

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extr. regi
P43405	KSYK_HUMAN	Homo sapiens	Tyrosine-protein kinase SYK	20.661177	T117	T28;S44;T47;T131;S202;T256;S295;T296;S297;S316;T317;S319;T323;T345;T348;S350;T352;T364;S379;T384;T484;T507;T525;T526;T530;T546;S579;T582;T629;T630;T631	30379171;29485866	MASSGMADSANHLPPFFGNITREEA EDYLVQGGMSDGLYLLROSRNYLG GFALSAVHGRKAHHYTIERELNGTY AIAGGRTHASPADLCHYHSQESDGL VCLLKKPFNRPGVQPKTGPFDLK ENLIREYVKQWNLQGGALEQAIHSQ KPQLEKLIATTAHEKMPWFHGKISR EESQIVLIGSKTNGKFLIRARDNNG SYALCLLHEGKVLHYRIDKDKTGKLS IPEGKKFDTLWQLVEHYSYKADGLL RVLTVPCQKIGTQGNVNFVGGRRQLP GSHPATWSAGGIISRIKSYSPKPGH RKSSPAQGNRQESTVSNFNYEPELA PWADKGFQREALPMDTEVYESPYA DPEEIRPKYEVLDKLLTLEDKELGS GNFCTVKKGYQMKVVKTVAVKIL KNEANDPALKDELLAEANVMQQLD NPYIVRMIGICEAESWMLVMEMAEL GPLNKYLQNRHVVDKNIHIELVHGV SMGMKYLEESNFVHRDLAARNVLL VTQHYAKISDFGLSKALRADENYYKA QTHGKWPVKWYAPWCINYYKFSSKS DVWSFGVLMWEAFSYGQKPYRGMK GSEVTAMLEKGERMGCPAGCPREM YDLMNLCWITYDVENRPGFAAVELR LRNYYYDVVN	True	True	4.787	5.0	2.567	2.379	1.898	5.0	2.95