

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membra
Q61818	RAI1_MOUSE	Mus musculus	Retinoic acid-induced protein 1	37.364315	NaN	S322;S328;T455;S551;S666;T679;S863;S875;S1047;T1051;S1105;S1332;S1343;S1400	22517741;28135057;34678516	MOSFRERCGFHGKQNYPTQSQET SRLENYRQPGQAGLSCDRQLLAKD YYSQPQYTYEGGDTGTPSGTVATAAA DKYHRGSKSLQGRPAFSPSYVQDSSP YPGRYSGEEGLQTVGGPQPPPPQPO PLPGAVSKYEENLMKKTIVVPPNRO YPEQGQPLPFRTHSLHVPVPPQPOQ LAYPKLORQKQNDLASPLPFPQGS HFPQHSQSFPSTSYAFTVQGGGQG AHSYKSC TAPSAQPHDRPMSANAN LAPGQRVONLHAYQPGRGLGYEQQQ QALQGRHHTQETLHYQNLAKYQHY GQQGGYCPDPTAVRTPEQYYQTF PSSSHSPARSVGRSPSYSTPSPLMP NLENFPYSQQPLSTGAFFTGITDHS HFMPLLNPSPTDAASSVDPOAGNCK PLQKEKLPDNLSDLSLQSLTALTSQ VENISNTVQQLLSKATMPQKKGVK NLVSRTPQEQHKSQHCSPEGSYSAE PAGTLPSEPPSSTPQSTHAEPODDY LSGSEDPLEERSFLYCSQARGSPARV NSNSKAKPEVSVTCSVTSRDDMSTK SDDSFQSLHSTLPLDSFSKFAVGER DCPRLLSALAQEDLASEILGLQEA VEKADKAWAEASSLPKDNKPPFSL ENHGACLDVAKTSWSQGPETPLP EPLQLDKGGSTKDFSPGLFEDPSVA FATTDPKTSSPLSFGTKPLLTATP DPTTAAFDCAFDPDTPASSVDGANPF AWPEENLGDACPRWGLHPGELTKG LEQGAKASDGVGKADAHEASACMG FQEDHAIGKPAALSGDFKQQAEG VKEEVGLLQCEVAKANQWLEES RHCCSSTDFGDLPLPPPPGRKEDLE AEEYSSELCELLGSPQRPQLQDPLS PKAPLMCTKEEAEEALDTKAGWVSP CHLSGEPVLLGSPVGAQSKVQSWF ESSLSHMKPGEEGPEMERAPGSSG TSQGS LAPKPNKPAVPEGPIAKKEPV PRGKSLRSRRVHRGLPEAEDSPCRV PALPKDLLLPECTGPPQQAEGAG APGRGLSEGLPRMCTRSALTALSEPQ TPGPPGLTTTTPPPDKLGGKQRAAFK SGKRVGKPSPKAASSPSNPAALPVAS DSSPMGSKTKEPDSMSPMGKDORS MVLRSRTKPPQVVFHAKRRRPSERI PDCRATKLLPANNHLPAYKVVSGP QKEGRMNQVRVVKPGTGKLSDR PLHTLKRKSAFMAPVPAKRSILIRS NNGSGDGREERAESSPGLLRMA SPQRARPRGSGEPPLPPPLEPPAAC MGLSTQSSLPASVARTKVLPRKGRG LKLEAIVQKITSPGLKLLACRVAGAP PGTPRSPALPERRPGGSPAGAEGL GGMGQMLPAASGADPLCRNPASRS LKGKLLNSKLLSSAADCPKAEAFMS PETLPSLGTARAPKKRSRGRGTGLG PSKGGLEKRPCPGQPLLLAPHDRAS STQGGGEDNSSGGGKPKTEELGPA SQPPEGRPCQPOTRAQKOPGQASYS SYSKRKRLSRGRGKTAHASPCKGRA TRRRQQVPLDPAEPEIRLKYISSC KRLRADSRTPAFSPFVRVEKRDAYTT ICTVNSPGDEPKHWPSSSAASS STSSSLEPAGASLTTFPGGSVLQQR PSLPLSSTMHLGPVVSALSTSLVC CLCQNPANFKDLGDLGPPYPEHCL PKKPKLKEKARLEGTLEASLPLER TLKGLECSASTAAAPTATITPTAL GRLSRPDGPADPAKQGPLRTSARGL SRRLQSCYCCDGGDGGEEVAQAD KSRKHECSKEAPTEPGGDTQEHVW HEACAVWTSGVYLVAGKLFGLQEA MKVAVDMPCTSCHEPGATISCSYKG CIHTYHYPCANDTGCTFIENFTLKC PKHKRLPL	None	None	None	None	None	None	None	