

Supporting Information

Atomistic Insight into Inhibition Mechanisms of Suppressors of Cytokine Signaling on Janus Kinase

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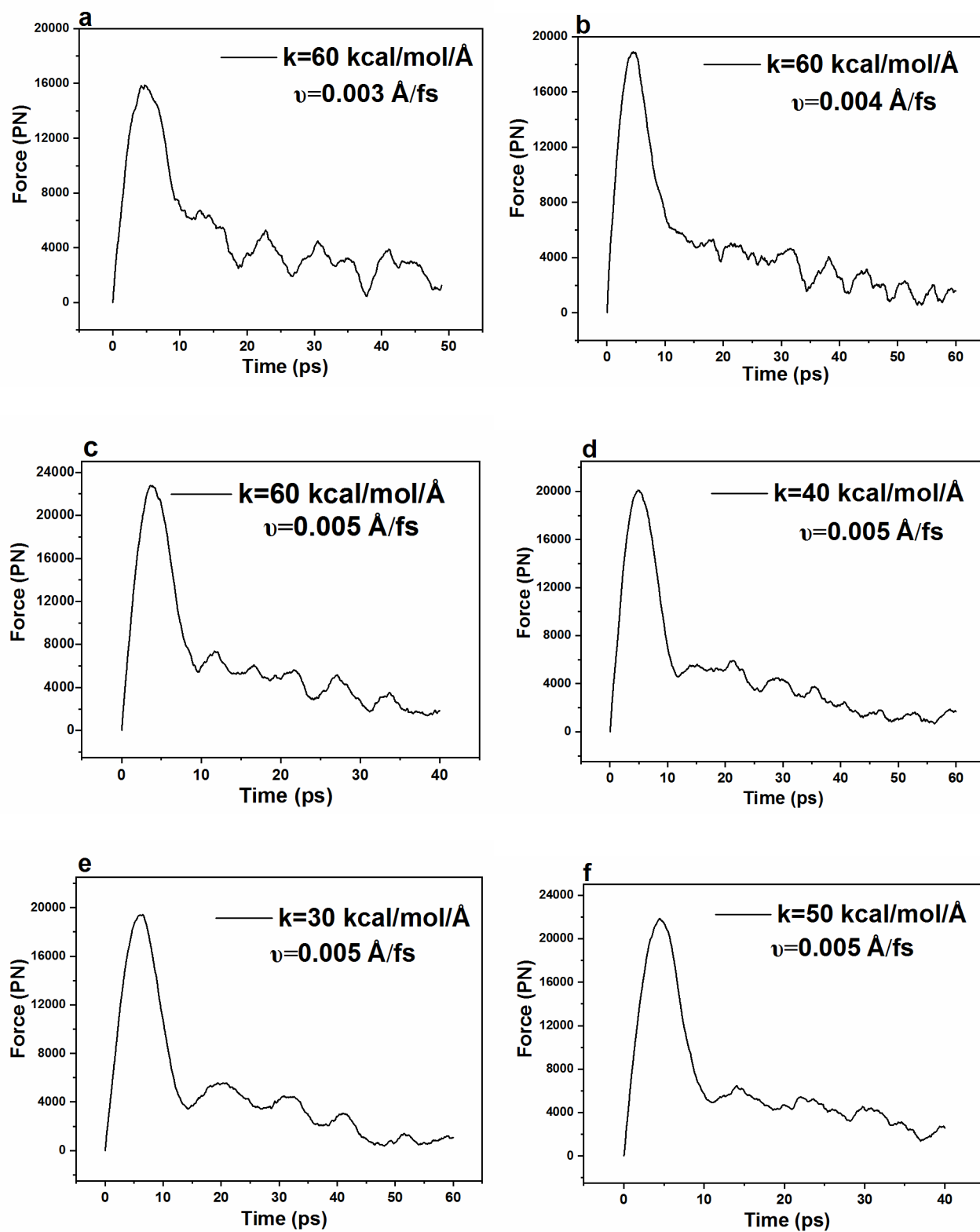
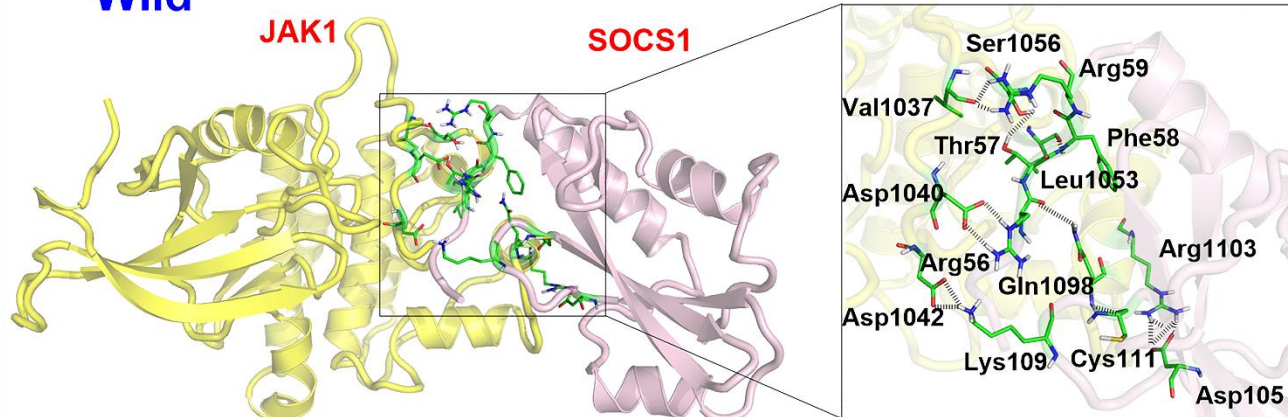
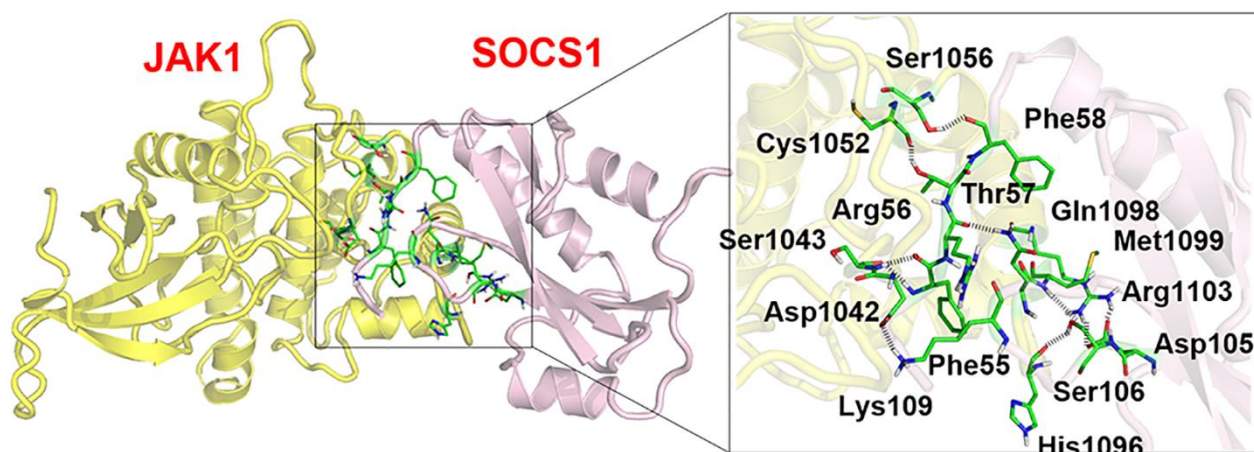


Fig S1. The test of a series of spring constants K and speed v .

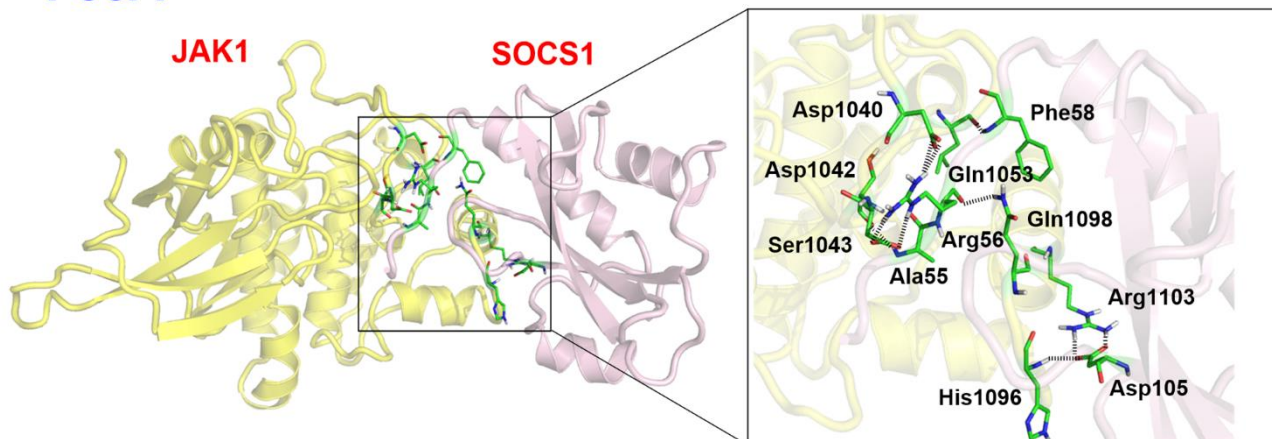
Wild



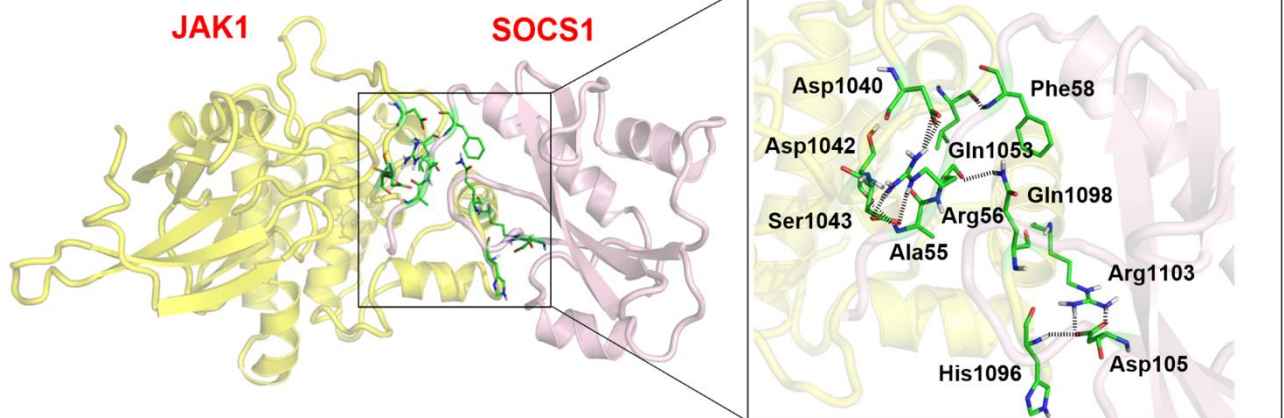
H54A



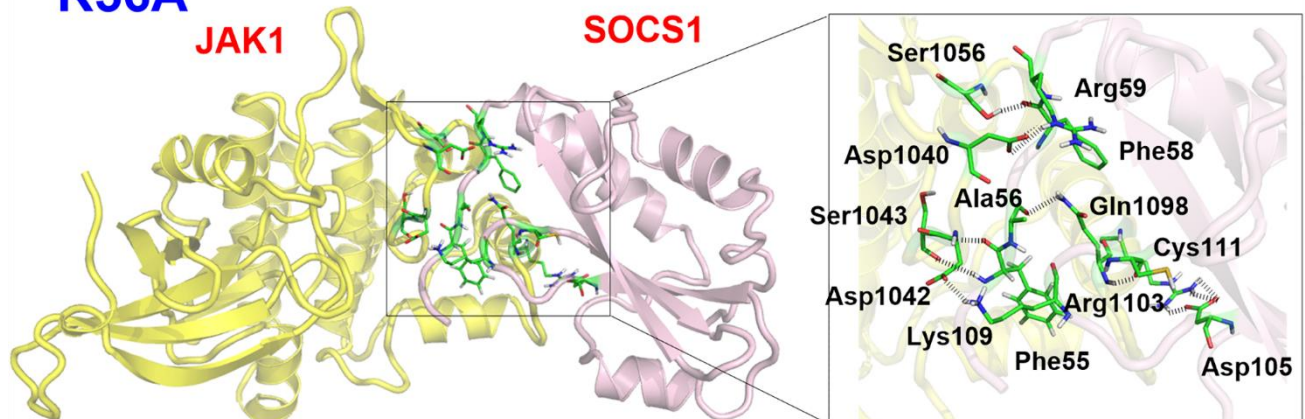
F55A



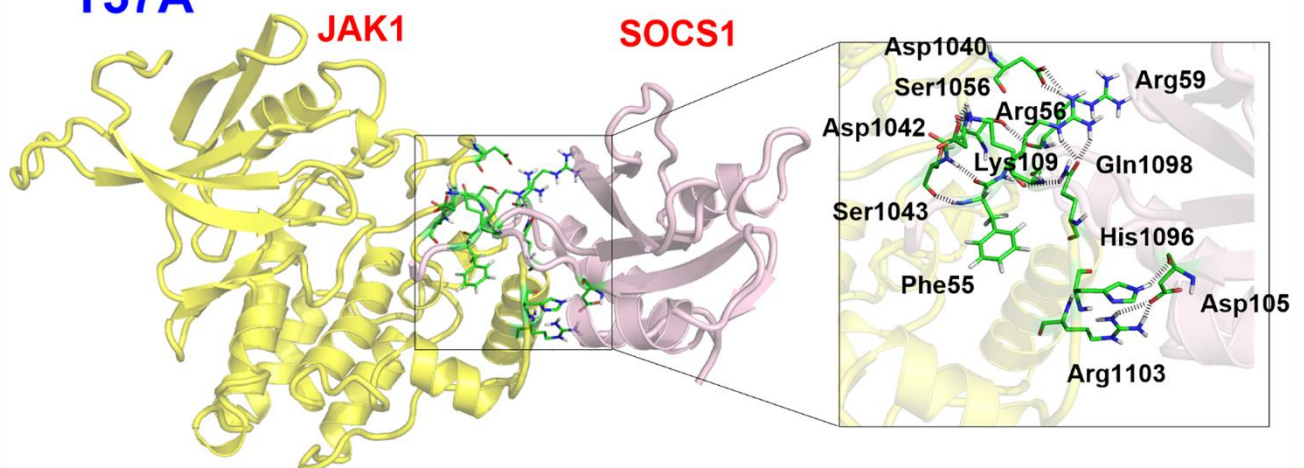
F55A



R56A



T57A



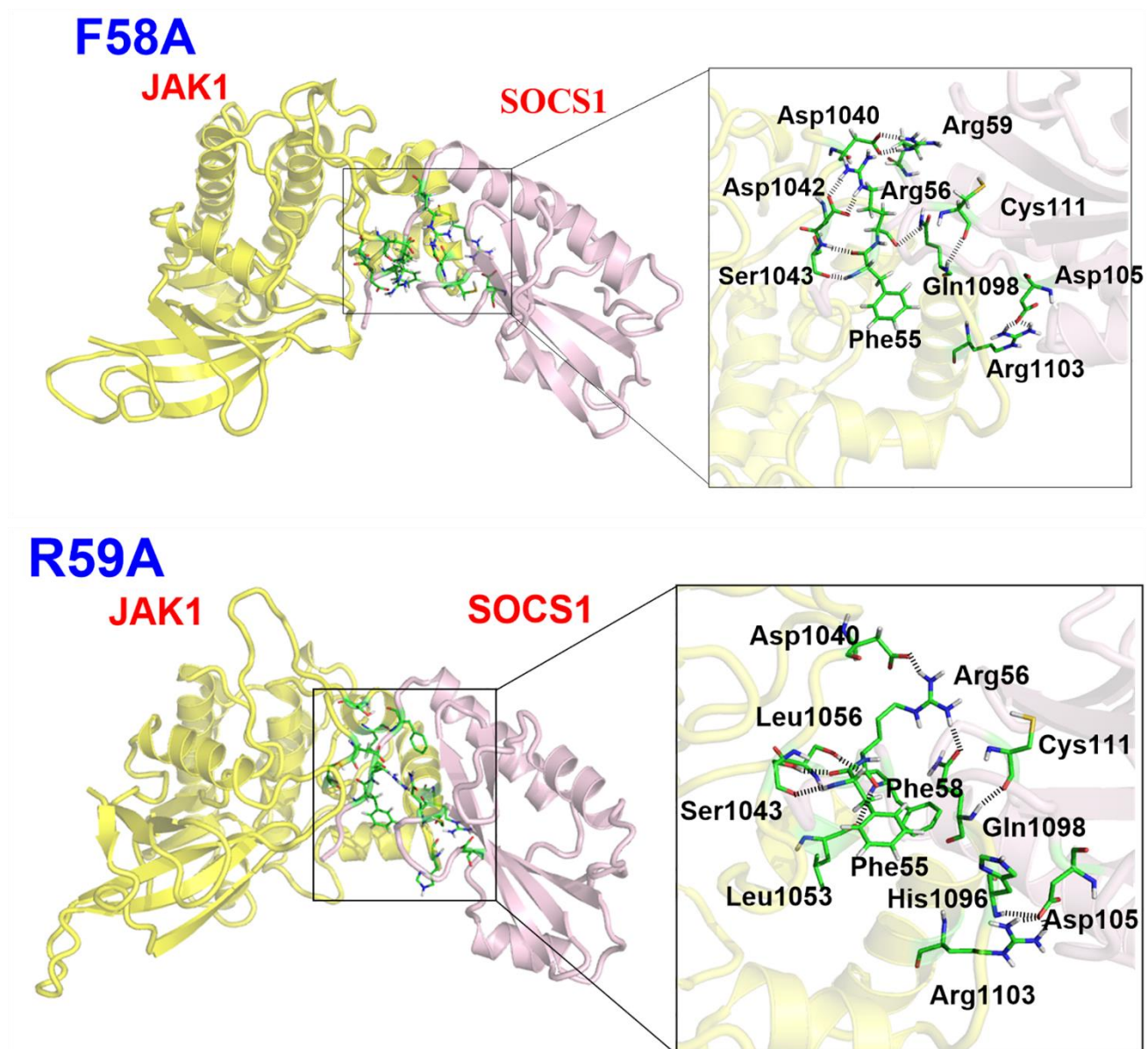
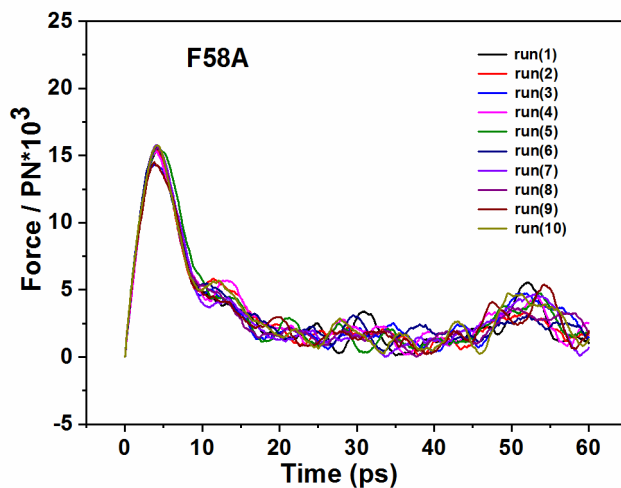
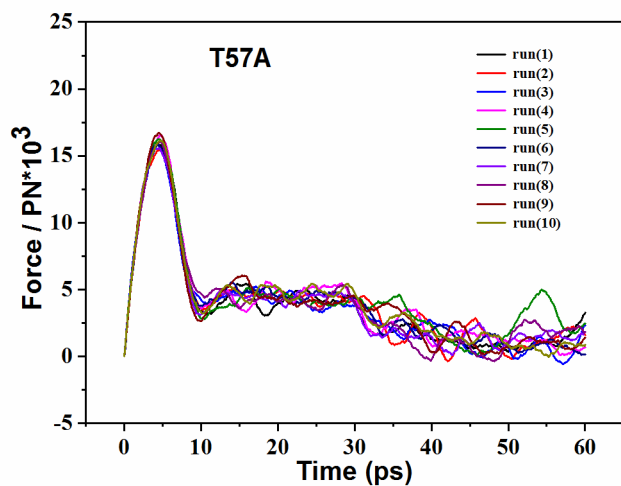
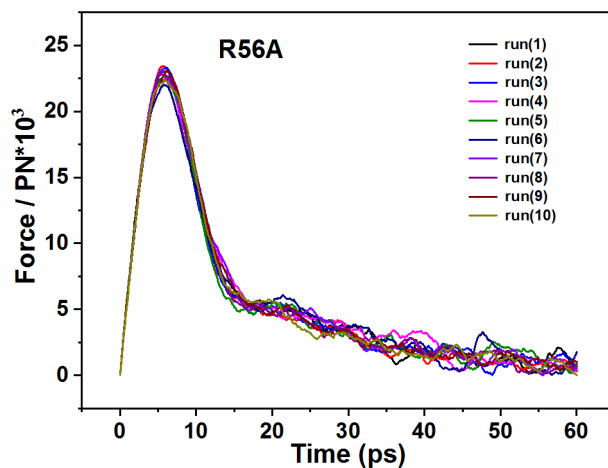
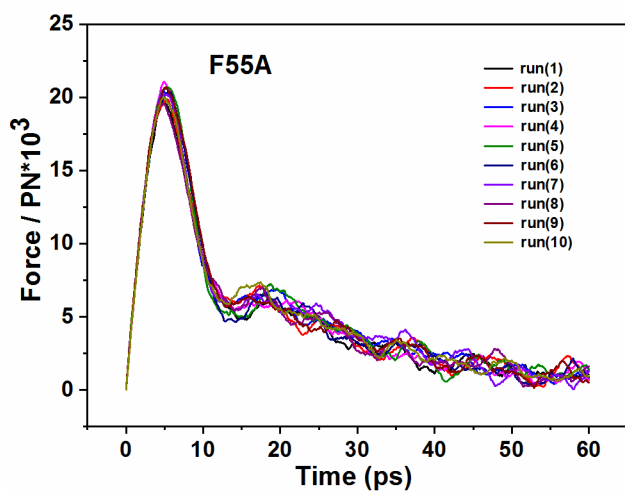
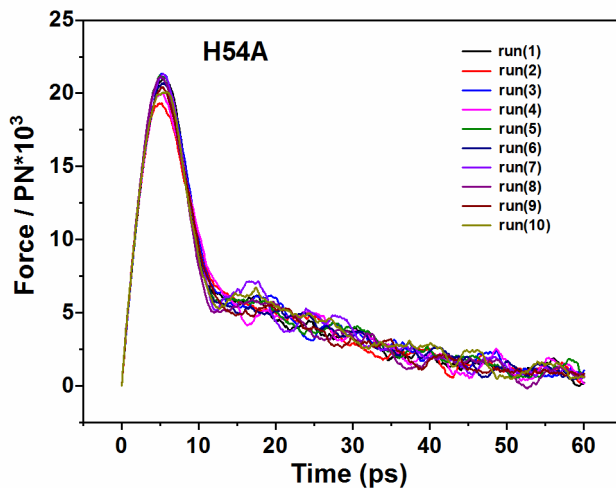
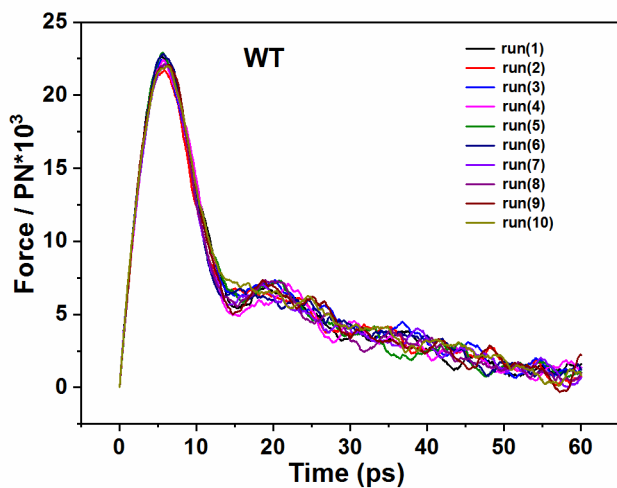


Fig S4. The details of the interaction between JAK1 and SOCS1 for wild and mutant systems



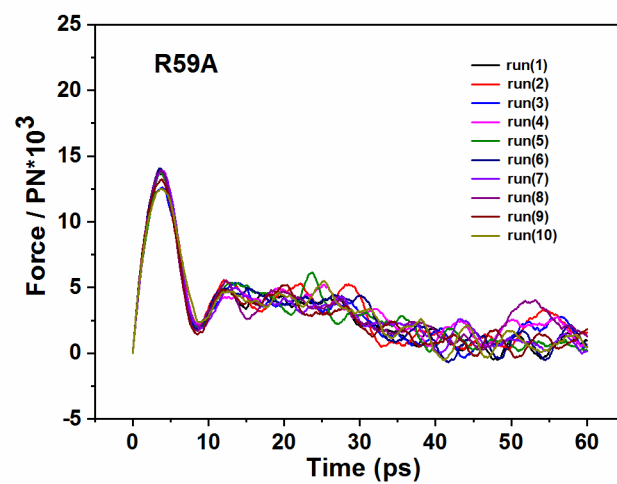


Fig S5. The SMD simulations of wild and mutant systems.

Table S1. The Binding Information of the Wild Systems

Binding couples			MD	
Wild	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Arg56@O	HE22-NE2@Gln1098	2.90	H-bond
	Ser1056@OG	HG1-OG1@Thr57	1.87	H-bond
	Val1037@O	HH22-NH2@Arg59	1.71	H-bond
	Val1037@O	HH12-NH1@Arg59	1.91	H-bond
	Asp105@OD2	HH12-NH1@ Arg1103	2.75	SB
	Cys111@O	H- N@Gln1098	2.84	H-bond
	Asp105@OD2	HH22- NH2@Arg1103	2.80	SB
	Leu1053@O	H- N@Phe58	2.87	H-bond
	Asp105@OD1	HH22- NH2@Arg1103	2.84	SB
	Asp1042@OD2	HZ3-NZ@Lys109	2.12	SB
	Asp1042@OD1	HZ3-NZ@Lys109	1.85	SB
	Asp105@OD1	HH12- NH1@Arg1103	2.87	SB
	Asp1040@OD2	HH21-NH2@Arg56	1.95	SB
	Asp1040@OD1	HE -NE@Arg56	1.85	SB

Table S2. The Binding Information of the H54A Systems

Binding couples			MD	
H54A	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Ser1043@O	H-O @Phe55	1.98	H-bond
	Phe55@O	H-O @Ser1043	2.05	H-bond
	Cys1052@OG	HG1-OG1@Thr57	2.19	H-bond
	Phe58@O	HG-OG @ Ser1056	1.83	H-bond
	Arg56@O	HE22-NE2@ Gln1098	2.03	H-bond
	Asp1042@OD2	HZ3-NZ @ Lys109	1.88	SB
	His1096@O	HG-OG @ Ser106	1.97	H-bond
	Ser106@OG	H-N @ Met1099	2.20	H-bond
	Asp105@OD1	HH22-NH2@ Arg1103	1.77	SB
	Asp105@OD2	HH12-NH1@ Arg1103	2.13	SB

Table S3. The Binding Information of the F55A Systems

F55A	Binding couples		MD	
	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Asp1040@OD2	HH11-NH1@Arg56	1.98	SB
	Asp1040@OD1	HH12-NH1@Arg56	2.98	SB
	Asp1042@OD2	HH21-NH2@ Arg56	1.93	SB
	Asp1042@OD1	HE-NE@Arg56	2.18	SB
	Ala55@O	H-O@Ser1043	2.01	H-bond
	Ser1043@O	H-O@Ala55	2.16	H-bond
	Arg56@O	HE22-NE@Gln1098	1.36	H-bond
	Lue1053@O	H-O@Phe58	2.07	H-bond
	Asp105@OD2	H-N@His1096	1.94	SB
	Asp105@OD1	HH22-NH2@Arg1103	1.68	SB
	Asp105@OD2	HH12-NH1@Arg1103	1.66	SB

Table S4. The Binding Information of the R56A Systems

Binding couples			MD	
R56A	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Asp1040@OD1	HH21-NH2@Arg59	2.26	SB
	Asp1040@OD1	HE-NE@ Arg59	1.81	SB
	Ser1043@O	H-O @Phe55	2.59	H-bond
	Phe55@O	H-O@ Ser1043	2.05	H-bond
	Phe58@O	H-O@ Ser1056	1.64	H-bond
	Ala56@OD2	HE22-NE2@Gln1098	2.38	H-bond
	Asp1042@OD1	HZ2-NZ@ Lys109	2.01	SB
	Lys111@O	H-N@ Gln1098	1.96	H-bond
	Asp105@OD2	HH12-NH1@Arg1103	1.83	SB
	Asp105@OD2	HH22-NH2@Arg1103	1.75	SB
	Asp105@OD1	HH22-NH2@Arg1103	2.61	SB

Table S5. The Binding Information of the T57A Systems

Binding couples			MD	
T57A	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Asp1040@OD2	HH11-NH1@ Arg56	2.11	SB
	Asp1040@OD1	HH11-NH1@ Arg56	2.17	SB
	Ser1043@O	H-O@Phe55	1.91	H-bond
	Phe55@O	H-O@Ser1043	1.67	H-bond
	Arg59@O	HG-OG@Ser1056	2.14	H-bond
	Gln1098@OE1	HE-NE@Arg56	2.01	H-bond
	Gln1098@OE1	HH21-NH2@Arg56	2.01	H-bond
	Arg56@O	HE22-NE@Gln1098	2.87	H-bond
	Asp1042@OD1	HZ3-NZ@Lys109	1.93	SB
	Asp105@O	HE2-NE@His1096	2.05	SB
	Asp105@OD2	HH12-NH1@Arg1103	2.39	SB
	Asp105@OD2	HH22-NH2@Arg1103	1.79	SB

Table S6. The Binding Information of the F58A Systems

F58A	Binding couples		MD	
	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Asp1040@OD1	Arg59@HH21-NH2	1.93	SB
	Asp1040@OD2	Arg59@HE-NE	2.00	SB
	Asp1042@OD2	Arg56@HH21-NH2	1.76	SB
	Asp1042@OD1	Arg56@HE-NE	1.92	SB
	Phe55@O	Ser1043@H-N	1.79	H-bond
	Ser1043@O	Phe55@H-N	1.97	H-bond
	Arg56@O	Gln1098@HE22-NE	2.44	H-bond
	Asp105@OD2	Arg1103@HH12-NH1	1.68	SB
	Asp105@OD1	Arg1103@HH22-NH2	2.27	SB
	Cys111@O	Gln1098@H-N	2.36	H-bond

Table S7. The Binding Information of the R59A Systems

Binding couples			MD	
R59A	DONOR	ACCEPTORH	Distance(Å)	Binding type
	Ser1043@O	H-O@ Phe55	1.94	H-bond
	Asp1040@OD2	HH11-NH1@Arg56	1.74	SB
	Phe55@O	HG-OG@ Ser1043	2.11	H-bond
	Leu1053@O	H-O@ Phe58	2.13	H-bond
	Phe58@O	HG-OG@ Leu1056	1.89	H-bond
	Gln1098@OE1	HH21-NH2 @Arg56	1.81	H-bond
	Asp105@OD2	H-N @ His1096	2.32	SB
	Asp105@OD1	HH22-NH2@ Arg1103	2.22	SB
	Asp105@OD1	HH12-NH1@ Arg1103	1.75	SB
	Cys111@O	H-N @ Gln1098	1.85	H-bond