	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plas mem
0	Q6PFD5	DLGP3_MOUSE	Mus musculus	Disks large-associated protein 3	40.905401	NaN	S58;S404;S407;S410;S414;S641;S643;S930;S933;S965	22517741;34678516;22645316	MRGYHGDRGSHPRPARFADQQHM DVGPAARAPYLLGSREAFSTEPRFCA PRAGLGHLSPEGPLSLSEGPSSVGPE GGPGGVGAGGGSSTFPRMYPGQGP FDTCEDCVGHPQGKGATRLPPTLLD QFEKQLPVQQDGFHTLPYQRGPAGP GPGPGSGAAPEARSESPSRIRHLVHS VQKLFAKSHSLEAPGKRDYNGPKAD GRGSSGGDSYSGPGSGGTPTSHHH HHHHHHHHHQSRHGKRSKSKDRK GDGRHQTKATGWWSSDDNLDSDS GFLGGRPPGEPGGPFCLDAPDGSYR DLSFKGRSGGSEGRCLACTGMSMS LDGQSVKRSAWHTMMVSQGRDGYP GAGPGKGLLGPETKAKARTYHYLQV PQDDWGGYPTGGKDGEIPCRRMRS GSYIKAMGDEESGDSDGSPKTSPKA LARRFASRRSSSVDTARINCCVPPRI HPRSSIPGYSRSLTTGQLSEEFNQQL EAVCGSVFGELESQAVDALDLPGCF RMRSHSYLRAIQAGCSQDDDCLPLL AAPASVSGRPGSSFNFRKAPPPIPPG SQAPPRISITAQSSTDSAHESFTAAE GPARRCSSADGLDGPTMGARTLELA PVPPRASPKPPTLIIKTIPGREELRSL ARQRKWRPSIGVQVETISDSDTENR SRREFHSIGVQVEEDKRRARFKRSN SVTAGVQADLELEGLAGLATVATED KALQFGRSFQRHASEPQPGPRAPTY SVFRTVHTQGQWAYREGYPLPYEPP ATDGSPGPTPVPAPGPGSGRRDSWM ERGSRSLPDSGRTSPCPRDGEWFIK MLRAEVEKLEHWCQQMEREAEDYE LPEEILEKIRSAVGSTQLLLSQKVQQ FFRLCQQSLDPTAFPVPTFQDLAGF WDLLQLSIEDVTLKFLELQQLKANS WKLLEPKEEKKVPPPIPKKPSRGRGV PVKERSLDSVDRQRQEARKRLLAAK RAASFRHSSATESADSIEIYIPEAQTR L	None	None	None	None	None	None	None	Non