

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
Q01974	ROR2_HUMAN	Homo sapiens	Tyrosine-protein kinase transmembrane receptor ROR2	23.451874	NaN	T646	29485866;38665916	MARGSALPRRPLLCPAVWAAAALL SVSRTSGEVEVLDPNDPLGPLDGDQD GPIPTLKGYFLNFLEPVNNITIVQGG TAILHCKVAGNPPPNVRWLKNDAPV VQEPRIIRKTEYGSRLRIQDLDTTD TGYQC VATNGMKTITATGVLFVRL GPTHSPNHNFDYHEDGFCQPYPYR GIACARFIGNRRTIYVDSLQMQGEIEN RITAAFTMIGTSTHLSDOCSQFAIPS FCHFVFLCDARSRTPKPRELRCRDE CEVLESDLCRQEYTIARSNPLILMRL QLPKCEALPMPESDAANCMRIGIP AERLGRYHQCYNGSGMDYRGTAST TKSGHOCQPWALQHPHSHHLSSTD FPELGGGHAYCRNPGGQMEGPWCF TQNKNVRMELCDVPSCSPRDSSKM GILYLVP SIAIPLVIACLFFLVCMCRN KQKASASTPQRRLMASPSQDMEM PLINQHKQAKLKEISLAVRFMEELG EDRFGKVYKGLFGPAPGEQTQAVA IKTLKDKAEGPLREEFRHEAMLRAR LQHPNVVCLLGVVTKDQPLSMIFSY CSHGDLHEFLVMRSPHSDVGSTDD DRTVKSALPPDFVHLVAQIAAGME YLSSHVVHKDLATRNVLYDKLNV KISDLGLFREYVADYKLLGNLLPI RWMAP EAIMYGKFSIDSDIWSYGVV LWEVFSYGLQPYCGYSNQDVVEMIR NRQVLPCPDDCPAWVYALMIECWN EFPSSRRPRFKDIHSRLRAWGNLSNY NSSAQTSGASNTTQTSSLSTSPVSN VSNARYVGPQKAPPPQPFIPMK GQIRPMVPPQLYVPVNGYQVPVAY GAYLPNFYVQIPMQMAPQVPPQ MVPKSSHHSGSGSTSTGYVTTAPS NTSMDRAALLSEGADDTQNAPEL GAQSTVQEAEEEEEGSVPETELLGD CDTLQVDEAQVQLEA	False	True	1.177	1.895	1.056	1.659	0.675	4.635	1.933