



# Differences in Enantioselective Hydroxylation of 2,2',3,6-Tetrachlorobiphenyl (CB45) and 2,2',3,4',6-Pentachlorobiphenyl (CB91) by Human and Rat CYP2B Subfamilies

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# Differences in enantioselective hydroxylation of 2,2',3,6-tetrachlorobiphenyl (CB45) and 2,2',3,4',6-pentachlorobiphenyl (CB91) by human and rat CYP2B subfamilies

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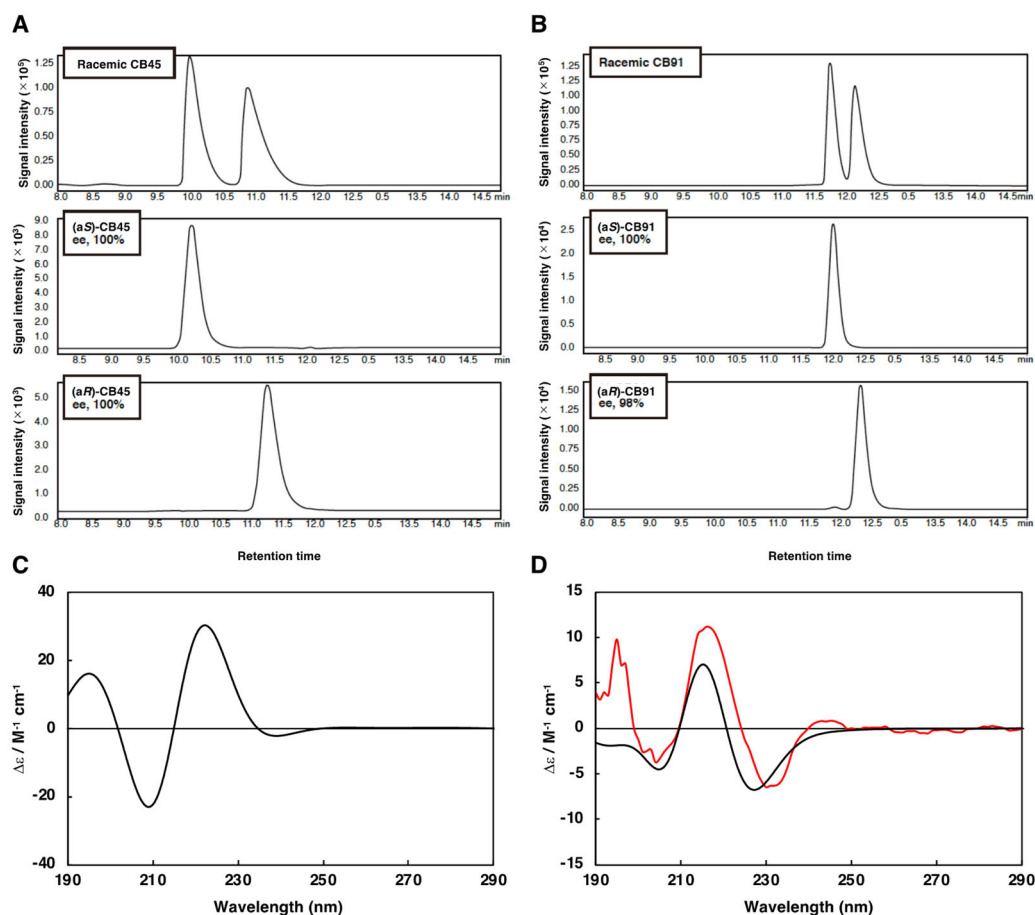
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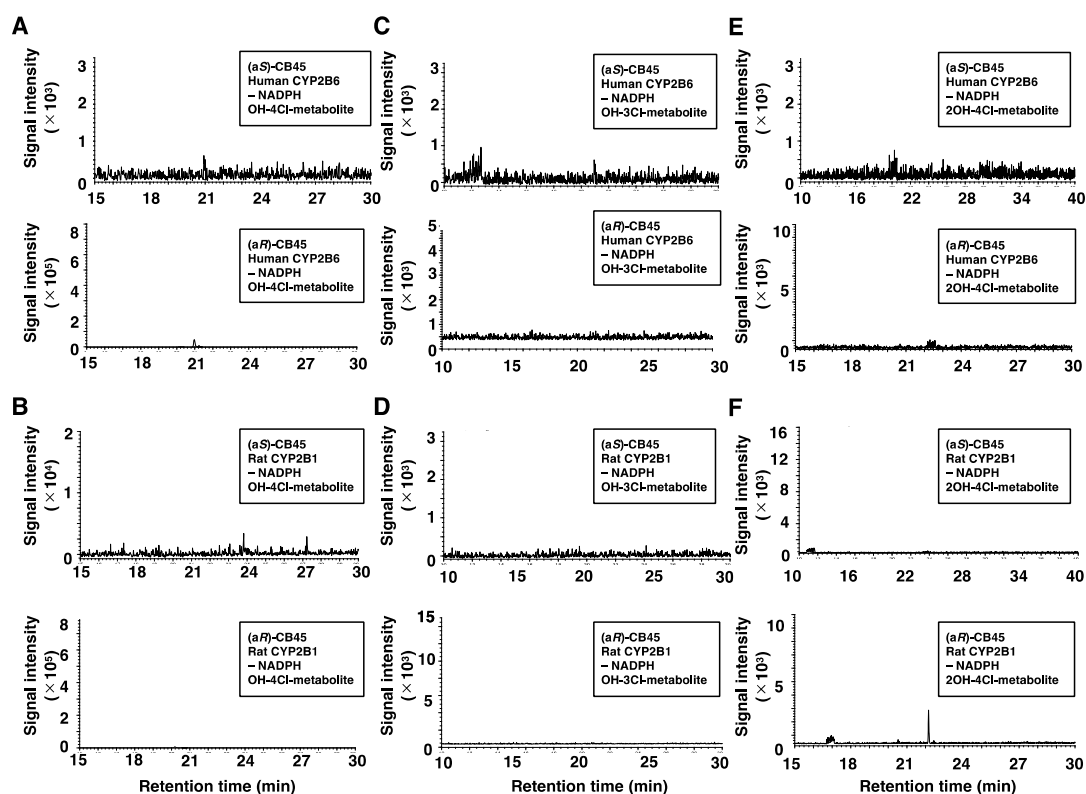
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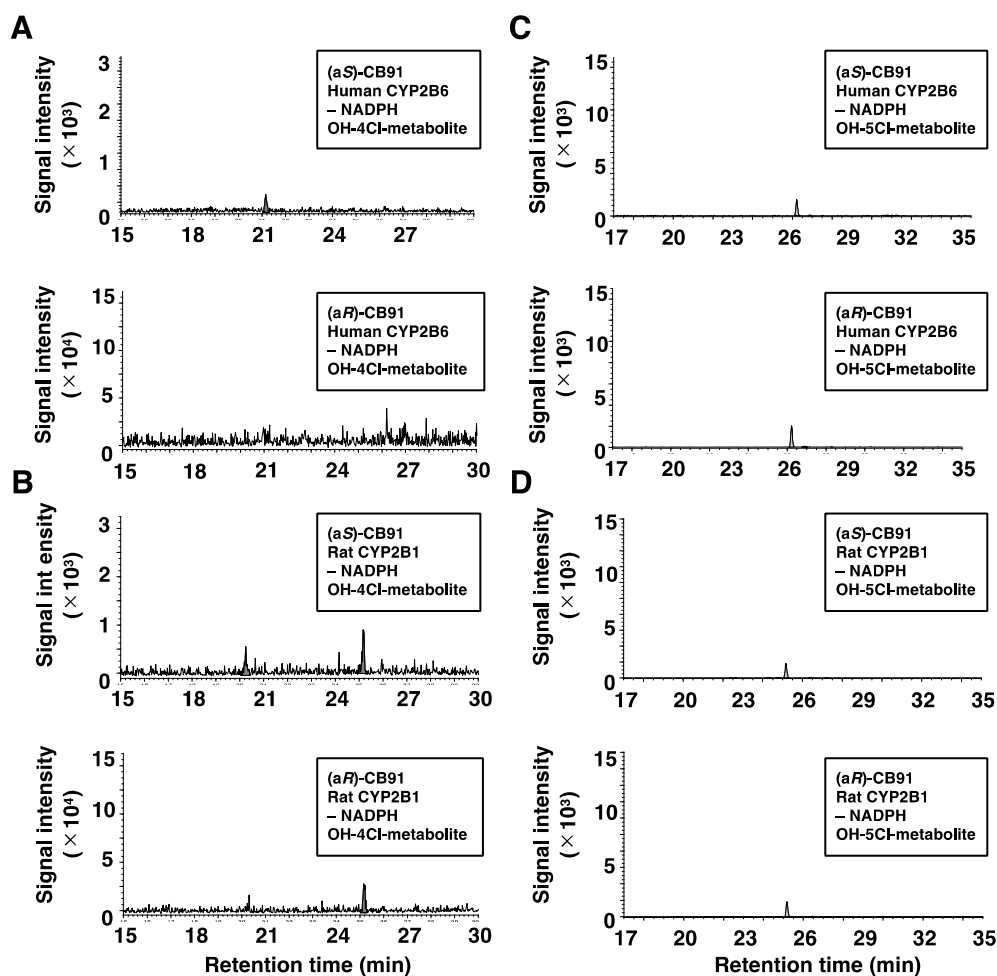
**Figure S1** Isolation of atropisomers from racemic 2,2',3,6-tetrachlorobiphenyl (CB45) (A) and 2,2',3,4',6-pentachlorobiphenyl (CB91) (B) and experimental (red) and theoretical (black) CD spectra of CB45 (C) and CB91 (D).

(a*S*)-CB45 and (a*S*)-CB91, and (a*R*)-CB45 and (a*R*)-CB91 were eluted in the first and second fractions, respectively. Enantiomeric excess (ee) values for each atropisomer were shown in the graphs. Experimental spectrum was obtained in hexane (ca. 7.7  $\mu\text{M}$ ) for the second fraction of CB91. Theoretical spectra for a*S* and a*R* atropisomers for CB45 and CB91 were calculated at the RI-CC2/aug-def2-TZVPP//B-LYP-D3/def2-TZVP level. The calculated rotational strengths in length gauge were scaled to one-half and were expanded by Gaussian functions and overlapped where the width of the band at 1/e height is fixed at 0.5 eV and the excitation energy was red-shifted by 0.6 eV.



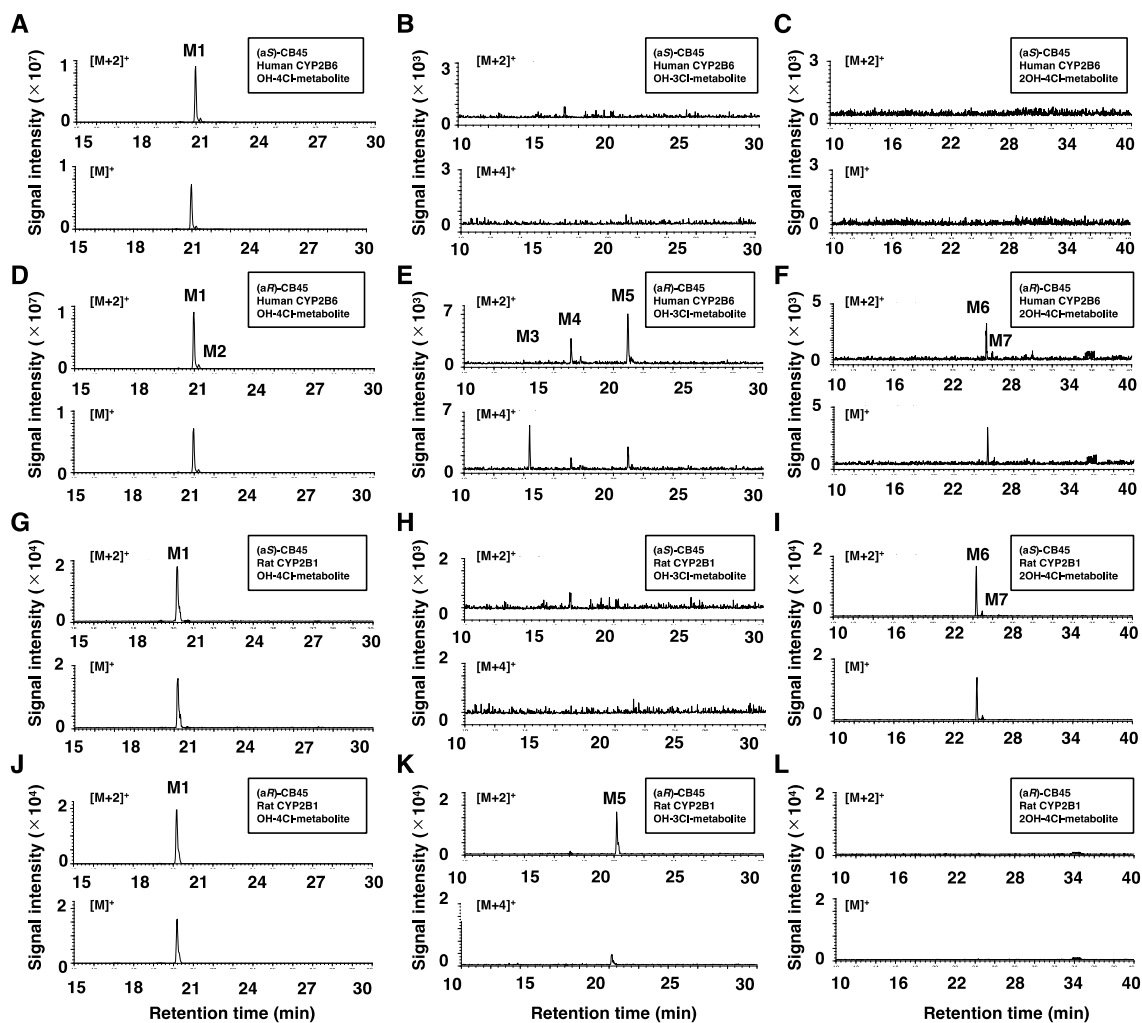
**Figure S2** Chromatograms of mono-hydroxylated tetrachloro (A, B), mono-hydroxylated trichloro (C, D), and di-hydroxylated tetrachloro (E, F) metabolites produced from 2,2',3,6-tetrachlorobiphenyl (CB45) by human CYP2B6 (A, C, E) and rat CYP2B1 (B, D, F), as analyzed by gas chromatography/ high-resolution mass spectrometry.

A, C, and E represent the metabolism of CB45 by human CYP2B6 without NADPH. B, D, and F show the metabolism of CB45 by rat CYP2B1 without NADPH. Upper and lower panels represent the metabolism of (aS)-CB45 and (aR)-CB45 as substrates, respectively.



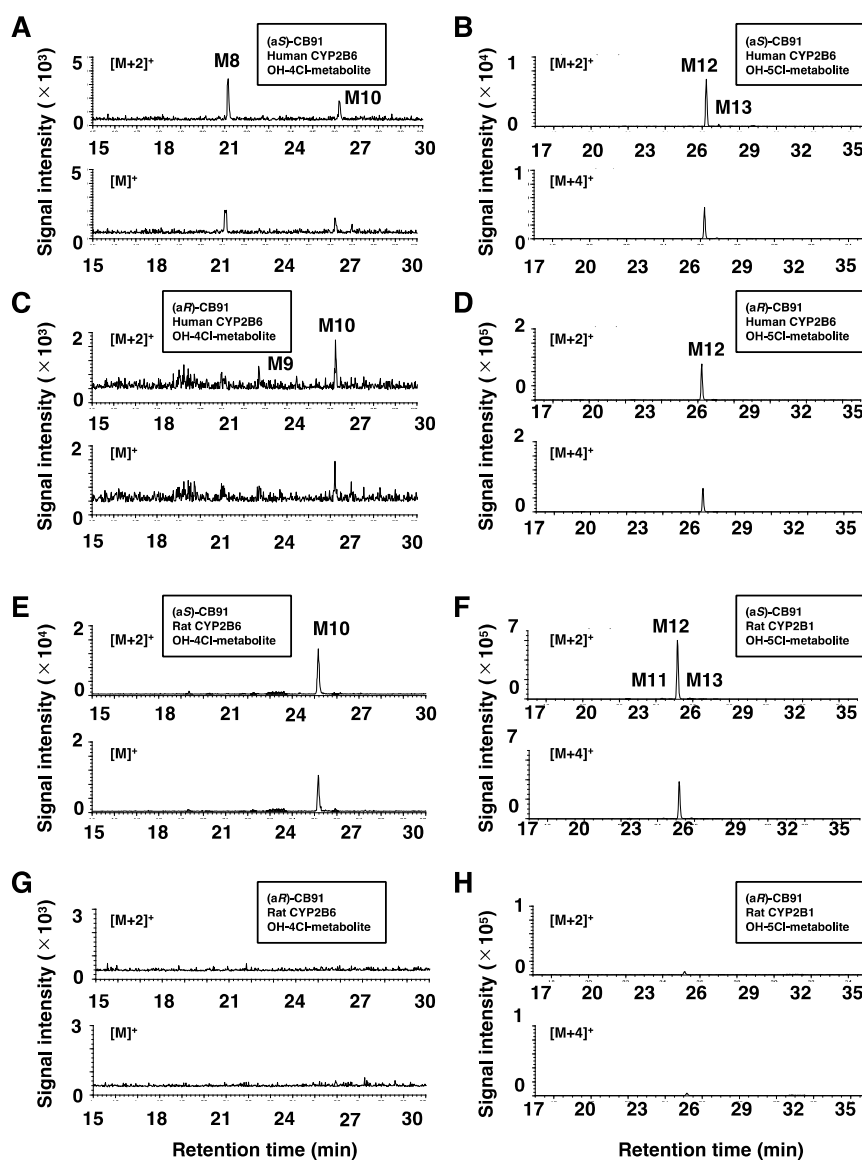
**Figure S3** Chromatograms of mono-hydroxylated tetrachloro (A, B) and pentachloro (C, D) metabolites produced from 2,2',3,4',6-pentachlorobiphenyl (CB91) by human CYP2B6 (A, C) and rat CYP2B1 (B, D), as analyzed by gas chromatography/high-resolution mass spectrometry.

A and C represent the metabolism of CB91 by human CYP2B6 without NADPH. B and D represent the metabolism of CB91 by rat CYP2B1 without NADPH. Upper and lower panels represent the metabolism of (aS)-CB91 and (aR)-CB91 as substrates, respectively.



**Figure S4** Isotope ratios of methylated 2,2',3,6-tetrachlorobiphenyl (CB45) metabolites produced by human CYP2B6 (A–F) and rat CYP2B1 (G–L).

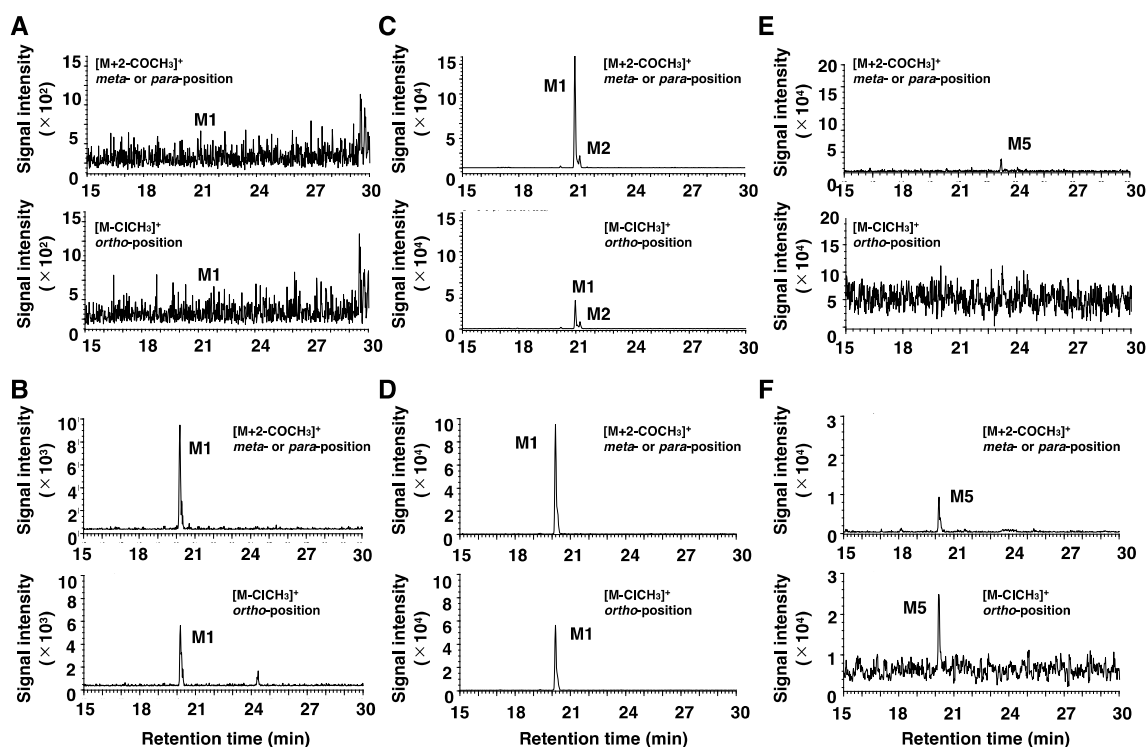
(aS)-CB45 (A–C, G–I) and (aR)-CB45 (D–F, J–L) were used as substrates. Hydroxylated tetrachloro (A, D, G, J), hydroxylated trichloro (B, E, H, K), and di-hydroxylated tetrachloro (C, F, I, L) metabolites were analyzed by gas chromatography/high-resolution mass spectrometry.



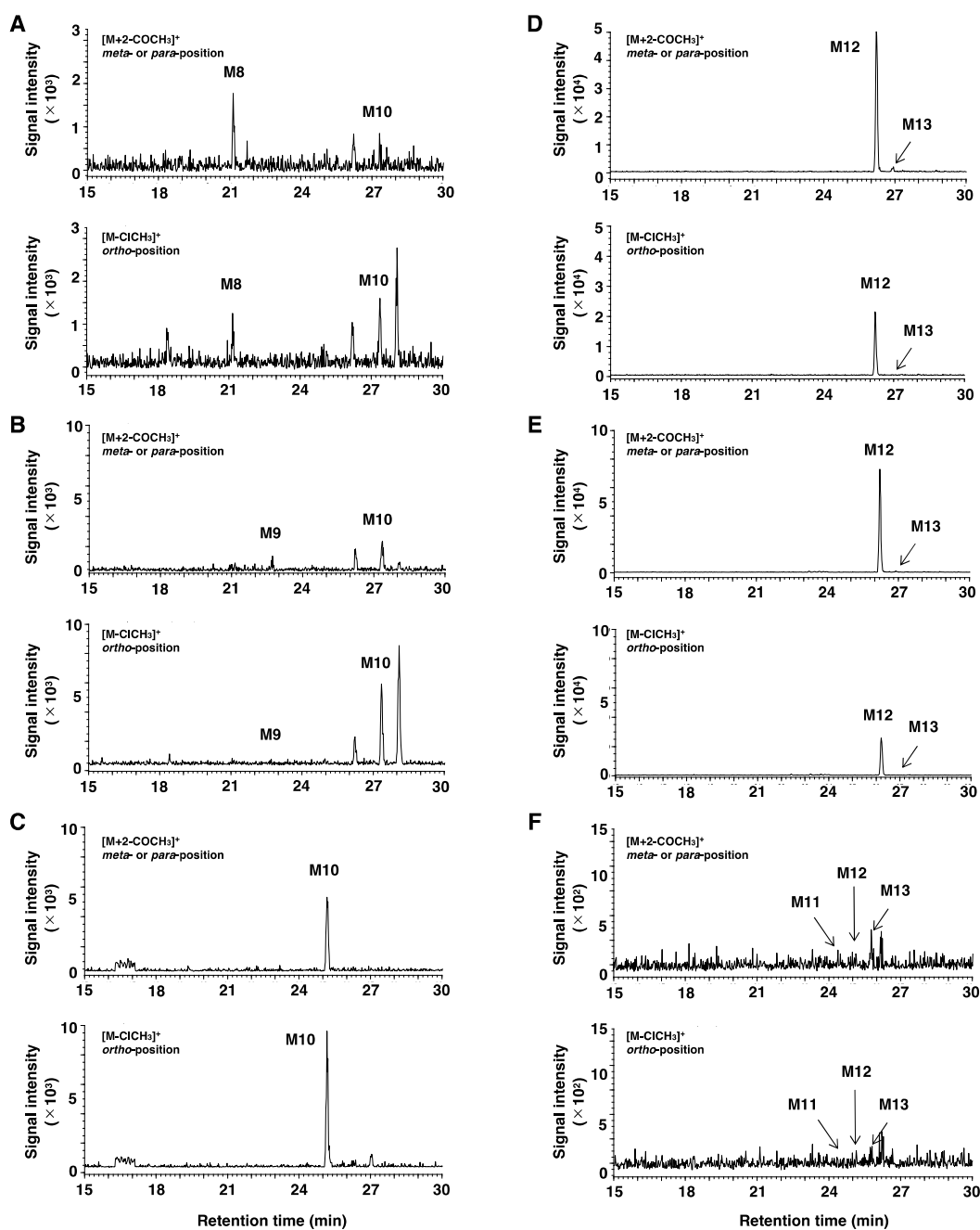
**Figure S5** Isotope ratios of methylated 2,2',3,4',6-pentachlorobiphenyl (CB91) metabolites produced by human CYP2B6 (A–D) and rat CYP2B1 (E–H).

(*aS*)-CB91 (A, B, E, F) and (*aR*)-CB91 (C, D, G, H) were used as substrates. Hydroxylated tetrachloro (A, C, E, G), and hydroxylated pentachloro (B, D, F, H) metabolites were analyzed by gas chromatography/high-resolution mass spectrometry.

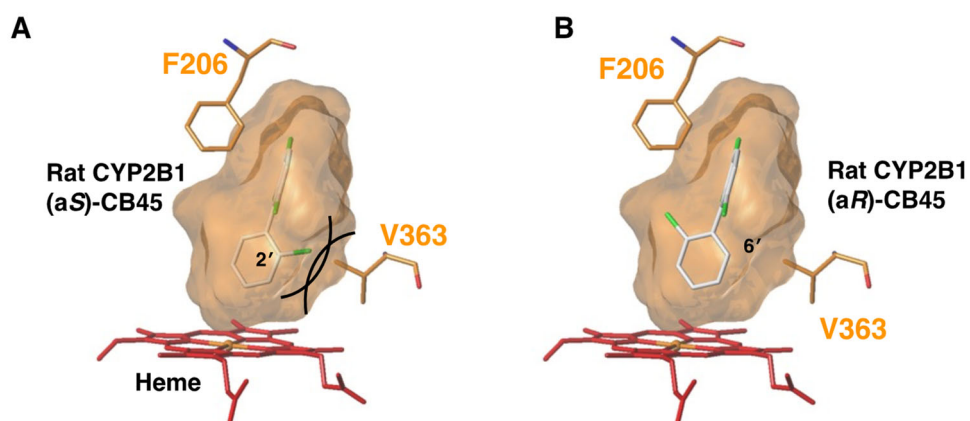




**Figure S6** Patterns of fragment ions ( $[M+2-COCH_3]^+$  and  $[M-CH_3Cl]^+$ ) of mono-hydroxylated tetrachloro metabolites of (aS)-2,2',3,6-tetrachlorobiphenyl (CB45) (A, B) and (aR)-CB45 (C, D) and mono-hydroxylated trichloro metabolites of (aR)-CB45 (E, F) produced on using human CYP2B6 (A, C, E) and rat CYP2B1 (B, D, F), as analyzed by gas chromatography/high-resolution mass spectrometry.



**Figure S7** Patterns of fragment ions ( $[M+2-COCH_3]^+$  and  $[M-CH_3Cl]^+$ ) of mono-hydroxylated tetrachloro and pentachloro metabolites of (aS)-2,2',3,4',6-pentachlorobiphenyl (CB91) (A, C, D, F) and (aR)-CB91 (B, E) produced on using human CYP2B6 (A, B, D, E) and rat CYP2B1 (C, F), as analyzed by gas chromatography/high-resolution mass spectrometry.



**Figure S8** Rat CYP2B1 preferentially metabolizes (a*R*)-2,2',3,6-tetrachlorobiphenyl (CB45).

Docking models of rat CYP2B1 toward (a*S*)-CB45 (A) and (a*R*)-CB45 (B) are described. The green stick in CB45 indicates a chlorine atom. Amino acids described in orange constitute the rat CYP2B1 substrate-binding cavity. Steric hindrance between chlorine at the 2'-position of (a*S*)-CB45 and a side chain of amino acids, such as V363, is observed in the rat CYP2B1 cavity.

**Table S1.** The analysis conditions for derivatized hydroxylated PCBs by gas chromatography/high-resolution mass spectrometry

	Condition
Gas chromatography	6890N (Agilent Technologies)
Mass spectrometry	JMS-800D (JEOL)
Column	HT8-PCB 60 m × 0.25 mm
Column temperature	130°C (1 min) → (20°C /min) → 210°C → (2°C /min) → 285°C → (30 °C /min) → 330 °C
Inlet mode	Splitless (Purge starting time; 1.5 min)
Inlet temperature	290°C
Injection volume	2 µL
Carrier gas	1.2 mL/min
Interface temperature	290°C
Ion source temperature	280°C
Ionization voltage	38 eV
Detection mode	SIM

**Table S2.** Monitored ions of derivatized hydroxylated PCBs by high-resolution gas chromatography/high-resolution mass spectrometry

Homologue	Derivatized homologue	Selected fragment ( <i>m/z</i> )
OH-TeCB <sup>*1</sup>	CH <sub>3</sub> O-TeCBs	321.9301 [M+2] <sup>+</sup> 319.9329 [M] <sup>+</sup> 278.9116 [M+2-COCH <sub>3</sub> ] <sup>+</sup> 269.9406 [M-ClCH <sub>3</sub> ] <sup>+</sup>
Di-OH-TeCB	Di-CH <sub>3</sub> O-TeCBs	351.9406 [M+2] <sup>+</sup> 340.9435 [M] <sup>+</sup>
OH-PeCB <sup>*2</sup>	CH <sub>3</sub> O-PeCBs	355.8911 [M+2] <sup>+</sup> 357.8882 [M+4] <sup>+</sup> 312.8727 [M+2-COCH <sub>3</sub> ] <sup>+</sup> 305.8942 [M+2-ClCH <sub>3</sub> ] <sup>+</sup>
OH-[ <sup>13</sup> C <sub>12</sub> ]-TeCB	CH <sub>3</sub> O-[ <sup>13</sup> C <sub>12</sub> ]-TeCB	333.9702 [M+2] <sup>+</sup> 331.9732 [M] <sup>+</sup>
OH-[ <sup>13</sup> C <sub>12</sub> ]-PeCB	CH <sub>3</sub> O-[ <sup>13</sup> C <sub>12</sub> ]-PeCB	367.9313 [M+2] <sup>+</sup> 365.9342 [M] <sup>+</sup>
[ <sup>13</sup> C <sub>12</sub> ]-2,3',4',5-TeCB <sup>*3</sup>		303.9597 [M+2] <sup>+</sup> 301.9626 [M] <sup>+</sup>

<sup>\*1</sup>Tetrachlorobiphenyl; <sup>\*2</sup>Pentachlorobiphenyl; <sup>\*3</sup>Syringe spike

**Table S3.** Retention times and retention time indexes for hydroxylated (OH)-metabolites of 2,2',3,6-tetrachlorobiphenyl (CB45) and 2,2',3,4',6-pentachlorobiphenyl (CB91).

PCB	Metabolite	Retention time (Retention time index <sup>*1</sup> )				Standard used for calibration curve
		Human CYP2B6		Rat CYP2B1		
		(a <i>S</i> )	(a <i>R</i> )	(a <i>S</i> )	(a <i>R</i> )	
CB45	M1	20.96 (0.7769)	20.98 (0.7779)	20.21 (0.7787)	20.21 (0.7786)	2-OH-2',3,3',4'- TeCB <sup>*3</sup>
	M2	- <sup>*2</sup>	21.23 (0.7870)	-	-	
	M3	-	14.41 (0.5340)	-	-	
	M4	-	17.18 (0.6371)	-	-	
	M5	-	20.99 (0.7782)	-	20.21 (0.7789)	4-OH-2,3',4-TrCB <sup>*5</sup>
	M6	-	25.40 (0.9415)	24.34 (0.9380)	-	2-OH-2',3,3',4'- TeCB
	M7	-	26.07 (0.9664)	24.91 (0.9599)	-	
CB91	M8	21.15 (0.7838)	-	-	-	2-OH-2',3,3',4'- TeCB
	M9	-	22.71 (0.8420)	-	-	
	M10	26.24 (0.9725)	26.24 (0.9727)	25.19 (0.9706)	-	4-OH-2,2',3',5- TeCB
	M11	-	-	24.34 (0.8929)	-	2-OH-3,3',5,5',6- PeCB <sup>*6</sup>
	M12	26.22 (0.9215)	26.22 (0.9271)	25.18 (0.9234)	-	
	M13	26.90 (1.026)	26.90 (1.026)	25.85 (1.027)	-	

<sup>\*1</sup>Retention time indexes were represented by the relative retention time to [<sup>13</sup>C<sub>12</sub>]-4-OH-2',3',4',5'-tetrachlorobiphenyl for M1–M10 and [<sup>13</sup>C<sub>12</sub>]-4-OH-2',3,4',5,5'-pentachlorobiphenyl for M11–M13.

<sup>\*2</sup>Not detected.

<sup>\*3</sup>Tetrachlorobiphenyl.

<sup>\*4</sup>M3 and M4 were not quantified.

<sup>\*5</sup>Trichlorobiphenyl.

<sup>\*6</sup>Pentachlorobiphenyl.

**Table S4.** Isotope ratios of hydroxylated (OH)-metabolites of 2,2',3,6-tetrachlorobiphenyl (CB45) and 2,2',3,4',6-pentachlorobiphenyl (CB91).

PCB	Metabolite (Cl number)	Isotope ratio				Theoretical ratio	Actual ratio (Standard)
		Human CYP2B6		Rat CYP2B1			
		(aS)	(aR)	(aS)	(aR)		
CB45	M1 (Tetra)	1:1.24 (M:M+2)	1:1.28 (M:M+2)	1:1.30 (M:M+2)	1:1.26 (M:M+2)	1:1.29 (M:M+2)	1:1.28 (4-OH-2,2',3',5-TeCB*2)
	M2 (Tetra)	-*1	1:1.28 (M:M+2)	-	-	1:1.29 (M:M+2)	
	M3 (Tri)	-	0.04:1 (M+2:M+4)	-	-	3.06:1 (M+2:M+4)	3.00:1 (4-OH-2,3',4'-TrCB*3)
	M4 (Tri)	-	2.26:1 (M+2:M+4)	-	-	3.06:1 (M+2:M+4)	
	M5 (Tri)	-	2.94:1 (M+2:M+4)	-	3.44:1 (M+2:M+4)	3.06:1 (M+2:M+4)	
	M6 (Tetra)	-	1:1.24 (M:M+2)	1:1.33 (M:M+2)	-	1:1.29 (M:M+2)	1:1.28 (4-OH-2,2',3',5-TeCB)
	M7 (Tetra)	-	1:1.19 (M:M+2)	1:1.06 (M:M+2)	-	1:1.29 (M:M+2)	
CB91	M8 (Tetra)	1:1.33 (M:M+2)	-	-	-	1:1.29 (M:M+2)	1:1.28 (4-OH-2,2',3',5-TeCB)
	M9 (Tetra)	-	1:1.27 (M:M+2)	-	-	1:1.29 (M:M+2)	
	M10 (Tetra)	1:1.12 (M:M+2)	1:1.28 (M:M+2)	1:1.24 (M:M+2)	-	1:1.29 (M:M+2)	
	M11 (Penta)	-	-	1.61:1 (M+2:M+4)	-	1.55:1 (M+2:M+4)	1.63:1 (3-OH-2,3',4,4',6-PeCB*4)
	M12 (Penta)	1.56:1 (M+2:M+4)	1.57:1 (M+2:M+4)	1.58:1 (M+2:M+4)	-	1.55:1 (M+2:M+4)	
	M13 (Penta)	1.44:1 (M+2:M+4)	1.60:1 (M+2:M+4)	1.79:1 (M+2:M+4)	-	1.55:1 (M+2:M+4)	

<sup>\*1</sup>Not detected.

<sup>\*2</sup>Tetrachlorobiphenyl.

<sup>\*3</sup>Trichlorobiphenyl.

<sup>\*4</sup>Pentachlorobiphenyl.