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## Supporting Information

Enantioselective Metabolism of Chiral Polychlorinated Biphenyl 2,2',3,4,4',5',6-Heptachlorobiphenyl (CB183) by Human and Rat CYP2B Subfamilies

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**Table S1** Monitored ions of derivatized hydroxylated PCBs observed in high-resolution gas chromatography/high-resolution mass spectrometry.

Homologue	Derivatized homologue	Selected fragment (m/z)	
ОН-НрСВ	CH <sub>3</sub> -HpCBs	423.8131 [M+2] <sup>+</sup>	
		425.8102 [M+4] <sup>+</sup>	
		380.7947 [M+2-COCH <sub>3</sub> ] <sup>+</sup>	
		373.8202 [M+2-CICH <sub>3</sub> ] <sup>+</sup>	
OH-[ <sup>13</sup> C <sub>12</sub> ]-HpCB	CH <sub>3</sub> -[ <sup>13</sup> C <sub>12</sub> ]-HpCB	433.8563 [M] <sup>+</sup>	

HpCB: heptachlorobiphenyl

**Table S2** Retention times of methylated products from 2,2',3,4,4',5',6-heptachlorobiphenyl (CB183) metabolites produced by human CYP2B6 and hydroxylated (OH)-CB183 standards.

MeOH-PCB*1		Retention time (min)			
		(+)*2	( <b>-</b> )*3	(±)	
Authentic	S1			33.3	
standard*4	S2			33.6	
CB183	M1	33.3	33.3		
metabolite	M2	33.7	33.7		

<sup>\*1</sup>Methylated hydroxylated polychlorinated biphenyl

<sup>\*2</sup>Retention times of methylated products of M1 and M2 that are produced by human CYP2B6 from (+)-CB183 as a substrate are represented.

<sup>\*3</sup>Retention times of methylated products of M1 and M2 that are produced by human CYP2B6 from (–)-CB183 as a substrate are represented.

<sup>\*4</sup>The authentic standards S1 and S2 are 3'-MeO-CB183 and 5-MeO-CB183, respectively.

**Table S3** Isotope ratios of methylated products from 2,2′,3,4,4′,5′,6-heptachlorobiphenyl (CB183) metabolites produced by human CYP2B6 and hydroxylated (OH)-CB183 standards.

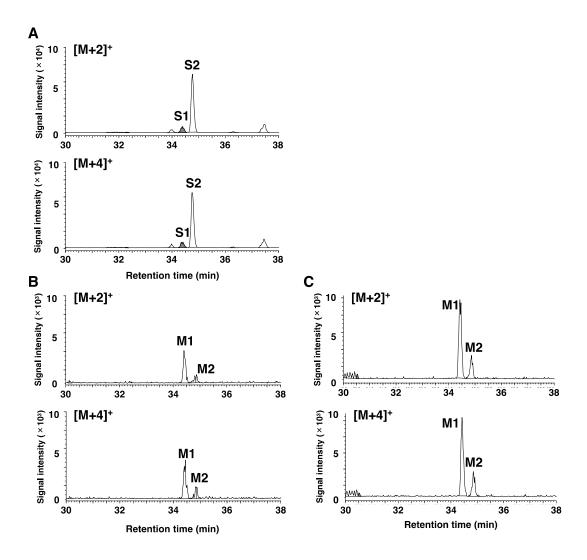
MeOH-PCB*1		Isotope ratio ([M+2]*:[M+4]*)			Theoretical
		(+)*2	(-)*3	(±)	ratio
Authentic	S1			1:1.02	
standard*4	S2			1:1.12	1:097
CB183	M1	1:1.10	1:0.97		1.097
metabolite	M2	1:1.04	1:1.09		

<sup>\*1</sup>Methylated hydroxylated polychlorinated biphenyl

<sup>\*2</sup>Isotope ratios of methylated products of M1 and M2 that are produced by human CYP2B6 from (+)-CB183 as a substrate are represented.

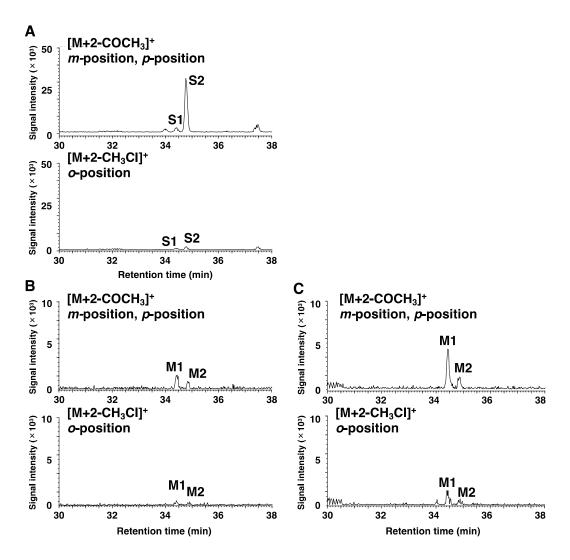
<sup>\*3</sup> Isotope ratios of methylated products of M1 and M2 that are produced by human CYP2B6 from (–)-CB183 as a substrate are represented.

<sup>\*4</sup>The authentic standards S1 and S2 are 3'-MeO-CB183 and 5-MeO-CB183, respectively.



**Figure S1** Isotope ratios ([M+2]<sup>+</sup> and [M+4]<sup>+</sup>) of hydroxylated heptachloro metabolites of CB183 by human CYP2B6 analyzed by high-resolution gas chromatography/ high-resolution mass spectrometry.

(A): Authentic standards S1 (3'-MeO-CB183) and S2 (5-MeO-CB183). (B) and (C): Hydroxylated heptachloro metabolites of (+)-CB183 and (-)-CB183, respectively.



**Figure S2** Pattern of fragment ions ([M+2-COCH<sub>3</sub>]<sup>+</sup> and [M+2-CH<sub>3</sub>Cl]<sup>+</sup>) of hydroxylated heptachloro metabolites of CB183 by human CYP2B6 analyzed by high-resolution gas chromatography/ high-resolution mass spectrometry.

(A) Authentic standards S1 (3'-MeO-CB183) and S2 (5-MeO-CB183). (B) and (C): Hydroxylated heptachloro metabolites of (+)-CB183 and (-)-CB183, respectively.

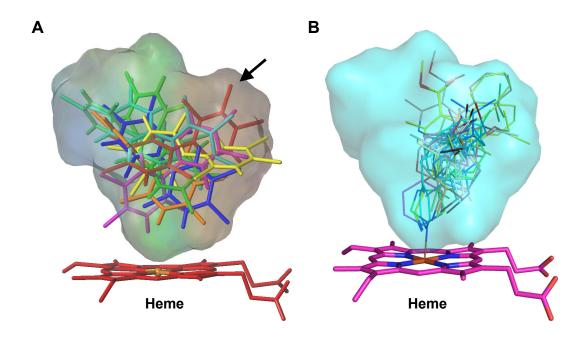
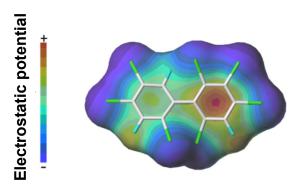


Figure S3 Docking models of CB183 with human CYP2B6.

(A): Eight conformations of CB183 were docked into the substrate-binding cavity of human CYP2B6 (PDB: 3IBD). The arrow indicates the plausible conformation of CB183 in the cavity. (B): Thirteen crystal structures of human CYP2B6 (mentioned under Materials and Methods) superimposed to show sixteen ligands in their substrate-binding cavities.



**Figure S4** Electrostatic potential map of CB183. Blue portion indicates region of high electron density.