

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
Q14289	FAK2_HUMAN	Homo sapiens	Protein-tyrosine kinase 2-beta	18.379595	NaN	S361;S375;S399;T402;T579;T580;T722;S762;T765;T819;T834;S839;T842;T849;S866;T881	30379171	MSGVSEPLSRVKLGLRRPEGPAEP MVVVVPDVEKEDVRLKVCFYNSNF NPGKNFKLVKCTVQTEIREIITSILLS GRIGPNIRLAECYGLRLKHKMSDEIH WLPQMTVGEVQDKYECLHVEAEW RYDLQIRYLPEDFMESLKEDRTTLLY FYQQLRNDYMQRYASKVSEGMALQ LGCELRFRFFKDMPHNALDKKSNF ELLEKEVGLDLFFPKQMENLKPKQ FRKMIQQTFFQYASLREEECVMKFF NTLAGFANIDQETRYRCELIQGWNTIV DLVIGPKGIRQLTSQDAKPTCLAEFK QIRSIRCLPLEEGQAVLQLGIEGAPQ ALSIKTSSLAEAEENMADLIDGYCRLQ GEHQGSLIHPKDGKRNSLPQIP MLNLEARRSHLSESCSIESDIYAEIP DETLRRPGGPQYGIAREDDVVLNRILG EGFFGEVYEGVYTNHKGEKINVAVK TCKKDCITLDNKEKFMSEAVIMKNL DHPHIVKLGIEEPTWIIMELYPYG ELGHYLERNKNSLKVLTLYLSLQIC KAMAYLESINCVHRDIAVRNIVASP ECVKLGDFGLSRYIEDEDYKASVTR LPIKWMSPESINFRFRFTASDVWWMF AVCMWEILSFGKQPPFWLENKDVIG VLEKGDRLPKPDLCPVLYTLMTRC WDYDPSDRPRFTLVCSLSDVYQME KDIAMEQERNARYRTPKILEPTAFQE PPPKPSRPKYRPPQTNLLAPKLQFQ VPEGLCASSPTLTSPMEYSPVNSLH TPPLHRHNVFKRHSMREEDFIQPS REEAQQLWEAEKVMRQILDKQKQK QMVEDYQWLRQEESLDPVMVYN DKSPLTPEKEVGYLEFTGPPQKPPRL GAQSIQPTANLDRDLDLVYLVNVMEL VRAVLELKNELCQLPPEGYVVVKN VGLTLRKLIGSVDDLPLSPSSRTEI EGTQKLLNKDLAELINKMRLAQQNA VTSLSECKRQMLTASHTLAVDAKN LLDAVDQAKVLANLAHPAE	None	None	None	None	None	None	None		

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