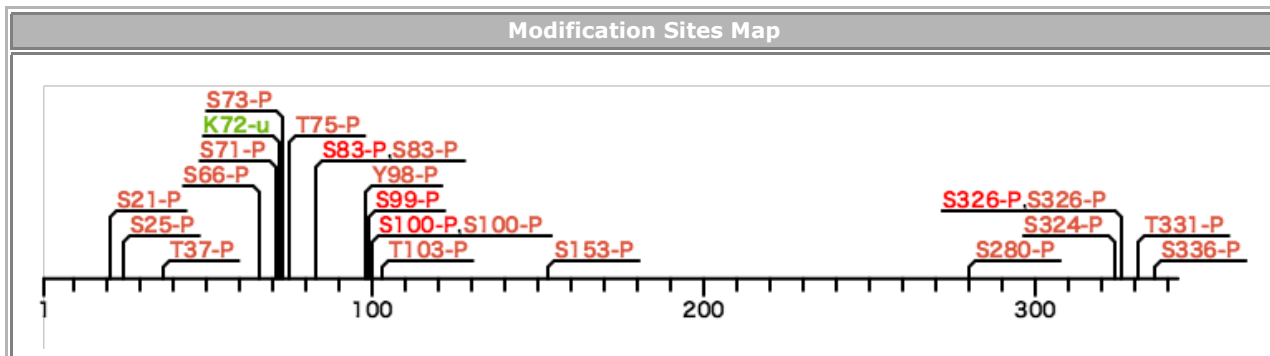


ID	Accession	GeneName	Chr.No.		Description
CCNY_HUMAN	Q8ND76	CCNY	10p11.21	35535953..35860852	Cyclin-Y



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
326	S	P	Lab	130327_HEK_CE_pphos.mgf[F015007]
326	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
326	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
326	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
326	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
326	S	P	Lab	130415_HEK_ME_tphos.mgf[F015010]
326	S	P	Lab	100510-lungc472.mgf[F017497]
326	S	P	Lab	100510-lungc480.mgf[F017498]
326	S	P	Lab	100510-lungc533.mgf[F017499]
326	S	P	Lab	100510-lungc858.mgf[F017501]
326	S	P	Lab	100510-lungc1006.mgf[F017502]
326	S	P	Lab	100510-lungc1016.mgf[F017503]
326	S	P	Lab	100510-lungc1046.mgf[F017504]
326	S	P	Lab	100510-lungc1059.mgf[F017505]
326	S	P	Lab	100510-lungc1060.mgf[F017506]
326	S	P	Lab	100510-lungc1067.mgf[F017507]
326	S	P	Lab	100510-lungc1077.mgf[F017508]
326	S	P	Lab	100510-lungc1105.mgf[F017510]
326	S	P	Lab	140320_Agarose_.mgf[F017423]
326	S	P	Lab	110218_pRMUGS_2.mgf[F017480]
326	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
326	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
326	S	P	Lab	100510-lungc126.mgf[F017483]
326	S	P	Lab	100510-lungc141.mgf[F017485]
326	S	P	Lab	100510-lungc220.mgf[F017487]
326	S	P	Lab	100510-lungc298.mgf[F017488]
326	S	P	Lab	100510-lungc343.mgf[F017490]
326	S	P	Lab	100510-lungc365.mgf[F017491]
326	S	P	Lab	100510-lungc391.mgf[F017493]
326	S	P	Lab	100510-lungc422.mgf[F017494]
326	S	P	Lab	100510-lungc429.mgf[F017495]

326	S	P	Lab	100510_lungc446.mgf[F017496]
326	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
326	S	P	Lab	100628_akimura_pOVSAHO_2.mgf[F017461]
326	S	P	Lab	100628_akimura_pOVSAHO_3.mgf[F017462]
326	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
326	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
326	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
326	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
326	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
326	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
326	S	P	Lab	110218_pOVSAYO_1.mgf[F017469]
326	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
326	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
326	S	P	Lab	110218_pRMG1_1.mgf[F017472]
326	S	P	Lab	110218_pRMG1_2.mgf[F017473]
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326	S	P	Lab	110218_pRMG2_4.mgf[F017478]
326	S	P	Lab	110218_pRMUGS_1.mgf[F017479]
326	S	P	Lab	100627_akimura_pOVISE_1.mgf[F017437]
326	S	P	Lab	100627_akimura_pOVISE_2.mgf[F017440]
326	S	P	Lab	110711_titania_LNCaP_AI_2.mgf[F017442]
326	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
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326	S	P	Lab	110711_titania_LNCaP_AI_4.mgf[F017445]
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326	S	P	Lab	100627_akimura_pOVTOKO_2.mgf[F017449]
326	S	P	Lab	100627_akimura_pOVTOKO_3.mgf[F017450]
326	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
326	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
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326	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
326	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
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326	S	P	Lab	140320_tita_C18_.mgf[F017426]
326	S	P	Lab	140320_HEK_ME_.mgf[F017427]
326	S	P	Lab	140320_HEK_SCE_.mgf[F017428]
326	S	P	Lab	140320_OVISE_ME_.mgf[F017429]
326	S	P	Lab	110711_titania_LNCaP_1.mgf[F017433]
326	S	P	Lab	110711_titania_LNCaP_2.mgf[F017434]

326	S	P	Lab	110711_titania_LNCaP_3.mgf[F017435]
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326	S	P	Lab	110711_titania_LNCaP_5.mgf[F017438]
326	S	P	Lab	110711_titania_LNCaP_6.mgf[F017439]
326	S	P	Lab	110711_titania_LNCaP_AI_1.mgf[F017441]
326	S	P	Lab	100520-GIST-IM1.mgf[F017509]
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326	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
326	S	P	Lab	100520-GIST-IM3.mgf[F017514]
326	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
326	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
326	S	P	Lab	100520-GIST-R2.mgf[F017517]
326	S	P	Lab	100520-GIST-R3.mgf[F017519]
326	S	P	Lab	100520-GIST-W1.mgf[F017521]
326	S	P	Lab	100520-GIST-W2.mgf[F017522]
326	S	P	Lab	100520-GIST-W3.mgf[F017524]
326	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
326	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
326	S	P	Paper	J Proteomics 2011, 75(4), 1343-1356
326	S	P	Paper	J Proteomics 2014, 96, 253-262
326	S	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668
326	S	P	Paper	Mol Cell Proteomics 2015, 14(6), 1599-1615
326	S	P	Paper	Sci Signal 2011, 4(164), rs3

Protein Sequence

MGNTTSCCVS SSPKLRRNAH **S**RLESYRPDT DLSRED**T**GCN LQHISDRENI DDLNMEFNPS DHPRAS**T**IFL **SKSQT**DVREK R
KSLFINHHP PGQIARK**YSS** CS**T**IFLDDST VSQPNLKYTI KCVAlAIYYH IKNRDPDGRM LLDIFDENLH PL**S**SKSEVPPD YDKHN
PEQKQ IYRFVRTLFS AAQLTAECAI VTLVYLERLL TYAEIDICPA NWKRIVLGAI LLASKVWDDQ AVWNVDYCQI LKDITVEDM
N ELERQFLELL QFNINVPSSV YAKYYFDL**S** LAEANNLSFP LEPLSRERAH KLEAISRLCE DKYKDLRRSA RKRS**SAS**ADNL **T**LP
RW**S**PAII

Backcolor of amino acid : Yellow -> site of modification, gray -> in front of processing