

Table S1. Summary of identified hMps1 phosphorylation sites.

Phosphorylation Site	Sequence	MASCOT Score	resembling Consensus	Reference
S7	DL <u>S</u> GR	39	Plk1	1, 6
T12	EL <u>T</u> ID	29	Plk1	1, 9
S15	ID <u>S</u> IM	8	--	1, 9
T33	DL <u>T</u> DE	69	Plk1	1, 5, 6, 7, 9
S37	EL <u>S</u> LN	43	Plk1	1, 5, 6, 9
S49/T51	DN <u>S</u> GTVN		Plk1/--	6
S80	PL <u>S</u> DA	32	--	1, 5, 6, 9
S281	LN <u>S</u> PD	58	Cdk1	1, 4, 6, 7
T288	VK <u>T</u> DD	17	--	1, 9
S317	KP <u>S</u> GN		--	6
S321	ND <u>S</u> CE	68	Plk1	1, 5, 7
S329	LK <u>S</u> VQ	23	--	1
S345/346	EK <u>S</u> SEL		Plk1/Aur-B	6
T351	II <u>T</u> DS		--	6
S353	TD <u>S</u> IT		--	4
T360	NK <u>T</u> ES	33	Plk1	1, 4, 5, 9
S362	TE <u>S</u> SL	17	--	1, 9
S363	ES <u>S</u> LL	24	Plk1	1, 4, 5, 9
T371	EET <u>K</u> E	56	Plk1	1, 4
S382	PE <u>S</u> NQ	50	--	1, 4
S393	RK <u>S</u> EC		Aur-B	5, 6, 7
T418	VN <u>T</u> QE	21	--	1
T423/T424	KH <u>T</u> TFE	9	Aur-B/--	1
S436	KQ <u>S</u> PP	39	Cdk1	1, 3, 4, 5, 6
T453	CK <u>T</u> PS	77	Cdk1	1, 3
S455	TP <u>S</u> SN	28	--	1, 7
T458	SN <u>T</u> LD	68	--	1
Y462	DD <u>Y</u> MS		--	5
T468	FR <u>T</u> PV		Cdk1	3, 4
T564	NQ <u>T</u> LD	58	Plk1	1, 8
S582	QH <u>S</u> DK		--	8
T675	PD <u>T</u> TS		--	2, 9
T676	DT <u>T</u> SV	50	Plk1	1, 2, 4, 5, 8, 9
S677	TT <u>S</u> VV		--	2, 9
S682	KD <u>S</u> QV	50	Aur-B	1, 8
T686	VG <u>T</u> VN	55	--	1, 2, 5, 8, 9
S742	QI <u>S</u> KL	46	Plk1	1, 8
T795	IQ <u>T</u> HP		--	8
T805	KG <u>T</u> TE		Aur-B	8
T806	GT <u>T</u> EE		--	8
Y811	MK <u>Y</u> VL		--	8
S821	LN <u>S</u> PN	46	Cdk1	1, 4, 5, 6, 7, 8
S824	PN <u>S</u> IL	8	--	1, 5, 7, 8
Y833/Y836/S837	TL <u>Y</u> EH <u>Y</u> SGG	55	--	1, 6
S845	NS <u>S</u> SS		Plk1	8
S847	SS <u>S</u> KT		--	8
T849	SK <u>T</u> FE		--	8

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