	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q6XUX3	DUSTY_HUMAN	Homo sapiens	Dual serine/threonine and tyrosine protein kinase	29.088996	S490	NaN	30620550;29485866	MEGDGVPWGSEPVSGPGPGGGGMI RELCRGFGRYRRYLGRLRQNLRETQ KFFRDIKCSHNHTCLSSLTGGGGAE RGPAGDVAETGLQAGQLSCISFPPKE EKYLQQIVDCLPCILILGQDCNVKCQ LLNLLLGVQVLPTTKLGSEESCKLRR LRFTYGTQTRVSLALPGQYELVHTLV AHQGNWETIPEEDLEVQENNEDAA HVLAELEVTMHHALLQEVDVVVAPC QGLRPTVDVLGDLVNDFLPVITYALH KDELSERDEQELQEIRKYFSFPVFFF KVPKLGSEIIDSSTRRMESERSPLYR QLIDLGYLSSSHWNCGAPGQDTKAQ SMLVEQSEKLRHLSTFSHQVLQTRL VDAAKALNLVHCHCLDIFINQAFDM QRDLQITPKRLEYTRKKENELYESL MNIANRKQEEMKDMIVETLNTMKE ELLDDATNMEFKDVIVPENGEPVGT REIKCCIRQIQELIISRLNQAVANKLI SSVDYLRESFVGTLERCLQSLEKSQ DVSVHITSNYLKQILNAAYHVEVTFH SGSSVTRMLWEQIKQIIQRITWVSPP AITLEWKRKVAQEAIESLSASKLAKSI CSQFRTRLNSSHEAFAASLRQLEAG HSGRLEKTEDLWLRVRKDHAPRLAR LSLESCSLQDVLLHRKPKLGQELGR GQYGVVYLCDNWGGHFPCALKSVV PPDEKHWNDLALEFHYMRSLPKHE RLVDLHGSVIDYNYGGGSSIAVLLIM ERLHRDLYTGLKAGLTLETRLQIALD VVEGIRFLHSQGLVHRDIKLKNVLLD KQNRAKITDLGFCKPEAMMSGSIVG TPIHMAPELFTGKYDNSVDVYAFGIL FWYICSGSVKLPEAFERCASKDHLW NNVRRGARPERLPVFDEECWQLME ACWDGDPLKRPLLGIVQPMLQGIM NRLCKSNSEQPNRGLDDST	None	None	None	None	None	None	None	None	None