

Self-Association of the Histidine Kinase CheA as Studied by Pulsed Dipolar ESR Spectroscopy

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Supplemental Table S1. Intersubunit Spin-Spin Separations within the CheA Δ 289 dimer.

Domains	Residue Pair	Cβ separations in crystal structure of CheAΔ289, (Å)	ESR Average Distance, (Å)	Full-width at half maximum from P(r) ,(Å)
P4-P4	A371-B371	65	50	23
	A387-B387	49	50	18
	A401-B401	52	50	25
	A458-B458*	101	62	31
	A496-B496*	90	67	29
	A508-B508	72	60	20
	A522-B522*	67	69	9
P5-P5	A545-B545	41	44	23
	A553-B553	64	63	25
	A568-B568	74	62	22
	A634-B634	31	40	15
	A639-B639	32	45	13
	A646-B646	60	58	14

A and B represent the two CheA subunits. Dipolar signals from some sites marked “*” were weak and contributed to only 40% of the total dipolar amplitude.

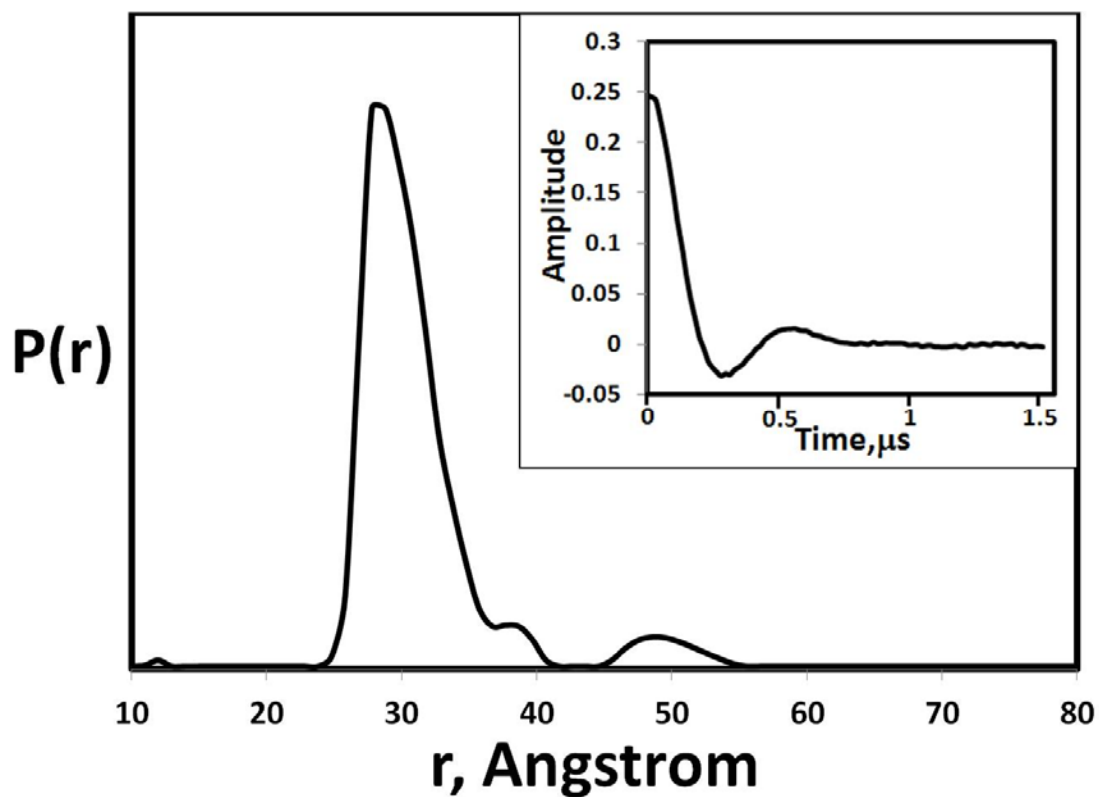


FIGURE S1. A Characteristic Intra-molecular Dipolar Signal from the CheA P3 Domain. Pairwise distance distribution for site E301C, which resides at the top of P3 domain with CheW. The distance distribution represents the width of the dimerization domain. **Inset:** Strong oscillating dipolar signal from site E301C after baseline correction.