## Electronic supplementary information

## Control of Stereoselectivity of Benzylic Hydroxylation Catalysed by Wild-Type Cytochrome P450BM3 Using Decoy Molecules <br> Kazuto Suzuki, Joshua Kyle Stanfield, a Osami Shoji, ${ }^{\text {ab }}$ Sota Yanagisawa, ${ }^{\text {a }}$ Hiroshi Sugimoto, ${ }^{\text {bc } \text { Yoshitsugu }}$ Shiroc and Yoshihito Watanabe*d

## Experimental Section

## Materials

Cytochrome P450BM3 was prepared according to methods described previously. ${ }^{1}$ The concentration of the enzyme was determined by CO difference spectra. ${ }^{2}$

Escherichia coli cells expressing P450BM3 suspended in 20 mM Tris- HCl ( pH 7.4 ) were disrupted using an ultrasonicator at $4^{\circ} \mathrm{C}$. After removing cell debris by centrifugation, the supernatant was applied to a CELLUFINE A-500 anion-exchange column (JNC). Weakly bound impurities were removed with 20 mM Tris- HCl containing $50 \mathrm{mM} \mathrm{KCl}(\mathrm{pH} 7.4)$ and tightly bound proteins including P450BM3 were eluted with Tris-buffer containing 250 mM KCl . Vividly red P450BM3 fractions were pooled and desalted by spin-centrifugation dialysis using an Amicon® Ultra Centrifuge Filter Ultracel® (Millipore,Co.) with a MWCO of 30 kDa , followed by further purification using a DEAE 650S anion-exchange column (TOSOH). P450BM3 was eluted with TrisHCl buffer over a KCl concentration gradient ranging from 0 to 120 mM . Eluted fractions were pooled and concentrated before applying to a Sephacryl S-300 gel-filtration column (GE Healthcare), equilibrated with 20 mM Tris buffer and $100 \mathrm{mM} \mathrm{KCl}(\mathrm{pH} 7.4)$ and the P450BM3 fraction was collected.

All chemical reagents were purchased from commercial sources and used without further purification. Ethylbenzene, propylbenzene, $(S)-(+)-1$-indanol, and cyclohexanepentanoic acid were purchased from Sigma-Aldrich Co. (St. Louis, MO). 1-phenylethyl alcohol, ( $R$ )-(+)-1-phenylethyl alcohol, 2-ethylphenol, 1-phenyl-1-propanol, 4-propylphenol, $(R)-(+)-1$-phenyl-propanol, indan, 1hydroxyindan, 2-hydroxyindan, 5-hydroxyindan, 1,2,3,4-tetrahydronaphthalene, 1,2,3,4-tetrahydro-1-naphthol, $(S)-(+)-1,2,3,4-$ tetrahydro-1-naphthol and BSTFA-TMCS (99:1) were obtained from Tokyo Chemical Industry Co. (Tokyo, Japan). The following chemicals were purchased from Wako Pure Chemical Industries, Ltd. (Osaka, Japan): acetophenone, phenylmethyl acetate.

## Measurement

Ultraviolet-visible spectra were recorded on a Shimadzu UV-2600 PC spectrophotometer. Gas
chromatography (GC) and GC-MS analysis were performed with Shimadzu GC-2014 ATF and Shimadzu GCMS-QP2010 SE both equipped with a cyclosil- $\beta$ column (Agilent Technologies, Inc., $30 \mathrm{~m} \times 0.25 \mathrm{~mm}) .1 \mathrm{H}$ NMR spectra were measured by A-400 spectrometer and ECA600 Delta (JEOL). ${ }^{1} \mathrm{H}$ NMR chemical shifts were reported versus tetramethylsilane (TMS) and referenced to residual solvent peaks (DMSO- $d_{6}$ ). ESI-TOF-MS spectra were measured by micrOTOF II (BRUKER ANALYTIC).

## Decoy molecules

The synthesis and characterisation of PFC9-Trp, PFC9-Phe, PFC9-Met were reported previously. ${ }^{1}$ Other decoy molecules were synthesised according to published procedures. ${ }^{3}$ C9-Phe, C9-Trp, Ph-C5-Phe, 5CHVA-Trp, 5CHVA-Phe, (R/S)-Ibuprofen-Phe, Z-Pro-Phe, Z-Pro-Met and Z-Gly-Phe were synthesised by the same method as described below.

General procedure for preparation of N -acyl amino acids ( $\mathrm{Ph}-\mathrm{C} 5-\mathrm{Phe}$ as an example):
To a mixture of 5-phenylpentanoic acid ( $356 \mathrm{mg}, 2.0 \mathrm{mmol}$ ), L-phenylalanine methylester hydrochloride ( $430 \mathrm{mg}, 2.0 \mathrm{mmol}$ ), 1-hydroxybenzotriazole monohydrate $(\mathrm{HOBt} \cdot \mathrm{H} 2 \mathrm{O}, 338 \mathrm{mg}, 2.5$ mmol ), 1-ethyl-3-(3-dimethylaminopropyl)carbodiimide hydrochloride (EDC $\cdot \mathrm{HCl}, 489 \mathrm{mg}, 2.5$ mmol) in dry DMF ( 5 mL ) was added $\mathrm{N}, \mathrm{N}$ '-diisopropylethylamine ( $0.65 \mathrm{~g}, 5 \mathrm{mmol}$ ) and stirred at room temperature for 13 h . The mixture was poured into water and extracted with ethyl acetate. The organic layer was washed with sat. $\mathrm{NaHCO}_{3}$ aq. and brine followed by drying over $\mathrm{MgSO}_{4}$. After removal of the solvent in vacuo, the residue was purified by column chromatography $\left(\mathrm{SiO}_{2}\right.$, hexane / $\mathrm{EtOAc}=7 / 3)$.

The solution of Ph-C5-Phe-Me in 1 M LiOH aq. / THF (4/1, 10 mL ) was stirred at $60^{\circ} \mathrm{C}$ for 3 h and then cooled to ambient temperature. 1 M HCl and EtOAc were added to the mixture and the aqueous phase was extracted ( $30 \mathrm{~mL} \times 2$ ) with EtOAc. The organic phases where combined, dried over $\mathrm{MgSO}_{4}$, filtered and concentrated in vacuo. The crude solid was recrystallised in hexane / EtOAc to give the product $\mathrm{Ph}-\mathrm{C} 5-\mathrm{Phe}$ ( $544 \mathrm{mg}, 84 \%$ ).

## Nonanoyl-L-Phenylalanine (C9-Phe):

${ }^{1} \mathrm{H}-\mathrm{NMR}\left(\mathrm{DMSO}_{6}\right) \delta: 12.66(1 \mathrm{H}, \mathrm{s}), 8.11(1 \mathrm{H}, \mathrm{d}, J=8.3 \mathrm{~Hz}), 7.28-7.17(5 \mathrm{H}, \mathrm{m}), 4.42(1 \mathrm{H}, \mathrm{td}, J$ $=9.0,4.2 \mathrm{~Hz}), 3.05(1 \mathrm{H}, \mathrm{dd}, J=13.7,4.4 \mathrm{~Hz}), 2.82(1 \mathrm{H}, \mathrm{dd}, J=13.7,10.2 \mathrm{~Hz}), 2.02(2 \mathrm{H}, \mathrm{t}, J=7.3$ $\mathrm{Hz}), 1.37(2 \mathrm{H}$, quin, $J=7.3 \mathrm{~Hz}), 1.29-1.10(10 \mathrm{H}, \mathrm{m}), 0.86(3 \mathrm{H}, \mathrm{t}, J=6.8 \mathrm{~Hz})$. ESI-MS: $m / z 328.19$ $\left([\mathrm{M}+\mathrm{Na}]^{+}\right), 350.17\left([\mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right), 633.40\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right), 655.38\left([2 \mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right), 938.60\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right)$, $960.58\left([3 \mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right)$.

## Nonanoyl-L-Tryptophan (C9-Trp):

${ }^{1} \mathrm{H}-$ NMR (DMSO-D ${ }_{6}$ ) $\delta: 12.55(1 \mathrm{H}, \mathrm{s}), 10.82(1 \mathrm{H}, \mathrm{s}), 8.03(1 \mathrm{H}, \mathrm{d}, \mathrm{J}=8.3 \mathrm{~Hz}), 7.52(1 \mathrm{H}, \mathrm{d}, \mathrm{J}=7.8$ $\mathrm{Hz}), 7.32(1 \mathrm{H}, \mathrm{d}, \mathrm{J}=8.3 \mathrm{~Hz}), 7.12(1 \mathrm{H}, \mathrm{d}, \mathrm{J}=2.0 \mathrm{~Hz}), 7.05(1 \mathrm{H}, \mathrm{t}, \mathrm{J}=7.3 \mathrm{~Hz}), 6.97(1 \mathrm{H}, \mathrm{t}, \mathrm{J}=7.3$ $\mathrm{Hz}), 4.46(1 \mathrm{H}, \mathrm{td}, \mathrm{J}=8.3,4.9 \mathrm{~Hz}), 3.15(1 \mathrm{H}, \mathrm{dd}, \mathrm{J}=14.6,4.9 \mathrm{~Hz}), 2.98(1 \mathrm{H}, \mathrm{dd}, \mathrm{J}=14.6,8.8 \mathrm{~Hz})$, $2.05(2 \mathrm{H}, \mathrm{t}, J=7.1 \mathrm{~Hz}), 1.40(2 \mathrm{H}$, quin, $J=7.2 \mathrm{~Hz}), 1.32-1.08(10 \mathrm{H}, \mathrm{m}), 0.85(3 \mathrm{H}, \mathrm{t}, \mathrm{J}=6.8 \mathrm{~Hz})$. ESI-MS: $m / z 367.21\left([\mathrm{M}+\mathrm{Na}]^{+}\right), 711.43\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right), 1055.64\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right)$

## (5-phenylpentanoyl)-L-phenylalanine (Ph-C5-Phe):

${ }^{1} \mathrm{H}$ NMR (DMSO- $d_{6}, 600 \mathrm{MHz}$ at $\left.80^{\circ} \mathrm{C}\right) \delta: 12.30(1 \mathrm{H}$, brs), $7.78(1 \mathrm{H}, \mathrm{d}, J=6.6 \mathrm{~Hz}), 7.26-7.13(10 \mathrm{H}$, m), $4.48(1 \mathrm{H}, \mathrm{td}, J=9.0,4.8 \mathrm{~Hz}), 3.06(1 \mathrm{H}, \mathrm{dd}, J=14.4,5.4 \mathrm{~Hz}), 2.87(1 \mathrm{H}, \mathrm{dd}, J=14.4,9.6 \mathrm{~Hz})$, $2.52(1 \mathrm{H}, \mathrm{t}, J=7.2 \mathrm{~Hz}), 2.09(2 \mathrm{H}, \mathrm{t}, J=7.2 \mathrm{~Hz}), 1.47(4 \mathrm{H}, \mathrm{m}) . \mathrm{ESI}-\mathrm{MS}: m / z 348.16\left([\mathrm{M}+\mathrm{Na}]^{+}\right)$ $370.14\left([\mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right) 673.33\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right) 695.32\left([2 \mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right) 324.16\left([\mathrm{M}-\mathrm{H}]^{-}\right) 739.54\left([2 \mathrm{M}-\mathrm{H}]^{-}\right.$ ) $649.32\left([2 \mathrm{M}-\mathrm{H}]^{-}\right)$

## (5-cyclohexylpentanoyl)-L-tryptophan (5CHVA-Trp):

${ }^{1} \mathrm{H}$ NMR (DMSO- $\left.d_{6}, 400 \mathrm{MHz}\right) \delta: 12.55(1 \mathrm{H}, \mathrm{brs}), 10.81(1 \mathrm{H}, \mathrm{s}), 8.03(1 \mathrm{H}, \mathrm{d}, J=7.6 \mathrm{~Hz}), 7.53(1 \mathrm{H}$, d, $J=7.6 \mathrm{~Hz}), 7.32(1 \mathrm{H}, \mathrm{d}, J=8.0 \mathrm{~Hz}), 7.12(1 \mathrm{H}, \mathrm{d}, J=2.0 \mathrm{~Hz}), 7.05(1 \mathrm{H}, \mathrm{t}, J=7.4 \mathrm{~Hz}), 6.75(1 \mathrm{H}, \mathrm{t}$, $J=7.2 \mathrm{~Hz}), 4.46(1 \mathrm{H}, \mathrm{td}, J=8.4,5.2 \mathrm{~Hz}), 3.15(1 \mathrm{H}, \mathrm{dd}, J=14.4,5.2 \mathrm{~Hz}), 2.98(1 \mathrm{H}, \mathrm{dd}, J=14.8,8.8$ $\mathrm{Hz}), 2.05(2 \mathrm{H}, \mathrm{m}, J=7.6,1.6 \mathrm{~Hz}), 1.62(5 \mathrm{H}, \mathrm{m}, J=9.6 \mathrm{~Hz}), 1.38(2 \mathrm{H}, \mathrm{m}, J=7.2 \mathrm{~Hz}), 1.21-1.05(8 \mathrm{H}$, m), $0.79(2 \mathrm{H}, \mathrm{q}, J=7.2 \mathrm{~Hz}) . \quad: \mathrm{MS} m / z 371.24\left([\mathrm{M}+\mathrm{H}]^{+}\right) 393.22\left([\mathrm{M}+\mathrm{Na}]^{+}\right) 741.47\left([2 \mathrm{M}+\mathrm{H}]^{+}\right)$ $763.45\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right) 1133.68\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right) 1503.91\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right) 369.21\left([\mathrm{M}-\mathrm{H}]^{-}\right) 739.44\left([2 \mathrm{M}-\mathrm{H}]^{-}\right)$ 1109.67 ([3M-H] $\left.{ }^{-}\right) 1479.90\left([4 \mathrm{M}-\mathrm{H}]^{-}\right)$

## (5-cyclohexylpentanoyl)-L-phenylalanine (5CHVA-Phe):

${ }^{1} \mathrm{H}$ NMR (DMSO- $\left.d_{6}, 400 \mathrm{MHz}\right) \delta: 12.78(1 \mathrm{H}, \mathrm{s}), 8.22(1 \mathrm{H}, \mathrm{d}, J=8.4 \mathrm{~Hz}), 7.39-7.29(5 \mathrm{H}, \mathrm{m}), 4.52$ $(1 \mathrm{H}, \mathrm{td}, J=13.2,4.8 \mathrm{~Hz}), 3.16(1 \mathrm{H}, \mathrm{dd}, J=13.6,4.4 \mathrm{~Hz}), 2.94(1 \mathrm{H}, \mathrm{dd}, J=14.0,10.0 \mathrm{~Hz}), 2.14(2 \mathrm{H}$, $\mathrm{t}, J=7.6 \mathrm{~Hz}), 1.74(5 \mathrm{H}, \mathrm{m}), 1.46(2 \mathrm{H}, \mathrm{q}, J=6.8 \mathrm{~Hz}), 1.34-1.20(8 \mathrm{H}, \mathrm{m}), 0.91(2 \mathrm{H}, \mathrm{q}, J=11.0 \mathrm{~Hz})$. ESI-MS: m/z $354.22\left([\mathrm{M}+\mathrm{Na}]^{+}\right) \quad 685.44\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right) \quad 1016.66\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right) 330.21\left([\mathrm{M}-\mathrm{H}]^{-}\right)$ $661.44\left([2 \mathrm{M}-\mathrm{H}]^{-}\right)$

## (2-(4-isobutylphenyl)propanoyl)-L-phenylalanine (R-Ibu-L-Phe):

${ }^{1} \mathrm{H}-\mathrm{NMR}\left(\mathrm{CDCl}_{3}\right) \delta: 7.82(1 \mathrm{H}, \mathrm{br} \mathrm{s}), 7.17-7.08(7 \mathrm{H}, \mathrm{m}), 6.82(2 \mathrm{H}, \mathrm{d}, J=7.3 \mathrm{~Hz}), 5.76(1 \mathrm{H}, \mathrm{d}, J=7.3$ $\mathrm{Hz}), 4.83(1 \mathrm{H}, \mathrm{q}, J=6.2 \mathrm{~Hz}), 3.51(1 \mathrm{H}, \mathrm{q}, J=7.0 \mathrm{~Hz}), 3.08(1 \mathrm{H}, \mathrm{dd}, J=13.9,5.1 \mathrm{~Hz}), 2.99(1 \mathrm{H}, \mathrm{dd}$, $J=13.9,6.1 \mathrm{~Hz}), 2.48(2 \mathrm{H}, \mathrm{d}, J=7.3 \mathrm{~Hz}), 1.91-1.84(1 \mathrm{H}, \mathrm{m}), 1.46(3 \mathrm{H}, \mathrm{d}, J=6.8 \mathrm{~Hz}), 0.92(6 \mathrm{H}, \mathrm{d}$, $J=6.8 \mathrm{~Hz})$. ESI-MS: $m / z 376.18\left([\mathrm{M}+\mathrm{Na}]^{+}\right), 729.38\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right), 1082.58\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right), 1104.57$ $\left([3 \mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right)$

## (2-(4-isobutylphenyl)propanoyl)-L-phenylalanine (S-Ibu-L-Phe):

${ }^{1} \mathrm{H}-\mathrm{NMR}\left(\mathrm{CDCl}_{3}\right) \delta: 8.06(1 \mathrm{H}, \mathrm{br} \mathrm{s}), 7.22-7.20(3 \mathrm{H}, \mathrm{m}), 7.10-7.07(4 \mathrm{H}, \mathrm{m})$, 6.96-6.94 (2H, m), 5.76
$(1 \mathrm{H}, \mathrm{d}, J=6.8 \mathrm{~Hz}), 4.74(1 \mathrm{H}, \mathrm{q}, J=6.3 \mathrm{~Hz}), 3.54(1 \mathrm{H}, \mathrm{q}, J=7.2 \mathrm{~Hz}), 3.15(1 \mathrm{H}, \mathrm{dd}, J=13.8,5.2$ $\mathrm{Hz}), 3.00(1 \mathrm{H}, \mathrm{dd}, J=14.0,6.7 \mathrm{~Hz}), 2.46(2 \mathrm{H}, \mathrm{d}, J=7.3 \mathrm{~Hz}), 1.91-1.80(1 \mathrm{H}, \mathrm{m}), 1.49(3 \mathrm{H}, \mathrm{d}, J=7.3$ $\mathrm{Hz}), 0.90(6 \mathrm{H}, \mathrm{d}, J=6.6 \mathrm{~Hz})$. ESI-MS: $m / z 376.19\left([\mathrm{M}+\mathrm{Na}]^{+}\right), 729.40\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right), 1082.60$ $\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right), 1104.58\left([3 \mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right)$

## ((benzyloxy)carbonyl)-L-prolyl-L-methionine (Z-Pro-Met):

${ }^{1} \mathrm{H}$ NMR (DMSO- $\left.d_{6}, 400 \mathrm{MHz}\right) \delta: 12.67(1 \mathrm{H}, \mathrm{s}), 8.25(1 \mathrm{H}, \mathrm{dd}, J=16.0,8.0 \mathrm{~Hz}), 7.37-7.28(5 \mathrm{H}, \mathrm{m})$, 5.09-4.94 ( $2 \mathrm{H}, \mathrm{m}$ ), $4.28(1 \mathrm{H}, \mathrm{dd}, J=8.4,3.6 \mathrm{~Hz}), 4.23(1 \mathrm{H}, \mathrm{d}, J=8.8 \mathrm{~Hz}), 3.49-3.39(2 \mathrm{H}, \mathrm{m}), 2.40-$ $2.33(2 \mathrm{H}, \mathrm{m}), 2.16-2.14(1 \mathrm{H}, \mathrm{m}), 2.04(1 \mathrm{H}, \mathrm{s}), 1.96(3 \mathrm{H}, \mathrm{s}), 1.84-1.80(4 \mathrm{H}, \mathrm{m})$. ESI-MS $m / z 403.14$ $\left([\mathrm{M}+\mathrm{Na}]^{+}\right) 783.29\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right) 1163.44\left([3 \mathrm{M}+\mathrm{Na}]^{+}\right) 379.13\left([\mathrm{M}-\mathrm{H}]^{-}\right) 759.29\left([2 \mathrm{M}-\mathrm{H}]^{-}\right)$

## (Benzyloxy)carbonyl)-L-prolyl-L-phenylalanine (Z-Pro-Phe):

${ }^{1} \mathrm{H}-\mathrm{NMR}\left(\mathrm{DMSO}_{6}\right) \delta: 12.70(1 \mathrm{H}, \mathrm{s}), 8.17(1 \mathrm{H}, \mathrm{dd}, J=39.0,7.8 \mathrm{~Hz}), 7.38-7.18(10 \mathrm{H}, \mathrm{m}), 5.10-$ $4.84(2 \mathrm{H}, \mathrm{m}), 4.50-4.39(1 \mathrm{H}, \mathrm{m}), 4.22(1 \mathrm{H}, \mathrm{d}, J=8.3 \mathrm{~Hz}), 3.45-3.35(2 \mathrm{H}, \mathrm{m}), 3.07-2.86(2 \mathrm{H}, \mathrm{m})$, 2.09-1.99 (1H, m), 1.77-1.68 (3H, m). ESI-MS: m/z $419.16\left([\mathrm{M}+\mathrm{Na}]^{+}\right), 441.14\left([\mathrm{M}-\mathrm{H}+2 \mathrm{Na}]^{+}\right)$, $859.30\left([2 \mathrm{M}-2 \mathrm{H}+3 \mathrm{Na}]^{+}\right)$

## (Benzyloxy)carbonyl)glycyl-L-phenylalanine (Z-Gly-Phe):

${ }^{1} \mathrm{H}-\mathrm{NMR}\left(\mathrm{DMSO}_{6}\right) \delta: 12.78(1 \mathrm{H}, \mathrm{s}), 8.12(1 \mathrm{H}, \mathrm{d}, J=7.8 \mathrm{~Hz}), 7.42-7.20(11 \mathrm{H}, \mathrm{m}), 5.02(2 \mathrm{H}, \mathrm{s})$, $4.44(1 \mathrm{H}, \mathrm{td}, J=8.1,5.2 \mathrm{~Hz}), 3.61(2 \mathrm{H}, \mathrm{qd}, J=16.7,6.4 \mathrm{~Hz}), 3.04(1 \mathrm{H}, \mathrm{dd}, J=13.9,5.1 \mathrm{~Hz}), 2.89$ $(1 \mathrm{H}, \mathrm{dd}, J=13.4,9.0 \mathrm{~Hz})$. ESI-MS: $m / z 379.12\left([\mathrm{M}+\mathrm{Na}]^{+}\right)$, $735.26\left([2 \mathrm{M}+\mathrm{Na}]^{+}\right)$

## Hydroxylation of ethylbenzene

The oxidation of ethylbenzene was carried out in 20 mM Tris- $\mathrm{HCl}(\mathrm{pH} 7.4)$ buffer containing 100 mM KCl at $25^{\circ} \mathrm{C}$ for 5 min in the presence of $0.5 \mu \mathrm{M} \mathrm{P450BM3}$,10 mM ethylbenzene, 5 mM NADPH and $100 \mu \mathrm{M}$ decoy molecule. Ethybenzene and decoy molecules were dissolved in DMSO and added to the reaction mixture. The final volume of the reaction mixture was 1 mL containing $1.5 \%(\mathrm{v} / \mathrm{v})$ DMSO. NADPH consumption was monitored by the absorption at 340 nm by UV-vis spectra. After 5 min reaction, dichloromethane was added to the reaction mixture to quench the reaction, and the products were extracted into the organic layer. Phenylmethyl acetate was also added to reaction mixture as an internal standard. Reaction extract was analysed by GC and GC-MS.

The conditions for GC were as follows: column temperature: $110^{\circ} \mathrm{C}(30 \mathrm{~min}) ; 10^{\circ} \mathrm{C} \mathrm{min}^{-1} ; 220^{\circ} \mathrm{C}$ $(15 \mathrm{~min})$, injection temperature: $250^{\circ} \mathrm{C}$, interface temperature: $250^{\circ} \mathrm{C}$, detector temperature: $250^{\circ} \mathrm{C}$, carrier gas: helium. The retention times of ethylbenzene, products and the internal standard were as follows: ethylbenzene ( 2.1 min ), phenylmethyl acetate $(11.1 \mathrm{~min}),(R)$-1-phenylethanol $(12.7 \mathrm{~min})$, ( $S$ )-1-phenylethanol ( 13.6 min ), 2-ethylphenol ( 23.2 min ) and 4-ethylphenol ( 27.0 min ). GC-MS condition: column temperature: $110^{\circ} \mathrm{C}(30 \mathrm{~min}) ; 20^{\circ} \mathrm{C} \mathrm{min}^{-1} ; 220^{\circ} \mathrm{C}(15 \mathrm{~min})$, injection temperature: $250^{\circ} \mathrm{C}$, interface temperature: $200^{\circ} \mathrm{C}$, ion source temperature: $200^{\circ} \mathrm{C}$, carrier gas: helium. The retention times of products and the internal standard were as follows: phenylmethyl acetate (11.7 $\min ),(R)$-1-phenylethanol (13.6 min), (S)-1-phenylethanol (14.6 min), 2-ethylphenol ( 24.7 min ) and 4-ethylphenol ( 28.2 min ). Reactions using each decoy molecule were performed at least three times. The unit of turnover rate is expressed as (nmol product) per min per (nmol P450).

## Hydroxylation of propylbenzene, indane, and tetralin

The oxidation of propylbenzene, indane, and tetralin were conducted in the same manner as ethylbenzene, but the products obtained from these substrates were derivatised to silylated alcohols by BSTFA-TMCS (99:1). The products of propylbenzene and indane oxidation were analysed by GC-MS, and those of tetralin oxidation were analysed by GC. The conditions for GC or GC-MS analysis and retention times of substrates, products and the internal standard were as follows:

Propylbenzene: GC-MS condition: column temperature: $90^{\circ} \mathrm{C}(30 \mathrm{~min}) ; 20^{\circ} \mathrm{C} \mathrm{min}^{-1} ; 220^{\circ} \mathrm{C}(10 \mathrm{~min})$, injection temperature: $250^{\circ} \mathrm{C}$, interface temperature: $200^{\circ} \mathrm{C}$, ion source temperature: $200^{\circ} \mathrm{C}$, carrier gas: helium. Retention times: propylbenzene ( 5.3 min ), ( $S$ )-1-phenylpropanol-BSTFA derivative $(15.0 \mathrm{~min}),(R)$-1-phenylpropanol-BSTFA derivative ( 15.5 min ), phenylmethyl acetate ( 29.5 min ), 4-propylphenol-BSTFA derivative ( 33.5 min ).

Indane: GC-MS condition: column temperature: $110^{\circ} \mathrm{C}(30 \mathrm{~min}) ; 20^{\circ} \mathrm{C} \mathrm{min}^{-1} ; 220^{\circ} \mathrm{C}(15 \mathrm{~min})$, injection temperature: $250^{\circ} \mathrm{C}$, interface temperature: $200^{\circ} \mathrm{C}$, ion source temperature: $200^{\circ} \mathrm{C}$, carrier gas: helium. Retention times: indane ( 4.7 min ), phenylmethyl acetate ( 11.9 min ), $(R)$-1-indanolBSTFA derivative ( 18.7 min ), $(S)$-1-indanol-BSTFA derivative ( 19.7 min ), 4-indanol-BSTFA derivative ( 23.4 min ), 1-indanone ( 28.7 min ), and 5-indanol-BSTFA derivative ( 30.7 min ). The peak of 1-indanone and 4-indanol-BSTFA derivative was assigned by GC-MS fragmentation.

Tetralin: GC condition: column temperature: $110^{\circ} \mathrm{C}(40 \mathrm{~min}) ; 30^{\circ} \mathrm{C} \mathrm{min}^{-1} ; 220^{\circ} \mathrm{C}(10 \mathrm{~min})$, injection temperature: $250^{\circ} \mathrm{C}$, interface temperature: $250^{\circ} \mathrm{C}$, detector temperature: $250^{\circ} \mathrm{C}$, carrier gas: helium. Retention times: tetralin ( 9.5 min ), phenylmethyl acetate ( 11.1 min ), ( $R$ )-1-tetralol-BSTFA derivative $(32.0 \mathrm{~min})$ and $(S)$-1-tetralol-BSTFA derivative $(34.3 \mathrm{~min})$. No further products were
detected by GC-MS analysis.

Table S1. Hydroxylation of ethylbenzene catalysed by WT P450BM3 in the presence of decoy molecules. ${ }^{\text {a }}$


${ }^{\text {a }}$ Reaction conditions: 10 mM ethylbenzene, 5 mM NADPH, $100 \mu \mathrm{M}$ decoy molecule, 100 mM KCl and $0.5 \mu \mathrm{M}$ P450BM3 in 20 mM Tris- HCl buffer ( pH 7.4 ) at $25^{\circ} \mathrm{C}$ for $5 \mathrm{~min}{ }^{\mathrm{b}}([$ Products $] /[\mathrm{NADPH}$ consumption] $) \times 100$.

Table S2. Hydroxylation of propylbenzene, indane, and tetralin.catalysed by WT P450BM3 in the presence of decoy molecules. ${ }^{\text {a }}$

| Reaction scheme | Decoy | Turnover rate | $\alpha-\mathrm{OH}$ | ee. | Coupling | Further |
| :---: | :---: | :---: | :---: | :---: | :---: | :---: |
|  | molecule | [/min/P450] | [\%] | [\%] (R/S) | efficiency ${ }^{\text {b }}$ [\%] | products |
| Propylbenzene | None | $165 \pm 8$ | >99 | $89(R)$ | 32 | 4-propylphenol (trace) |
|  | PFC9-Phe | $305 \pm 12$ | >99 | $94(R)$ | 32 | 4-propylphenol (trace) |
|  | 5CHVA-Phe | $479 \pm 15$ | >99 | 95 (R) | 66 | 4-propylphenol (trace) |
|  | (R)-Ibu-Phe | $308 \pm 13$ | 96 | 86 (R) | 51 | 4-propylphenol (4\%) |
|  | Z-Pro-Phe | $175 \pm 13$ | 97 | 81 (R) | 42 | 4-propylphenol (3\%) |
| Indane | None | $36 \pm 2$ | >99 | 16 (S) | 8 | indanone (trace) |
|  | PFC9-Phe | $216 \pm 8$ | 96 | 6 (R) | 24 | 4-indanol (4\%), 5-indanol (trace), indanone (trace) |
|  | 5CHVA-Phe | $435 \pm 27$ | 97 | 53 (R) | 44 | 4-indanol (3\%), 5-indanol (trace), indanone (trace) |
|  | (R)-Ibu-Phe | $314 \pm 5$ | 90 | 45 (S) | 39 | 4-indanol (7\%), 5-indanol (3\%), indanone (trace) |
|  | Z-Pro-Phe | $302 \pm 10$ | 90 | 56 (S) | 43 | 4-indanol (7\%), 5-indanol (3\%), indanone (trace) |
| Tetralin | None | $20 \pm 1$ | >99 | 55 (S) | 7 | - |
|  | PFC9-Phe | $212 \pm 16$ | >99 | 69 (S) | 17 | - |
|  | 5CHVA-Phe | $320 \pm 31$ | >99 | 13 (S) | 49 | - |
|  | (R)-Ibu-Phe | $332 \pm 40$ | >99 | $89(S)$ | 66 | - |
|  | Z-Pro-Phe | $260 \pm 15$ | >99 | 96 (S) | 82 | - |

[^0] ([Products] $][$ NADPH consumption $]) \times 100$.


Fig. S1 Chromatograms of products in benzylic hydroxylation reactions by P450BM3 with 5CHVAPhe (solid line) and Z-Pro-Phe (dashed line). (a): Ethylbenzene hydroxylation, (b): propylbenzene hydroxylation, (c): indane hydroxylation, (d): tetralin hydroxylation.

## Crystallisation of P450BM3 with 5CHVA-L-Trp, Data Collection, and Refinement.

Buffer containing purified P450BM3 was exchanged with 50 mM Tris- HCl ( pH 7.4 ) containing 200 $\mu \mathrm{M}$ of 5-cyclohexylvaleroyl-L-tryptophan and $1.0 \%(\mathrm{v} / \mathrm{v})$ dimethyl sulfoxide and concentrated to 30 $\mathrm{mg} / \mathrm{mL}$ by centrifugation using Amicon Ultra filter units (Millipore,Co.). An aliquot of the concentrated P450BM3 solution ( $1 \mu \mathrm{M}$ ) was mixed with $1 \mu \mathrm{M}$ of a reservoir solution composed of 50 mM Tris- $\mathrm{HCl}(\mathrm{pH} 8.5), 260 \mathrm{mM} \mathrm{MgCl}$, and $17 \%$ (w/v) PEG8000. Cocrystals were grown by sitting-drop vapour diffusion method at $20^{\circ} \mathrm{C}$. Crystals were harvested and flash-cooled in liquid nitrogen. X-ray diffraction data sets were collected at SPring-8 (Hyogo, Japan) on the beamline BL26B2 equipped with a MAR225 CCD detector at $1.0 \AA$ wavelength and 100 K . The program HKL2000 ${ }^{4}$ was used for integration of diffraction intensities and scaling. The structure was solved by molecular replacement with MolRep. ${ }^{5}$ The structure of P450BM3 with $N$-perfluorononanoyl-Ltryptophan (3WSP) was used as a search model. Model building and refinement were performed using COOT ${ }^{6}$ and REFMAC5. ${ }^{7}$ The 5-cyclohexylvaleroyl-L-tryptophan model was generated employing a Dundee PRODRG server ${ }^{8}$ and used in the refinement with COOT and REFMAC5. The final refinement statistics are summarised in Table S3.

Table S3 Data collection and refinement statistics

| PDB code | 5XHJ |
| :---: | :---: |
| Data collection |  |
| Wavelength ( $\AA$ ) | 1.000 |
| Space group | $P 21$ |
| Cell dimensions |  |
| $a, b, c(\AA)$ | 58.653, 146.779, 63.642 |
| $\alpha, \beta, \gamma\left({ }^{\circ}\right)$ | 90.000, 97.251, 90.000 |
| Resolution ( $\AA$ ) | 50.00-2.00 (2.00-2.07) |
| No. of total observed reflections | 245648 |
| No. of unique reflections | 69112 |
| $R_{\text {merge }}{ }^{a, b}$ (\%) | 3.3 (14.7) |
| Completeness ${ }^{\text {a }}$ (\%) | 96.7 (95.9) |
| $I / \sigma(I){ }^{a}$ | 35.2 (7.4) |
| Redundancy ${ }^{a}$ | 3.6 (3.5) |
| Refinement statistics |  |
| Resolution range ( $\AA$ ) | 20.00-2.00 |
| No. of monomer/asymmetric unit | 2 |
| $R_{\text {work }} / R_{\text {free }}{ }^{c, d}(\%)$ | 17.8/20.6 |
| RMSD bond length ${ }^{e}(\AA)$ | 0.008 |
| RMSD bond angles ${ }^{e}\left({ }^{\circ}\right)$ | 1.3159 |
| No. of atoms | 8040 |
| Average $B$-factor ( $\AA^{2}$ ) | 30 |

${ }^{a}$ The values in parentheses are for the highest resolution shell.
${ }^{b} R_{\text {merge }}=\Sigma_{h k l} \Sigma_{i}\left|I_{i}(h k l)-<\mathrm{I}(\mathrm{hkl})>\right| / \Sigma_{h k l} \Sigma_{i} I_{i}(h k l)$, where $<I(h k l)>$ is the average intensity of the $i$ observations.
${ }^{c} R_{\text {work }}=\Sigma_{h k \mid}\left|F_{\text {obs }}(h k l)-F_{\text {calc }}(h k l)\right| \Sigma_{h k l} F_{\text {obs }}(h k l)$, where $F_{\text {obs }}$ and $F_{\text {calc }}$ are the observed and calculated structure factors, respectively.
${ }^{d} R_{\text {free }}$ was calculated with $5 \%$ of the reflections that were not included in the refinement.
${ }^{{ }^{e}}$ r. m. s. d. $=$ root mean square deviation


Fig. S2 Superimposed structures of the I helix and residues contained in the I helix (Phe261, Leu262, Ile263, Ala264, Gly265, His266, Glu267 and Thr268).


Fig. S3 Cavities of the active site depicted by surface model. (Left) 5CHVA-Trp- and (right) Z-ProPhe bound P450BM3

## Docking simulation

The docking simulations were performed by AutoDockFR and AutoGridFR. ${ }^{9}$ To prepare the active species compound I, receptor structures were edited by Maestro (Schrödinger Release 2016-2: Maestro, version 10.6, Schrödinger, LLC, New York, NY, 2016). ${ }^{10}$ Indane model structure was obtained from Pubchem database. ${ }^{11}$ Structures of receptor and ligand for simulations were prepared by AutoDock Tools. ${ }^{12}$ Phe87, Leu262, Ile263, His266, Glu267 and Thr268, located around haem, were set as flexible residues. Partial charge of compound I species and Cys 400 were derived from previous research. ${ }^{13}$ The docking pockets were generated by AutoGridFR and pockets at the haem distal side were chosen for each simulation. After each simulation, results were visualised and hydrogen atoms of indane were added by Pymol. ${ }^{14}$

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[^0]:    ${ }^{\text {a }}$ Reaction conditions: 10 mM substrate, 5 mM NADPH, $100 \mu \mathrm{M}$ decoy molecule, 100 mM KCl and $0.5 \mu \mathrm{M}$ P450BM3 in 20 mM Tris- HCl buffer ( pH 7.4 ) at $25^{\circ} \mathrm{C}$ for 5 min . ${ }^{\text {b }}$

