

Electronic Supplementary Information

Theoretical evaluation of a bulky ortho-thioalkyl-azobenzene as an alternative to photocontrol structural cytotoxic effect of metal-free and disulfide oxidized hSOD1 in pathogenesis of ALS

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Table S1. Optimization energies for different functionals through studies of DFT for 1A and 2A molecules.

Functional	<i>trans</i> ortho-TIABP, 1A (Kcal)	<i>trans</i> ortho-TMABP, 2A (Kcal)
B3LYP	-4,1399e ⁻¹⁸	-3.8121e ⁻¹⁸
CAM-B3LYP	-4,1386e ⁻¹⁸	-3.8109e ⁻¹⁸
M06-2X	-4,1381e ⁻¹⁸	-3.8106e ⁻¹⁸
PBE	-4,1370e ⁻¹⁸	-3.8095e ⁻¹⁸

Table S2. Theoretical absorption wavelength (λ) and energy (E) for the calibration of functionals for ortho-S-alkyl substituents (*trans* ortho-TIABP (1A) and *trans* ortho-TMABP (2A), respectively). Theoretical values were calculated using the CPCM solvent model (Dimethyl sulfoxide) and the 6-31++G (d,p) basis set.

Molecule	Functional	Excited State	Transition (CI)	λ (nm)/E (eV)
1A	B3LYP	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.63) H -1 $\rightarrow$ L (0.70)	529/2.36 444/2.79
	CAM-B3LYP	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.50) H -1 $\rightarrow$ L (0.69)	467/2.65 347/3.57
	M06-2X	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.61) H -1 $\rightarrow$ L (0.68)	540/2.29 420/2.95
	PBE	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.61) H -1 $\rightarrow$ L (0.70)	511/2.42 421/2.95
2A	B3LYP	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.64) H -1 $\rightarrow$ L (0.70)	533/2.32 449/2.76
	CAM-B3LYP	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.63) H -1 $\rightarrow$ L (0.67)	454/2.72 353/3.44
	M06-2X	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.66) H -1 $\rightarrow$ L (0.69)	523/2.37 461/2.69
	PBE	$n \rightarrow \pi^*$ $\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.67) H -1 $\rightarrow$ L (0.69)	499/2.49 450/2.76

Table S3. Absorption properties, including vertical transition energies (λ), oscillator strengths (f) and molecular orbitals involved in the excitation features of the ortho-O-alkyl azobenzenes (ortho-OABPs) and their respective hydrazobenzenes. All calculations were made on Gaussian09.

Azo compound	Isomer	Excited State	Transition (CI)	λ (nm)/E (eV)	f
3A	E	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.66)	539/2.29	0.295
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.69)	358/3.46	0.035
3B	E	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.47)	339/3.66	0.464
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.42)	331/3.74	0.083
3C	Z	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.69)	498/2.49	0.014
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.70)	382/3.18	0.003
3D	Z	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.69)	378/3.28	0.113
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.70)	365/3.40	0.047
4A	E	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.65)	541/2.29	0.319
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.56)	351/3.54	0.080
4B	E	$\pi \rightarrow \pi^*$	H $\rightarrow$ L (0.68)	333/3.73	0.060
		$n \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.52)	336/3.69	0.464
4C	Z	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.69)	472/2.63	0.146
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.70)	378/2.63	0.032
4D	Z	$n \rightarrow \pi^*$	H $\rightarrow$ L (0.70)	389/3.19	0.153
		$\pi \rightarrow \pi^*$	H -1 $\rightarrow$ L (0.70)	329/3.19	0.200

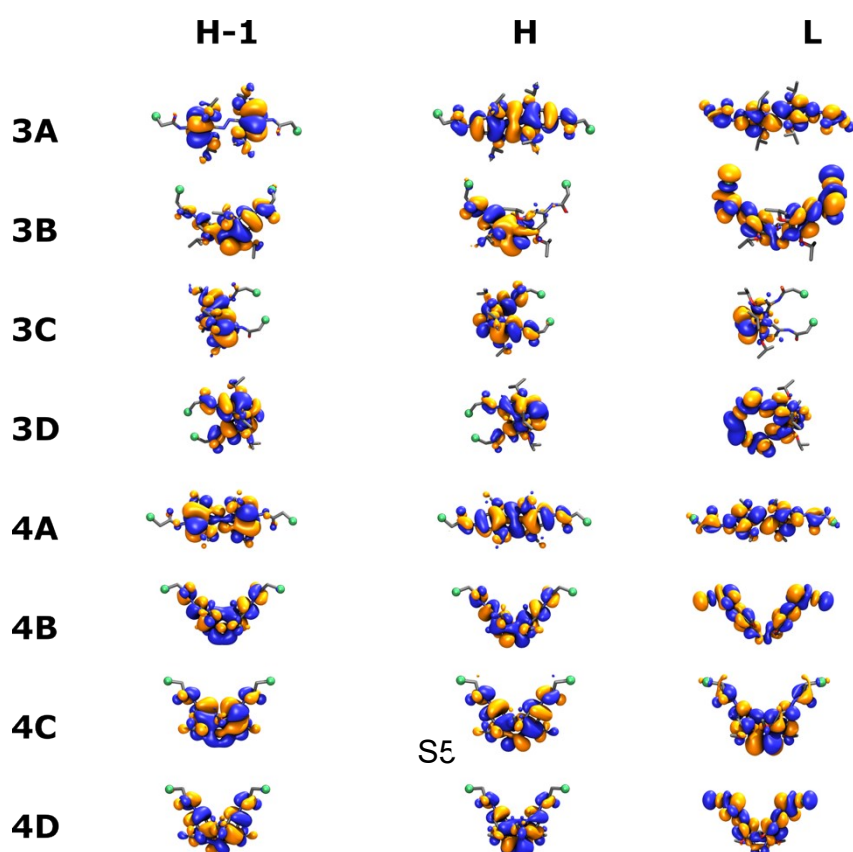


Figure S1. Molecular orbital surface for LUMO (L), HOMO (H) and HOMO-1(H-1) of ortho-O-alkyl azobenzenes (ortho-OABPs) and its corresponding hydrazobenzenes calculated with B3LYP/6-31G(d,p). Solvent was simulated with C-PCM model. 3A and 3B represent ortho-OIABP and their corresponding hydrazobenzenes in the *trans* conformation, while 3C and 3D represent ortho-OIABP and their corresponding hydrazobenzenes in the *cis* conformation. 4A and 4B represent ortho-OMABP and their corresponding hydrazobenzenes in the *trans* conformation, while 4C and 4D represent ortho-OMABP and their corresponding hydrazobenzenes in the *cis* conformation. Representations were obtained from VMD.

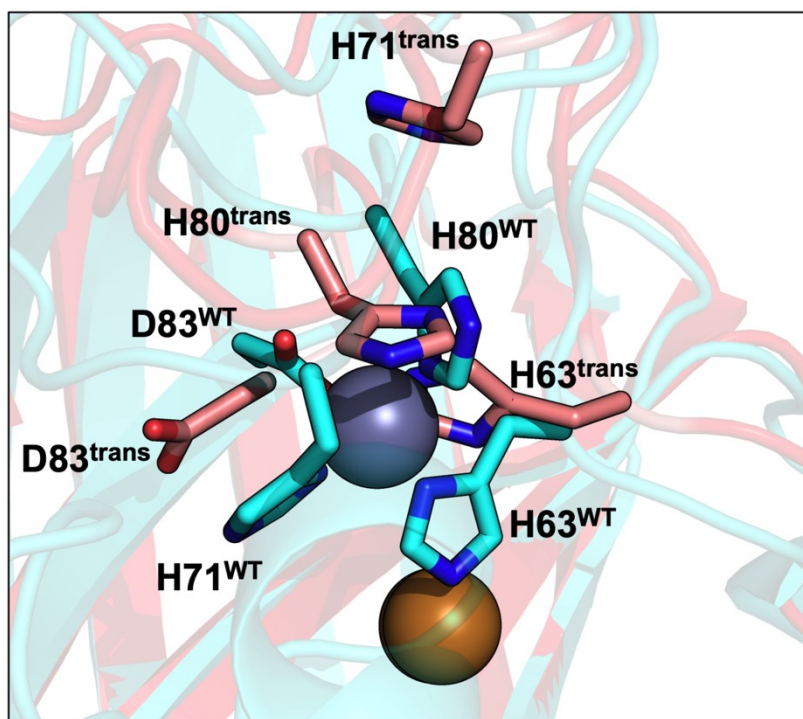


Figure S2. Metal-binding site of apo-hSOD1^{S-S} and apo-hSOD1^{S-Strans}. The position of the amino acid residues that are part of the Zn(II) coordination site in the apo-hSOD1^{S-S} and apo-hSOD1^{S-S} derivatized with one ortho-TABP in *cis* conformation is shown in cyan. The position of the amino acid residues that are part of the Zn(II) coordination site after the transition of the ortho-TABP from *cis* to *trans* conformation is shown in red. Representation obtained from VMD.

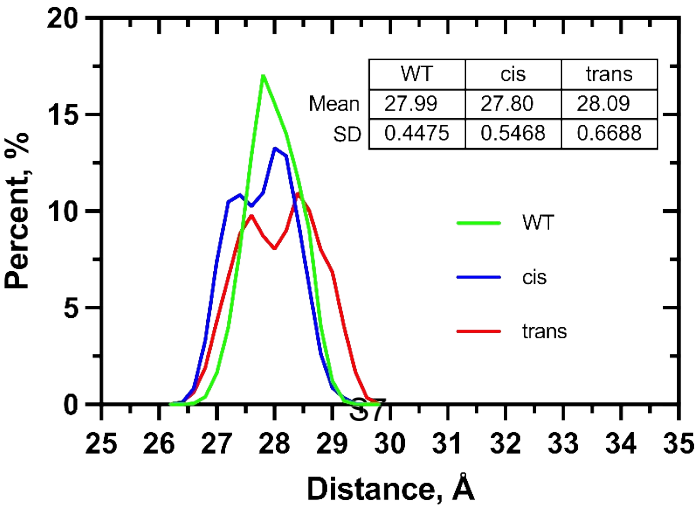


Figure S3. Probability distribution of the distances between the centres of mass of each monomer in apo-hSOD1^{S-S}, apo-hSOD1^{S-Scis} and apo-hSOD1^{S-Strans}, calculated by the adaptive biasing force method (ABF).