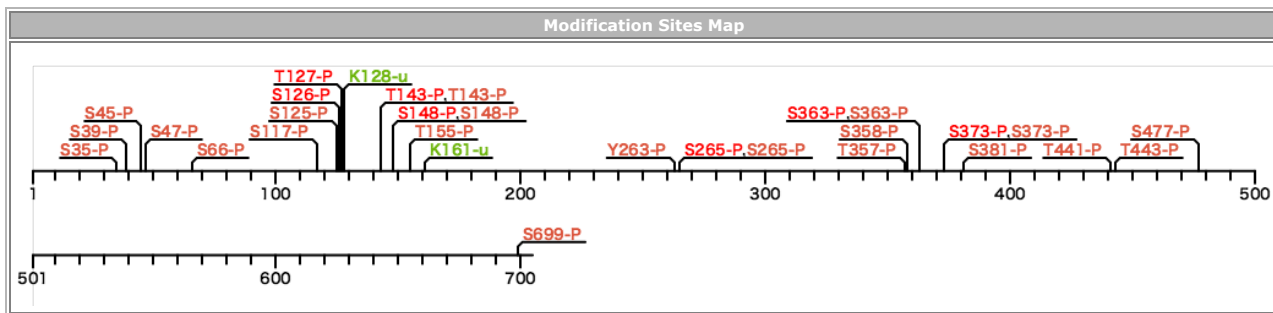


ID	Accession	GeneName	Chr.No.	Description
CASC3_HUMAN	O15234	CASC3	17q21.1 38296571..38328436	Protein CASC3



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
143	T	P	Lab	140320_Agarose_.mgf[F017423]
143	T	P	Lab	110218_pOVKATE_3.mgf[F017465]
143	T	P	Lab	110218_pOVSAYO_1.mgf[F017469]
143	T	P	Lab	110218_pOVSAYO_3.mgf[F017471]
143	T	P	Lab	110218_pRMG1_3.mgf[F017474]
143	T	P	Lab	100627_akimura_pOVISe_2.mgf[F017440]
143	T	P	Lab	100520-GIST-IM2.mgf[F017512]
143	T	P	Paper	Sci Signal 2011, 4(179), rs5

**Protein Sequence**

MADRRRQRAS QDTEDEESGA SGSDSGGSPL RGGGSCSGSA GGGGSGSLPS QRGGRTGALH LRRVESGGAK SAESECESE DGIEGDAVLV DYESAEDSEG EEEYSEEE  
 N SKVELKSEAN DAVN SSTKKEE KGEEKPDTKS TVTGERQSGD GQESTEPVEN KVGKKGPKHL DDEDRKNPA YIPRKLFFH HDLRGQTQEE EVRPKGRQRK LWKDEGRW  
 EH DKFREDEQAP KSRQELIALY GDIRSAHNP DDIKPRRIRK PRYGSPPQRD PNWNGERLNK SHRHQGLGGT LPPRTFINRN AAGTGRMSAP RNYSRSGGFK EGRAGFRPV  
 E AGGQHGGRSQ ETVKHEISYR SRRLEQTSVR DPSPEADAPV LGSPEKEEAA SEPPAAAPDA APPPPDRPIE KKSYSRARRT RTKVGDAVKL AEEVPPPPPEG LIPAPPVPET TPT  
 PPTKTGT WEAPVDSSTS GLEQDVAQLN IAEQNWSPGQ PSFLQPRELR GMPNHIHMG A GPPQFNRM E EMGVQGGRAK RYSSQRQRPV PEPPAPPVHI SIMEGHYDP LQF  
 QGPIYTH GDSPAPLPPQ GMLVQPGMNL PHPLPHQPT PAPLPNPLY PPPVMSMPGQ PPPQQLLAPT YFSAVGMNF GNPSYPYAPG ALPPPPPHL YPNTQAPSQV YGGVTY  
 YNPA QQVQPKPSP PRRTQPVTI KPPPPEVVS R GS

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