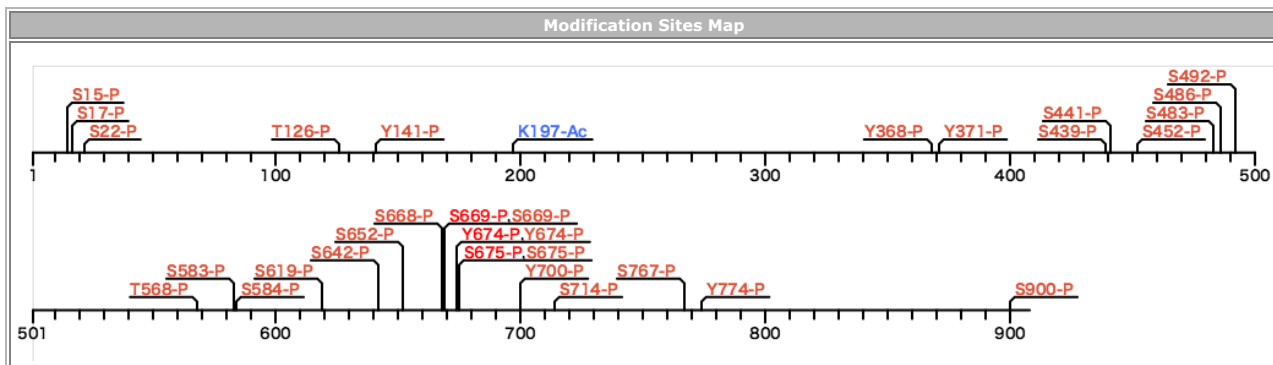


ID	Accession	GeneName	Chr.No.		Description
CBL_HUMAN	P22681	CBL	11q23.3	119076752..119178859	E3 ubiquitin-protein ligase CBL



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
674	Y	P	Lab	100407-nagata-pMCAS-10mg.mgf[F014984]
674	Y	P	Lab	100407-nagata-pMCAS-10mg.mgf[F014984]
674	Y	P	Lab	100407-nagata-pOVCAR3-10mg-MGF.mgf[F014985]
674	Y	P	Lab	100626-nagata-pOVCAR3-10mg.mgf[F014996]
674	Y	P	Lab	100626-nagata-pOVISE-10mg.mgf[F014998]
674	Y	P	Lab	100626-nagata-pOVMANA-10mg.mgf[F015000]
674	Y	P	Paper	Cell Rep 2014, 8(5), 1583-1594

Protein Sequence	
MAGNVKSSG AGGGSGSGGS GSGGLIGLMK DAFQPHHHH HHLSPHPPGT VDKMVEKCW KLMDKVRLC QNPKLALKNS PPYILDLLPD TYQHLRTILS RYEGKMETLG ENEYFRVFM E NLMKKTKQTI SLFKEGKERM YEENSQPRRN LTKLSLIFSH MLAELKGIFP SGLFQGDTRF ITKADAAEFW RKAFGEKTI V PWKSRFQALH EVHPISSGLE AMAL KSTIDL TCNDYISVFE FDIFTRLFQP WSSLLRNWNS LAVTHPGYMA FLTIDEVKAR LQKFIHKPGS YIFRLSCTRL GQWAIGYVTA DGNILQTIPIH NKPLFQALID GFREGFYLF P DGRNQNPDLT GLCEPTQDH IKVTQEYEL YCEMGSTFQL CKICAENDKD VKIEPCGHLM CTSCLTSWQE SEGQGCPCFR CEIKGTEPIV VDFPDRGSG SLLLRQAEGA P SPNYDDDDD ERADDTLFMM KELAGAKVER PPSPFMAPQ ASLPPVPPRL DLLPQRVCVP SSASALGTAS KAASGSLHKD KPLPVPPTLR DLPPPPPPDR PYSVGAESRP QRR PLPCIPG DCPSRDKLP VPSSRLGDSW LPRPIKVPV SAPSSSDPWT GRELTNRHSL PFSLPQMEP RPDVPRLGST FSLDTSMSMN SPLVGPEDC HPKIKPSSA NAIYSL AARP LPVPKLPPE QCEGEEDTEY MTPSSRPLRP LDTSQSSRAC DCDQQIDSC T YEAMYNISQ APSITESSTF GEGNLAHAHA NTGPPESENE DDGYDVPKPP VPAVLARRT L SDISNASSSF GWLSLDGDP T TNVTEGSQVP ERPPKPPRR INSEKAGSC QQSGSPAASA ATASPLSSE IENLMSQGY S YQDIQKALVI AQNNIEMAKN ILREFVSISS P AHVA	

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