

## Supporting Information

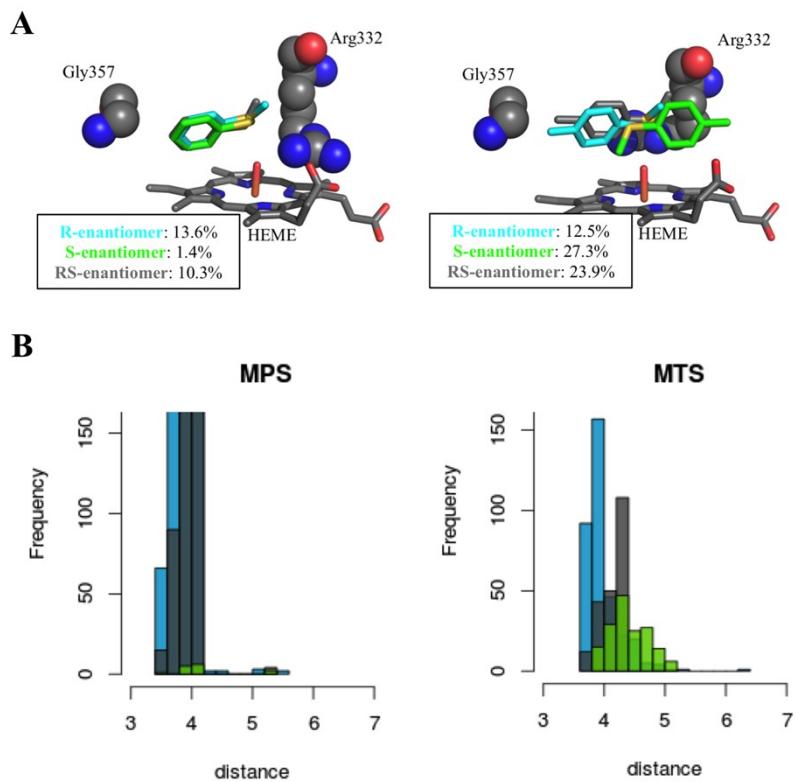
# Asymmetric sulfoxidation by engineering the heme pocket of a dye-decolorizing peroxidase: An experimental and computational study

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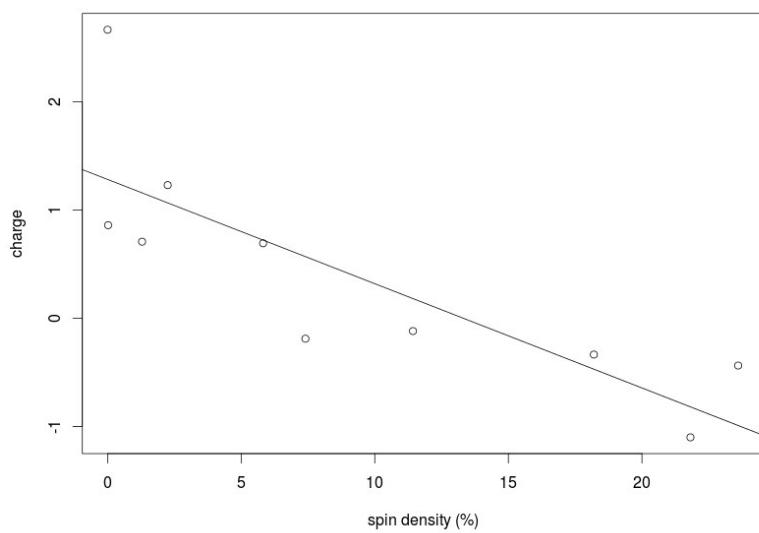
**Table S1** X-ray data collection and refinement statistics of L357G and F359G variants of *A. auricula-judae* DyP

	L357G	F359G
<b>Data collection</b>		
Beamline	ID23-1 (ESRF)	BL13-XALOC (ALBA)
Wavelength (Å)	0.9792	0.9795
Space group	C2	C2
Unit cell parameters (Å, °)	a=100.9, b=61.0, c=81.0; β=95.0	a=104.9, b=56.1, c=82.7; β=97.03
Resolution range (Å)	44.73 - 1.95 (2.05 - 1.95)	46.63 - 1.10 (1.17 - 1.10)
Total reflections	239,457 (35,489)	997,210 (135,507)
Unique reflections	35,827 (5,247)	184,770 (26,363)
Multiplicity	6.7 (6.8)	5.4 (5.1)
Completeness (%)	99.2 (99.9)	98.4 (96.4)
Mean I/σ(I)	12.6 (2.3)	9.5 (2.4)
R <sub>merge</sub> <sup>a</sup>	0.073 (0.729)	0.088 (0.726)
CC <sub>1/2</sub> (%) <sup>b</sup>	99.8 (90.2)	99.7 (77.2)
Mosaicity (°)	0.16	0.15
Wilson B factor (Å <sup>2</sup> )	33.21	7.78
<b>Refinement</b>		
R <sub>work</sub> /R <sub>free</sub>	0.198 (0.367) / 0.23 (0.415)	0.149 (0.231) / 0.158 (0.246)
Working reflections	33,925 (5,206)	182,839 (25,500)
Testing reflections	1,863 (308)	1,897 (280)
Protein atoms (non H)	3,298	3,310
Heme group	43	43
Water molecules	127	576
Mean B factors (Å <sup>2</sup> ):		
-Protein atoms (non H)	46.2	11.9
-Heme group	41.7	7.8
-Water molecules	47.5	24.7
Deviations from ideality:		
-rmsd bond lengths (Å)	0.007	0.006
-rmsd angles (°)	0.876	0.987
Ramachandran statistics:		
-Favoured (%)	97.5	98.9
-Outliers (%)	0	0
PDB code	5IKG	5IKD

Statistics for the highest-resolution shell are shown in parenthesis. <sup>a</sup> $R_{\text{merge}} = \frac{\sum_{hkl} \sum_i |I_i(hkl) - \langle I(hkl) \rangle|}{\sum_{hkl} \sum_i I_i(hkl)}$ , where  $I_i(hkl)$  is the intensity measured for the  $i$ th reflection and  $\langle I(hkl) \rangle$  is the average intensity of all reflections with indices  $hkl$ . <sup>b</sup> $CC_{1/2}$  is the correlation coefficient between two random half datasets (Karplus P.A and Diederichs K., Science, 2012, 336, 1030-1033).



**Fig. S1.** L357G variant substrate oxidation and positioning analysis: **A)** pro-*R*, pro-*S* and pro-*RS* MPS (left) and MTS (right) positioning on the L357G variant. Spin density population on the substrate, from QM/MM calculations, is indicated in each image. **B)** Frequency distance (in Å) distribution of MPS and MTS S atom to Arg332 CZ atom in the L357G variant for the selected reactive structures: last 10 kcal·mol<sup>-1</sup> interaction energies and distances to the heme group below 5 Å. Pro-*S* substrate positions are colored in green, pro-*R* in cyan and pro-*RS* in gray.



**Fig. S2.** Correlation between the substrate spin density and the charge distribution on 10 representative points.