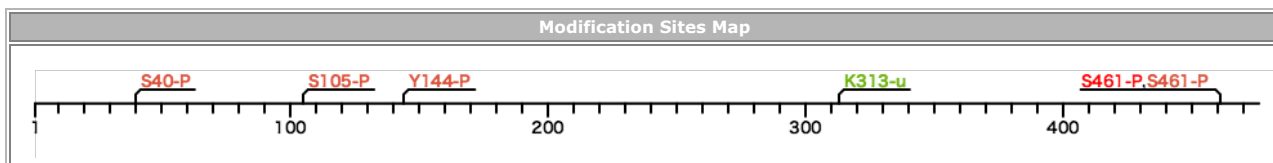


ID	Accession	GeneName	Chr.No.	Description
CBLC_HUMAN	Q9ULV8	CBLC	19q13.32 45281126..45303891	E3 ubiquitin-protein ligase CBL-C



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
461	S	P	Lab	110218_pOVMANA_1.mgf[F017466]
461	S	P	Lab	110218_pOVMANA_2.mgf[F017467]
461	S	P	Lab	110218_pOVMANA_3.mgf[F017468]
461	S	P	Lab	110218_pOVSAYO_2.mgf[F017470]
461	S	P	Lab	110218_pOVSAYO_3.mgf[F017471]
461	S	P	Lab	100627_akimura_pOVICE_1.mgf[F017437]
461	S	P	Lab	100627_akimura_pOVICE_2.mgf[F017440]
461	S	P	Lab	100627_akimura_pOVICE_3.mgf[F017443]
461	S	P	Lab	100627_akimura_pRMG1_3.mgf[F017453]
461	S	P	Lab	100628_akimura_pMCAS_1.mgf[F017454]
461	S	P	Lab	100628_akimura_pMCAS_2.mgf[F017455]
461	S	P	Lab	100628_akimura_pMCAS_3.mgf[F017456]
461	S	P	Lab	140320_tita_C18_.mgf[F017426]
461	S	P	Lab	140320_OVICE_SCE_.mgf[F017431]
461	S	P	Paper	J Proteomics 2014, 96, 253-262
461	S	P	Paper	Mol Cell Proteomics 2012, 11(9), 651-668

Protein Sequence	
MALAVAPWGR QWEEARALGR AVRMLQRLEE QCVDPRLSV S PPSLRDLLPR TAQLLREVAH SRRAAGGGGP GGPGGSGDFL LIYLANLEAK SRQVAALLPP RGR R SA NDEL FRAGSRLRRQ LAKLAIIFSH MHAELHALFP G G K Y CGHMYQ LTKAPAHTFW RESCGARCVL PWAEFESLLG TCHPVEPGCT ALALRTTIDL TCSGHVSIFE FDV FTRLFQP WPTLLKNWQL LAVNHPGYMA FLT Y DEVQER LQACRDKPGS YIFRPSCTRL GQWAIGYVSS DGSILQTIPA NKPLSQVLE GQKDGFLYP DG K THNPDLT ELGQAEPQQR IHVSEELQLL YWAMDSTFEL CKICAESNKD VKIEPCGHLL CSCCLAAWQH SDSQTCPCFR CEIKGWEAVS IYQFHGQATA EDSGNSSDQE GRELE LGQVP LSAPLPPRP DLPPRKPRNA QPKVRLKGN S PPAALGPQD PAP	

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