	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	Q9Y2H0	DLGP4_HUMAN	Homo sapiens	Disks large-associated protein 4	26.386321	T547	S206;S207;S378;S381;S388;S405;S415;S421;S580;S581;S609;S611;S665;S744;T915;S973	28411811;28657654;37217939	MKGLGDSRPRHLSDSLDPPHEPLFA GTDRNPYLLSPTEAFAREARFPGQN TLPGDGLFPLNNQLPPPSSTFPRIHY NSHFEVPEESPFPSHAQATKINRLPA NLLDQFEKQLPIHRDGFSTLQFPRG EAKARGESPGRIRHLVHSVQRLFFT KAPSLEGTAGKVGGNGSKKGGMED GKGRRAKSKERAKAGEPKRRSRSNI SGWWSSDDNLDGEAGAFRSSGPAS GLMTLGRQAERSQPRYFMHAYNTIS GHMLKTTKNNTTELTAPPPPPAPPA TCPSLGVGTDTNYVKRGSWSTLTLS HAHEVCQKTSATLDKSLLKSKSCHQ GLAYHYLQVPGGGGEWSTTLLSPRE TDAAAEGPIPCRRMRSGSYIKAMGD EDSDESGGSPKPSPKTAARRQSYLR ATQQSLGEQSNPRRSLDRLDSVDML LPSKCPSWEEDYTPVSDSLNDSSCIS QIFGQASLIPQLFGHEQQVREAELSD QYEAACESACSEAESTAAETLDLPLP SYFRSRSHSYLRAIQAGCSQEEDSVS LQSLSPPPSTGSLSNSRTLPSSSCLV AYKKTPPPVPPRTTSKPFISVTVQSST ESAQDTYLDSQDHKSEVTSQSGLSN SSDSLDSSTRPPSVTRGGVAPAPEAP EPPPKHAALKSEQGTLTSSESHPEA APKRKLSSIGIQVDCIQPVPKEEPSPA TKFQSIGVQVEDDWRSSVPSHSMSS RRDTDSDTQDANDSSCKSSERSLPD CTPHPNSISIDAGPRQAPKIAQIKRN LSYGDNSDPALEASSLPPPDPWLET SSSSPAEPAQPGACRRDGYWFLKLL QAETERLEGWCCQMDKETKENNLS EEVLGKVLSAVGSAQLLMSQKFQQF RGLCEQNLNPDANPRPTAQDLAGF WDLLQLSIEDISMKFDELYHLKANS WQLVETPEKRKEEKKPPPPVPKKPA KSKPAVSRDKASDASDKQRQEARKR LLAAKRAASVRQNSATESADSIEIYV PEAQTRL	None	None	None	None	None	None	None	Non