

trans-Di- μ -acetato- κ^4 O:O'-bis[2-(5-phenylisoxazolin-3-yl)phenyl- κ^2 C¹,N]-dipalladium(II)

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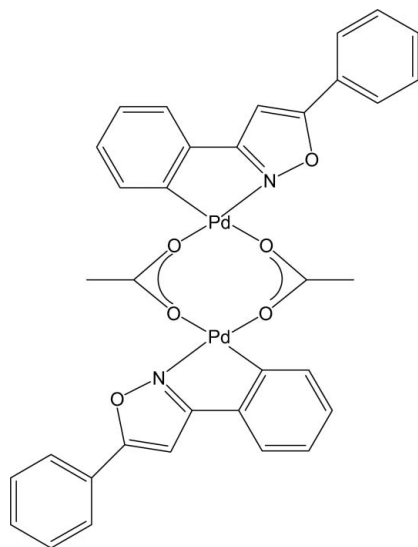
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Key indicators: single-crystal X-ray study; $T = 298$ K; mean $\sigma(\text{C}-\text{C}) = 0.013$ Å; R factor = 0.049; wR factor = 0.143; data-to-parameter ratio = 15.3.

The title compound, $[\text{Pd}_2(\text{C}_{15}\text{H}_{10}\text{NO})_2(\text{C}_2\text{H}_3\text{O}_2)_2]$, crystallized from a dichloromethane/*n*-hexane solution with two crystallographically independent dimeric molecules in the asymmetric unit. Each molecule may be described as a dimer with an *anti* configuration and the cyclometallated fragments in the characteristic open-book disposition, linked by two bridging acetate ligands.

Related literature

For a related palladacycle bridged by acetate ligands, see: Schultz *et al.* (2004). For related literature, see: Dupont *et al.* (2005).



Experimental

Crystal data

$[\text{Pd}_2(\text{C}_{15}\text{H}_{10}\text{NO})_2(\text{C}_2\text{H}_3\text{O}_2)_2]$
 $M_r = 771.37$

Monoclinic, $P2_1/c$
 $a = 14.8160$ (6) Å
 $b = 24.2339$ (10) Å
 $c = 19.6397$ (8) Å
 $\beta = 103.233$ (1)°

$V = 6864.4$ (5) Å³
 $Z = 8$

Mo $K\alpha$ radiation
 $\mu = 1.09$ mm⁻¹
 $T = 298$ (2) K
 $0.28 \times 0.20 \times 0.15$ mm

Data collection

Bruker SMART APEXII
diffractometer
Absorption correction: multi-scan
(*SADABS*; Sheldrick, 1996)
 $T_{\min} = 0.750$, $T_{\max} = 0.854$

80599 measured reflections
12098 independent reflections
8103 reflections with $I > 2\sigma(I)$
 $R_{\text{int}} = 0.038$

Refinement

$R[F^2 > 2\sigma(F^2)] = 0.049$
 $wR(F^2) = 0.143$
 $S = 1.09$
12098 reflections

793 parameters
H-atom parameters constrained
 $\Delta\rho_{\text{max}} = 0.81$ e Å⁻³
 $\Delta\rho_{\text{min}} = -0.58$ e Å⁻³

Data collection: *APEX2* (Bruker, 2005); cell refinement: *SAINT* (Bruker, 1997); data reduction: *SAINT*; program(s) used to solve structure: *SHELXS97* (Sheldrick, 2008); program(s) used to refine structure: *SHELXL97* (Sheldrick, 2008) and *PLATON* (Spek, 2003); molecular graphics: *SHELXTL* (Sheldrick, 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: CF2190).

References

- Bruker (1997). *SAINT*. Bruker AXS Inc., Madison, Wisconsin, USA.
Bruker (2005). *APEX2*. Bruker AXS Inc., Madison, Wisconsin, USA.
Dupont, J., Consorti, C. S. & Spencer, J. (2005). *Chem. Rev.* **105**, 2527–2571.
Schultz, T., Schmees, N. & Pfaltz, A. (2004). *Appl. Organomet. Chem.* **18**, 595–601.
Sheldrick, G. M. (1996). *SADABS*. University of Göttingen, Germany.
Sheldrick, G. M. (2008). *Acta Cryst.* **A64**, 112–122.
Spek, A. L. (2003). *J. Appl. Cryst.* **36**, 7–13.