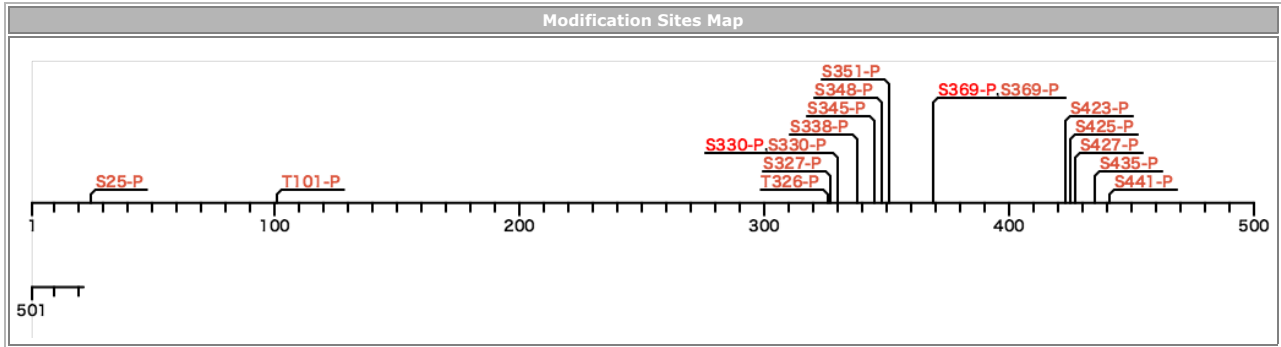


ID	Accession	GeneName	Chr.No.	Description
CCNL2_HUMAN	Q96S94	CCNL2	1p36.33 1321091..1334708	Cyclin-L2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
369	S	P	Lab	140326_OVISE_NES_tita_2_.mgf[F017520]
369	S	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
369	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
369	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
369	S	P	Lab	140326_OVISE_NES_tita_.mgf[F017518]
369	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
369	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
369	S	P	Paper	J Proteomics 2011, 75(4), 1343-1356
369	S	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668
369	S	P	Paper	Mol Cell Proteomics 2015, 14(6), 1599-1615
369	S	P	Paper	Proc Natl Acad Sci USA 2014, 111(21), E2182-E2190

Protein Sequence	
MAAAAAAGA AGSAAPAAA GAPGSGGAPS GSQVGLIGDR LYSGLVLTLE NCLLPDDKLR FTPSMSSGLD TDTETDLRVV GCELIQAAGI LLRLPQVAMA TGGVLFQRF YTK	SFVKHSM EHVSMACVHL ASKIEEAPRR IRDVINVFHR LRQLRDKKKP VPLLLDQDYV NLKNQIIKAE RRVLKELGFC VHVKHHPHKII VMYLQVLECE RNQHLVQTSW NYMND
SLRTD VVRFQPEST ACACIYLAAR TLEIPLNRP HWFLLFGATE EEIQEICKI LQLYARKVD LTHLEGEVEK RKHAIEEAKA QARGLLPGGT QVLDGTSGFSPAPKLVESPK E	GKGSKPSPL SVKNTKRRLE GAKKAKADSP VNGLPKGRES RSRSRSEQS YSRSPRSAS PKRRKSDSGS TSGGSKSQSR SRSRSDSPPR QAPRSAPYKG SEIRGSRKSK D
CKYPQKPHK SRSRSSRSR SRSRERADNP GKYYKKSHYY RDQRRERSRS YERTGRRYER DHPGHSRHR	

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