$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.032$
 $wR(F^2) = 0.108$
 $S = 1.05$
6647 reflections
358 parameters

14 restraints
H-atom parameters constrained
 $\Delta\rho_{\max} = 1.47$ e Å⁻³
 $\Delta\rho_{\min} = -2.78$ e Å⁻³

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: *SHELXTL* (Sheldrick, 2008) and *Mercury* (Macrae *et al.*, 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: BV2164).

References

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supporting information

Acta Cryst. (2010). E66, m1704 [https://doi.org/10.1107/S1600536810048750]

Chlorido(4,4',4''-tri-*tert*-butyl-2,2':6',2''-terpyridine)platinum(II) chloride toluene monosolvate

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S1. Comment

The bond distances and angles around the platinum atom in the title structure are all similar to the structures of the perchlorate (Lai, *et al.*, 1999) and tetrafluoroborate salts (Batrice *et al.* 2010) of the $[(\text{tbtrpy})\text{PtCl}]^+$ complex. The cations in these structures all pack in head-to-tail dimers. Interestingly, the interplanar (Pt, Cl and N atoms) distance between the two cations seems to be related to the size of the anion with the Cl^- , BF_4^- , and ClO_4^- being 3.283, 3.390, and 3.536 Å, respectively. In addition to a smaller counterion, the structure of the title complex contains a toluene molecule. This suggests that the solvent molecule may also influence the ability of these types of complexes to interact significantly with each other (Bailey, *et al.*, 1995).

The short Pt(1)—Pt(1') distance, 3.2774 (3) Å, of the title complex is similar to the intermolecular Pt—Pt distance in the structures of $[(\text{trpy})\text{PtCl}]\text{Cl}$, 3.397 Å, (Sengul, 2004) and $[(\text{trpy})\text{PtCl}]\text{ClO}_4$, 3.269 Å, (Bailey *et al.*, 1995). This indicates that the bulky *tert*-Butyl groups of the tbtrpy ligand do not appear to restrict the ability of this complex to form suitable M—M and/or π — π interactions between the two molecules of the dimer.

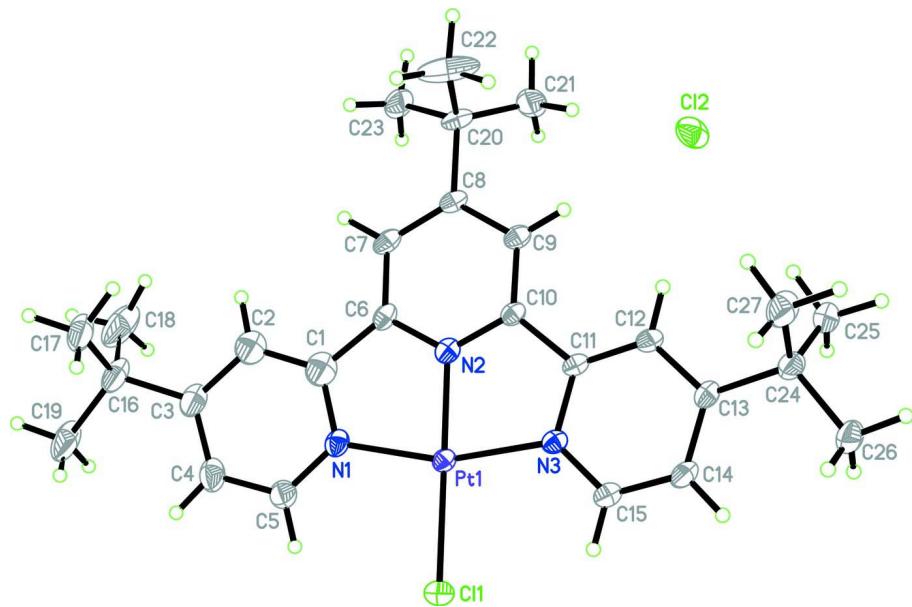
The solvent molecule is disordered and occupies two positions with a ratio of 0.553 (6):0.447 (6).

S2. Experimental

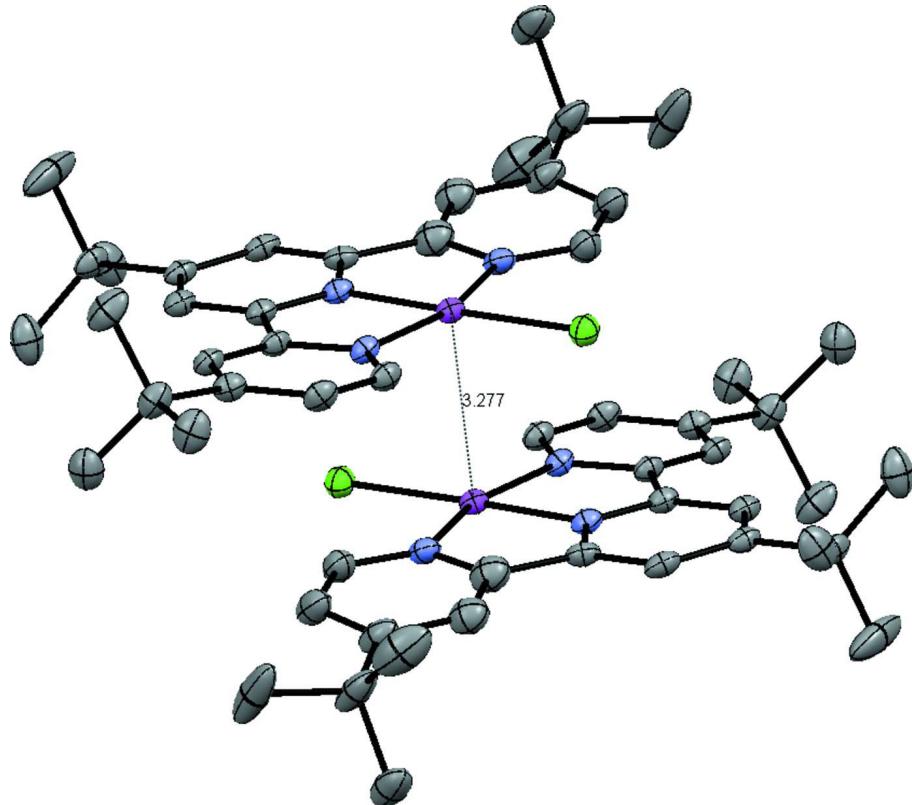
$[(\text{tbtrpy})\text{PtCl}]\text{Cl}$ was synthesized according to modifications on a published procedure (Howe-Grant *et al.*, 1980). This $[(\text{tbtrpy})\text{PtCl}]^+$ complex was reacted with various aromatic thiol ligands (SAr). Crystals of the title compound were grown from the slow evaporation of an acetonitrile/toluene solution containing $[(\text{tbtrpy})\text{Pt}(\text{SAr})]\text{Cl}$ and $[(\text{tbtrpy})\text{PtCl}]\text{Cl}$.

S3. Refinement

H atoms attached to C atoms were placed in idealized positions (C—H = 0.95–0.98 Å) and allowed to ride on their parent atoms. All H atoms were constrained so that $U_{\text{iso}}(\text{H})$ were equal to 1.2 U_{eq} or 1.5 U_{eq} of their respective parent atoms. The solvent molecule is disordered and occupies two positions with a ratio of 0.553 (6): 0.447 (6). Aromatic C atoms were fitted to a regular hexagon with default distances 1.390 Å (AFIX 66) and refined anisotropically. Both CH_3 groups were refined anisotropically with fixed C—C distances as 1.51 Å. The largest peak in the final Fourier difference map (1.51 e Å^{−3}) was located 1.40 Å from the several disordered C atoms of the solvent.

**Figure 1**

View of title complex (50% probability displacement ellipsoids) without the toluene solvate.

**Figure 2**

Mercury (Macrae, *et al.*, 2008) rendition of head-to-tail packing with Pt—Pt' distance in 3.2774 (3) Å

Chlorido(4,4',4''-tri-tert-butyl-2,2':6',2''-terpyridine)platinum chloride toluene monosolvate*Crystal data*

$M_r = 759.70$

Monoclinic, $P2_1/c$

Hall symbol: -P 2ybc

$a = 9.4418 (3)$ Å

$b = 20.0002 (7)$ Å

$c = 17.2321 (6)$ Å

$\beta = 91.948 (1)^\circ$

$V = 3252.19 (19)$ Å³

$Z = 4$

$F(000) = 1520$

$D_x = 1.552$ Mg m⁻³

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

Cell parameters from 9878 reflections

$\theta = 2.4\text{--}27.2^\circ$

$\mu = 4.51$ mm⁻¹

$T = 100$ K

Plate, yellow

0.26 × 0.21 × 0.09 mm

Data collection

Bruker SMART APEXII CCD
diffractometer

Radiation source: fine-focus sealed tube

Graphite monochromator

ω scans

Absorption correction: multi-scan
(SADABS; Bruker, 2007)

$T_{\min} = 0.387$, $T_{\max} = 0.677$

37919 measured reflections

6647 independent reflections

6232 reflections with $I > 2\sigma(I)$

$R_{\text{int}} = 0.030$

$\theta_{\max} = 26.4^\circ$, $\theta_{\min} = 1.6^\circ$

$h = -11 \rightarrow 11$

$k = -24 \rightarrow 24$

$l = -21 \rightarrow 21$

Refinement

Refinement on F^2

Least-squares matrix: full

$R[F^2 > 2\sigma(F^2)] = 0.032$

$wR(F^2) = 0.108$

$S = 1.05$

6647 reflections

358 parameters

14 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

$w = 1/[\sigma^2(F_o^2) + (0.080P)^2 + 5.P]$
where $P = (F_o^2 + 2F_c^2)/3$

$(\Delta/\sigma)_{\max} = 0.008$

$\Delta\rho_{\max} = 1.47$ e Å⁻³

$\Delta\rho_{\min} = -2.78$ e Å⁻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 (Å²)

| | <i>x</i> | <i>y</i> | <i>z</i> | $U_{\text{iso}}^*/U_{\text{eq}}$ | Occ. (<1) |
|-----|---------------|--------------|--------------|----------------------------------|-----------|
| Pt1 | 0.588936 (16) | 0.429628 (7) | 0.502253 (9) | 0.01709 (9) | |
| Cl1 | 0.80289 (11) | 0.48538 (5) | 0.50311 (6) | 0.0234 (2) | |
| N1 | 0.5430 (5) | 0.43245 (16) | 0.3872 (2) | 0.0212 (8) | |
| N2 | 0.4112 (3) | 0.38083 (19) | 0.50120 (19) | 0.0192 (8) | |

| | | | | |
|------|-------------|--------------|------------|-------------|
| N3 | 0.5802 (4) | 0.40887 (18) | 0.6163 (2) | 0.0176 (7) |
| C1 | 0.4201 (6) | 0.3997 (3) | 0.3663 (3) | 0.0343 (7) |
| C2 | 0.3788 (6) | 0.3916 (3) | 0.2892 (3) | 0.0343 (7) |
| H2A | 0.2930 | 0.3690 | 0.2759 | 0.041* |
| C3 | 0.4636 (7) | 0.4170 (2) | 0.2303 (3) | 0.0317 (11) |
| C4 | 0.5847 (6) | 0.4531 (3) | 0.2542 (3) | 0.0346 (12) |
| H4A | 0.6420 | 0.4734 | 0.2166 | 0.041* |
| C5 | 0.6209 (5) | 0.4591 (2) | 0.3320 (3) | 0.0276 (10) |
| H5A | 0.7043 | 0.4831 | 0.3468 | 0.033* |
| C6 | 0.3424 (5) | 0.3720 (2) | 0.4324 (2) | 0.0189 (8) |
| C7 | 0.2114 (5) | 0.3402 (2) | 0.4302 (3) | 0.0210 (9) |
| H7A | 0.1619 | 0.3328 | 0.3820 | 0.025* |
| C8 | 0.1530 (4) | 0.3190 (2) | 0.4998 (2) | 0.0190 (8) |
| C9 | 0.2314 (4) | 0.3270 (2) | 0.5698 (3) | 0.0197 (8) |
| H9A | 0.1957 | 0.3110 | 0.6171 | 0.024* |
| C10 | 0.3625 (4) | 0.3590 (2) | 0.5688 (2) | 0.0171 (8) |
| C11 | 0.4619 (4) | 0.3728 (2) | 0.6359 (2) | 0.0177 (8) |
| C12 | 0.4448 (4) | 0.3510 (2) | 0.7101 (2) | 0.0194 (8) |
| H12A | 0.3620 | 0.3267 | 0.7222 | 0.023* |
| C13 | 0.5482 (5) | 0.3639 (2) | 0.7686 (2) | 0.0204 (8) |
| C14 | 0.6643 (5) | 0.4025 (2) | 0.7476 (3) | 0.0232 (9) |
| H14A | 0.7348 | 0.4144 | 0.7858 | 0.028* |
| C15 | 0.6773 (5) | 0.4234 (2) | 0.6722 (3) | 0.0211 (9) |
| H15A | 0.7579 | 0.4489 | 0.6592 | 0.025* |
| C16 | 0.4265 (8) | 0.4015 (3) | 0.1449 (3) | 0.0423 (15) |
| C17 | 0.4511 (8) | 0.3261 (3) | 0.1326 (3) | 0.0464 (15) |
| H17A | 0.3892 | 0.3006 | 0.1661 | 0.070* |
| H17B | 0.5503 | 0.3153 | 0.1457 | 0.070* |
| H17C | 0.4296 | 0.3147 | 0.0782 | 0.070* |
| C18 | 0.2729 (9) | 0.4188 (3) | 0.1272 (4) | 0.058 (2) |
| H18A | 0.2554 | 0.4653 | 0.1421 | 0.087* |
| H18B | 0.2119 | 0.3891 | 0.1565 | 0.087* |
| H18C | 0.2520 | 0.4133 | 0.0715 | 0.087* |
| C19 | 0.5205 (11) | 0.4413 (4) | 0.0910 (3) | 0.072 (3) |
| H19A | 0.5057 | 0.4892 | 0.0992 | 0.109* |
| H19B | 0.4959 | 0.4299 | 0.0369 | 0.109* |
| H19C | 0.6201 | 0.4302 | 0.1024 | 0.109* |
| C20 | 0.0069 (5) | 0.2843 (2) | 0.4993 (3) | 0.0242 (9) |
| C21 | -0.0735 (5) | 0.3042 (3) | 0.5714 (3) | 0.0385 (13) |
| H21A | -0.1653 | 0.2814 | 0.5708 | 0.058* |
| H21B | -0.0885 | 0.3527 | 0.5713 | 0.058* |
| H21C | -0.0181 | 0.2914 | 0.6181 | 0.058* |
| C22 | 0.0365 (7) | 0.2091 (3) | 0.5021 (5) | 0.061 (2) |
| H22A | -0.0532 | 0.1848 | 0.5053 | 0.092* |
| H22B | 0.0972 | 0.1988 | 0.5478 | 0.092* |
| H22C | 0.0843 | 0.1956 | 0.4550 | 0.092* |
| C23 | -0.0820 (5) | 0.3035 (3) | 0.4267 (3) | 0.0364 (12) |
| H23A | -0.1783 | 0.2859 | 0.4310 | 0.055* |

| | | | | |
|------|--------------|-------------|-------------|----------------------|
| H23B | -0.0389 | 0.2846 | 0.3807 | 0.055* |
| H23C | -0.0857 | 0.3523 | 0.4222 | 0.055* |
| C24 | 0.5345 (5) | 0.3346 (2) | 0.8503 (3) | 0.0256 (9) |
| C25 | 0.3910 (5) | 0.3546 (3) | 0.8826 (3) | 0.0318 (11) |
| H25A | 0.3847 | 0.3380 | 0.9359 | 0.048* |
| H25B | 0.3143 | 0.3352 | 0.8502 | 0.048* |
| H25C | 0.3824 | 0.4035 | 0.8825 | 0.048* |
| C26 | 0.6550 (5) | 0.3573 (3) | 0.9056 (3) | 0.0318 (11) |
| H26A | 0.6420 | 0.3378 | 0.9571 | 0.048* |
| H26B | 0.6544 | 0.4061 | 0.9095 | 0.048* |
| H26C | 0.7458 | 0.3424 | 0.8858 | 0.048* |
| C27 | 0.5403 (6) | 0.2585 (3) | 0.8428 (3) | 0.0352 (12) |
| H27A | 0.5393 | 0.2382 | 0.8946 | 0.053* |
| H27B | 0.6274 | 0.2456 | 0.8173 | 0.053* |
| H27C | 0.4580 | 0.2429 | 0.8117 | 0.053* |
| Cl2 | 0.14340 (13) | 0.24444 (6) | 0.74948 (7) | 0.0315 (3) |
| C1A | 0.9381 (7) | 0.0032 (4) | 0.6429 (3) | 0.067 (4) 0.553 (6) |
| C2A | 0.9481 (12) | 0.0561 (3) | 0.6952 (5) | 0.081 (5) 0.553 (6) |
| H2AA | 0.9471 | 0.1008 | 0.6768 | 0.097* 0.553 (6) |
| C3A | 0.9594 (12) | 0.0434 (4) | 0.7745 (4) | 0.068 (4) 0.553 (6) |
| H3AA | 0.9662 | 0.0795 | 0.8103 | 0.081* 0.553 (6) |
| C4A | 0.9608 (9) | -0.0221 (5) | 0.8015 (3) | 0.050 (4) 0.553 (6) |
| H4AA | 0.9686 | -0.0307 | 0.8557 | 0.060* 0.553 (6) |
| C5A | 0.9509 (11) | -0.0749 (3) | 0.7492 (5) | 0.049 (3) 0.553 (6) |
| H5AA | 0.9518 | -0.1197 | 0.7676 | 0.059* 0.553 (6) |
| C6A | 0.9395 (10) | -0.0623 (3) | 0.6699 (4) | 0.059 (3) 0.553 (6) |
| H6AA | 0.9327 | -0.0984 | 0.6342 | 0.070* 0.553 (6) |
| C7A | 0.9269 (10) | 0.0104 (4) | 0.5583 (4) | 0.0343 (7) 0.553 (6) |
| H7AA | 0.8482 | -0.0170 | 0.5378 | 0.051* 0.553 (6) |
| H7AB | 1.0154 | -0.0045 | 0.5357 | 0.051* 0.553 (6) |
| H7AC | 0.9097 | 0.0574 | 0.5449 | 0.051* 0.553 (6) |
| C1B | 0.9217 (9) | 0.0531 (4) | 0.6741 (5) | 0.067 (4) 0.447 (6) |
| C2B | 0.9072 (14) | -0.0135 (5) | 0.6522 (5) | 0.081 (5) 0.447 (6) |
| H2BA | 0.8806 | -0.0245 | 0.6001 | 0.097* 0.447 (6) |
| C3B | 0.9319 (16) | -0.0640 (4) | 0.7064 (7) | 0.068 (4) 0.447 (6) |
| H3BA | 0.9220 | -0.1095 | 0.6914 | 0.081* 0.447 (6) |
| C4B | 0.9709 (13) | -0.0479 (5) | 0.7826 (6) | 0.050 (4) 0.447 (6) |
| H4BA | 0.9877 | -0.0824 | 0.8197 | 0.060* 0.447 (6) |
| C5B | 0.9853 (13) | 0.0187 (5) | 0.8046 (4) | 0.049 (3) 0.447 (6) |
| H5BA | 1.0120 | 0.0297 | 0.8567 | 0.059* 0.447 (6) |
| C6B | 0.9607 (12) | 0.0692 (4) | 0.7504 (5) | 0.059 (3) 0.447 (6) |
| H6BA | 0.9706 | 0.1147 | 0.7654 | 0.070* 0.447 (6) |
| C7B | 0.8956 (12) | 0.1064 (5) | 0.6175 (6) | 0.0343 (7) 0.447 (6) |
| H7BA | 0.9856 | 0.1203 | 0.5959 | 0.051* 0.447 (6) |
| H7BB | 0.8520 | 0.1446 | 0.6431 | 0.051* 0.447 (6) |
| H7BC | 0.8318 | 0.0901 | 0.5757 | 0.051* 0.447 (6) |

Atomic displacement parameters (\AA^2)

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|-----|--------------|--------------|--------------|--------------|--------------|--------------|
| Pt1 | 0.01831 (12) | 0.01521 (12) | 0.01768 (12) | -0.00046 (5) | -0.00049 (7) | -0.00152 (5) |
| C11 | 0.0202 (5) | 0.0217 (5) | 0.0284 (5) | -0.0019 (4) | 0.0007 (4) | 0.0000 (4) |
| N1 | 0.030 (2) | 0.0149 (18) | 0.0188 (19) | 0.0005 (13) | 0.0009 (16) | -0.0026 (13) |
| N2 | 0.023 (2) | 0.0140 (18) | 0.0207 (19) | 0.0014 (13) | -0.0033 (15) | 0.0005 (12) |
| N3 | 0.0180 (17) | 0.0145 (16) | 0.0199 (17) | -0.0012 (14) | -0.0036 (14) | 0.0012 (14) |
| C1 | 0.0421 (19) | 0.0285 (16) | 0.0320 (16) | 0.0018 (14) | -0.0020 (14) | 0.0033 (13) |
| C2 | 0.0421 (19) | 0.0285 (16) | 0.0320 (16) | 0.0018 (14) | -0.0020 (14) | 0.0033 (13) |
| C3 | 0.052 (3) | 0.020 (2) | 0.022 (2) | -0.005 (2) | -0.006 (2) | 0.0014 (19) |
| C4 | 0.058 (3) | 0.024 (2) | 0.022 (2) | -0.006 (2) | 0.006 (2) | -0.0009 (19) |
| C5 | 0.035 (3) | 0.023 (2) | 0.025 (2) | -0.0054 (19) | 0.0051 (19) | -0.0019 (18) |
| C6 | 0.022 (2) | 0.0164 (19) | 0.0178 (19) | 0.0026 (16) | -0.0045 (16) | -0.0031 (15) |
| C7 | 0.024 (2) | 0.0129 (19) | 0.026 (2) | 0.0012 (16) | -0.0086 (17) | -0.0027 (16) |
| C8 | 0.017 (2) | 0.0133 (19) | 0.026 (2) | 0.0001 (16) | -0.0061 (16) | -0.0029 (16) |
| C9 | 0.018 (2) | 0.016 (2) | 0.025 (2) | 0.0008 (16) | -0.0044 (16) | -0.0028 (16) |
| C10 | 0.018 (2) | 0.0146 (19) | 0.0185 (19) | 0.0003 (15) | -0.0031 (15) | -0.0009 (15) |
| C11 | 0.0162 (19) | 0.0140 (19) | 0.023 (2) | -0.0018 (15) | -0.0022 (16) | -0.0024 (15) |
| C12 | 0.020 (2) | 0.0142 (19) | 0.024 (2) | -0.0014 (16) | -0.0033 (16) | -0.0012 (16) |
| C13 | 0.022 (2) | 0.020 (2) | 0.019 (2) | 0.0005 (16) | -0.0052 (16) | -0.0010 (16) |
| C14 | 0.025 (2) | 0.021 (2) | 0.023 (2) | -0.0040 (18) | -0.0066 (17) | -0.0028 (17) |
| C15 | 0.021 (2) | 0.019 (2) | 0.024 (2) | -0.0023 (16) | -0.0040 (18) | -0.0015 (16) |
| C16 | 0.085 (5) | 0.024 (3) | 0.018 (2) | -0.020 (3) | -0.005 (3) | 0.000 (2) |
| C17 | 0.085 (5) | 0.030 (3) | 0.025 (3) | -0.018 (3) | 0.009 (3) | -0.006 (2) |
| C18 | 0.098 (6) | 0.037 (3) | 0.037 (3) | -0.009 (3) | -0.031 (4) | 0.002 (3) |
| C19 | 0.150 (8) | 0.052 (4) | 0.016 (3) | -0.061 (5) | 0.006 (4) | 0.000 (3) |
| C20 | 0.017 (2) | 0.019 (2) | 0.036 (2) | -0.0056 (17) | -0.0070 (18) | -0.0045 (18) |
| C21 | 0.023 (2) | 0.060 (4) | 0.032 (3) | -0.010 (2) | -0.003 (2) | 0.005 (2) |
| C22 | 0.038 (3) | 0.020 (3) | 0.124 (7) | -0.009 (2) | -0.016 (4) | -0.002 (3) |
| C23 | 0.021 (2) | 0.057 (4) | 0.031 (3) | -0.009 (2) | -0.005 (2) | -0.010 (2) |
| C24 | 0.031 (2) | 0.024 (2) | 0.021 (2) | -0.0055 (18) | -0.0064 (18) | 0.0030 (18) |
| C25 | 0.034 (3) | 0.042 (3) | 0.019 (2) | -0.008 (2) | -0.0038 (19) | 0.000 (2) |
| C26 | 0.036 (3) | 0.038 (3) | 0.021 (2) | -0.006 (2) | -0.0103 (19) | 0.006 (2) |
| C27 | 0.051 (3) | 0.026 (3) | 0.027 (2) | -0.004 (2) | -0.015 (2) | 0.010 (2) |
| Cl2 | 0.0288 (6) | 0.0344 (6) | 0.0309 (6) | -0.0016 (5) | -0.0036 (4) | 0.0147 (5) |
| C1A | 0.078 (8) | 0.071 (8) | 0.051 (6) | -0.039 (8) | -0.012 (6) | 0.013 (5) |
| C2A | 0.108 (12) | 0.087 (10) | 0.047 (6) | -0.017 (9) | -0.014 (7) | -0.031 (6) |
| C3A | 0.049 (7) | 0.065 (7) | 0.090 (11) | -0.009 (5) | 0.001 (6) | -0.052 (8) |
| C4A | 0.022 (4) | 0.087 (12) | 0.043 (6) | 0.032 (5) | 0.017 (4) | 0.013 (7) |
| C5A | 0.042 (6) | 0.052 (6) | 0.055 (6) | 0.007 (4) | 0.007 (5) | 0.009 (5) |
| C6A | 0.042 (6) | 0.049 (6) | 0.086 (9) | -0.014 (4) | 0.018 (6) | -0.024 (5) |
| C7A | 0.0421 (19) | 0.0285 (16) | 0.0320 (16) | 0.0018 (14) | -0.0020 (14) | 0.0033 (13) |
| C1B | 0.078 (8) | 0.071 (8) | 0.051 (6) | -0.039 (8) | -0.012 (6) | 0.013 (5) |
| C2B | 0.108 (12) | 0.087 (10) | 0.047 (6) | -0.017 (9) | -0.014 (7) | -0.031 (6) |
| C3B | 0.049 (7) | 0.065 (7) | 0.090 (11) | -0.009 (5) | 0.001 (6) | -0.052 (8) |
| C4B | 0.022 (4) | 0.087 (12) | 0.043 (6) | 0.032 (5) | 0.017 (4) | 0.013 (7) |
| C5B | 0.042 (6) | 0.052 (6) | 0.055 (6) | 0.007 (4) | 0.007 (5) | 0.009 (5) |

| | | | | | | |
|-----|-------------|-------------|-------------|-------------|--------------|-------------|
| C6B | 0.042 (6) | 0.049 (6) | 0.086 (9) | -0.014 (4) | 0.018 (6) | -0.024 (5) |
| C7B | 0.0421 (19) | 0.0285 (16) | 0.0320 (16) | 0.0018 (14) | -0.0020 (14) | 0.0033 (13) |

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

| | | | |
|----------------------|-------------|----------|-----------|
| Pt1—N2 | 1.941 (4) | C21—H21A | 0.9800 |
| Pt1—N3 | 2.013 (4) | C21—H21B | 0.9800 |
| Pt1—N1 | 2.015 (4) | C21—H21C | 0.9800 |
| Pt1—Cl1 | 2.3070 (10) | C22—H22A | 0.9800 |
| Pt1—Pt1 ⁱ | 3.2774 (3) | C22—H22B | 0.9800 |
| N1—C5 | 1.333 (6) | C22—H22C | 0.9800 |
| N1—C1 | 1.371 (7) | C23—H23A | 0.9800 |
| N2—C10 | 1.340 (5) | C23—H23B | 0.9800 |
| N2—C6 | 1.344 (5) | C23—H23C | 0.9800 |
| N3—C15 | 1.338 (6) | C24—C27 | 1.528 (7) |
| N3—C11 | 1.381 (5) | C24—C26 | 1.529 (6) |
| C1—C2 | 1.381 (7) | C24—C25 | 1.536 (7) |
| C1—C6 | 1.483 (7) | C25—H25A | 0.9800 |
| C2—C3 | 1.408 (8) | C25—H25B | 0.9800 |
| C2—H2A | 0.9500 | C25—H25C | 0.9800 |
| C3—C4 | 1.402 (8) | C26—H26A | 0.9800 |
| C3—C16 | 1.531 (7) | C26—H26B | 0.9800 |
| C4—C5 | 1.378 (7) | C26—H26C | 0.9800 |
| C4—H4A | 0.9500 | C27—H27A | 0.9800 |
| C5—H5A | 0.9500 | C27—H27B | 0.9800 |
| C6—C7 | 1.391 (6) | C27—H27C | 0.9800 |
| C7—C8 | 1.402 (6) | C1A—C2A | 1.3900 |
| C7—H7A | 0.9500 | C1A—C6A | 1.3900 |
| C8—C9 | 1.403 (6) | C1A—C7A | 1.465 (6) |
| C8—C20 | 1.544 (6) | C2A—C3A | 1.3900 |
| C9—C10 | 1.393 (6) | C2A—H2AA | 0.9500 |
| C9—H9A | 0.9500 | C3A—C4A | 1.3900 |
| C10—C11 | 1.490 (5) | C3A—H3AA | 0.9500 |
| C11—C12 | 1.365 (6) | C4A—C5A | 1.3900 |
| C12—C13 | 1.404 (6) | C4A—H4AA | 0.9500 |
| C12—H12A | 0.9500 | C5A—C6A | 1.3900 |
| C13—C14 | 1.398 (6) | C5A—H5AA | 0.9500 |
| C13—C24 | 1.534 (6) | C6A—H6AA | 0.9500 |
| C14—C15 | 1.375 (7) | C7A—H7AA | 0.9800 |
| C14—H14A | 0.9500 | C7A—H7AB | 0.9800 |
| C15—H15A | 0.9500 | C7A—H7AC | 0.9800 |
| C16—C18 | 1.511 (11) | C1B—C2B | 1.3900 |
| C16—C19 | 1.530 (8) | C1B—C6B | 1.3900 |
| C16—C17 | 1.541 (8) | C1B—C7B | 1.461 (6) |
| C17—H17A | 0.9800 | C2B—C3B | 1.3900 |
| C17—H17B | 0.9800 | C2B—H2BA | 0.9500 |
| C17—H17C | 0.9800 | C3B—C4B | 1.3900 |
| C18—H18A | 0.9800 | C3B—H3BA | 0.9500 |

| | | | |
|------------|-------------|---------------|-----------|
| C18—H18B | 0.9800 | C4B—C5B | 1.3900 |
| C18—H18C | 0.9800 | C4B—H4BA | 0.9500 |
| C19—H19A | 0.9800 | C5B—C6B | 1.3900 |
| C19—H19B | 0.9800 | C5B—H5BA | 0.9500 |
| C19—H19C | 0.9800 | C6B—H6BA | 0.9500 |
| C20—C22 | 1.529 (7) | C7B—H7BA | 0.9800 |
| C20—C21 | 1.530 (7) | C7B—H7BB | 0.9800 |
| C20—C23 | 1.532 (7) | C7B—H7BC | 0.9800 |
| | | | |
| N2—Pt1—N3 | 80.87 (14) | C22—C20—C8 | 106.3 (4) |
| N2—Pt1—N1 | 81.25 (15) | C21—C20—C8 | 110.2 (4) |
| N3—Pt1—N1 | 162.06 (16) | C23—C20—C8 | 110.8 (4) |
| N2—Pt1—Cl1 | 178.70 (11) | C20—C21—H21A | 109.5 |
| N3—Pt1—Cl1 | 99.11 (10) | C20—C21—H21B | 109.5 |
| N1—Pt1—Cl1 | 98.72 (12) | H21A—C21—H21B | 109.5 |
| C5—N1—C1 | 119.1 (4) | C20—C21—H21C | 109.5 |
| C5—N1—Pt1 | 127.4 (3) | H21A—C21—H21C | 109.5 |
| C1—N1—Pt1 | 113.4 (3) | H21B—C21—H21C | 109.5 |
| C10—N2—C6 | 123.7 (4) | C20—C22—H22A | 109.5 |
| C10—N2—Pt1 | 118.5 (3) | C20—C22—H22B | 109.5 |
| C6—N2—Pt1 | 117.8 (3) | H22A—C22—H22B | 109.5 |
| C15—N3—C11 | 118.5 (4) | C20—C22—H22C | 109.5 |
| C15—N3—Pt1 | 127.4 (3) | H22A—C22—H22C | 109.5 |
| C11—N3—Pt1 | 114.0 (3) | H22B—C22—H22C | 109.5 |
| N1—C1—C2 | 121.2 (5) | C20—C23—H23A | 109.5 |
| N1—C1—C6 | 114.4 (4) | C20—C23—H23B | 109.5 |
| C2—C1—C6 | 124.4 (5) | H23A—C23—H23B | 109.5 |
| C1—C2—C3 | 120.2 (5) | C20—C23—H23C | 109.5 |
| C1—C2—H2A | 119.9 | H23A—C23—H23C | 109.5 |
| C3—C2—H2A | 119.9 | H23B—C23—H23C | 109.5 |
| C4—C3—C2 | 116.8 (5) | C27—C24—C26 | 108.7 (4) |
| C4—C3—C16 | 122.9 (5) | C27—C24—C13 | 107.4 (4) |
| C2—C3—C16 | 120.2 (5) | C26—C24—C13 | 112.0 (4) |
| C5—C4—C3 | 120.3 (5) | C27—C24—C25 | 109.0 (4) |
| C5—C4—H4A | 119.9 | C26—C24—C25 | 110.1 (4) |
| C3—C4—H4A | 119.9 | C13—C24—C25 | 109.6 (4) |
| N1—C5—C4 | 122.3 (5) | C24—C25—H25A | 109.5 |
| N1—C5—H5A | 118.8 | C24—C25—H25B | 109.5 |
| C4—C5—H5A | 118.8 | H25A—C25—H25B | 109.5 |
| N2—C6—C7 | 119.1 (4) | C24—C25—H25C | 109.5 |
| N2—C6—C1 | 113.1 (4) | H25A—C25—H25C | 109.5 |
| C7—C6—C1 | 127.8 (4) | H25B—C25—H25C | 109.5 |
| C6—C7—C8 | 119.3 (4) | C24—C26—H26A | 109.5 |
| C6—C7—H7A | 120.3 | C24—C26—H26B | 109.5 |
| C8—C7—H7A | 120.3 | H26A—C26—H26B | 109.5 |
| C7—C8—C9 | 119.3 (4) | C24—C26—H26C | 109.5 |
| C7—C8—C20 | 120.6 (4) | H26A—C26—H26C | 109.5 |
| C9—C8—C20 | 120.1 (4) | H26B—C26—H26C | 109.5 |

| | | | |
|---------------|-----------|---------------|-----------|
| C10—C9—C8 | 119.1 (4) | C24—C27—H27A | 109.5 |
| C10—C9—H9A | 120.5 | C24—C27—H27B | 109.5 |
| C8—C9—H9A | 120.5 | H27A—C27—H27B | 109.5 |
| N2—C10—C9 | 119.4 (4) | C24—C27—H27C | 109.5 |
| N2—C10—C11 | 112.9 (4) | H27A—C27—H27C | 109.5 |
| C9—C10—C11 | 127.7 (4) | H27B—C27—H27C | 109.5 |
| C12—C11—N3 | 121.2 (4) | C2A—C1A—C6A | 120.0 |
| C12—C11—C10 | 125.2 (4) | C2A—C1A—C7A | 124.9 (5) |
| N3—C11—C10 | 113.6 (4) | C6A—C1A—C7A | 115.1 (5) |
| C11—C12—C13 | 120.8 (4) | C3A—C2A—C1A | 120.0 |
| C11—C12—H12A | 119.6 | C3A—C2A—H2AA | 120.0 |
| C13—C12—H12A | 119.6 | C1A—C2A—H2AA | 120.0 |
| C14—C13—C12 | 116.6 (4) | C4A—C3A—C2A | 120.0 |
| C14—C13—C24 | 122.7 (4) | C4A—C3A—H3AA | 120.0 |
| C12—C13—C24 | 120.7 (4) | C2A—C3A—H3AA | 120.0 |
| C15—C14—C13 | 120.5 (4) | C3A—C4A—C5A | 120.0 |
| C15—C14—H14A | 119.7 | C3A—C4A—H4AA | 120.0 |
| C13—C14—H14A | 119.7 | C5A—C4A—H4AA | 120.0 |
| N3—C15—C14 | 122.2 (4) | C4A—C5A—C6A | 120.0 |
| N3—C15—H15A | 118.9 | C4A—C5A—H5AA | 120.0 |
| C14—C15—H15A | 118.9 | C6A—C5A—H5AA | 120.0 |
| C18—C16—C19 | 109.3 (6) | C5A—C6A—C1A | 120.0 |
| C18—C16—C3 | 109.6 (5) | C5A—C6A—H6AA | 120.0 |
| C19—C16—C3 | 111.1 (5) | C1A—C6A—H6AA | 120.0 |
| C18—C16—C17 | 110.2 (5) | C2B—C1B—C6B | 120.0 |
| C19—C16—C17 | 109.3 (6) | C2B—C1B—C7B | 120.3 (5) |
| C3—C16—C17 | 107.4 (4) | C6B—C1B—C7B | 119.7 (5) |
| C16—C17—H17A | 109.5 | C1B—C2B—C3B | 120.0 |
| C16—C17—H17B | 109.5 | C1B—C2B—H2BA | 120.0 |
| H17A—C17—H17B | 109.5 | C3B—C2B—H2BA | 120.0 |
| C16—C17—H17C | 109.5 | C4B—C3B—C2B | 120.0 |
| H17A—C17—H17C | 109.5 | C4B—C3B—H3BA | 120.0 |
| H17B—C17—H17C | 109.5 | C2B—C3B—H3BA | 120.0 |
| C16—C18—H18A | 109.5 | C3B—C4B—C5B | 120.0 |
| C16—C18—H18B | 109.5 | C3B—C4B—H4BA | 120.0 |
| H18A—C18—H18B | 109.5 | C5B—C4B—H4BA | 120.0 |
| C16—C18—H18C | 109.5 | C4B—C5B—C6B | 120.0 |
| H18A—C18—H18C | 109.5 | C4B—C5B—H5BA | 120.0 |
| H18B—C18—H18C | 109.5 | C6B—C5B—H5BA | 120.0 |
| C16—C19—H19A | 109.5 | C5B—C6B—C1B | 120.0 |
| C16—C19—H19B | 109.5 | C5B—C6B—H6BA | 120.0 |
| H19A—C19—H19B | 109.5 | C1B—C6B—H6BA | 120.0 |
| C16—C19—H19C | 109.5 | C1B—C7B—H7BA | 109.5 |
| H19A—C19—H19C | 109.5 | C1B—C7B—H7BB | 109.5 |
| H19B—C19—H19C | 109.5 | H7BA—C7B—H7BB | 109.5 |
| C22—C20—C21 | 109.1 (5) | C1B—C7B—H7BC | 109.5 |

| | | | |
|-------------|-----------|---------------|-------|
| C22—C20—C23 | 111.5 (5) | H7BA—C7B—H7BC | 109.5 |
| C21—C20—C23 | 108.9 (4) | H7BB—C7B—H7BC | 109.5 |

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