

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	extracellular region
Q96F46	I17RA_HUMAN	Homo sapiens	Interleukin-17 receptor A	19.553086	T69	S708;S736	35289036;38665916	MGAARSPPSAVPCPLGLLLLLLGVL APGGASLRLLDHRALVCSQPGLNCT VKNSTCLDDSWIHPRNLTPSSPKDL QIQLHFAHTQGGDLFPVAHIEWTLQ TDASILYLEGAELSVLQLNTERLCV RFEFLSKLRHHHRRRWRFTFSHFVV DPDQEYEVTVHHLKPIPDGDPNHQ SKNFLVPDCEHARMKVTTPCMSSG SLWDPNITVETLEAHQLRVSFILWN ESTHYQILLTSPHMHNSCFEHEM HHIPAPRPEEFHQRSNVTLTLRNLK GCCRHQVQIQPFSSCLNDCLRHSA TVSCPEMPDTPPEPIPDYMLVWYWF ITGISILLVGSVILLIVCMTWRLAGPG SEKYSDDTKYTDGLPAADLIPPLKP RKVWHIYSADHPLYVDVVLKFAQFLL TACGTEVALDLEEQAISEAGVMTW VGRQKQEMVESNSKIIVLCSRGTRA KWQALLGRGAPVRLRCDHKGKPVGD LFTAAMNMILPDFKRFACFGTYVVC YFSEVSCDGDVPDFGAAPRYPLMD RFEVYFRIQDLEMFQPMHRMVRG ELSGDNYLRSPPGGRQLRAALDRFRD WQVRCPDWFECENLYSADDQDAPS LDEEVFEPLPFGTGIVKRAPLVRE PGSQACLAIDPLVGEEGGAAVAKLEP HLQPRGQAPQPLHTLVLAEEGAL VAAVEPGPLADGAAVRLALAGEGEA CPLLGSPGAGRNSVLFPLVDPEDSPL GSSTPMASPDLLPEDVREHLEGLML SLFEQSLSCQAQGGCSRPMVLTDP HTPYEEQRQSVQSDQGYISRSSPQ PPEGLTEMEEEEEEEQDPGKPALPL SPEDLESLSLQRQLLFRQLQKNSG WDTMGSESEGPSA	False	True	3.033	3.384	1.928	2.197	0.951	4.7	4.476