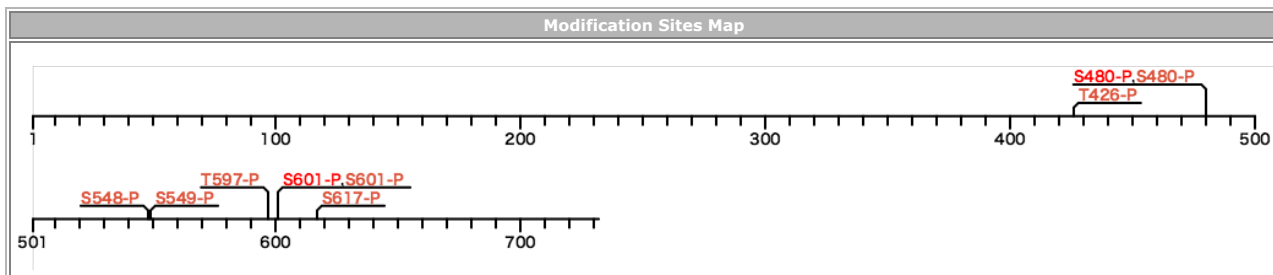


ID	Accession	GeneName	Chr.No.	Description
CCNT2_HUMAN	O60583	CCNT2	2q21.3	135675805..135716912 Cyclin-T2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
601	S	P	Lab	140320_tita_C18_.mgf[F017426]
601	S	P	Lab	140320_tita_SDB_.mgf[F017430]
601	S	P	Lab	140320_OVISE_SCE_.mgf[F017431]
601	S	P	Paper	Cancer Res 2009, 69(6), 2663-2668
601	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
601	S	P	Paper	Mol Cell 2008, 31(3), 438-448
601	S	P	Paper	Mol Cell Proteomics 2009, 8(7), 1751-1764
601	S	P	Paper	Mol Cell Proteomics 2010, 9(6), 1167-1181
601	S	P	Paper	Mol Cell Proteomics 2015, 14(6), 1599-1615
601	S	P	Paper	Sci Signal 2009, 2(84), ra46
601	S	P	Paper	Sci Signal 2011, 4(179), rs5

Protein Sequence	
MASGRGASSR WFFTREQLN TPSRRCGVEA DKELSCRQQA ANLIQEMGQR LNVSQLTINT AIVYMHRYM HHSFTKFNKN IISSTALFLA AKVEEQARKL EHVIVAHAC LH PLEPLDLD KDAYLQQTQ ELVILEITIML QTLGFEITIE HPHTDVVKCT QLVASKDLA QTSYFMATNS LHLTTFCLQY KPTVIACVCI HBLACKWSNWE IPVSTDGKHW WEYVDP TVTL ELLDELTHEF LQILEKTPNR LKKIRNWRAN QAARKPKVDG QVSETPLGGS SLVQNSILVD SVTGVPTNPS FQKPSTSAFP APVPLNSGNI SVQDSHTSDN LSMLATGMP S TSYGLSSHQ WPQHQSART EQLYSQKQET SLGGSQYNIN FQQGPSISLH SGLHHRPDKI SDHSSVKQEY THKAGSSKHH GPISLPGIIPQKMSLDKYR EKRRKLETLDL DVRDHVIAAQ VEQHQKQGS QAASSSSVTS PIKMKIPIAN TEKYMADKKE KSGSLKLRIP IPPTDKSASK EELKMKIKVS SSERHSSSDE GSGKSKHSSP HISRDHKEKH KE HPSSRHHT SSHKSHSHS GSSSGGSKHS ADGIPPTVLRSPVGLSSDGI SSSSSSRKR LHVNDASHNH HSKMSKSSKS SGSSSSSSSS VKQYISSHNS VFNHPLPPP PV TYQVGYGH LSTLVKLDKK PVETNGPDAN HEYSTSSQHM DYKDTFDMLD SLLSAQGMN	

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