

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	extra regio
Q9EQH3	VPS35_MOUSE	Mus musculus	Vacuolar protein sorting-associated protein 35	26.459305	NaN	S7;S783;T791	34887587;33300544;37507081;37453647;28528544	MPTTQQSPQDEQEKLLEDAIQAVKV QSFQMKRCLDKNKLMDALKHASN MLGELRTSMLSPKSYELYMAISDE LHYLEVYLTDEFAGRKRKADLYELV QYAGNIIPRLYLLITVGVVYKSFQPS RKDILKDLVEMCRGVQHPLRGLFLR NYLLQCTRNILPDEGEPTDEETGDI SDSMDFVLLNFAEMNKLWVRMQH QGHSRDREKRERERQELRILVGTNL VRLSQLEGVNVVERYKQIVLTGILEQV VNCRDALAQEYLMECHQVFPDEFH LQTLNPFRLRACAEHLQNVNVKNIIIA LIDRLALFAHREDGPGIPAEIKLFDIF SQQVATVIQSRQDMPSEDVVSLQVS LINLAMKCYPRVDYVDKVLTTVEI FNKLNLEHIATSSAVSKELTRLLKIP VDTYNNILTVLKLFHPLFEYFDYE SRKSMSCYVLSNVLDYNTIIVSQDQ VDSIMNLVSTLIQDQDPQVEDPDP EDFADEQSLVGRFIHLRSDDPDQDQ YLILNTARKHFGAGGNQIRFTLPPL VFAAYQLAFRYKENSQMDDKWEKK CQKIFSFAHQTSALIKAEELPLRL FLQGALAAAGEIGFENHETVAYEFMS QAFSLYEDEISDSKAQLAAITLIIGTF ERMKCFSEENHEPLRTQCALAASKL LKKPDQGRAVSTCAHLFWSGRNTD KNGEELHGGKRVMECLKKALKIAN QCMDPSLQVQLFIEILNRYIFYEKE NDAVTIQVLNQLIQKIREDLPNLESS EETEIQINKHFHNTLEHLRSRESPE SEGPIYEGLLI	True	False	4.705	2.675	3.404	2.049	2.818	2.392	1.997