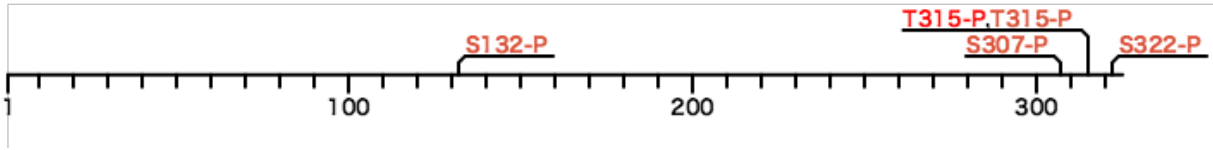
ID	Accession	GeneName	Chr.No.		Description
CCNH_HUMAN	P51946	CCNH	5q14.3	86687311..86708836	Cyclin-H

Modification Sites Map



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
315	T	P	Lab	140326_OVISE_NES_tita_3_.mgf[F017523]
315	T	P	Lab	140320_Mag_new_.mgf[F017424]
315	T	P	Lab	100627_akimura_pOVISE_3.mgf[F017443]
315	T	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
315	T	P	Lab	140326_GIST_NES_tita_.mgf[F017511]
315	T	P	Paper	Cell Rep 2014, 8(5), 1583-1594
315	T	P	Paper	J Proteome Res 2013, 12(1), 260-271
315	T	P	Paper	J Proteomics 2011, 75(4), 1343-1356
315	T	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668
315	T	P	Paper	Proc Natl Acad Sci USA 2014, 111(21), E2182-E2190

Protein Sequence

MYHNSSQKRH WTFSSEEQLA RLRADANRKF RCKAVANGKV LPNDPVFLEP HEEMTLCKYY EKRLLEFCSV FKPAMPRSV
V GTACMYFKRF YLNNSVMEYH PRIIMLTCAF LACKVDEFNV SSPQFVGNLR ESPLGQEKAL EQILEYELLL IQQLNFHLIV
HNPYRPFEGF LIDLKTRYPI LENPEILRKT ADDFLNRIAL TDAYLLYTPS QIALTAILSS ASRAGITMES YLSESLMLKE NRT
CLSQLLD IMKSMRNLVK KYEPPRSEEV AVLKQKLERC HSAELALNVI TKKRKGYEDD DYVSKKSKHE EEEWTDDDLV E
S

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