

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
Q6P5F6	S39AA_MOUSE	Mus musculus	Zinc transporter ZIP10	24.474581	NaN	T538;T555;S593	33300544	MKVHIHTKFCLICLLTFIFHHCNHC HEDHDHGPEELHRHHRGMTESESS KFSVQDAENEKKYIEKLFDRYGEN GRLSFFGLEKLLTNLGLGEIKVVEIN HEDLGHHDVSHLDILAVQEGKHFH SHTHQHFHNHLNAENHTTTSVTSK RNHKCDPEKEAAELPIKADDKHLHD RNHRFHHRHRLHHHLDHNTTRHV HNSVAHSEHGEPGHSPSPETNKT QEQSEVKSVKVRRKEKGRKKENSE VNTPGFLPNHDHSEQYEHNRVHKL DRVHSPGHPHAHLPEHSGHELGHG HOELDPDNEGELRHTRKREAPHVR KSAIYTPSHKDQSEDDROHECLNV TQLLKHFGLGPNSPISPDLFYLCPA LLYQIDSRLCIEHFDKLLVEDLNKDK TLVPEDKTNIGASAWICGHSITVISLL SLLGVILVPIINQGCFFKLLTFLVALA VGTMSGDALLHLLPHSQGGHDHSH QHTHGHGSHGHEKFELEYDAV LKGLVALGGIYLLFIIIEHCIRMFKHYK QQRGKQWFMKQSTEESTIGRKLK DHKLNSTPDADWLQKPLAGTDDS VVEDRLNETELTDLEAQQESPPKN YLGVEEEKIMDHSUSDGLHTIHEHE VHVTSHNHHDEDKAVLRKHSHQW HHRHAHSHGPGCHSGSDLKETGIA NIAWMVMGDGIHNFSDGLAIGAAF SAGLTGGISTSIAVFCHLPHLGDGDF AVLLKAGMTVKQAIVYNLLSAMMAY IGMLIGTAVGQYANNITLWIFAITAG MFLYVALVDMLEPEMLHGDGDHEEH GFCPVGQFLLQNLGGLLFGFAIMLVIA LYEDKIVFDIQF	False	True	1.838	1.747	1.474	1.61	1.68	5.0	1.828