

1,3,5-Tris(chloromethyl)benzene

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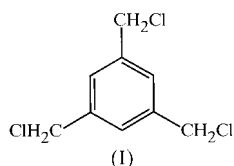
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The title compound, C₉H₉Cl₃, is being used as a platform for new tripodal receptors. Two molecules make up the asymmetric unit; weak intermolecular hydrogen bonding is observed between methylene H atoms and the chlorine of an adjacent molecule. There are also C—H... π interactions.

Comment

Two molecules make up the asymmetric unit of the title compound, (I). Weak intermolecular hydrogen bonding is observed (full details in Table 1) between a methylene H atom H18B and chlorine Cl4 of an adjacent molecule. There are also C—H... π interactions with H...ring-centroid separations of 3.07–3.31 Å (Table 1).



Experimental

The title compound was prepared following literature methods (Cochrane *et al.*, 1968) and was recrystallized from toluene.

Crystal data

C₉H₉Cl₃
M_r = 223.51
Triclinic, P1
a = 8.7124 (12) Å
b = 9.0659 (5) Å
c = 12.7901 (15) Å
 α = 87.537 (8)°
 β = 86.992 (10)°
 γ = 87.873 (8)°
V = 1007.3 (2) Å³

Z = 4
D_x = 1.474 Mg m⁻³
Mo K α radiation
Cell parameters from 25 reflections
 θ = 10.4–14.9°
 μ = 0.85 mm⁻¹
T = 163 (2) K
Fragment, colourless
0.30 × 0.30 × 0.25 mm

Data collection

Nonius CAD-4 diffractometer
 ω - θ scans
Absorption correction: ψ scan
(Siemens, 1994)
T_{min} = 0.722, T_{max} = 0.809
6800 measured reflections
3943 independent reflections
3047 reflections with I > 2 σ (I)

R_{int} = 0.025
 θ_{max} = 26.0°
h = -9 → 10
k = -11 → 11
l = -15 → 15
3 standard reflections
frequency: 120 min
intensity decay: 1%

Refinement

Refinement on F²
R[F² > 2 σ (F²)] = 0.032
wR(F²) = 0.088
S = 1.041
3943 reflections
217 parameters
H-atom parameters constrained

w = 1/[\sigma²(F_o²) + (0.0424P)²
+ 0.2112P]
where P = (F_o² + 2F_c²)/3
(Δ/σ)_{max} = 0.001
 $\Delta\rho_{max}$ = 0.31 e Å⁻³
 $\Delta\rho_{min}$ = -0.34 e Å⁻³

Table 1

Hydrogen-bonding geometry (Å, °).

D—H...A	D—H	H...A	D...A	D—H...A
C18—H18B...Cl4 ⁱ	0.99	2.72	3.682 (2)	163
C7—H7A...CG1 ⁱⁱ	0.99	3.07	3.509 (2)	108
C7—H7B...CG1 ⁱⁱ	0.99	3.12	3.509 (2)	105
C17—H17A...CG2 ⁱⁱⁱ	0.99	3.31	3.729 (2)	107

Symmetry codes: (i) 1 - x, 2 - y, 1 - z; (ii) 1 - x, 2 - y, 2 - z; (iii) 1 - x, 1 - y, 1 - z.

Data collection: CAD-4-PC (Nonius, 1993); cell refinement: CAD-4-PC (Nonius, 1993); data reduction: XCAD4 (Harms, 1995); program(s) used to solve structure: SHELXS86 (Sheldrick, 1990); program(s) used to refine structure: SHELXL97 (Sheldrick, 1997); molecular graphics: SHELXTL (Siemens, 1994); software used to prepare material for publication: SHELXTL (Siemens, 1994).

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