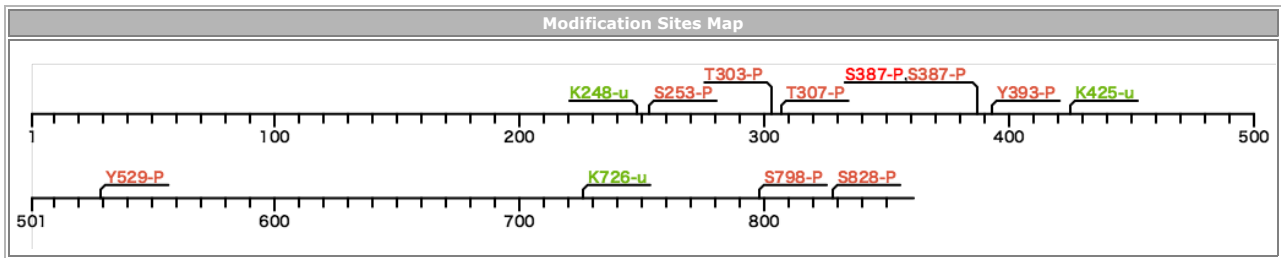


ID	Accession	GeneName	Chr.No.		Description
AGO2_HUMAN	Q9UKV8	EIF2C2	8q24.3	141541264..141645718	Protein argonaute-2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
---------	------------	------	----------	--------

Protein Sequence	
MYSGAGPALA PPAPPPPIQG YAFKPPRPD FGTSGRITKL QANFFEMDIP KIDIHYELD IKPEKPRRV NREIVEHMQ HFKTQIFGDR KPVFDGRKNL YTAMPLIGR DKVEL EVTLP GEGKDRIFKV SIKWVSCVSL QALHDALSGR LPSVPFETIQ ALDVMRHLR SMRYTPVGRS FFTASEGCSN PLGGGREVWF GFHQSVRPSL WKMMLNIDVS ATAFYKA QPV IEFVCEVLD KSIIEEQKPL TD SQRVKFTK EIKGLKVEIT HCGQMKRKYR VCNVTRRPAS HQTFPLQES GQ T VECTVAQ YFKDRHKLVL RYPHLPCLQV GQEQKHTYLP LEVCNIVAGQ RCIKKLTDNQ TSTMIRATAR SAPDRQEIS KLMRSASFNT DP YVREFGIM VKDEMVDVTG RVLQPPSILY GGRNKAIATP VQGVWDMRNL QFHTGIEIKV WA IACFAPQR QCTEVHLKSF TEQLRKISR AGMPIQGQPC FCKYAQGADS VEPMFRHLKN TYAGLQLVV ILPGKTPVYA EVKRVGDTVL GMATQCVQMK NVQRTTPQTL SNLC LKINVK LGGVNNILL QGRPPVFQQP VIFLGADVTH PPAGDGKPKS IAAVVGSDMA HPNRYCATVR VQHRQEIIQ DLAAMVRELL IQFYKSTRFK PTRIIFYRDG VSEGQFQ QVL HHELLAIREA CIKLEKDYQP GITFIVVQKR HHTRLFCTDK NERVGKSGNI PAGTTVDTKI THPTEFDYFL CSHAGIQGTS RPSHYHVLWD DNRFSDELQ ILTYQLCHTY V RCTRSV SIP APAYYAHVA FRARYHLVDK EHDSAEGSHT SGQSNGRDHQ ALAKAVQVHQ DTLRMTMYF	

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