Supplementary Table 1. Crystal data collection statistics.

\mathbf{X} reviewed a path $(\mathbf{\hat{\lambda}})$	0.0700		
X-ray wavelength (A)	0.9792		
Space group	P2 ₁ 2 ₁ 2 ₁		
Unit cell dimensions	a = 31.6 Å, b = 43.6 Å, c = 65.6 Å,		
	$\alpha = \beta = \gamma = 90^{\circ}$		
Resolution* (Å)	26.2 – 1.25	(1.27 - 1.25)	
No. of unique reflections	23934	(749)	
Completeness	92.5%	(59.3%)	
R-merge	0.053	(0.48)	
CC1/2 (Å ²)	-	(0.82)	
l/σ	10.9	(2.04)	
Redundancy	6.1	(3.1)	
Wilson plot B-factor (Å ²)	10.9		
Molecules per asymmetric unit	1		
No. of protein residues	92		

* Numbers in parenthesis are shown for the highest resolution shell.

Supplementary Table 2. Structure refinement statistics.

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Resolution range (Å)	26.2 - 1.25	(1.28 – 1.25)		
Reflections	23883	(1228)		
σcutoff	None			
R-value (all) (%)	12.3			
R-value (R-work) (%)	12.2	(20.7)		
Free R-value (%)	14.9	(24.6)		
Rms deviations from ideal				
geometry				
bond length (Å)	0.011			
angle (degrees)	1.53			
chiral (Å)	0.096			
No. of atoms				
protein	771			
water	148			
Mean B-factor (Ų)				
all atoms	17.3			
protein atoms	14.3			
protein main chain	11.9			
protein side chain	16.3			
water	32.7			
Molprobity Ramachandran plot statistics				
Residues in favored regions (%)	100.0			
Residues in allowed regions (%)	100.0			
Residues in disallowed region (%)	0.0			

Supplementary Figure Legend

Fig. 1. Malate dehydrogenase (MDH) refolding time course. MDH was heat denatured and diluted into chaperone mixtures including the indicated J-protein and Hsp70, as well as Hsp104 and Sse1. Aliquots were removed at the indicated times and enzymatic activity determined. Activity of non-denatured enzyme was taken as 100%. Hsp70: Ssa1 (Hsp70 WT) or Ssa1 lacking the EEVD (Hsp70_{Δ EEVD}) (a) Sis1, Xdj1, and Ydj1 with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 1c. (b) Sis1 and Sis1 with Xdj1 J-domain (J_{xdj1}Sis1) or Ydj1 J-domain (J_{ydj1}Sis1) with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 3c. (d) Sis1 WT and variants with indicated substitutions with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 4b. (e) Sis1 WT, E50A and R73A with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 5c. (f) Sis1 WT and variants having indicated lysines substituted by asparagine with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 5c. (f) Sis1 WT and variants having indicated lysines substituted by asparagine with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 5c. (f) Sis1 WT and variants having indicated lysines substituted by asparagine with Hsp70 WT or Hsp70_{Δ EEVD}. Activity at 90 min, Fig. 6c.





Time (minutes)

- K199/K202/K214N:E50A
- K199/K202/K214N:R73A