

Supplementary Online Material for

The Structure of the *Staphylococcus aureus* Sortase-Substrate Complex Reveals how the Universally Conserved LPXTG Sorting Signal is Recognized

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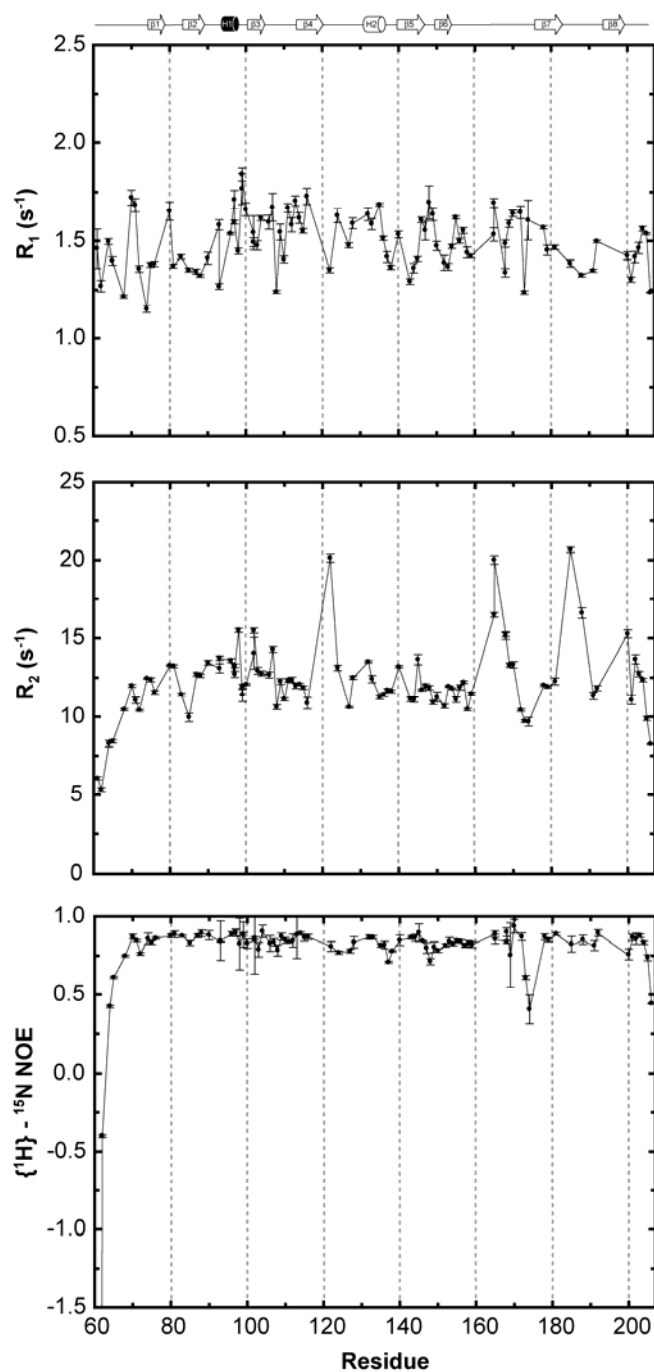


Fig. S1: Backbone amide ^{15}N spin relaxation data. The connecting lines represent only the trends of the data and are not necessarily representing residues with no data available (due to low signal or overlapped resonance).

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Table S1: Model-free parameters for the SrtA-LPAT* complex

Residue		Model	S^2	S_f^2	τ_e (ps)	R_{ex} (s^{-1})
A61		1	0.496 (\pm 0.009)			
K62		5	0.350 (\pm 0.011)	0.849 (\pm 0.015)	570 (\pm 17)	
Q64		5	0.628 (\pm 0.019)	0.898 (\pm 0.011)	923 (\pm 47)	
I65		5	0.652 (\pm 0.011)	0.825 (\pm 0.008)	1217 (\pm 66)	
D68		4	0.760 (\pm 0.005)		15 (\pm 1)	
S70		1	0.981 (\pm 0.007)			
K71		1	0.928 (\pm 0.013)			
V72		1	0.831 (\pm 0.004)			
G74	β 1	3	0.740 (\pm 0.011)			3.3 (\pm 0.2)
Y75	β 1	3	0.884 (\pm 0.008)			1.4 (\pm 0.2)
I76	β 1	1	0.912 (\pm 0.006)			
D80		1	1.000 (\pm 0.002)			
I83	β 2	1	0.916 (\pm 0.005)			
E85	β 2	1	0.869 (\pm 0.006)			
G90		1	1.000 (\pm 0.006)			
T93	H1	1	1.000 (\pm 0.008)			
Q96	H1	1	1.000 (\pm 0.002)			
L97	H1	1	1.000 (\pm 0.008)			
N98	H1	3	0.961 (\pm 0.010)			3.2 (\pm 0.2)
G100		1	0.930 (\pm 0.004)			
F103	β 3	3	0.936 (\pm 0.015)			1.5 (\pm 0.3)
A104	β 3	1	1.000 (\pm 0.004)			
E106		1	1.000 (\pm 0.008)			
N107		3	1.000 (\pm 0.026)			2.1 (\pm 0.4)
E108		3	0.807 (\pm 0.005)			0.5 (\pm 0.2)
S109		1	0.997 (\pm 0.008)			
L110		1	0.904 (\pm 0.006)			
D111		1	1.000 (\pm 0.004)			
D112		1	1.000 (\pm 0.007)			
Q113	β 4	1	0.997 (\pm 0.007)			
N114	β 4	1	0.936 (\pm 0.005)			
I115	β 4	1	0.980 (\pm 0.006)			
S116	β 4	1	1.000 (\pm 0.011)			
F122		3	0.867 (\pm 0.009)			9.4 (\pm 0.3)
D124		1	1.000 (\pm 0.007)			
N127		1	0.876 (\pm 0.004)			
Y128		1	1.000 (\pm 0.005)			
L133	H2	1	1.000 (\pm 0.009)			
A136	H2	1	0.951 (\pm 0.005)			
K138		1	0.907 (\pm 0.005)			
S140	β 5	3	0.981 (\pm 0.011)			1.1 (\pm 0.2)
Y143	β 5	1	0.856 (\pm 0.007)			
F144	β 5	1	0.888 (\pm 0.011)			
K145	β 5	3	0.904 (\pm 0.008)			2.5 (\pm 0.3)
V146	β 5	1	0.970 (\pm 0.005)			
G147		1	0.974 (\pm 0.009)			
N148		1	0.925 (\pm 0.011)			
E149	β 6	1	0.898 (\pm 0.010)			

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Residue		Model	S^2	S_f^2	τ_e (ps)	R_{ex} (s^{-1})
T150	$\beta 6$	2	0.926 (± 0.013)		31 (± 12)	
K152	$\beta 6$	1	0.848 (± 0.009)			
Y153	$\beta 6$	3	0.877 (± 0.010)			1.1 (± 0.1)
K154	$\beta 6$	1	0.935 (± 0.004)			
T156		1	0.962 (± 0.007)			
S157		1	0.989 (± 0.006)			
I158		1	0.851 (± 0.006)			
R159		1	0.922 (± 0.005)			
V168		3	0.866 (± 0.014)			4.4 (± 0.2)
L169		1	1.000 (± 0.005)			
D170		1	1.000 (± 0.005)			
Q172		1	0.839 (± 0.006)			
K173		5	0.750 (± 0.008)	0.826 (± 0.008)	471 (± 97)	
G174		2	0.780 (± 0.023)		400 (± 101)	
Q178	$\beta 7$	1	0.990 (± 0.004)			
L179	$\beta 7$	1	0.947 (± 0.005)			
L181	$\beta 7$	1	0.940 (± 0.007)			
D185		3	0.880 (± 0.011)			9.9 (± 0.2)
N188		3	0.851 (± 0.005)			6.1 (± 0.3)
T191		1	0.893 (± 0.005)			
G192		1	0.952 (± 0.004)			
F200	$\beta 8$	3	0.915 (± 0.013)			3.9 (± 0.3)
V201	$\beta 8$	1	0.835 (± 0.008)			
A202		3	0.924 (± 0.019)			2.1 (± 0.4)
T203		1	0.997 (± 0.007)			
E204		1	1.000 (± 0.004)			
V205		5	0.752 (± 0.013)	0.894 (± 0.009)	2106 (± 565)	
K206		5	0.595 (± 0.004)	0.795 (± 0.002)	848 (± 12)	

Note: For model 5: $S^2 = S_f^2 S_s^2$, and $\tau_e = \tau_s$. For models 1 to 4 $\tau_e = \tau_f$.