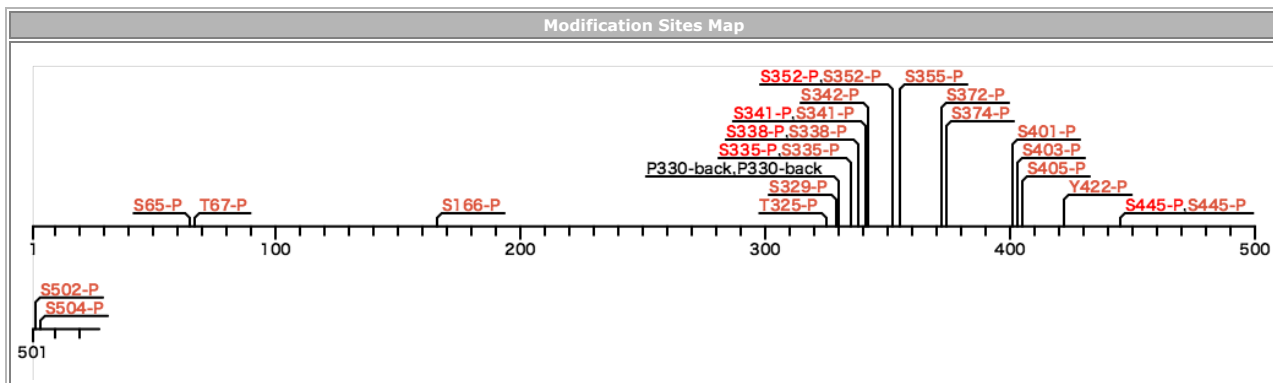


ID	Accession	GeneName	Chr.No.		Description
CCNL1_HUMAN	Q9UK58	CCNL1	3q25.31	156864297..156878549	Cyclin-L1



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
352	S	P	Lab	130415_HEK_CE_tphos.mgf[F015009]
352	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
352	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
352	S	P	Lab	140320_Agarose_.mgf[F017423]
352	S	P	Lab	110218_pRMUGS_3.mgf[F017481]
352	S	P	Lab	110218_pRMUGS_4.mgf[F017482]
352	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
352	S	P	Lab	100628_akimura_pOVSAHO_1.mgf[F017460]
352	S	P	Lab	100628_akimura_pOVSAHO_2.mgf[F017461]
352	S	P	Lab	100628_akimura_pOVSAHO_3.mgf[F017462]
352	S	P	Lab	110218_pOVKATE_1.mgf[F017463]
352	S	P	Lab	110218_pOVKATE_2.mgf[F017464]
352	S	P	Lab	110218_pOVKATE_3.mgf[F017465]
352	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
352	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
352	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
352	S	P	Lab	110218_pOVSAYO_1.mgf[F017469]
352	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
352	S	P	Lab	110218_pRMG2_1.mgf[F017475]
352	S	P	Lab	110218_pRMG2_2.mgf[F017476]
352	S	P	Lab	110711_titania_LNcAP_AI_2.mgf[F017442]
352	S	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
352	S	P	Lab	110711_titania_LNcAP_AI_3.mgf[F017444]
352	S	P	Lab	110711_titania_LNcAP_AI_4.mgf[F017445]
352	S	P	Lab	110711_titania_LNcAP_AI_5.mgf[F017446]
352	S	P	Lab	100627_akimura_pOVTOKO_1.mgf[F017447]
352	S	P	Lab	110711_titania_LNcAP_AI_6.mgf[F017448]
352	S	P	Lab	100627_akimura_pOVTOKO_2.mgf[F017449]
352	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
352	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
352	S	P	Lab	100627_akimura_pRMG1_3.mgf[F017453]
352	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
352	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
352	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
352	S	P	Lab	100628_akimura_pOVCAR3_1.mgf[F017457]
352	S	P	Lab	100628_akimura_pOVCAR3_2.mgf[F017458]
352	S	P	Lab	100628_akimura_pOVCAR3_3.mgf[F017459]
352	S	P	Lab	140320_tita_C18_.mgf[F017426]
352	S	P	Lab	140320_HEK_SCE_.mgf[F017428]
352	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
352	S	P	Lab	110711_titania_LNcAP_1.mgf[F017433]
352	S	P	Lab	110711_titania_LNcAP_2.mgf[F017434]
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352	S	P	Lab	110711_titania_LNcaP_6.mgf[F017439]
352	S	P	Lab	110711_titania_LNcaP_AI_1.mgf[F017441]
352	S	P	Lab	100520-GIST-IM1.mgf[F017509]
352	S	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
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352	S	P	Lab	100520-GIST-IM2.mgf[F017512]
352	S	P	Lab	140326_GIST_NES_tita_2_.mgf[F017513]
352	S	P	Lab	100520-GIST-IM3.mgf[F017514]
352	S	P	Lab	140326_GIST_NES_tita_3_.mgf[F017516]
352	S	P	Lab	100520-GIST-R2.mgf[F017517]
352	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
352	S	P	Lab	100520-GIST-R3.mgf[F017519]
352	S	P	Lab	100520-GIST-W1.mgf[F017521]
352	S	P	Lab	100520-GIST-W2.mgf[F017522]
352	S	P	Lab	100520-GIST-W3.mgf[F017524]
352	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
352	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
352	S	P	Paper	J Proteomics 2011, 75(4), 1343-1356
352	S	P	Paper	J Proteomics 2014, 96, 253-262
352	S	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668
352	S	P	Paper	Mol Cell Proteomics 2015, 14(6), 1599-1615

Protein Sequence

MASGPHSTAT AAAAASSAAP SAGGSSSGTT TTTTTTGGI LIGDRLYSEV SLTIDHSLIP EERLSPITPSM QDGLDLPSET DLRILGCELI QAAGILLRLP QVAMATGQVL FHRFF
 YKSF VKHSFEIVAM ACINLASKIE EAPRRIRDVI NVFHHLRQLR GKRTPLIL DQNYINTKNQ VIKARRVLK ELGFCVHVKH PHKIIVMYLQ VLECERNQTL VQTAWNYMN
 D SLRTNVFVRF QPETIACACI YLAARALQIP LPTRPHWLL FGTTEEEIQE ICIELRLYT RKKPNYELLE KEVEKRKVAL QEAKLKAKGL NPDGTPALST LGGFSPASPKSSPRE
 VKAEE KSPISINVKT VKKEPEDRQQ ASKSPYNGVR KDSKRSRNSR SASRSRSTR SRSRSHTPRR HYNRRRSRSG TYSSRSRSRSHSESPPRH HNHGSPHLKA KHTRD
 DLKSS NRHGHKRKKS RRSQSQRD HSDAAKKRRH ERGHHRDRRE RRSFERSHK SKHHGGSRSG HGRHR

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