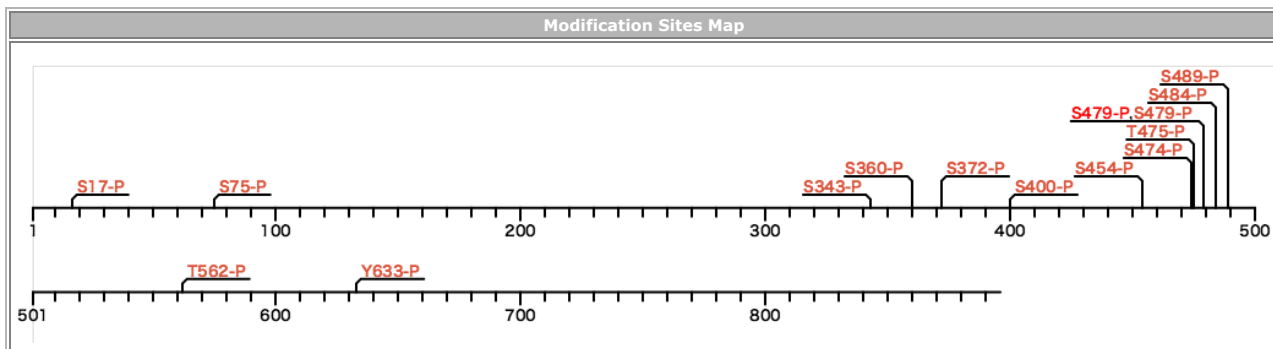


ID	Accession	GeneName	Chr.No.		Description
C19L2_HUMAN	Q2TBE0	CWF19L2	11q22.3	107197071..107328572	CWF19-like protein 2



Click a modification site to display information in detail.

Site no	Amino acid	Type	Division	Detail
479	S	P	Lab	130327_HEK_ME_pphos.mgf[F015008]
479	S	P	Paper	Cell 2006, 127(3), 635-648
479	S	P	Paper	Cell Rep 2014, 8(5), 1583-1594
479	S	P	Paper	Sci Signal 2009, 2(84), ra46

Protein Sequence

MATSM^SAAASG RFESA^SIEE RKEQTRNARA EVLRQAKANF EKEERRKELK RLRGEDTWML PDVNERIEQF SQEH^SVK^SKKK^S KDKH^SSKKAK KE^SKKK^SSKK^SQ KYEKNNESSD
SSSSSEDEWV EAVPSQTPDK EKAWKVKDEK SGKDDTQIIK RDEWMTVD^SFM SVKTVSSSSL KAEKETMRKI EQEKNQALEQ SKLMERELNP YWKDGGTGLP PEDCSVSSIT
KVS^SVVEDGGL SWLRKSYLRM KEQAEQSRN FEDIVAERYG SMEIFQSKLE DAEKA^SASTKE DYRRERWRKP TYSDKAQNCQ ESRESDLVKY GNSSRD^SRYAT TDTAKNSNNE
KF^SIGDEKDKR PG^SLET^SCRRE SNPRQNQEF^S FG^SNLRAKFLR PS^SDDELSFH SKGRKFEPLS SSSALVAQGS^S LC^SSGFRKPTK NSEERLTSWS RSDGRGDKKH SNQK^SPSETST D
EHQHVPE^SDP REK^SQDEVLR DDP^SPKKEHLR DTK^STFAG^SSP ERES^SIHL^SSV DEKNKLGAKI IKAEMMG^SNME LAEQLKVQLE KANKFKETIT QIPK^SSGVEN EDQ^SQEVILVR TDQS
GRVW^SPV N^STPGK^SLESQ GGR^SRRKQ^SMVS THEERERVRY FHDDDNLSL^SN DLV^SKNEKMG^ST AENQ^SNKLFMR MASK^SFMGKTD GD^SY^STLDDMF VSKAAERERL GEE^SENQR^SKK AIA
EHRSLAA QMEKCLYCFD SSQFPKHLIV AIGVKVYLCL PN^SVRSLTEGH CLIVPLQHHR AATLLDEDI^SW EEIQMFRKSL VKMFEDKGLD CIFLETNMSM KKQYHMVYEC IPLPKEV
GDM APIYFKKAIM ESDEEWSMNK KLIDLSSKDI RKS^SVPRGLPY FSVDFGLHGG FAHVIEDQH^SK FPHYFGKEII GGMLDIEPRL WRKGIRESFE DQRK^SKALQFA QWWKPYDFT
K SKN

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