

Supplementary Materials for

Intramolecular Heme Ligation of the CYP2C9-

R108H Mutant Demonstrates Pronounced

Conformational Flexibility of the B-C Loop

Region: Implications for Substrate Binding

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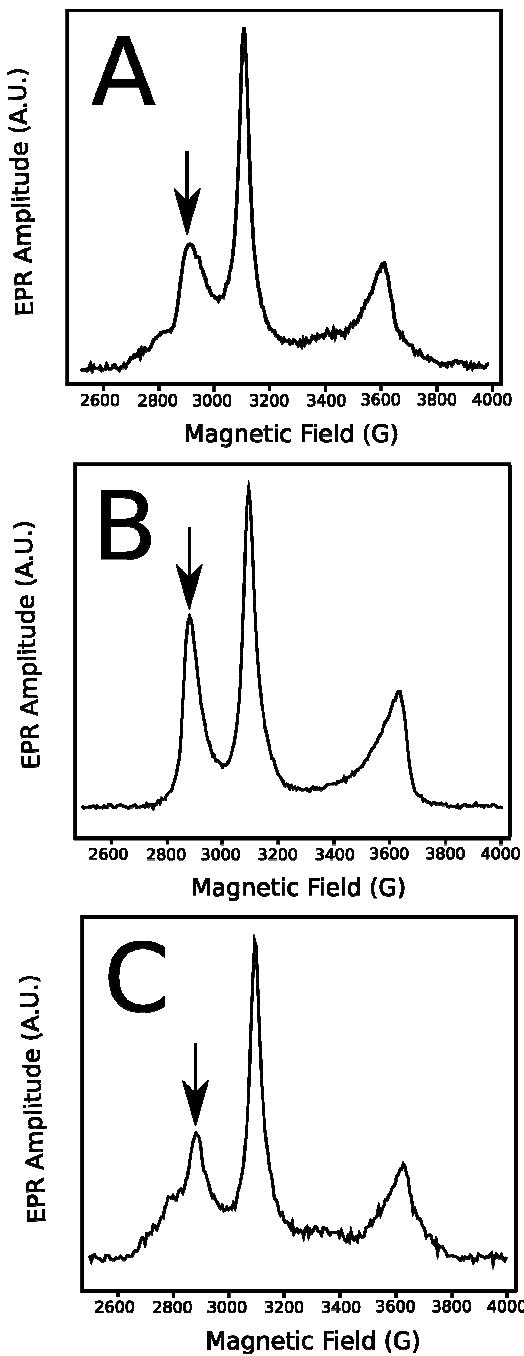


Figure S1. Spin echo-detected EPR spectra are shown for A) CYP2C9, B) the R108H mutant and C) CYP2C9 with 100 mM imidazole. The spectra were measured with the spacing between the two microwave pulses at $\tau=160$ ns in order to emphasize the principal g-values as a result of the orientation dependence of the ESEEM and the phase memory relaxation of the sample. The g-values of the EPR peaks under the various conditions were similar. The arrow denotes the position where the HYSCORE spectra were measured.

Table S1. Additional Force field parameters used for CYP2C9^a

Bond	Energy (kcal mol ⁻¹ Å ⁻²)	Distance (Å) ^b	Reference
Fe-S	87.589	2.377	(I)
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Angle ^c	Energy (kcal mol ⁻¹ rad ⁻²)	Reference	
Fe-S-C _β	21.646	(I)	
N _P -Fe-S	13.277	(I)	
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Dihedral ^c	Energy (kcal mol ⁻¹)	Reference	
C _β -S-Fe-N _P	0.034 ^d	(I)	

^a Fe, heme iron; S, sulfur of C435; C_β, C_β of C435; N_P, pyrrole heme nitrogen.

^b Equilibrium distance value from (I).

^c The angles and dihedrals, used in the simulations, were calculated using the values measured from the 1R9O CYP2C9 X-ray crystal structure (2).

^d Autenrieth et al., 2004 obtained energy values for cytochrome c of 0.04 kcal mol⁻¹(3)

Table S2. Additional force field parameters for R108H CYP2C9^a

Bond	Energy (kcal mol ⁻¹ Å ⁻²)	Distance (Å)	Reference
Fe-N _ε	65 ^b	2.05 ^c	(3)

Angle	Energy (kcal mol ⁻¹ rad ⁻²)	Angle (degrees)	Reference
N _P -Fe-N _ε	50	90°	(4)

Improper Dihedral	Energy (kcal mol ⁻¹ rad ⁻²)	Angle (degrees)	Reference
N _ε -C _δ -C _ε -Fe	29.4	0°	(3)

^a N_ε, C_δ and C_ε correspond to atoms on H108 of the mutant.

^b Henry et al., 1985 obtained similar values for hemoglobin of 60 kcal mol⁻¹ Å⁻².

^c Intermediate value between 2-2.1 Å that was used previously in (1, 3, 4).

References

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