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4,6-Dimethyl-2-*p*-tolylpyrimidine

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Key indicators: single-crystal X-ray study; T = 296 K; mean σ (C–C) = 0.002 Å; *R* factor = 0.042; *wR* factor = 0.131; data-to-parameter ratio = 14.0.

The molecule of the title compound, $C_{13}H_{14}N_2$, is located on a crystallographic mirror plane. The aromatic rings make a dihedral angle of 3.4 (2)°. The H atoms of the methyl groups on the benzene ring are disordered over two positions; their site-occupation factors were fixed at 0.5. In the crystal, intermolecular $C-H\cdots\pi$ contacts form infinite chains perpendicular to the *b* axis.

Related literature

The title compound was derived from the reaction of *p*-tolylmercutic chlorides and 4,6-dimethyl-2-iodopyrimidine. For general background to theuse of organomercury compounds in cross-coupling reactions, see: Beletskaya *et al.* (2001); Braga *et al.* (2004). For a related structure, see: Santoni *et al.* (2008). For the synthesis, see: Xu *et al.* (2009*a*,*b*).



Experimental

Crystal data C₁₃H₁₄N₂

 $M_r = 198.26$

organic compounds

Z = 4

Mo $K\alpha$ radiation

 $0.35 \times 0.25 \times 0.22$ mm

7934 measured reflections 1089 independent reflections 777 reflections with $I > 2\sigma(I)$

 $\mu = 0.07 \text{ mm}^{-1}$

T = 296 K

 $R_{\rm int} = 0.025$

Orthorhombic, *Pnma* a = 7.2086 (10) Å b = 12.4668 (18) Å c = 12.4335 (18) Å $V = 1117.4 (3) \text{ Å}^3$

Data collection

| Bruker SMART APEX CCD area- |
|--|
| detector diffractometer |
| Absorption correction: multi-scan |
| (SADABS; Sheldrick, 1996) |
| $T_{\rm min} = 0.976, T_{\rm max} = 0.985$ |

Refinement

 $R[F^2 > 2\sigma(F^2)] = 0.042$ 78 parameters $wR(F^2) = 0.131$ H-atom parameters constrainedS = 1.06 $\Delta \rho_{max} = 0.19$ e Å⁻³1089 reflections $\Delta \rho_{min} = -0.14$ e Å⁻³

Table 1

Hydrogen-bond geometry (Å, °).

| $D - H \cdot \cdot \cdot A$ | D-H | $H \cdot \cdot \cdot A$ | $D \cdot \cdot \cdot A$ | $D - \mathbf{H} \cdots A$ |
|-----------------------------|------|-------------------------|-------------------------|---------------------------|
| $C8-H8\cdots Cg1^i$ | 0.93 | 2.79 | 3.638 (2) | 152 |
| | . 1 | | | |

Symmetry code: (i) -x - 1, $y + \frac{1}{2}$, -z. Cg1 is the centroid of the pyrimidine ring.

Data collection: *SMART* (Bruker, 2004); cell refinement: *SAINT* (Bruker, 2004); 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: *SHELXL97* and *PLATON* (Spek, 2009).

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Supplementary data and figures for this paper are available from the IUCr electronic archives (Reference: SI2211).

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supporting information

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4,6-Dimethyl-2-p-tolylpyrimidine

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S1. Comment

The organomercury compounds have a number of notable advantages over other organometallic compounds commonly used in cross-coupling reactions, including higher selectivity of reactions, extra stability and easy availability by a direct mercuration (Beletskaya *et al.*, 2001; Braga *et al.*, 2004). We have recently reported ferrocene-heterocycles were obtained from the coupling reaction(Xu *et al.*, 2009*a*,b). Here we report the crystal structure of the title compound, derived from the reaction of *p*-tolylmercutic chlorides and 4,6-dimethyl-2-iodopyrimidine.

Due to the molecular mirror symmetry m of the title compound (Fig.1), and coincidence with the crystallographic mirror plane m (space group *Pnma*), the atoms C1, C2, C5, C8, H8 are half occupied and the H atoms of the methyl groups in the benzene ring are disordered over two positions; their site-occupation factors were fixed at 0.5. The aromatic rings have very small angles between their planes (dihedral angle is 3.4 (2)°) due to the absence of H—H repulsion (Santoni *et al.*, 2008). Fig.2 shows that in the crystal there exist intermolecular C—H… π interactions (Table 1, *Cg*1 is the centroid of the pyrimidine ring).

S2. Experimental

The title compound was obtained from the coupling reaction of *p*-tolylmercutic chlorides and 4,6-dimethyl-2-iodopyrimidine as described in literature (Xu *et al.*, 2009*b*) and recrystallized from ethanol at room temperature to give the desired crystals suitable for single-crystal X-ray diffraction.

S3. Refinement

H atoms attached to C atoms of the title compound were placed in geometrically idealized positions and treated as riding with C—H distances constrained to 0.93–0.96 Å, and with $U_{iso}(H)=1.2-1.5U_{eq}(C)$.



Figure 1

The molecular structure of the title compound with displacement ellipsoids at the 30% probability level, the disordered H atoms are omitted (Symmetry code A: -x + 2, -y, -z).



Figure 2

Partial view of the crystal packing showing the formation of the infinite chain of molecules formed by the C—H $\cdots\pi$ interactions.

4,6-Dimethyl-2-p-tolylpyrimidine

Crystal data

 $C_{13}H_{14}N_2$ $M_r = 198.26$ Orthorhombic, *Pnma* a = 7.2086 (10) Å b = 12.4668 (18) Å c = 12.4335 (18) Å $V = 1117.4 (3) \text{ Å}^3$ Z = 4F(000) = 424 $D_x = 1.179 \text{ Mg m}^{-3}$ Mo K α radiation, $\lambda = 0.71073 \text{ Å}$ Cell parameters from 1634 reflections $\theta = 2.3-23.3^{\circ}$ $\mu = 0.07 \text{ mm}^{-1}$ T = 296 KBlock, colourless $0.35 \times 0.25 \times 0.22 \text{ mm}$ Data collection

| Bruker SMART APEX CCD area-detector diffractometer | 7934 measured reflections 1089 independent reflections |
|---|--|
| Radiation source: fine-focus sealed tube | 777 reflections with $I > 2\sigma(I)$ |
| Graphite monochromator | $R_{\rm int} = 0.025$ |
| phi and ω scans | $\theta_{\rm max} = 25.5^\circ, \theta_{\rm min} = 2.3^\circ$ |
| Absorption correction: multi-scan | $h = -8 \rightarrow 8$ |
| (SADABS; Sheldrick, 1996) | $k = -15 \rightarrow 15$ |
| $T_{\min} = 0.976, \ T_{\max} = 0.985$ | $l = -15 \rightarrow 14$ |
| Refinement | |
| Refinement on F^2 | Secondary atom site location: difference Fourier |
| Least-squares matrix: full | map |
| $R[F^2 > 2\sigma(F^2)] = 0.042$ | Hydrogen site location: inferred from |
| $wR(F^2) = 0.131$ | neighbouring sites |
| S = 1.06 | H-atom parameters constrained |
| 1089 reflections | $w = 1/[\sigma^2(F_o^2) + (0.0557P)^2 + 0.291P]$ |
| 78 parameters | where $P = (F_o^2 + 2F_c^2)/3$ |

78 parameterswhere $P = (F_o^2 + 0)^2$ 0 restraints $(\Delta/\sigma)_{max} < 0.001$ Primary atom site location: structure-invariant $\Delta \rho_{max} = 0.19 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{min} = -0.14 \text{ e} \text{ Å}^{-3}$ $\Delta \rho_{min} = -0.14 \text{ e} \text{ Å}^{-3}$

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*, and *R*-factors based on ALL data will be even larger.

Fractional atomic coordinates and isotropic or equivalent isotropic displacement parameters $(Å^2)$

| | x | У | Ζ | $U_{ m iso}$ */ $U_{ m eq}$ | Occ. (<1) |
|-----|-------------|--------------|--------------|-----------------------------|-----------|
| C1 | 0.8988 (4) | 0.2500 | 0.6728 (2) | 0.0709 (8) | |
| H1A | 0.8781 | 0.2870 | 0.7394 | 0.106* | 0.50 |
| H1B | 0.9952 | 0.2856 | 0.6332 | 0.106* | 0.50 |
| H1C | 0.9355 | 0.1774 | 0.6874 | 0.106* | 0.50 |
| C2 | 0.7231 (3) | 0.2500 | 0.60770 (17) | 0.0499 (6) | |
| C3 | 0.6383 (2) | 0.34507 (13) | 0.57708 (13) | 0.0544 (5) | |
| H3 | 0.6934 | 0.4100 | 0.5953 | 0.065* | |
| C4 | 0.4741 (2) | 0.34546 (12) | 0.52012 (13) | 0.0522 (5) | |
| H4 | 0.4201 | 0.4105 | 0.5009 | 0.063* | |
| C5 | 0.3884 (3) | 0.2500 | 0.49107 (16) | 0.0449 (5) | |
| C6 | 0.2078 (3) | 0.2500 | 0.43367 (17) | 0.0468 (5) | |
| C7 | -0.0325 (2) | 0.34504 (13) | 0.36089 (13) | 0.0529 (5) | |
| | | | | | |

supporting information

| C8 | -0.1188 (3) | 0.2500 | 0.33466 (18) | 0.0549 (6) |
|-----|--------------|--------------|--------------|------------|
| H8 | -0.2331 | 0.2500 | 0.3000 | 0.066* |
| C9 | -0.1170 (3) | 0.45227 (14) | 0.33570 (16) | 0.0732 (6) |
| H9A | -0.0974 | 0.5001 | 0.3951 | 0.110* |
| H9B | -0.2477 | 0.4438 | 0.3235 | 0.110* |
| H9C | -0.0598 | 0.4814 | 0.2724 | 0.110* |
| N1 | 0.13292 (18) | 0.34589 (10) | 0.41062 (10) | 0.0509 (4) |

Atomic displacement parameters $(Å^2)$

| | U^{11} | U^{22} | U^{33} | U^{12} | U^{13} | U^{23} |
|----|-------------|-------------|-------------|-------------|--------------|-------------|
| C1 | 0.0703 (17) | 0.0703 (18) | 0.0720 (17) | 0.000 | -0.0169 (14) | 0.000 |
| C2 | 0.0541 (14) | 0.0531 (13) | 0.0425 (11) | 0.000 | -0.0007 (10) | 0.000 |
| C3 | 0.0600 (11) | 0.0446 (9) | 0.0584 (10) | -0.0043 (8) | -0.0036 (8) | -0.0048 (7) |
| C4 | 0.0588 (10) | 0.0390 (9) | 0.0589 (10) | 0.0022 (7) | -0.0028 (8) | 0.0002 (7) |
| C5 | 0.0507 (12) | 0.0407 (11) | 0.0433 (11) | 0.000 | 0.0027 (10) | 0.000 |
| C6 | 0.0545 (13) | 0.0430 (12) | 0.0428 (11) | 0.000 | 0.0018 (10) | 0.000 |
| C7 | 0.0550 (10) | 0.0541 (10) | 0.0496 (9) | 0.0045 (8) | 0.0009 (7) | 0.0034 (7) |
| C8 | 0.0510 (13) | 0.0608 (15) | 0.0528 (13) | 0.000 | -0.0047 (11) | 0.000 |
| C9 | 0.0702 (12) | 0.0598 (12) | 0.0896 (14) | 0.0091 (10) | -0.0128 (10) | 0.0094 (10) |
| N1 | 0.0546 (8) | 0.0449 (8) | 0.0532 (8) | 0.0031 (6) | -0.0025(6) | 0.0021 (6) |

Geometric parameters (Å, °)

| C1—C2 | 1.503 (3) | C5—C6 | 1.485 (3) | |
|------------------------|-------------|------------------------|-------------|--|
| C1—H1A | 0.9600 | C6—N1 ⁱ | 1.3426 (16) | |
| C1—H1B | 0.9600 | C6—N1 | 1.3426 (16) | |
| C1—H1C | 0.9600 | C7—N1 | 1.343 (2) | |
| C2-C3 ⁱ | 1.387 (2) | C7—C8 | 1.378 (2) | |
| С2—С3 | 1.387 (2) | С7—С9 | 1.502 (2) | |
| C3—C4 | 1.380 (2) | $C8$ — $C7^{i}$ | 1.378 (2) | |
| С3—Н3 | 0.9300 | C8—H8 | 0.9300 | |
| C4—C5 | 1.3885 (19) | С9—Н9А | 0.9600 | |
| C4—H4 | 0.9300 | С9—Н9В | 0.9600 | |
| $C5-C4^i$ | 1.3885 (19) | С9—Н9С | 0.9600 | |
| | | | | |
| C2—C1—H1A | 109.5 | C4 ⁱ —C5—C6 | 121.00 (11) | |
| C2—C1—H1B | 109.5 | N1 ⁱ —C6—N1 | 125.8 (2) | |
| H1A—C1—H1B | 109.5 | N1 ⁱ —C6—C5 | 117.07 (10) | |
| C2—C1—H1C | 109.5 | N1—C6—C5 | 117.07 (10) | |
| H1A—C1—H1C | 109.5 | N1—C7—C8 | 121.12 (16) | |
| H1B—C1—H1C | 109.5 | N1—C7—C9 | 116.67 (15) | |
| C3 ⁱ —C2—C3 | 117.4 (2) | C8—C7—C9 | 122.20 (16) | |
| C3 ⁱ —C2—C1 | 121.28 (11) | $C7$ — $C8$ — $C7^i$ | 118.7 (2) | |
| C3—C2—C1 | 121.28 (11) | С7—С8—Н8 | 120.7 | |
| C4—C3—C2 | 121.48 (16) | C7 ⁱ —C8—H8 | 120.7 | |
| С4—С3—Н3 | 119.3 | С7—С9—Н9А | 109.5 | |
| С2—С3—Н3 | 119.3 | С7—С9—Н9В | 109.5 | |
| | | | | |

| C3—C4—C5 | 120.81 (16) | H9A—C9—H9B | 109.5 |
|-----------------------------------|--------------|---------------------------|--------------|
| C3—C4—H4 | 119.6 | С7—С9—Н9С | 109.5 |
| С5—С4—Н4 | 119.6 | Н9А—С9—Н9С | 109.5 |
| $C4$ — $C5$ — $C4^i$ | 118.0 (2) | H9B—C9—H9C | 109.5 |
| C4—C5—C6 | 121.00 (11) | C6—N1—C7 | 116.62 (15) |
| | | | |
| C3 ⁱ —C2—C3—C4 | 1.2 (3) | C4 ⁱ —C5—C6—N1 | 178.65 (17) |
| C1—C2—C3—C4 | -178.02 (19) | $N1-C7-C8-C7^{i}$ | 0.3 (3) |
| C2—C3—C4—C5 | -0.3 (3) | C9—C7—C8—C7 ⁱ | -179.77 (14) |
| C3-C4-C5-C4 ⁱ | -0.7 (3) | N1 ⁱ C6N1C7 | 0.6 (3) |
| C3—C4—C5—C6 | 177.47 (16) | C5—C6—N1—C7 | -178.53 (15) |
| C4C5C6N1 ⁱ | -178.65 (17) | C8—C7—N1—C6 | -0.4 (3) |
| $C4^{i}$ — $C5$ — $C6$ — $N1^{i}$ | -0.6 (3) | C9—C7—N1—C6 | 179.62 (16) |
| C4—C5—C6—N1 | 0.6 (3) | | |
| | | | |

Symmetry code: (i) x, -y+1/2, z.

Hydrogen-bond geometry (Å, °)

| D—H···A | <i>D</i> —Н | Н…А | D····A | <i>D</i> —H··· <i>A</i> |
|-------------|-------------|------|-----------|-------------------------|
| С8—Н8…Сg1іі | 0.93 | 2.79 | 3.638 (2) | 152 |

Symmetry code: (ii) -x-1, y+1/2, -z.