	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	Q9UJ98	STAG3_HUMAN	Homo sapiens	Cohesin subunit SA-3	19.376537	NaN	S1203	28411811;30379171	MSSPLQRAVGDTKRALSASSSSSAS LPFDDRDSNHTSEGNGDSLLADEDT DFEDSLNRNVKKRAAKRPPKTTPVA KHPKKGSRVVHRHSRKQSEPPAND LFNAVKAAKSDMQSLVDEWLDSYK QDQDAGFLELVNFFIQSCGCKGIVTP EMFKKMSNSEIIQHLTEQFNEDSGD YPLIAPGPSWKKFQGSFCEFVRTLVC QCQYSLLYDGFPMDDLISLLTGLSDS QVRAFRHTSTLAAMKLMTSLVKVAL QLSVHQDNNQRQYEAERNKGPGQR APERLESLLEKRKELQEHQEEIEGM MNALFRGVFVHRYRDVLPEIRAICIE EIGCWMQSYSTSFLTDSYLKYIGWTL HDKHREVRLKCVKALKGLYGNRDLT TRLELFTSRFKDRMVSMVMDREYD VAVEAVRLLILILKNMEGVLTDADCE SVYPVVYASHRGLASAAGEFLYWKL FYPECEIRMMGGREQRQSPGAQRTF FQLLLSFFVESELHDHAAYLVDSLW DCAGARLKDWEGLTSLLLEKDQNL GDVQESTLIEILVSSARQASEGHPPV GRVTGRKGLTSKERKTQADDRVKLT EHLIPLLPQLLAKFSADAEKVTPLLQ LLSCFDLHIYCTGRLEKHLELFLQQL QEVVVKHAEPAVLEAGAHALYLLCN PEFTFFSRADFARSQLVDLLTDRFQ QELEELLQSSFLDEDEVYNLAATLK RLSAFYNTHDLTRWELYEPCCQLLQ KAVDTGEVPHQVILPALTLVYFSILW TLTHISKSDASQKQLSSLRDRMVAF CELCQSCLSDVDTEIQEQAFVLLSDL LLIFSPQMIVGGRDFLRPLVFFPEAT LQSELASFLMDHVFIQPGDLGSGDS QEDHLQIERLHQRRRLLAGFCKLLL YGVLEMDAASDVFKHYNKFYNDYG DIIKETLTRARQIDRSHCSRILLLSLK QLYTELLQEHGPQGLNELPAFIEMR DLARRFALSFGPQQLQNRDLVVML HKEGIQFSLSELPPAGSSNQPPNLAF LELLSEFSPRLFHQDKQLLLSYLEKC LQHVSQAPGHPWGPVTTYCHSLSPV ENTAETSPQVLPSSKRRRVEGPAKP NREDVSSSQEESLQLNSIPPTPTLTS TAVKSRQPLWGLKEMEEEDGSELD FAQGQPVAGTERSRFLGPQYFQTPH NPSGPGLGNQLMRLSLMEEDEEEE LEIQDESNEERQDTDMQASSYSSTS ERGLDLLDSTELDIEDF	None	None	None	None	None	None	None	None	None