

**Sigma Bond Metathesis with Pentamethylcyclopentadienyl Ligands in Sterically
Crowded (C₅Me₅)₃M Complexes**

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X-ray Data Collection, Structure Solution and Refinement for $[(C_5Me_5)_2La(SePh)]_2$, **1b**.

A colorless crystal of approximate dimensions 0.22 x 0.24 x 0.26 mm was mounted on a glass fiber and transferred to a Bruker SMART1K diffractometer. The SMART¹ program package was used to determine the unit-cell parameters and for data collection (25 sec/frame scan time for a sphere of diffraction data). The raw frame data was processed using SAINT² and SADABS³ to yield the reflection data file. Subsequent calculations were carried out using the SHELXTL⁴ program. The diffraction symmetry was *mmm* and the systematic absences were consistent with the orthorhombic space groups *Pnn2* and *Pnmm*. It was later determined that the noncentrosymmetric space group *Pnn2* was correct.

The structure was solved using the coordinates of the cerium sulfur analogue and refined on F^2 by full-matrix least-squares techniques. The analytical scattering factors⁵ for neutral atoms were used throughout the analysis. Hydrogen atoms were included using a riding model. The molecule is located on a two-fold rotation axis.

At convergence, $wR2 = 0.0460$ and $Goof = 1.057$ for 265 variables refined against 6015 data (0.75\AA), $R1 = 0.0187$ for those 5560 data with $I > 2.0\sigma(I)$. The structure was refined as a twin with the Flack⁶ parameter/BASF = 0.232(9).

¹ SMART Software Users Guide, Version 5.1, Bruker Analytical X-Ray Systems, Inc.; Madison, WI 1999.

² SAINT Version 6.36a, Bruker Analytical X-Ray Systems, Inc.; Madison, WI 1999.

³ Sheldrick, G. M. SADABS, Version 2008/1, Bruker Analytical X-Ray Systems, Inc.; Madison, WI 2008.

⁴ Sheldrick, G. M. SHELXTL Version 6.12, Bruker Analytical X-Ray Systems, Inc.; Madison, WI 2001.

⁵ International Tables for X-Ray Crystallography 1992, Vol. C., Dordrecht: Kluwer Academic Publishers.

⁶ H. D. Flack, *Acta. Cryst.*, 1983, **A39**, 876.

Table 1. Crystal data and structure refinement for **1b**.

Identification code	tjm4 (1b)	
Empirical formula	C ₅₂ H ₇₀ La ₂ Se ₂	
Formula weight	1130.82	
Temperature	153(2) K	
Wavelength	0.71073 Å	
Crystal system	Orthorhombic	
Space group	<i>Pnn2</i>	
Unit cell dimensions	a = 10.3729(10) Å	α = 90°.
	b = 12.0187(12) Å	β = 90°.
	c = 19.4835(19) Å	γ = 90°.
Volume	2429.0(4) Å ³	
Z	2	
Density (calculated)	1.546 Mg/m ³	
Absorption coefficient	3.266 mm ⁻¹	
F(000)	1128	
Crystal color	colorless	
Crystal size	0.26 x 0.24 x 0.22 mm ³	
Theta range for data collection	1.99 to 28.30°	
Index ranges	-13 ≤ h ≤ 13, -15 ≤ k ≤ 16, -25 ≤ l ≤ 25	
Reflections collected	25662	
Independent reflections	6015 [R(int) = 0.0258]	
Completeness to theta = 28.30°	100.0 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.5336 and 0.4839	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	6015 / 1 / 265	
Goodness-of-fit on F ²	1.057	
Final R indices [I > 2σ(I) = 5560 data]	R1 = 0.0187, wR2 = 0.0445	
R indices (all data, 0.75 Å)	R1 = 0.0221, wR2 = 0.0460	
Absolute structure parameter	0.232(9)	
Largest diff. peak and hole	0.487 and -0.599 e.Å ⁻³	

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y+1,z

X-ray Data Collection, Structure Solution and Refinement for $[(C_5Me_5)_2La(SePh)(NCCMe_3)]_2$, **2b**.

A colorless crystal of approximate dimensions 0.09 x 0.19 x 0.34 mm was mounted on a glass fiber and transferred to a Bruker SMART APEX II diffractometer. The APEX2⁷ program package was used to determine the unit-cell parameters and for data collection (25 sec/frame scan time for a sphere of diffraction data). The raw frame data was processed using SAINT⁸ and SADABS⁹ to yield the reflection data file. Subsequent calculations were carried out using the SHELXTL⁴ program. There were no systematic absences nor any diffraction symmetry other than the Friedel condition. The centrosymmetric triclinic space group $P\bar{1}$ was assigned and later determined to be correct.

The structure was solved by direct methods and refined on F^2 by full-matrix least-squares techniques. The analytical scattering factors⁵ for neutral atoms were used throughout the analysis. Hydrogen atoms were included using a riding model. The molecules were dimers and were located about inversion centers. There were 3.5 molecules of benzene solvent present per dimeric formula-unit. At convergence, $wR2 = 0.0583$ and $Goof = 1.023$ for 828 variables refined against 17502 data (0.74Å), $R1 = 0.0237$ for those 15175 data with $I > 2.0\sigma(I)$.

⁷ APEX2 Version 2.2-0 Bruker AXS, Inc.; Madison, WI 2007.

⁸ SAINT Version 7.46a, Bruker AXS, Inc.; Madison, WI 2007

⁹ Sheldrick, G. M. SADABS, Version 2007/4, Bruker AXS, Inc.; Madison, WI 2007

Table 2. Crystal data and structure refinement for **2b**.

Identification code	tjm30 (2b)	
Empirical formula	C ₆₂ H ₈₈ La ₂ N ₂ Se ₂ • 3.5(C ₆ H ₆)	
Formula weight	1570.46	
Temperature	153(2) K	
Wavelength	0.71073 Å	
Crystal system	Triclinic	
Space group	<i>P</i> $\bar{1}$	
Unit cell dimensions	<i>a</i> = 14.9927(9) Å	α = 88.5675(7)°.
	<i>b</i> = 15.3140(9) Å	β = 88.8528(7)°.
	<i>c</i> = 16.9117(10) Å	γ = 79.9654(7)°.
Volume	3821.8(4) Å ³	
<i>Z</i>	2	
Density (calculated)	1.365 Mg/m ³	
Absorption coefficient	2.098 mm ⁻¹	
<i>F</i> (000)	1606	
Crystal color	colorless	
Crystal size	0.34 x 0.19 x 0.09 mm ³	
Theta range for data collection	1.75 to 28.50°	
Index ranges	-20 ≤ <i>h</i> ≤ 19, -20 ≤ <i>k</i> ≤ 20, -22 ≤ <i>l</i> ≤ 22	
Reflections collected	44479	
Independent reflections	17502 [R(int) = 0.0210]	
Completeness to theta = 28.50°	90.4 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.8337 and 0.5357	
Refinement method	Full-matrix least-squares on <i>F</i> ²	
Data / restraints / parameters	17502 / 0 / 828	
Goodness-of-fit on <i>F</i> ²	1.023	
Final R indices [<i>I</i> > 2σ(<i>I</i>) = 15175 data]	R1 = 0.0237, wR2 = 0.0558	
R indices (all data)	R1 = 0.0305, wR2 = 0.0583	
Largest diff. peak and hole	0.853 and -0.464 e.Å ⁻³	

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y,-z+1 #2 -x+2,-y+1,-z #3 -x+1,-y+1,-z+1

X-ray Data Collection, Structure Solution and Refinement for $[(C_5Me_5)_2Ce(SPh)]_2$, **3**,
Figure S1.

A purple crystal of approximate dimensions 0.19 x 0.21 x 0.32 mm was mounted on a glass fiber and transferred to a Bruker SMART APEX II diffractometer. The APEX2¹⁰ program package was used to determine the unit-cell parameters and for data collection (20 sec/frame scan time for a sphere of diffraction data). The raw frame data was processed using SAINT¹¹ and SADABS³ to yield the reflection data file. Subsequent calculations were carried out using the SHELXTL⁴ program. The diffraction symmetry was *mmm* and the systematic absences were consistent with the orthorhombic space groups *Pnn2* and *Pnmm*. It was later determined that the noncentrosymmetric space group *Pnn2* was correct.

The structure was solved by direct methods and refined on F^2 by full-matrix least-squares techniques. The analytical scattering factors⁵ for neutral atoms were used throughout the analysis. Hydrogen atoms were included using a riding model. The molecule was located on a two-fold rotation axis.

At convergence, $wR2 = 0.0584$ and $Goof = 1.046$ for 264 variables refined against 5782 data (0.75\AA), $R1 = 0.0207$ for those 5498 data with $I > 2.0\sigma(I)$. The absolute structure was assigned by refinement of the Flack parameter.⁶

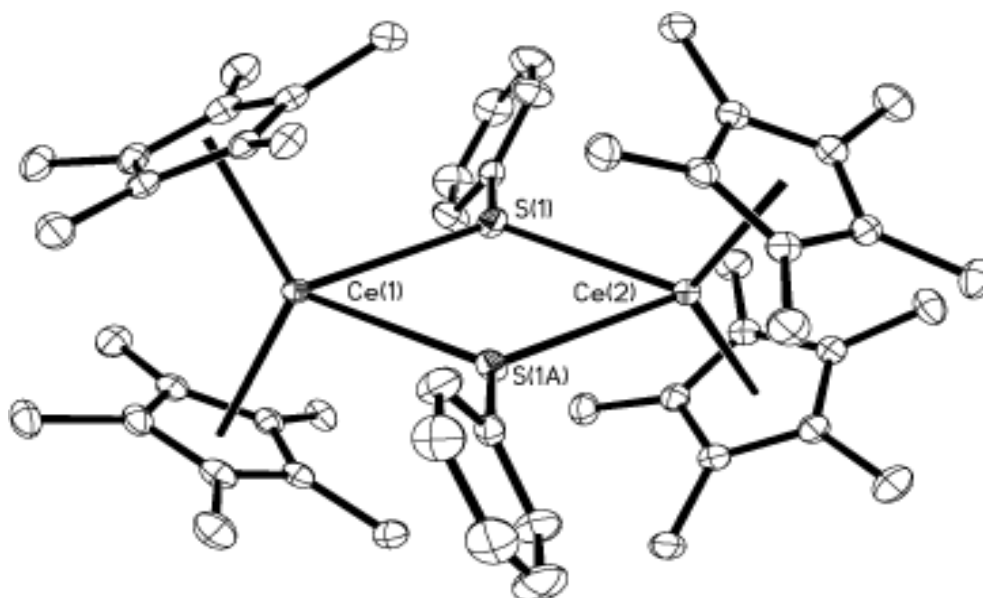


Figure S1. Thermal ellipsoid plot of $[(C_5Me_5)_2Ce(SPh)]_2$, **3**, drawn at the 50% probability level. Hydrogen atoms have been omitted for clarity.

¹⁰ APEX2 Version 2008.3-0, Bruker AXS, Inc.; Madison, WI 200/8.

¹¹ SAINT Version 7.53a, Bruker AXS, Inc.; Madison, WI 2007.

Table 3. Crystal data and structure refinement for **3**.

Identification code	tjm29 (3)	
Empirical formula	C ₅₂ H ₇₀ Ce ₂ S ₂	
Formula weight	1039.44	
Temperature	103(2) K	
Wavelength	0.71073 Å	
Crystal system	Orthorhombic	
Space group	<i>Pnn2</i>	
Unit cell dimensions	a = 10.3287(7) Å	α = 90°.
	b = 11.7979(8) Å	β = 90°.
	c = 19.4736(13) Å	γ = 90°.
Volume	2373.0(3) Å ³	
Z	2	
Density (calculated)	1.455 Mg/m ³	
Absorption coefficient	2.015 mm ⁻¹	
F(000)	1060	
Crystal color	purple	
Crystal size	0.32 x 0.21 x 0.19 mm ³	
Theta range for data collection	2.02 to 28.28°	
Index ranges	-13 ≤ h ≤ 13, -15 ≤ k ≤ 15, -25 ≤ l ≤ 25	
Reflections collected	26891	
Independent reflections	5782 [R(int) = 0.0247]	
Completeness to theta = 28.28°	99.5 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.7008 and 0.5649	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	5782 / 1 / 264	
Goodness-of-fit on F ²	1.046	
Final R indices [I > 2σ(I) = 5498 data]	R1 = 0.0207, wR2 = 0.0571	
R indices (all data, 0.75 Å)	R1 = 0.0218, wR2 = 0.0584	
Absolute structure parameter	0.001(14)	
Largest diff. peak and hole	1.152 and -0.276 e.Å ⁻³	

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y+1,z

X-ray Data Collection, Structure Solution and Refinement for $[(C_5Me_5)_2Pr(SPh)]_2$, **5**, Figure S2.

A yellow crystal of approximate dimensions 0.13 x 0.15 x 0.18 mm was mounted on a glass fiber and transferred to a Bruker SMART APEX II diffractometer. The APEX2⁷ program package was used to determine the unit-cell parameters and for data collection (10 sec/frame scan time for a sphere of diffraction data). The raw frame data was processed using SAINT⁸ and SADABS³ to yield the reflection data file. Subsequent calculations were carried out using the SHELXTL¹² program. The diffraction symmetry was *mmm* and the systematic absences were consistent with the orthorhombic space groups *Pnn2* and *Pnnm*. It was later determined that the noncentrosymmetric space group *Pnn2* was correct.

The structure was solved using the coordinates of the analogous cerium complex and refined on F^2 by full-matrix least-squares techniques. The analytical scattering factors⁵ for neutral atoms were used throughout the analysis. Hydrogen atoms were included using a riding model. The molecule was located on a two-fold rotation axis.

At convergence, $wR2 = 0.0930$ and $Goof = 1.028$ for 264 variables refined against 5403 data (0.77\AA), $R1 = 0.0334$ for those 4399 data with $I > 2.0\sigma(I)$. The absolute structure was assigned by refinement of the Flack parameter.⁶

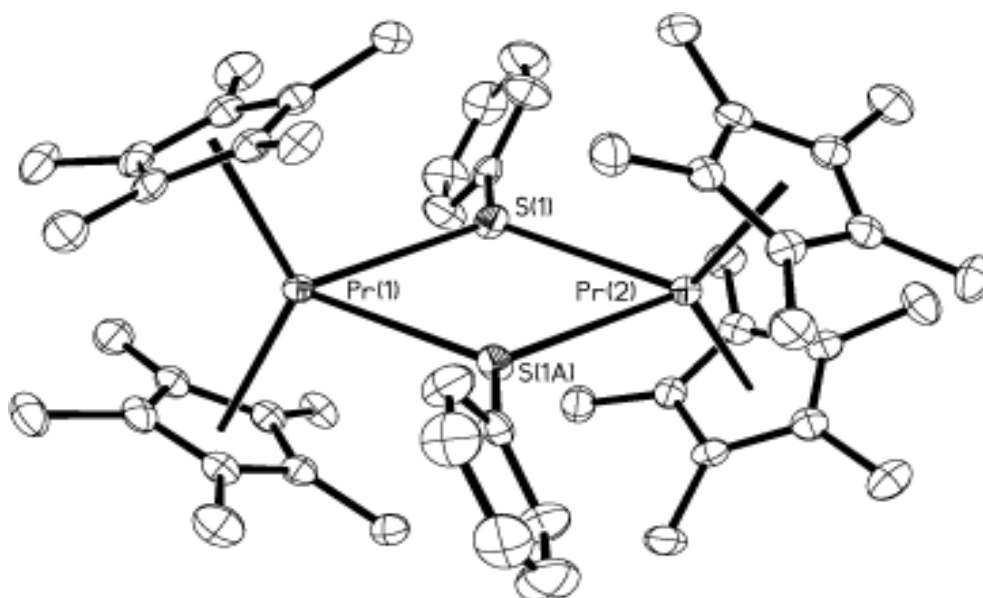


Figure S2. Thermal ellipsoid plot of $[(C_5Me_5)_2Pr(SPh)]_2$, **5**, drawn at the 50% probability level. Hydrogen atoms have been omitted for clarity.

¹² Sheldrick, G. M. SHELXTL, Version 2008/3, Bruker AXS, Inc.; Madison, WI 2008.

Table 4. Crystal data and structure refinement for $[(C_5Me_5)_2Pr(SPh)]_2$, **5**.

Identification code	tjm35 (5)	
Empirical formula	$C_{52} H_{70} Pr_2 S_2$	
Formula weight	1041.02	
Temperature	148(2) K	
Wavelength	0.71073 Å	
Crystal system	Orthorhombic	
Space group	$Pnn2$	
Unit cell dimensions	$a = 10.3504(6)$ Å	$\alpha = 90^\circ$.
	$b = 11.7647(7)$ Å	$\beta = 90^\circ$.
	$c = 19.4542(12)$ Å	$\gamma = 90^\circ$.
Volume	$2368.9(2)$ Å ³	
Z	2	
Density (calculated)	1.459 Mg/m ³	
Absorption coefficient	2.153 mm ⁻¹	
F(000)	1064	
Crystal color	yellow	
Crystal size	0.18 x 0.15 x 0.13 mm ³	
Theta range for data collection	2.02 to 27.48°	
Index ranges	$-13 \leq h \leq 13, -15 \leq k \leq 15, -25 \leq l \leq 25$	
Reflections collected	25686	
Independent reflections	5403 [R(int) = 0.0477]	
Completeness to theta = 27.48°	99.8 %	
Absorption correction	Semi-empirical from equivalents	
Max. and min. transmission	0.7671 and 0.6979	
Refinement method	Full-matrix least-squares on F ²	
Data / restraints / parameters	5403 / 1 / 264	
Goodness-of-fit on F ²	1.028	
Final R indices [$I > 2\sigma(I)$ = 4399 data]	R1 = 0.0334, wR2 = 0.0859	
R indices (all data, 0.77Å)	R1 = 0.0434, wR2 = 0.0930	
Absolute structure parameter	-0.01(3)	
Largest diff. peak and hole	2.663 and -0.492 e.Å ⁻³	

Symmetry transformations used to generate equivalent atoms:

#1 -x+1,-y+1,z