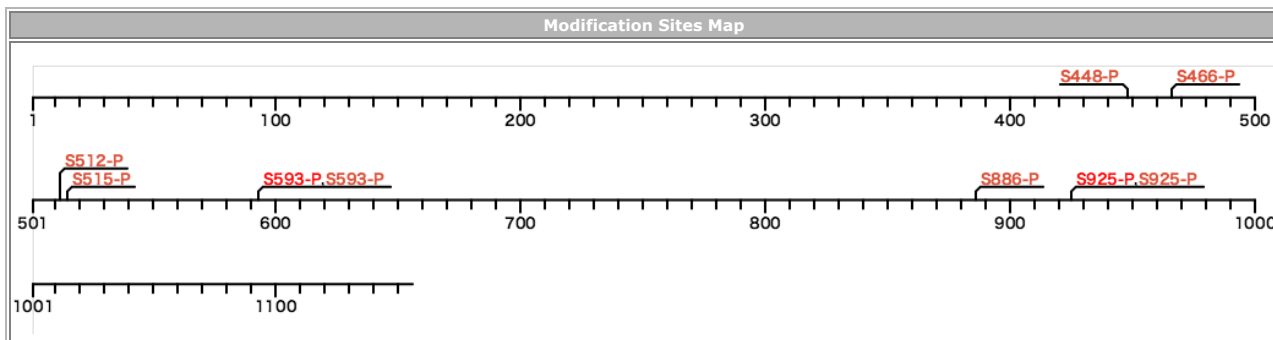


ID	Accession	GeneName	Chr.No.	Description	
CAR11_HUMAN	Q9BXL7	CARD11	7p22.2	2945775..3083579	Caspase recruitment domain-containing protein 11



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
593	S	P	Lab	100627_akimura_pRMG1_1.mgf[F017451]
593	S	P	Lab	100627_akimura_pRMG1_2.mgf[F017452]
593	S	P	Paper	J Proteome Res 2013, 12(1), 260-271
593	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MPGGGPEMDD YMETLKDEED ALWENVECNR HMLSRYNPA KLTPLYRQCK VIDEQDEDEV LNPMLPSKI NRAGRLLDIL HTKGQRGYVV FLESLEFYYP ELYKLVGTGKE PTR RFSTIV EEGHEGLTHF LMNEVIKQQ QMKAKDLQRC ELLARLRQLE DEKKQMTLTR VELLTFQERY YKMKEERDSY NDELVKVKDD NYNLAMRYAQ LSEKNNMAMV RSRD LQLEID QLKHLRLNKME EECKLERNQS LKLKNDIENR PKKEQVLELE RENEMLKTKN QELQSIIQAG KRSLPDSDKA ILDILEHDRK EALEDRQELV NRIYNLQEEA RQAEELRD KY LEEKEDLELK CSTLGKDCM YKHRMNTVML QLEEVERERD QAFHSRDEAQ TOYSQCLIEK DKYRKQIREL EEKNDEMRIE MVRREACIVN LESKLRRLSK DSNNLDQSLP RNLVPTIISQ DFGDASPRTN GQEADDSSTS EESPEDSKYF LPYHPPQRRM NLKGIQLQRA KSPISLKRKTS DFQAKGHEEE GTDASPSSCG SLPITNSFTK MQPPRSRSSI MSI TAEPPGN DSIVRRYKED APHRSTVEED NDGGFDALD LDDDSHERYS FGPSSIHSSS SSHQSEGLDA YDLEQVNLMF RKFSLERPFR PSVTSVGHVR GPGPSVQHTT LNGD SLTSQL TLLGGNARGS FVHSVKPGSL AEKAGLREGH QLLLLEG CIR GERQSVPLDT CTKEEAHWIT QRCSGPVT LH YKVNHEGYRK LVKDMEDGLI TSGDSFYIRL NLNISSQ LDA CTMSLKCDDV VHVVRTMYQD RHEWL CARVD PFDHDLDMG TIPSYSRAQQ LLLVQLQRLM HRGSREEVDG THHTLRALRN TLQPPEALST SDPRVSPRLS RASFLFGQ LL QFVSRSENKY KRMNSNERVR IISGSP LGS L ARSSLDATKL L TEKQEELDP ESELGKNLSL IPYSLVRAFY CERRRPVLFT PTVLAKTLVQ RLLNSGGAME FTICKSDIVT RDE FLRRQKT ETIYSREKN PNAFECIAPA NIEVAANKN HCLLEAGIGC TRDLIKSNIY PIVLFIRVCE KNIKRFKLL PRPETEEEF L RVCRLKEKEL EALPCLYATV EPDMWGSVEE LLRVVKDKIG EEQRKTIWVD EDQ

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