

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
Q60943	I17RA_MOUSE	Mus musculus	Interleukin-17 receptor A	25.685855	NaN	S736	39627609	MAIRRCWPRVVPGPALGWLLLLNV LAPGRASPRLLDFPAPVCAQEGLS RVKNSTCLDDSWIHPKNLTPSSPKN IYINLSVSSSTQHGEIVLVHVEWTLQ TDASILYLEGAELSVLQLNTNERLCV KFQFLSMLQHHRKRWRFSFHFV DPGQYEYEVTVHHLPKPIPDGDPNHK SKIIFVPDCEDSKMKMTTSCVSSGSL WDPNITVETLDTQHRLRVDFTLWNES TPYQVLLSFSDSENHSCFDVVKQIF APRQEEFHQRANVTFTLSKFHWCC HHHVQVQPFSSCLNDCLRHAVTV PCPVISNTTVPKPVADYIPLWVYGLIT LIAILLVGSVIVLIICMTWRLSGADQE KHGDDSKINGILPVADLTPPPLRPRK VWIVYSADHPLYVEVVLKFAQFLITA CGTEVALDLLEEQVISEVGVMTWVS ROKQEMVESNSKIIILCSRGTQAKW KAILGWAEPVQLRCDHWKPAAGDLF TAAMNMILPDFKRPACFGTYVVCYF SGICSERDVPDLFNITSRYPLMDRFE EYVFRIQDLEMFEPGRMHVRELTG DNYLQSPSGRQLKEAVLRFQEWQT QCPDWFERENLCLADGQDLPSLDE EVFEDPLLPPGGIVKQQLVRELPS DGCLVVDVCVSEESRMAKLDPOL WPQRELVAHTLQSMVLPAEQVPAA HVVEPLHLPDGSAAAQLPMTEDSE ACPLLGVQRNSILCLPVDSDDLPLCS TPMMSPDHLQGDAREQLESMLLSV LQQLSGQPLESWPREVVLEGCTP SEEEQRQSVQSDQGYISRSSQPPE WLTETEELELGEVPEVSLSPPELRLR KLQRQLFFWELEKNPGWNSLEPRR PTPEEQNPS	False	True	2.35	2.804	2.093	1.787	1.043	5.0	3.377