

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	e. ri
Q8WXE1	ATRIP_HUMAN	Homo sapiens	ATR-interacting protein	27.244452	S339;T343;S352;T353;S358	NaN	28411811;38253038	MAGTSAPGSKRRSEPPAPRPGPPPG TGHPPSKRARGFSAAAAPDDDDPFG AHGDFTADDDLEELDTLASQALSQCP AAARDVSSDHKVRHLLDGMKSNPS GKNRETVPIKDNFELEVLQAQYKEL KEKMKVMEEVLIKNGEIKILRDSL HQTESVLEEQRSHFLLQEQKTAAL SDKEKEFSKQLQSLQSEIQFKDAEM NELRTKLQTSERANKLAAPSVSHVS PRKNPSVVIKPEACSPQFGKTSFPTK ESFSANMSLPHPCQTESGYKPLVGR EDSKPHSLRGDSIKQEEAQKSFVDS WRORSNTQGSILINLLLKQPLIPGSS LSLCHLLSSSESAPAGTLPQPPGFGS TLAGMSGLRTTGSYDGSFSLSALRE AQNLAFGLNLVARNECSRDPGPAE GGRAFPPLCQLPGAVHFLPLVQFFIG LHCQALQDLAAAKRSGAPGDSPTHS SCVSSGVETNPEDSVCILEGFSVTAL SILQHLVCHSGAVVSLLSGVGADSA AGEGNRSLVHRLSDGDMTSALRGV ADDQGHPLLKMLLHLLAFSSAATG HLQASVLTQCLKVLVLAENTSCDF LPRFQCVFQVLPKCLSPETPLPSVLL AVELLSLLADHDQLAPQLCSHSEGC LLLLLYMYITSRPDRVALETQWLQLE QEVVWLLAKLGVQSPPLPVTGSNCQ CNVEVVRLTVMHHRQWLTVRRAG GPPRTDQRRITVRCLRDVTVLLHGL SQDKLFMMHCVEVLHQFDQVMP GVSMIRGLPDVTDCEEAAALDDLCA AETDVEDPEVECG	True	False	2.656	4.747	1.509	0.915	0.685	1.05	1.