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THE CRYSTAL AND MOLECULAR STRUCTURE OF BIS[1,2-BIS(DIPHENYLPHOSPHINO)ETHANE] DICHLORORUTHENIUM(II)

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The title compound, *trans*-[Ru(dppe)₂Cl₂], has $\overline{1}$ symmetry such that the chloride atoms are exactly *trans*. The coordination geometry is distorted octahedral owing, in part, to the restricted bite distance of the chelating dppe ligands. Important interatomic parameters are Ru-Cl 2.436(1), Ru-P(1), 2.389(1) Ru-P(2) 2.369(1) Å and P(1)-Ru-P(2) 82.1(1)°. The compound crystallizes as a dichloromethane solvate in the monoclinic space group C2/c with unit cell dimensions a = 27.995(3), b = 13.85(1), c = 13.265(3) Å, $\beta = 105.38(2)°$ and Z = 4. The structure was refined by a full-matrix least-squares procedure to final R = 0.038 for 3522 reflections with $I \ge 2.5\sigma(I)$.

Keywords: Ruthenium(II), diphosphine, X-ray structure.

INTRODUCTION

The crystal structure of the title compound, trans-[Ru(dppe)₂Cl₂], was determined as a part of a continuing investigation of metal phosphine complexes.¹⁻³ Particular interest in the present system arises as a result of our study of the photochemistry of derivatives based on trans-[Ru(dppe)₂Cl₂]. Further interest in these compounds is related to their potential use as catalysts; for example, trans-[Ru(dppe)₂Cl₂] has been shown to catalytically hydrogenate olefins.⁴

EXPERIMENTAL

Crystallography

Trans-[Ru(dppe)₂Cl₂] is a known compound⁵ and crystals suitable for the X-ray analysis were grown as a CH₂Cl₂ solvate by the slow evaporation of a solution of the compound. Intensity data for 5693 reflections were measured on an Enraf-Nonius CAD4F diffractometer with the use of graphite monochromatized MoKa radiation and the ω : 2 θ scan technique such that θ_{max} was 25°. The data were corrected for Lorentz and polarization effects and for absorption.⁶ There were 4390 unique reflections (R_{ama1} 0.019) and of these 3522 satisfied the $I \ge 2.5\sigma(I)$ criterion of observability. Crystal data and refinement details are given in Table I.

Formula	C ₅₃ H ₅₀ Cl ₄ P ₄ Ru			
M _r	1053.7			
Crystal system	monoclinic			
Space group	C2/c			
<i>a</i> , Å	27.995(3)			
b, Å	13.85(1)			
c, Å	13.265(3)			
β(°)	105.38(4)			
Vol. (Å ³)	4959(2)			
Ζ	4			
$D_{c} (g cm^{-3})$	1.411			
F(000)	2328			
μ, cm^{-1}	7.44			
Transmission factors (max./min.)	0.964; 0.785			
θ limits (°)	1.5-25.0			
No. of data collected	5693			
No. of unique data	4390			
No. of unique reflections used				
with $I \ge 2.5\sigma(I)$	3522			
R	0.038			
g	0.004			
R _w	0.043			
ρ_{max} (e Å ⁻³)	0.76			

TABLE I Crystal and refinement details for *trans*-[Ru(dppe)₂Cl₂].CH₂Cl₂.



FIGURE 1 Molecular structure and crystallographic numbering scheme for trans-[Ru(dppe)₂Cl₂].

The structure was solved from the interpretation of the Patterson synthesis and refined by a full-matrix least-squares procedure based on $F.^6$ Non-hydrogen atoms were refined with anisotropic thermal parameters and hydrogen atoms were included in the model at their calculated positions. The solvent CH_2Cl_2 molecule was found to

be disordered about a crystallographic 2-fold axis and was refined with three sites for the chloride atoms each with a fractional occupancy factor. Full details are given in Table S(5) of the supplementary material. Refinement with a weighting scheme of the form $w = [\sigma^2(F) + g(F)^2]^{-1}$ converged with final R 0.038, R_w 0.043, and g 0.004.

TABLE II

Fractional atomic coordinates (×10⁵ for Ru, P, Cl; ×10⁴ others) for trans-[Ru(dppe)₂Cl₂].

Atom	x/a	<i>y/b</i>	<i>z/c</i>
Ru	25000(-)	25000(-)	50000(-)
P(1)	18105(3)	33012(6)	38559(6)
P(2)	29487(3)	34740(6)	41012(7)
CI	24965(3)	10996(6)	38870(6)
C(11)	2052(1)	4302(3)	3184(3)
C(21)	2581(1)	4584(2)	3745(3)
C(121)	1381(1)	3901(2)	4487(3)
C(122)	1466(1)	4845(3)	4856(3)
C(123)	1153(2)	5278(3)	5388(3)
C(124)	762(2)	4777(4)	5546(4)
C(125)	671(2)	3844(4)	5178(4)
C(126)	979(1)	3399(3)	4665(3)
C(131)	1392(1)	2741(3)	2697(3)
C(132)	1600(2)	2116(3)	2101(3)
C(133)	1300(2)	1683(3)	1215(3)
C(134)	807(2)	1866(4)	892(4)
C(135)	597(2)	2503(4)	1451(4)
C(136)	890(2)	2942(3)	2338(3)
C(221)	3566(1)	3927(2)	4756(3)
C(222)	3661(2)	4236(3)	5793(3)
C(223)	4144(2)	4495(4)	6342(4)
C(224)	4519(2)	4475(4)	5885(4)
C(225)	4433(2)	4184(4)	4873(5)
C(226)	3960(2)	3924(3)	4298(4)
C(231)	3038(1)	3058(3)	2845(3)
C(232)	2906(2)	3607(3)	1940(3)
C(233)	3012(3)	3283(4)	1043(4)
C(234)	3266(3)	2420(3)	1033(4)
C(235)	3398(2)	1877(3)	1930(3)
C(236)	3277(1)	2179(3)	2824(3)

Fractional atomic coordinates are listed in Table II and the numbering scheme employed is shown in Figure 1 (drawn at 25% probability ellipsoids with ORTEP⁷). The refinement was performed with the SHELX76⁹ program system installed on a SUN 4/280 computer system. Scattering factors for Ru (corrected for f' and f'') were the *International Tables for X-ray Crystallography*⁸ and those for the remaining atoms were those incorporated in SHELX76.⁶ Listings of thermal parameters, hydrogen atom parameters, all bond distances and angles, details of the disordered CH₂Cl₂ molecule and the observed and calculated structure factors are available from the Editor.

RESULTS AND DISCUSSION

The crystal structure determination confirms the stoichiometry of the compound and the *trans* disposition of the chloride atoms. A diagram of the molecular structure is shown in Figure 1 and selected interatomic parameters are given in Table III. The ruthenium atom is situated on a crystallographic centre of inversion (located at 1/4, 1/4,1/2) so that half the molecule comprises the asymmetric unit. In addition there is disordered CH₂Cl₂ in the crystal lattice as discussed in the Experimental section.

The ruthenium atom exists in a distorted octahedral geometry with the maximum deviation from the ideal geometry arising as a result of the restricted bite distance of the diphosphine ligand; P(1)-Ru-P(2) 82.1(1)°. The Ru-Cl bond distance of 2.436(1) Å lies within the narrow range of Ru-Cl bond distances found in related *trans*-ruthenium(diphosphine)₂Cl₂ complexes. Thus Ru-Cl distances of $2 \times 2.426(1)$ Å (molecule has 1 symmetry) has been reported for *trans*-[Ru(dppm)₂Cl₂]⁹ [dppm is bis(diphenylphosphino)methane] and Ru-Cl distances of 2.443(4) and 2.446(4) Å were found in *trans*-[Ru(PhMePCH₂CH₂PPhMe)₂Cl₂].¹⁰ The Ru-Cl distances in *trans*-[Ru(dppe)₂Cl₂] may also be compared to the Ru-Cl distances found in *trans*-[Ru(PHMe₂)₄Cl₂] of 2 × 2.440(1) Å (molecule has 1 symmetry).¹¹

TABLE III Selected bond distances (Å) and angles (°) for *trans*-[Ru(dppe)₂Cl₂].

Distance			
Distances			
Ru-Cl	2.436(1)	Ru-P(1)	2.389(1)
Ru-P(2)	2.369(1)	P(1)-C(11)	1.868(3)
P(1)-C(121)	1.836(3)	P(1)-C(131)	1.839(4)
P(2)-C(21)	1.841(3)	P(2)-C(221)	1.827(3)
P(2)-C(231)	1.841(4)	C(11)-C(21)	1.522(5)
Angles			
Cl-Ru-P(1)	96.4(1)	Cl-Ru-P(2)	93.8(1)
ClRuP(1')	83.6(1)	Cl-Ru-P(2')	86.2(1)
P(1)-Ru-P(2)	82.1(1)	P(1)-Ru-P(2')	97.9(1)
Ru-P(1)-C(11)	108.3(1)	Ru-P(1)-C(121)	115.9(1)
Ru-P(1)-C(131)	124.3(1)	C(11)-P(1)-C(121)	104.2(2)
C(11)-P(1)-C(131)	98.3(2)	C(121)-P(1)-C(131)	102.9(2)
Ru-P(2)-C(21)	105.7(1)	Ru-P(2)-C(221)	121.1(1)
Ru-P(2)-C(231)	119.6(1)	C(21)-P(1)-C(221)	103.2(1)
C(21)-P(1)-C(231)	103.3(2)	C(221)-P(1)-C(231)	101.5(2)
P(1)-C(11)-C(21)	112.7(2)	P(2)-C(21)-C(11)	108.4(2)

The dppe ligands in *trans*-[Ru(dppe)₂Cl₂] coordinate in a bidentate fashion, there being a slight asymmetry in the Ru-P bond distances: Ru-P(1) 2.389(1) and 2.369(1) Å. The Ru-P(1) distance is significantly longer than other Ru-P bonds in the complexes cited above whereas the Ru-P(2) bond distance, while at the upper limit of Ru-P bond distances, has a precedent. In *trans*-[Ru(dppm)₂Cl₂]⁹ the Ru-P bond distances are 2.340(1) and 2.367(1) Å and in *trans*-[Ru(PhMePCH₂CH₂PPhMe)₂Cl₂] the Ru-P bonds were found to be 2.319(4), $2 \times 2.346(4)$ and 2.361(4) Å.¹⁰ In the *trans*-[Ru(PH₂Me)₄Cl₂] compound the two unique Ru-P bond distances are at the lower end of the range at 2.323(1) and 2.331(1) Å.¹¹ Bond lengths and angles associated with

trans-[Ru(dppe)₂Cl₂]

the dppe ligand show no abnormal features. In conclusion there does not appear to be any systematic variation in the Ru-Cl and Ru-P bond distances as the nature of the diphosphine is varied; any differences that do occur in the ruthenium atom geometry may be related to the bite distance of the diphosphine.

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