

Appendix

Chapter 4

Table S1: Details of the binding energies and quantum mechanical descriptor values for the 45 complex systems studied with B3LYP/TZVP level of theory.

Complex	MOC	ρ	OWBO	WBI	E_I (kJ/mol)
HSH-FNF2	0.0043	0.0037	0.0011	0.0017	-0.10
ClH-FOF	0.0046	0.0070	-0.0005	0.0058	-0.19
BrH-FNF2	0.0056	0.0055	0.0023	0.0035	-0.69
H3CH-OMe2	0.0093	0.0061	0.0015	0.0023	-1.20
H3CH-NHMe2	0.0061	0.0066	-0.0020	0.0053	-1.51
MeHPH-NH3	-0.0093	0.0082	0.0164	0.0070	-1.84
HSH-BrH	0.0065	0.0051	0.0042	0.0054	-1.98
HOPh-FMe	0.0121	0.0062	0.0032	0.0018	-2.55
H2PH-OH2	-0.0008	0.0080	0.0059	0.0037	-2.64
FH-FNF2	0.0108	0.0131	0.0196	0.0092	-3.80
BrH-BrH	0.0173	0.0096	0.0377	0.0215	-4.11
FH-FOF	0.0128	0.0150	0.0142	0.0118	-4.75
FH-ClOCl	0.0294	0.0143	0.0073	0.0249	-5.55
HSH-PH2Me	0.0002	0.0096	0.0119	0.0207	-5.87
MeSH-NH2Ph	0.0109	0.0153	0.0147	0.0202	-8.00
MeOH-PH3	-0.0087	0.0125	0.0142	0.0175	-9.00
MeSH-OHMe	0.0052	0.0168	0.0174	0.0157	-9.55
MeHNH-OMe2	0.0179	0.0148	0.0106	0.0077	-10.67
MeHNH-NMe3	0.0263	0.0175	0.0155	0.0150	-10.87
MeHNH-NH3	-0.0254	0.0171	0.0202	0.0167	-12.53
ClH-SH2	0.0389	0.0187	0.0469	0.0569	-13.78
PhOH-PH2Me	0.0082	0.0160	0.0308	0.0375	-16.17
HSH-NH2Me	0.0072	0.0259	0.0676	0.0506	-16.99
MeOH-PMe3	0.0245	0.0181	0.0199	0.0464	-17.40
ClH-PH2Me	0.0363	0.0219	0.1173	0.0851	-18.52
FH-FH	0.0205	0.0263	0.0305	0.0248	-20.38
BrH-OH2	0.0137	0.0303	0.0701	0.0488	-21.67
MeOH-OH2	0.0225	0.0259	0.0364	0.0243	-21.86
HOH-OH2	0.0168	0.0256	0.0369	0.0243	-22.60
MeOH-OHMe	0.0267	0.0281	0.0400	0.0284	-22.62
FH-SH2	0.0559	0.0245	0.0491	0.0627	-22.68
HOH-OHEt	0.0224	0.0276	0.0398	0.0282	-23.17
FH-OHCl	0.0186	0.0345	0.0566	0.0393	-24.20
ClH-OH2	0.0190	0.0332	0.0621	0.0511	-26.32
HOH-NMe3	0.0454	0.0353	0.0656	0.0483	-28.34
HOH-NH3	-0.0065	0.0297	0.0550	0.0452	-28.92
MeOH-NH3	-0.0131	0.0307	0.0582	0.0476	-28.96
FH-PH2Me	0.0553	0.0277	0.0773	0.0890	-29.35
HOH-NHMe2	0.0339	0.0347	0.0655	0.0535	-29.71
HOH-NH2Me	0.0165	0.0328	0.0618	0.0525	-30.03
MeOH-NH2Me	0.0068	0.0336	0.0685	0.0543	-30.11
FH-OH2	0.0320	0.0434	0.0790	0.0612	-40.66
FH-OHMe	0.0469	0.0496	0.0945	0.0754	-44.21
ClH-NH3	0.0281	0.0540	0.1737	0.1383	-44.89
FH-OMe2	0.0536	0.0531	0.1034	0.0800	-45.03

Table S2: Details of the binding energies and quantum mechanical descriptor values for the 45 complex systems studied with B3LYP/aug-TZVP level of theory.

Complex	MOC	ρ	OWBO	WBI	E_I (kJ/mol)
HSH-FNF2	-0.002	0.00372	0.0031	0.0016	0.0
ClH-FOF	-0.005	0.00702	-0.0040	0.0079	0.0
BrH-FNF2	0.001	0.00551	0.0064	0.0037	-0.5
H3CH-OMe2	-0.047	0.00600	-0.0031	0.0039	-1.1
H3CH-NHMe2	-0.037	0.00649	-0.0098	0.0076	-1.3
MeHPH-NH3	-0.057	0.00784	0.0188	0.0071	-1.3
HSH-BrH	0.002	0.00511	0.0035	0.0048	-1.9
HOPh-FMe	-0.031	0.00612	0.0013	0.0025	-2.5
H2PH-OH2	-0.056	0.00774	0.0079	0.0041	-2.3
FH-FNF2	0.013	0.01307	0.0074	0.0084	-3.7
BrH-BrH	0.016	0.00966	0.0132	0.0213	-4.0
FH-FOF	0.008	0.01499	0.0092	0.0118	-4.6
FH-ClOCl	0.040	0.01436	0.0055	0.0232	-5.4
HSH-PH2Me	-0.030	0.00957	0.0079	0.0187	-5.8
MeSH-NH2Ph	-0.092	0.01518	0.0188	0.0213	-7.8
MeOH-PH3	0.014	0.01254	0.0116	0.0188	-8.9
MeSH-OHMe	-0.053	0.01664	0.0175	0.0164	-9.1
MeHNH-OMe2	-0.090	0.01477	0.0099	0.0086	-10.5
MeHNH-NMe3	-0.162	0.01747	0.0368	0.0212	-10.8
MeHNH-NH3	-0.156	0.01668	0.0189	0.0169	-11.4
ClH-SH2	0.021	0.01866	0.0756	0.0584	-13.5
PhOH-PH2Me	0.087	0.01607	0.0245	0.0390	-16.0
HSH-NH2Me	-0.098	0.02569	0.0750	0.0513	-16.4
MeOH-PMe3	0.029	0.01812	0.0871	0.0606	-17.6
ClH-PH2Me	0.026	0.02199	0.1178	0.0836	-18.7
FH-FH	-0.021	0.02606	0.0289	0.0233	-19.3
BrH-OH2	-0.089	0.03005	0.0787	0.0503	-20.0
MeOH-OH2	-0.130	0.02558	0.0346	0.0246	-20.3
HOH-OH2	-0.135	0.02530	0.0349	0.0237	-21.7
MeOH-OHMe	-0.124	0.02795	0.0407	0.0284	-21.9
FH-SH2	0.043	0.02452	0.0445	0.0632	-22.9
HOH-OHEt	-0.106	0.02753	0.0346	0.0276	-23.4
FH-OHCl	-0.128	0.03439	0.0499	0.0373	-23.9
ClH-OH2	-0.090	0.03304	0.0839	0.0522	-24.9
HOH-NMe3	-0.184	0.03527	0.0848	0.0566	-29.1
HOH-NH3	-0.150	0.02949	0.0527	0.0453	-28.5
MeOH-NH3	-0.151	0.03050	0.0614	0.0494	-27.9
FH-PH2Me	0.043	0.02782	0.0717	0.0888	-30.1
HOH-NHMe2	-0.152	0.03472	0.0569	0.0536	-30.2
HOH-NH2Me	-0.139	0.03265	0.0556	0.0521	-30.1
MeOH-NH2Me	-0.130	0.03350	0.0667	0.0551	-29.6
FH-OH2	-0.135	0.04334	0.0790	0.0606	-39.8
FH-OHMe	-0.123	0.04962	0.0896	0.0737	-44.4
ClH-NH3	-0.079	0.05415	0.1821	0.1423	-44.7
FH-OMe2	-0.112	0.05311	0.0920	0.0799	-45.9

Table S3: Details of the binding energies and quantum mechanical descriptor values for the 45 complex systems studied with B97D/aug-TZVP level of theory.

Complex	MOC	ρ	OWBO	WBI	E_I (kJ/mol)
HSH-FNF2	-0.002	0.00360	0.0030	0.0018	-2.2
ClH-FOF	-0.002	0.00697	-0.0039	0.0090	-2.7
BrH-FNF2	0.001	0.00540	0.0063	0.0041	-3.5
H3CH-OMe2	-0.044	0.00583	-0.0026	0.0039	-3.9
H3CH-NHMe2	-0.015	0.00639	-0.0099	0.0081	-4.8
MeHPH-NH3	-0.044	0.00779	0.0198	0.0086	-4.0
HSH-BrH	0.004	0.00502	0.0039	0.0055	-5.1
HOPh-FMe	-0.037	0.00598	0.0057	0.0030	-7.2
H2PH-OH2	-0.052	0.00761	0.0082	0.0047	-3.8
FH-FNF2	0.012	0.01313	0.0078	0.0093	-4.7
BrH-BrH	0.020	0.00967	0.0145	0.0240	-8.3
FH-FOF	0.008	0.01515	0.0097	0.0132	-4.7
FH-ClOCl	0.038	0.01449	0.0063	0.0249	-6.7
HSH-PH2Me	-0.030	0.00956	0.0085	0.0208	-9.6
MeSH-NH2Ph	-0.075	0.01523	0.0214	0.0238	-14.8
MeOH-PH3	0.002	0.01258	0.0327	0.0200	-11.7
MeSH-OHMe	-0.043	0.01671	0.0200	0.0184	-11.7
MeHNH-OMe2	-0.092	0.01470	0.0107	0.0091	-14.8
MeHNH-NMe3	-0.146	0.01753	0.0371	0.0219	-20.3
MeHNH-NH3	-0.153	0.01680	0.0189	0.0187	-13.9
ClH-SH2	0.023	0.01889	0.0776	0.0637	-16.3
PhOH-PH2Me	0.081	0.01617	0.0256	0.0411	-20.5
HSH-NH2Me	-0.075	0.02614	0.0780	0.0578	-21.6
MeOH-PMe3	0.016	0.01826	0.0869	0.0622	-22.0
ClH-PH2Me	0.021	0.02224	0.1195	0.0887	-22.5
FH-FH	-0.021	0.02648	0.0302	0.0254	-16.8
BrH-OH2	-0.078	0.03065	0.0833	0.0561	-21.3
MeOH-OH2	-0.133	0.02596	0.0357	0.0269	-20.1
HOH-OH2	-0.140	0.02568	0.0365	0.0259	-20.4
MeOH-OHMe	-0.128	0.02836	0.0423	0.0307	-23.5
FH-SH2	0.029	0.02490	0.0463	0.0661	-23.2
HOH-OHEt	-0.110	0.02795	0.0363	0.0298	-24.0
FH-OHCl	-0.135	0.03508	0.0519	0.0401	-22.6
ClH-OH2	-0.081	0.03365	0.0876	0.0575	-25.1
HOH-NMe3	-0.193	0.03581	0.0866	0.0591	-35.8
HOH-NH3	-0.154	0.03004	0.0549	0.0489	-29.2
MeOH-NH3	-0.156	0.03107	0.0631	0.0534	-30.1
FH-PH2Me	0.021	0.02815	0.0734	0.0907	-30.7
HOH-NHMe2	-0.152	0.03532	0.0597	0.0572	-35.4
HOH-NH2Me	-0.145	0.03326	0.0574	0.0560	-33.0
MeOH-NH2Me	-0.137	0.03413	0.0686	0.0592	-34.4
FH-OH2	-0.140	0.04431	0.0815	0.0646	-37.6
FH-OHMe	-0.132	0.05073	0.0922	0.0778	-43.7
ClH-NH3	-0.060	0.05528	0.1852	0.1516	-49.7
FH-OMe2	-0.131	0.05422	0.0946	0.0834	-46.5

Table S4: Details of the binding energies and quantum mechanical descriptor values for the 45 complex systems studied with *wB97XD/aug-TZVP* level of theory.

Complex	MOC	ρ	OWBO	WBI	E_I (kJ/mol)
HSH-FNF2	-0.005	0.00354	0.0025	0.0014	-1.5
ClH-FOF	-0.006	0.00682	-0.0036	0.0072	-1.3
BrH-FNF2	-0.002	0.00529	0.0057	0.0033	-2.3
H3CH-OMe2	-0.041	0.00588	-0.0034	0.0035	-3.6
H3CH-NHMe2	-0.035	0.00643	-0.0096	0.0068	-4.9
MeHPh-NH3	-0.045	0.00776	0.0185	0.0065	-4.8
HSH-BrH	0.004	0.00501	0.0030	0.0041	-4.7
HOPh-FMe	-0.031	0.00596	0.0017	0.0022	-6.9
H2PH-OH2	-0.047	0.00764	0.0080	0.0038	-4.3
FH-FNF2	0.014	0.01286	0.0065	0.0079	-4.3
BrH-BrH	0.017	0.00949	0.0291	0.0186	-6.9
FH-FOF	0.010	0.01479	0.0087	0.0111	-4.3
FH-ClOCl	0.038	0.01412	0.0049	0.0206	-5.9
HSH-PH2Me	-0.011	0.00949	0.0070	0.0167	-9.4
MeSH-NH2Ph	-0.100	0.01502	0.0160	0.0195	-15.5
MeOH-PH3	0.027	0.01242	0.0308	0.0169	-11.7
MeSH-OHMe	-0.048	0.01648	0.0136	0.0154	-13.5
MeHNH-OMe2	-0.077	0.01468	0.0091	0.0081	-16.1
MeHNH-NMe3	-0.147	0.01737	0.0363	0.0197	-20.7
MeHNH-NH3	-0.120	0.01658	0.0175	0.0160	-15.2
ClH-SH2	0.021	0.01834	0.0723	0.0531	-16.1
PhOH-PH2Me	0.103	0.01594	0.0227	0.0361	-21.0
HSH-NH2Me	-0.094	0.02537	0.0722	0.0477	-22.3
MeOH-PMe3	0.038	0.01800	0.0197	0.0457	-22.2
ClH-PH2Me	0.037	0.02175	0.1135	0.0780	-21.9
FH-FH	-0.016	0.02596	0.0281	0.0227	-18.9
BrH-OH2	-0.084	0.02968	0.0746	0.0478	-23.4
MeOH-OH2	-0.115	0.02541	0.0336	0.0238	-22.9
HOH-OH2	-0.118	0.02517	0.0341	0.0228	-23.4
MeOH-OHMe	-0.111	0.02774	0.0393	0.0272	-25.9
FH-SH2	0.046	0.02421	0.0426	0.0577	-23.5
HOH-OHEt	-0.095	0.02734	0.0337	0.0263	-26.6
FH-OHCl	-0.122	0.03420	0.0490	0.0354	-24.7
ClH-OH2	-0.081	0.03275	0.0821	0.0504	-28.0
HOH-NMe3	-0.161	0.03491	0.0579	0.0471	-36.9
HOH-NH3	-0.119	0.02930	0.0510	0.0431	-31.5
MeOH-NH3	-0.119	0.03027	0.0592	0.0473	-32.1
FH-PH2Me	0.058	0.02767	0.0690	0.0829	-31.1
HOH-NHMe2	-0.120	0.03438	0.0553	0.0509	-36.6
HOH-NH2Me	-0.107	0.03237	0.0535	0.0495	-34.6
MeOH-NH2Me	-0.097	0.03318	0.0641	0.0526	-35.6
FH-OH2	-0.126	0.04308	0.0780	0.0587	-41.0
FH-OHMe	-0.115	0.04921	0.0879	0.0711	-46.6
ClH-NH3	-0.067	0.05355	0.1800	0.1376	-49.5
FH-OMe2	-0.101	0.05257	0.0902	0.0768	-49.3

Table S5: Details of the binding energies and quantum mechanical descriptor values for the 45 complex systems studied with MP2/aug-TZVP level of theory.

Complex	MOC	ρ	OWBO	WBI	E_I (kJ/mol)
HSH-FNF2	-0.005	0.00368	-0.0012	0.0012	-1.1
ClH-FOF	-0.018	0.00696	-0.0024	0.0066	-0.8
BrH-FNF2	-0.002	0.00538	0.0064	0.0027	-1.8
H3CH-OMe2	-0.052	0.00601	-0.0048	0.0034	-2.6
H3CH-NHMe2	-0.052	0.00656	-0.0096	0.0065	-2.7
MeHPH-NH3	-0.079	0.00776	0.0068	0.0050	-2.0
HSH-BrH	-0.006	0.00500	0.0006	0.0030	-3.1
HOPh-FMe	-0.035	0.00610	-0.0006	0.0024	-5.2
H2PH-OH2	-0.070	0.00768	0.0096	0.0031	-2.6
FH-FNF2	0.003	0.01274	0.0032	0.0060	-2.9
BrH-BrH	0.004	0.00934	0.0254	0.0146	-4.7
FH-FOF	-0.001	0.01461	0.0044	0.0085	-3.1
FH-ClOCl	0.023	0.01385	0.0032	0.0161	-3.3
HSH-PH2Me	-0.043	0.00941	0.0062	0.0132	-6.0
MeSH-NH2Ph	-0.206	0.01513	0.0125	0.0172	-12.3
MeOH-PH3	0.004	0.01243	0.0130	0.0143	-9.1
MeSH-OHMe	-0.086	0.01643	0.0084	0.0127	-10.4
MeHNH-OMe2	-0.117	0.01468	0.0244	0.0109	-13.2
MeHNH-NMe3	-0.263	0.01757	0.0026	0.0185	-15.6
MeHNH-NH3	-0.190	0.01648	0.0201	0.0132	-11.0
ClH-SH2	-0.006	0.01804	0.0694	0.0434	-10.5
PhOH-PH2Me	0.062	0.01593	0.0283	0.0313	-17.8
HSH-NH2Me	-0.153	0.02525	0.0604	0.0404	-15.0
MeOH-PMe3	0.028	0.01793	0.0750	0.0473	-17.5
ClH-PH2Me	0.005	0.02142	0.1043	0.0660	-14.9
FH-FH	-0.033	0.02533	0.0220	0.0176	-15.8
BrH-OH2	-0.124	0.02922	0.0675	0.0406	-18.1
MeOH-OH2	-0.175	0.02519	0.0339	0.0199	-18.7
HOH-OH2	-0.175	0.02484	0.0304	0.0190	-19.2
MeOH-OHMe	-0.177	0.02762	0.0348	0.0232	-21.6
FH-SH2	0.029	0.02383	0.0408	0.0488	-18.2
HOH-OHEt	-0.162	0.02714	0.0291	0.0224	-22.1
FH-OHCl	-0.172	0.03381	0.0469	0.0302	-21.6
ClH-OH2	-0.129	0.03222	0.0736	0.0419	-21.2
HOH-NMe3	-0.317	0.03511	0.0408	0.0490	-30.1
HOH-NH3	-0.177	0.02899	0.0489	0.0370	-25.4
MeOH-NH3	-0.180	0.03008	0.0576	0.0409	-25.8
FH-PH2Me	0.037	0.02723	0.0645	0.0727	-24.2
HOH-NHMe2	-0.233	0.03438	0.0534	0.0456	-29.8
HOH-NH2Me	-0.181	0.03220	0.0523	0.0435	-27.9
MeOH-NH2Me	-0.178	0.03314	0.0639	0.0467	-28.8
FH-OH2	-0.174	0.04236	0.0721	0.0502	-34.6
FH-OHMe	-0.174	0.04867	0.0827	0.0624	-39.6
ClH-NH3	-0.132	0.05295	0.1728	0.1216	-37.8
FH-OMe2	-0.188	0.05225	0.0882	0.0684	-42.1

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Figure S1: *RMSD equilibration of the backbone atoms of the interfacial residues relative to the starting structure in the three systems; LongAB, LatB and LatA.*

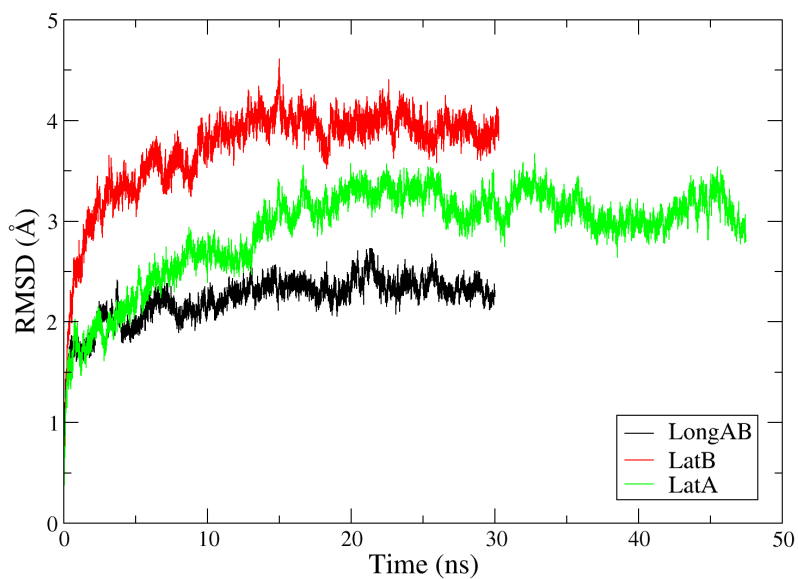


Figure S2: *An all-atom model of the LongAB system with subunit assignment.*

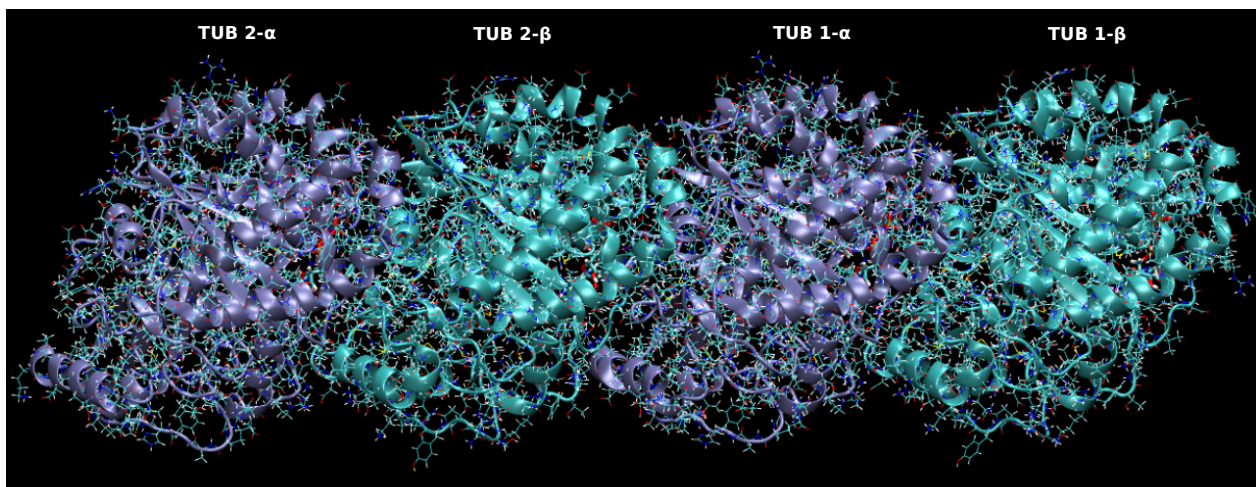


Table S6: Energy of hydrogen bonds in the LongAB interface in kJ/mol. SS# is snapshot number.

TUB 1- α	TUB 2- β	SS1	SS2	SS3	SS4	SS5	SS6	SS7	SS8	$E_{average}$	SD
Arg2	Glu71	-62.5	-33.4	-42.6	-42.3	-38.3	-20.1	-58.7	-23.2	-40.1	15
Glu434	Arg401	-30.0	-17.5	-39.3	-59.0	-32.4	-44.3	-38.1	0.0	-32.6	18
Tyr262	Arg401	-46.7	-45.0	-32.7	-45.5	0.0	-13.3	-25.6	-47.6	-32.0	18
Arg243	Asp76	-37.3	-17.3	-39.4	-35.7	0.0	-24.4	-42.8	-42.1	-29.9	15
Thr349	Val181	-44.6	-34.6	-13.8	-14.3	-43.6	-21.5	-21.0	-13.0	-25.8	13
Asp438	Arg401	-26.3	-55.1	-57.1	0.0	0.0	0.0	0.0	-52.7	-23.9	27
Val260	His406	-31.7	-34.1	-12.9	-8.8	-12.1	-41.6	-15.5	-20.0	-22.1	12
Gln133	Gly98	-16.8	-39.1	-20.7	-22.8	-21.2	-28.2	-10.7	-16.0	-21.9	9
Thr257	Gly100	-22.7	-35.5	-17.6	-27.2	-16.3	-13.2	-14.3	-24.2	-21.4	8
Lys352	Thr180	-18.5	-16.8	-22.1	0.0	-30.4	-16.3	-32.1	-12.9	-18.7	10
Asn249	Gln11	-22.6	-20.8	-27.5	-23.1	0.0	-28.6	0.0	-26.3	-18.6	12
Asn329	Lys176	-26.4	-18.2	-26.1	-19.4	-24.4	0.0	0.0	-7.6	-15.3	11
Lys163	Glu411	-34.2	0.0	-40.1	0.0	0.0	-43.7	0.0	0.0	-14.8	21
Asn258	Val181	0.0	-14.8	-15.6	-29.7	-8.8	-9.4	-22.4	-10.2	-13.9	9
Lys352	Asp179	0.0	0.0	0.0	-20.0	-12.2	-33.7	-12.9	0.0	-9.9	12
Asn249	Glu71	0.0	0.0	-31.3	0.0	0.0	-22.8	0.0	0.0	-6.8	13
Asp345	Arg400	0.0	0.0	0.0	0.0	0.0	-28.0	-23.6	0.0	-6.4	12
Asn258	Asn101	0.0	-20.6	0.0	0.0	-14.3	0.0	-12.9	0.0	-6.0	9
Arg2	Gln96	-3.5	0.0	0.0	-15.2	0.0	0.0	0.0	-23.8	-5.3	9
Val260	Trp407	0.0	-9.5	0.0	-4.3	-17.7	-6.7	0.0	0.0	-4.8	6
Gln133	Ser97	0.0	0.0	-4.7	0.0	0.0	0.0	0.0	-30.4	-4.4	11
Leu132	Gln96	0.0	0.0	0.0	0.0	0.0	-18.5	-14.8	0.0	-4.2	8
Asn258	Val182	0.0	-13.8	-13.5	0.0	0.0	0.0	0.0	0.0	-3.4	6
Lys326	Tyr210	0.0	0.0	0.0	0.0	0.0	-17.9	0.0	0.0	-2.2	6
Thr130	Gln96	0	0	0	0	-17	0	0	0	-2.1	6
Asn258	Thr180	0.0	0.0	0.0	-16.1	0.0	0.0	0.0	0.0	-2.0	6
Val437	Arg401	0.0	0.0	0.0	0.0	0.0	-11.6	0.0	0.0	-1.4	4
Val353	Asp179	0.0	-5.0	-4.4	0.0	0.0	0.0	0.0	0.0	-1.2	2
Lys336	Lys176	0.0	0.0	-2.3	0.0	0.0	0.0	0.0	-4.2	-0.8	2
Glu254	Asn101	0.0	0.0	0.0	0.0	0.0	0.0	-3.5	0.0	-0.4	1
Total Energy		-424	-431	-464	-384	-289	-444	-349	-354	-392	59

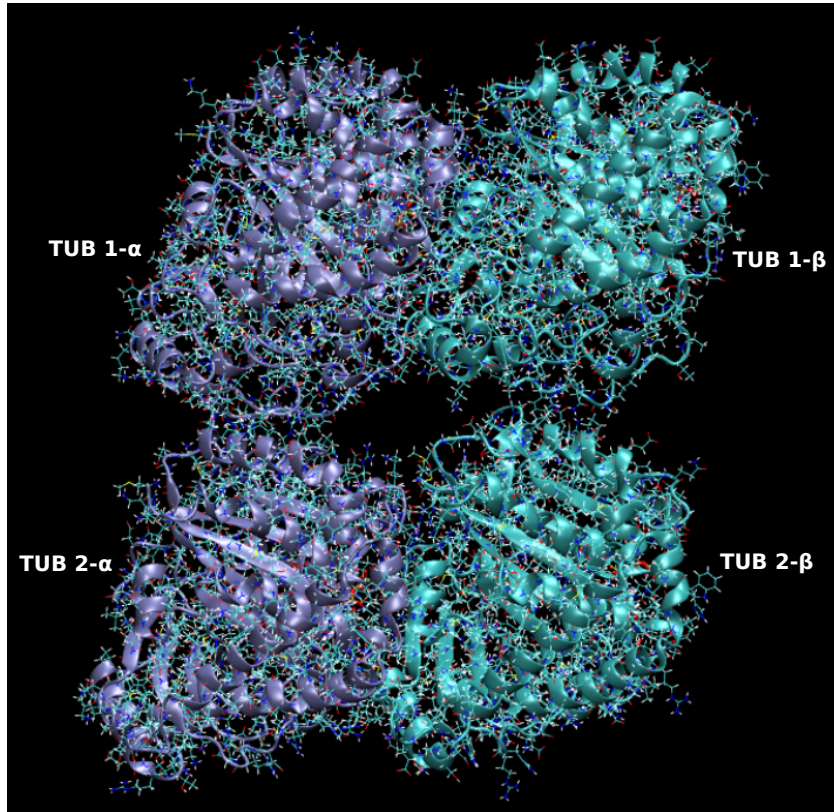
Figure S3: An all-atom model of the LatB system with subunit assignment.

Table S7: Energy of hydrogen bonds in the LatB interface in kJ/mol. SS# is snapshot number.

α - α Interactions											
TUB 1- α	TUB 2- α	SS1	SS2	SS3	SS4	SS5	SS6	SS7	SS8	$E_{average}$	SD
Arg215	Glu90	0.0	-44.0	-36.7	-49.3	-64.7	-51.4	-40.6	-55.3	-42.7	19
Lys338	Asp127	-24.5	-13.8	-33.6	-34.1	-26.2	-41.9	-28.9	-24.7	-28.5	8
Glu297	Arg121	-26.1	-21.9	-40.0	0.0	-62.8	0.0	-23.6	-22.2	-24.6	20
Glu297	Lys124	-20.0	-29.2	-15.9	-38.3	0.0	-32.1	-35.8	-17.1	-23.5	13
Glu284	Ser54	0.0	-32.6	-24.3	0.0	-33.4	-24.5	-38.7	-24.8	-22.3	15
Gln372	Glu55	-28.0	0.0	-13.7	-28.0	-15.9	-20.9	-16.9	-24.2	-18.4	9
Tyr282	Ser48	0.0	-31.8	-19.3	-20.9	-8.1	0.0	0.0	-13.2	-11.7	12
His283	Phe49	-7.5	-8.8	-28.2	0.0	-12.8	-5.8	-7.9	-13.5	-10.5	8
Ala278	Asn50	-20.3	0.0	0.0	-10.0	-14.9	0.0	0.0	0.0	-5.7	8
Arg373	Ser54	-8.1	0.0	-25.4	0.0	0.0	0.0	0.0	0.0	-4.2	9
Glu284	Lys60	0.0	0.0	0.0	0.0	0.0	-8.5	-23.6	0.0	-4.0	8
Leu286	Ser54	0.0	-2.9	-7.6	0.0	-9.3	-4.7	0.0	-6.1	-3.8	4
Gln285	Gly57	-15.1	0.0	0.0	0.0	0.0	0.0	-11.9	0.0	-3.4	6
Gln372	Thr56	0.0	0.0	0.0	0.0	0.0	-24.6	0.0	0.0	-3.1	9
His283	Asn50	0.0	-11.3	0.0	0.0	0.0	0.0	0.0	0.0	-1.4	4
Glu290	Gln128	0	0	0	0	-9	0	0	0	-1.1	3
Lys370	Thr51	0.0	0.0	0.0	0.0	0.0	0.0	-7.1	0.0	-0.9	3
Total Subunit Energy		-149	-196	-245	-180	-257	-214	-235	-201	-210	35
β - β Interactions											
TUB 1- β	TUB 2- β	SS1	SS2	SS3	SS4	SS5	SS6	SS7	SS8	$E_{average}$	SD
Arg308	Asp116	-46.7	-58.9	0.0	-37.5	-69.8	-57.4	-54.3	-33.7	-44.8	22
Glu290	Arg88	-51.6	-35.3	-32.2	-55.4	-22.6	-30.3	-30.3	-53.9	-38.9	13
Arg308	Asp120	-29.2	-26.9	-27.3	-25.6	-37.7	-27.6	-61.5	-58.2	-36.7	15
Lys299	Asp90	-11.0	-37.8	-25.5	-34.6	-31.0	-34.7	-31.3	-55.0	-32.6	12
Asp297	Lys124	-31.8	0.0	-31.3	0.0	-32.1	0.0	-29.0	-54.4	-22.3	20
Tyr342	Asp120	0.0	-9.9	0.0	-10.8	-25.1	-8.1	-41.7	-51.8	-18.4	19
Ser280	Arg88	-14.5	-21.7	-6.3	-22.9	0.0	-51.6	0.0	-16.1	-16.6	17
Lys338	Lys124	-6.4	-20.8	-15.5	-24.9	-25.2	0.0	0.0	-25.4	-14.8	11
Lys338	Ser126	-24.0	-23.6	-7.2	0.0	-14.0	0.0	0.0	-27.5	-12.0	12
Lys338	Arg123	0.0	0.0	0.0	0.0	0.0	-35.9	-19.9	-4.1	-7.5	13
Asn334	Glu127	0.0	-19.0	0.0	-32.7	0.0	0.0	0.0	0.0	-6.5	12
Asn335	Glu128	0.0	-6.8	0.0	0.0	0.0	0.0	0.0	0.0	-0.9	2
Total Subunit Energy		-215	-261	-145	-244	-257	-246	-268	-380	-252	65
Total Energy		-365	-457	-390	-425	-514	-460	-503	-581	-462	70

Table S8: Energy of hydrogen bonds in the LatA interface in kJ/mol. SS# is snapshot number.

α - β Interactions											
TUB 1- α	TUB 2- β	SS1	SS2	SS3	SS4	SS5	SS6	SS7	SS8	$E_{average}$	SD
Asp47	Arg284	-44.6	-17.9	-41.5	-40.4	-32.1	-54.1	-71.1	-36.0	-42.2	15.7
Lys124	Asp297	-24.7	-19.4	-39.2	-42.9	-29.9	-25.5	-27.4	-44.8	-31.7	9.4
Gln85	Ser280	-31.4	-23.8	0.0	-29.4	-34.9	-23.1	-22.4	-23.4	-23.6	10.6
Asp46	Arg278	-26.7	-10.3	-39.2	-39.6	-14.3	-19.8	-33.9	0.0	-23.0	14.4
Asp127	Asn334	-25.0	-24.3	-27.9	-16.4	-21.9	-21.8	-28.5	-15.0	-22.6	4.9
Asp120	Lys338	-39.2	0.0	-20.4	-36.6	-26.5	0.0	-25.3	-31.4	-22.4	15.1
Gln128	Gln293	-6.5	-20.3	-27.8	-36.5	-10.6	-28.8	-18.1	0.0	-18.6	12.3
Asp47	Gln282	-32.0	-8.2	-17.8	0.0	-29.7	0.0	-8.6	-36.0	-16.5	14.5
Glu55	Arg284	-39.8	0.0	-45.9	0.0	0.0	-19.8	-13.0	0.0	-14.8	18.9
Arg121	Asp297	0.0	-69.3	0.0	0.0	-35.3	0.0	0.0	0.0	-13.1	25.9
Asp47	Arg278	0.0	-18.5	0.0	0.0	0.0	-34.0	-16.8	-25.5	-11.9	13.7
Ser54	Lys372	0.0	0.0	0.0	0.0	-21.1	-18.6	-13.5	-24.3	-9.7	10.8
Lys124	Gln293	0.0	-12.8	-17.0	0.0	-15.6	0.0	0.0	-9.9	-6.9	7.7
His88	Gln281	0.0	-9.3	-19.8	-16.2	0.0	0.0	0.0	0.0	-5.7	8.3
Ser54	Arg284	-11.1	0.0	0.0	0.0	-32.0	0.0	0.0	0.0	-5.4	11.4
Arg123	Asn334	0.0	-15.8	0.0	-5.3	0.0	-1.9	-16.1	0.0	-4.9	7.1
Thr56	Gly370	-7.7	-16.7	-4.4	-7.5	0.0	0.0	0.0	0.0	-4.5	6.0
Gly59	Arg284	0.0	0.0	-19.2	-16.3	0.0	0.0	0.0	0.0	-4.4	8.2
Phe53	Ser374	0.0	0.0	0.0	0.0	0.0	0.0	0.0	-31.1	-3.9	11.0
Asn50	Gln282	0.0	0.0	0.0	-9.6	0.0	-20.4	0.0	0.0	-3.8	7.5
Phe87	Ser280	0.0	0.0	-24.5	0.0	0.0	0.0	0.0	0.0	-3.1	8.7
Gln128	Thr287	0.0	0.0	0.0	0.0	0.0	0.0	0.0	-18.8	-2.4	6.7
Ser54	Met373	0.0	0.0	0.0	0.0	0.0	-15.1	0.0	0.0	-1.9	5.3
Gln128	Glu290	0.0	0.0	0.0	0.0	0.0	0.0	0.0	-14.6	-1.8	5.2
Total Subunit Energy		-289	-267	-344	-297	-304	-283	-295	-311	-299	23
β - α Interactions											
TUB 1- β	TUB 3- α	SS1	SS2	SS3	SS4	SS5	SS6	SS7	SS8	$E_{average}$	SD
Arg88	Glu279	-45.6	-26.5	-36.3	-47.6	-40.4	-40.1	-45.3	0.0	-35.2	15.7
Lys124	Glu284	-25.6	0.0	-36.6	-32.4	-30.2	-38.7	-35.2	-17.2	-27.0	12.9
Ile86	Tyr282	-7.5	-29.9	0.0	-31.9	-38.9	-23.1	-25.2	-27.1	-22.9	12.9
Asp90	Lys280	-36.6	0.0	-27.4	-19.7	-39.1	-25.8	0.0	0.0	-18.6	16.5
Asn54	Glu284	-37.0	0.0	-24.6	0.0	0.0	-19.8	-14.5	-19.5	-14.4	13.6
Glu127	Thr334	0.0	-34.3	0.0	-33.4	-43.0	0.0	0.0	0.0	-13.8	19.3
Asp90	Ala281	-17.4	-14.8	0.0	0.0	-22.6	-17.9	-30.9	0.0	-12.9	11.7
Glu127	Thr337	0.0	-22.9	0.0	-39.0	0.0	0.0	0.0	-20.4	-10.3	15.2
Glu55	Gln285	-5.4	-13.1	-11.9	-9.3	-6.7	-20.2	-9.1	0.0	-9.5	6.0
Thr33	His283	0.0	0.0	-18.3	-5.2	0.0	0.0	0.0	-11.6	-4.4	7.0
Asp120	Lys338	-18.4	0.0	0.0	0.0	0.0	0.0	0.0	0.0	-2.3	6.5
Arg88	Ala281	0.0	0.0	0.0	-8.4	0.0	-5.0	0.0	0.0	-1.7	3.2
Total Subunit Energy		-193	-141	-155	-227	-221	-191	-160	-96	-173	44
Total Energy		-482	-408	-499	-523	-525	-474	-455	-406	-472	46

Figure S4: *An all-atom model of the LatA system with subunit assignment.*

