	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q9P218	COKA1_HUMAN	Homo sapiens	Collagen alpha-1(XX) chain	26.273286	S396;T398	NaN	31637018	MSSGDPAHLGLCLWLWLGATLGRE QVQASGLLRLAVLPEDRLQMKWRE SEGSGLGYLVQVKPMAGDSEQEVIL TTKTPKATVGGLSPSKGYTLQIFELT GSGRFLLARREFVIEDLKSSSLDRSS QRPLGSGAPEPTPSHTGSPDPEQAS EPQVAFTPSQDPRTPAGPQFRCLPPV PADMVFLVDGSWSIGHSHFQQVKD FLASVIAPFEIGPDKVQVGLTQYSGD AQTEWDLNSLSTKEQVLAAVRRLRY KGGNTFTGLALTHVLGQNLQPAAGL RPEAAKVVILVTDGKSQDDVHTAAR VLKDLGVNVFAVGVKNADEAELRLL ASPPRDITVHSVLDFLQLGALAGLLS RLICQRLQGGSPRQGPAAAPALDTLP APTSLVLSQVTSSSIRLSWTPAPRHP LKYLIVWRASRGGTPREVVVEGPAAS TELHNLASRTEYLVSVFPIYEGGVGE GLRGLVTTAPLPPPRALTLAAVTPRT VHLTWQPSAGATHYLVRCSPASPKG EEEEREVQVGRPEVLLDGLEPGRDY EVSVQSLRGPEGSEARGIRARTPTLA PPRHLGFSDVSHDAARVFWEGAPRP VRLVRVTYVSSEGGHSGQTEAPGNA TSATLGPLSSSTTYTVRVTCLYPGGG SSTLTGRVTTKKAPSPSQLSMTELPG DAVQLAWVAAAPSGVLVYQITWTPL GEGKAHEISVPGNLGTAVLPGLGRH TEYDVTILAYYRDGARSDPVSLRYTP STVSRSPPSNLALASETPDSLQVSWT PPLGRVLHYWLTYAPASGLGPEKSV SVPGARSHVTLPDLQAATKYRVLVSA IYAAGRSEAVSATGQTACPALRPDGS LPGFDLMVAFSLVEKAYASIRGVAM EPSAFGGTPTFTLFKDAQLTRRVSDV YPAPLPPEHTIVFLVRLLPETPREAFA LWQMTAEDFQPLLGVLLDAGKKSLT YFHRDPRAALQEATFDPQEVRKIFF GSFHKVHVAVGRSKVRLYVDCRKVA ERPLGEMGSPPAAGFVTLGRLAKAR GPRSSSAAFQLQMLQIVCSDTWADE DRCCELPASRDGETCPAFVSACSCS SETPGPPGPQGPPGLPGRNGTPGEQ GFPGPRGPPGVKGEKGDHGLPGLQ GHPGHQGIPGRVGLQGPKGMRGLE GTAGLPGPPGPRGFQGMAGARGTS GERGPPGTVGPTGLPGPKGERGEKG EPQSLATLYQLVSQASHVSKFDSFH ENTRPPMPILEQKLEPGTEPLGSPGT RSKALVPGEWGRGGRHLEGRGEPG AVGQMGSPGQQGASTQGLWE	None	None	None	None	None	None	None	None	None