

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	ext. reg.
P43403	ZAP70_HUMAN	Homo sapiens	Tyrosine-protein kinase ZAP-70	29.956363	S497	T248;S289;T292;T315;T319;T492;T493	27655845;29351928	MPDPA AHL PFFYGSISR AEAEEHLK LAGMADGLFLLRQCLRSLGGYVLSL VHDVRFHHFPIERQLNGTYAIAAGGK AHC GPAELCEFYSRDPDGLPCNLRK PCNRPSGLEPQPGVDFDCLRDAMVR DYVRQ TWKLEGEALEQAHSQAPQVE KLIATTAHERMPWYHSSLTREEAER KLYSGAQTDGKFLLRPRKEQGTAL SLIYGKTVYHYLISQDKAGKYCIPEGT KFDTLWQLVEYLKADGLIYCLKEA CPNSSASNASGAAAPTLP AHPSTLT HPQRRIDTLNSDGYTPEPARITSPDK PRPMPMDTSVYESPYS DPEELKDKK LFLKRDNLLIADIELGCGNFGSVRQG VYRMRKKQIDVAIKVLKQGT E KADT EEMMREAQIMHQLDNPYIVRLIGVC QAEALMLVMEMAGGGPLHKFLV GK REEIPVSNVAELLHQVSMGMKYLEE KNFVHRDLAARNVLLVNRHYAKISD FGLSKALGADDSYYTARSAGKWPLK WYAP E C I N F R K F S S R S D V W S Y G V T M WEALSYGQKPYKKMKGPEVMAFIE QGKRMECPPECPPELYALMSDCWIY KWEDRPDFTLVEQRM RACYSLASK VEGPPGSTQKAEAAACA	True	True	4.68	3.327	2.09	2.152	1.826	5.0	2.7