

Supplementary Figure S1. Alternate conformation of the free Cys95 in molecule B of Pa-SOD. Electron density maps $(2F_o-F_c)$ and (F_o-F_c) are contoured at 1.2σ and 3σ level respectively.



Supplementary Figure S2. Comparison of loop I structure of eukaryotic CuZnSODs. (*a*) Molecules A (Olive green) and B (Yellow) of Pa-SOD with So-SOD (Orchid), (*b*) Pa-SOD with Hs-SOD (Deep sky blue). The loop I structure in Sc, Sm, Bt and XI-SODs is the same as in Hs-SOD. (*c*) In Sc and Sm-SODs (Gold) an additional water molecule Wt4 is present. For clarity, side-chains are not shown except for Ser10 in Pa-SOD and the residues are labeled according to Pa-SOD numbering.



Supplementary Figure S3. Comparison of metal binding residues with secondary shell stabilization residues of Pa-SOD (Olive green) and Sm-SOD (Gold). The water molecule Wm (orange) in Pa-SOD superimposes with the higher occupancy copper site of Sm-SOD.



Supplementary Figure S4. Comparison of water Ws2 and Arg114 of copper unbound Pa-SOD and the early bound state So-SOD.



Supplementary Figure S5. Movements of the structural water and the C-terminal Greek key loop between copper unbound Pa-SOD (Olive green) and the early copper bound state So-SOD (Orchid).



Supplementary Figure S6. Comparison of active site water molecules of copper unbound Pa-SOD with early chelation state So-SOD. Views of *(a)* Superposition of Cu unbound Pa-SOD (Olive green) and the early copper bound state So-SOD (Orchid) and *(d)* Superposition of molecules A and B of Pa-SOD. Additional five-membered ring water structures are observed for molecule B of Pa-SOD.



Supplementary Figure S7. Comparison of dimeric interfaces of Pa-SOD (Olive green) and So-SOD (Orchid).

Conserved secondary shell interactions in solution and crystal structures of CuZnSODs

- (i) backbone N of His62 with carbonyl O of 135 (Lys/Ser/Thr/Val)
- (ii) backbone N of His70 and carbonyl O of 134 (Lys/Leu/Thr/Pro)
- (iii) backbone N of His79 with the side chain $O^{\delta 2}$ of Asp82
- (iv) backbone N of Asp82 with carbonyl O of His79
- (v) backbone N of His119 and carbonyl O of Gly43
- (vi) carbonyl O of His119 with $N^{\delta 1}$ of His42 [except Sc-SOD where His is replaced by
- Arg (Djinović et al., 1992)]
- (vii) carbonyl O of His79 also with N^{ϵ} of Arg78
- (viii) carbonyl O of His62 with water molecule
- (ix) carbonyl O of His70 with water molecule
- (x) $N^{\epsilon 2}$ of His45 with side chain $O^{\delta 1}$ or $O^{\delta 2}$ of Asp123
- (xi) $N^{\delta 1}$ of His119 with carbonyl O of Gly140
- (xii) N^{ϵ_2} of His70 with side chain O^{δ_1} and O^{δ_2} of Asp123
- (xiii) $N^{\delta 1}$ of His47 with carbonyl O of Gly60
- (xiv) $N^{\epsilon 2}$ of His79 with carbonyl O of 68 (Lys/Arg/Gln) through water molecule.

(xv) Residues His45 and His47 are part of β -strand 4f and interact with residues Val117 and Ala/Ser/Thr115 in the β -strand 7g. A slight movement in the carbonyl O of His47 is found for most of the SOD structures studied so far and thus, the distance between carbonyl O of His47 with N of 115 is >3.0 Å for all the structures.

1. Salt Bride			
	i.	Arg78	Asp100
		$N^{\eta 1}$ Arg78	$O^{\delta 1}$ Asp100
		$N^{\eta 2}$ Arg78	$O^{\delta 2}$ Asp100
2. Main-chain - main-chain interaction			
	i.	N (Gly/Ser/Asn/Lys)67	O Asn64
	ii.	N Gly81	O Phe63
	iii.	N Asp82	O His79
3. Main-chain - side-chain interaction			
	i.	N Gly71	$O^{\delta 2}$ Asp82
	ii.	N (Lys/Arg/Gln)68	$O^{\delta 1}$ Asn64
	iii.	O (Lys/Arg/Gln)68	$N^{\delta 2}$ Asn64
	iv.	O His79	N ^ε Arg78
	v.	O Pro73	$N^{\eta 1}$ Arg78
	vi.	O (Ala/Val)80	$N^{\eta 2}$ Arg78
	vii.	N His79	O ⁸² Asp82
4. Water mediated interactions			
	i.	N Gly84	Water 14 (PA numbering)
	ii.	O Gly71	Water 14, Ws1
	iii.	O ⁸¹ Asp123	Water 14, Ws1
	Iv	N Ala/Leu/Thr/Asp/Glu/Phe66	Water 13, Ws5
	v.	N Ala/Val80	Water 13, Ws5
	Vi	N Asn64	Water/I3, Ws2
	vii.	O ⁸¹ Asn64	Water 13, Ws5
	viii.	O Arg78	Water 87, Ws6
	ix.	Water 13, Ws5	Water 87, Ws6

Supplementary Table 1. Conserved hydrogen bonding interactions in the zinc binding loop