

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q7Z5J4	RAI1_HUMAN	Homo sapiens	Retinoic acid-induced protein 1	12.580955	T330;S343;S345;S1772	S339;S345;T472;S568;S683;T696;S805;S880;S892;S1064;T1068;S1122;S1352;S1358;S1374;S1431	34019948;35132862;35138101;33214551;35289036;37340703;23301498;31492838;32574038;30379171	MQSFRERCGFH GKQ QNYQ QTSQET SRLENYRQPSQAGLSCDRQLLAKD YYNPQYPYSYEGGAGTPSGTAAAVAA DKYHRGSKALPTQQGLQGRPAFPGY GVQDSSPYPGRYAGEESLQAWGAPQ PPPPQPQLPAGVAKYDENLMKTA VPPSRQYAEQGAQVPFRTHSLHVQQ PPPPQQPLAYPKLQRQKLQNDIASPL FFPQGTHTFPQHSQSFPSTSYSSSV QGGGQGAHSYKSC TAPTAQPHDRPL TASSSLAPGQRVQNLHAYQSGRLSY DQQQQQQQQQQQQQQALQSRHHA QETLHYQNLAKYQHYGQQGGQGYCQ PDAAVRTPEQYYQTFSPSSSHSPARS VGRSPSYSTPSPLMPNLENFPYSQ QPLSTGAFPAGITDHSHFMPLLNPS PTDATSSVDTQAGNCKPLQKDKLPE NLLSDLSLQSLTALTSQVENISNTVQ QLLLSKAAVPQKKGKVLVSRTEQ HKSQHCSPEGSGYSAEPAGTPLSEP PSSTPQSTHAEPQEADYLSGSEDPLE RSFLYCNQARGSPARVNSNSKAKPE SVSTCSVTSPDDMSTKSDSFQSLH GSLPLDSFSK FVAGERDCPRLLLSAL AQEDLASEILGLQEAI GEKADKAWA EAPSLVKDSSKPPFSLENHSACLDS VAKSAWPRPGEPEALPDSLQLDKGG NAKDFSPGLFEDPSVAFATPDPKTT GPLSFGTKPTLGVPAPDPTTAAFDCE PDTTAAASSADSANPFAWPEENLGDA CPRWGLHPGELTKGLEQGGKASDGI SKGDTHEASACLGFQEEDPPGEKVA SLPGDFKQEEVGGVKEEAGGLLQCP EVAKADRWLED SRHCCSTADFGDL PLLPTSRKEDLEAEEYSSLCCELLG SPEQRPGMQDPLSPKAPLIC TKEEV EEVLDSKAGWGPCHLSGESVILLG PTVGTESKVQSWFESSLSHMKPGEE GPDGERAPGDSTTSDASLAQKPNKP AVPEAPIAKKEPVPRGKSLRRRVHR GLPEAEDSPCRAPVLPKDLLLPESCT GPPQGQMEGAGAPGRGASEGLPRM CTRSLTALSEPRTPGPPGLTTTPAPP DKLGKQRAAFKSGKRVGKPSPKAA SSPSNPAALPVASDSSPMGSKTKET DSPSTPGKDQRSMLRSRTKTQEIFH SKRRRPSEGRLPNCRATKLLDNSH LPATFKVSSSPQKEGRVSRARVPKP GAGSKLSDRPLHALKRKSAFMAPVP TKKRNVLRSRSSSSSNASGNGGDG KEERPEGSPTLFRMSSPKKAKPTK GNGEPATKLPPEPDA CLKLASRAA FQGAMTKVLP PPRKGRGLKLEAIVQ KITSPSLKKFACKAPGASPGNPLSPS LSDKDRGLKGAGGSPVGV EGLVNV GTGQKLPTSGADPLCRNPTNRS LKG KLMNSKLSSTDCFKTEAFTSPEAL

								QPGGTALAPKKRSRKGRAGAHGLSK GPLEKRPYLGALLLTPRDRASGTQG ASEDNSSGGGGKKPKMEELGLASQP PEGRPCQPQTRAQKQPGHTNYSSYS KRRKRLTRGRAKNTTSSPCKGRAKRR RQQQVLPLDPAEPEIRLKYISSCKRL RSDSRTPAFSPFVVRVEKRDFTTICT VVNSPGDAPKPHRKPSSSSSSSSSS SFLSLLAAGASLALPLGGGILQPRPSL PLSSTMHLGPVVSKALSTSLVCCL CQNPNANFKDLGDLGCGPYYPEHCLPK KKPKLKEKVRPEGTCEEASLPLERTL KGPECAAAATAGKPPRPDGPADPAK QGPLRTSARGLSRRLQSCYCCDGRE DGGEEAAPADKGRKHECSKEAPAEP GGEAQEHVWEACAVWTGGVYLVA GKLFGLQEAMKVAVDMMCSQCQA GATIGCCCHKGLHTYHYPCASDAGC IFIEENFSLKCPKHRLP
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