	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	ex re
0	Q15334	L2GL1_HUMAN	Homo sapiens	Lethal(2) giant larvae protein homolog 1	19.981646	S123	S663;T958;S967;S985	30379171;23301498;34725712;29351928	MMKFRFRRQGADPQREKLKQELFA FNKTVEHGFPNQPSALAFDPELRIM AIGTRSGAVKIYGAPGVEFTGLHRDA ATVTQMHFLTGQGRLLSLLDDSSLH LWEIVHHNGCAHLEEALSFQLPSRP GFDGASAPLSLTRVTVVLLVAASDIA ALGTEGSSVFFLDVTTLTLLEGQTLA PGEVLRSVPDDYRCGKALGPVESLQ GHLRDPTKILIGYSRGLLVIWNQASQ CVDHIFLGNQQLESLCWGRDSSTVV SSHSDGSYAVWSVDAGSFPTLQPTV ATTPYGPFPCKAINKILWRNCESGG HFIIFSGGMPRASYGDRHCVSVLRA ETLVTLDFTSRIIDFFTVHSTRPEDEF DDPQALAVLLEEELVVLDLQTPGWP AVPAPYLAPLHSSAITCSAHVASVPA KLWARIVSAGEQQSPQPVSSALSWPI TGGRNLAQEPSQRGLLLTGHEDGTV RFWDASGVALRPLYKLSTAGLFQTD CEHADSLAQAAEDDWPPFRKVGCF DPYSDDPRLGVQKVALCKYTAQMVV AGTAGQVLVLELSDVPVEQAVSVAII DLLQDREGFTWKGHERLSPRTGPLP WPAGFQPRVLVQCLPPAAVTAVTLH TEWSLVAFGTSHGFGLFDYQRKSPV LARCTLHPNDSLAMEGPLSRVKSLK KSLRQSFRRIRKSRVSGKKRAANASS KLQEANAQLAEQACPHDVEMTPVQ RRIEPRSADDSLSGVVRCLYFADTFL RDGAHHGPTMWAGTNSGSVFAYAL EVPAAAVGGEKRPEQAVEAVLGKEV QLMHRAPVVAIAVLDGRGRPLPEPY EASRDLAQAPDMQGGHAVLIASEEQ FKVFTLPKVSAKTKFKLTAHEGCRV RKVALATFASVACEDYAETCLACLTN LGDVHVFSVPGLRPQVHYSCIRKEDI SGIASCVFTRHGQGFYLISPSEFERF SLSARNITEPLCSLDINWPRDATQAS YRIRESPKLSQANGTPSILLAPQSLD GSPDPAHSMGPDTPEPPEAALSPMS IDSATSADTTLDTTGDVTVEDVKDFL GSSEESEKNLRNLAEDEAHACAILIK	False	False	3.152	3.173	1.579	1.357	4.183	4.193	1.