

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnacsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
F7A4A7	OTOGL_MOUSE	Mus musculus	Otogelin-like protein	24.474581	NaN	NaN	33300544	MVPWRALSLPILLVSLRQYVVCASSVL SETSESEFYENEQRALLAVQFEATS PRYFFHEAINWGESKIKGSCSHECL NGAFCSKTGTDCQIFQALGTRCQI VPNMGSGRDGICKTWGQYHFETFD GIYYFPGSCSYFAKDCGNLEPQYT VWVHNSPKCLGVSYSYRISLFFS NQEIRIYGHEIRKNGISLSLPQTLG QVHLEKVADYILVKTTFGFLAWDGI SGIYLKLEEHRGKSCGLCANYNGIQ SDDFVILQEDYTEDIAMFANSWLVL TPDDTKCVPTPSDFPNPCSSGMPAF EAIFFKCQILLQFPFLSCHEYIDPYLY IASCVNDLCKTDDDETYCRAATEYA RACSHAGFPIQDWRDDFPACTDKCD DSFVHRDCISCCPPSCTFEKQCLGS NLHCLDGCYCADGLIMDNGTCISLE SCPCSFHGLAYSVGSKIEQECTECVC VGGVWNCTEHDPCVQCSVVGDSHF TTFDGRHYSFIGLCQYILVKGTKDR FTITLQKAHCEQNLGLVCLQSITLILE DDFNKQVTLSRGGQIVTSPNQGFLL NGIVEIQTLSLIFILLRTTFGLKILFAI DGERIYIQLSSAWKRRRLGLCGTFN GNIRDDFLSPSGMIEGTPQLHAHAW RVSSTCFAPVHVPMDVPCNINQONI GYAAHCDVIHQELFAPCHVYVSPGL YYQLCRHDACKCGSPCLCNALAHYA YLCGQRGVPIDFRAHISFCAVVCQKG MLYHHCSSLCLRSTSLSSPEQCKD DCAEGCNCPEGKFYEETLNFVPIY HCRCHYRGSYIYQPGELIPTPSGLCQC SNGTVKCELATPSTVHACPEGKEY FDCRFDPALPAGGINCETTANLA MNFTCAPSSPCISGCVCAAGMAEHK GKCYVPESCPCIWKDWEYSSGEVIST PCYTCVRRGMFNCTYPCPAVCTV YGDRHYHSFDGLEDYISDCQVFLIK STDDSDISVISQNKKCFDNDIVCSKS VLISIGDTEIYLNDAFYKQKRSQFFLE SRPEYQLWKAGFYIVYFPEEDITILW DEKTTIHKVGPQWKNKLAGLCCGNF DKCTSNDMTTSNIEVRNAQVFGD SWALGQCEDLMEALKPCEAHQNKF PYAKRECSILYSDVFAPCRNVIDVTS FAKNCHEDTCNCLGGDCECLCTS VAAYAYKCCQEGVPVHWRSPVICAL DCEYYNQGLGEGPYMLASYGQSGLV LGNMTRSRSVSLPRSNRGNLFFI FMITPGLFKEKTSALVLSLESAERP NYFLYVHDNDTSLKLWRANSEFH QRATFFHHQGLWIPGYSAFELYSKK GYFIVFMGSSVKASKYDDSEEFKQS SSFSIEEIQAVVPYRRMCEWRYEPCA TPCFKTCSDPEALACTFLPPVEGCLP YCPKNMILDETTLKCVHPEDCIPLFP TEPALPPDITPSDMTPTPGLECEPQQ FDPVYNCQYICLNMEWTFYNWSL NCPKDLEMPDCGFRGWVQVNTDI CCPEWECPCRCMSLSELSIITFDGN SAALSSMASYLVRVPGIIVVHIDK CSMNQNGHALKPPASFRISGLCFK KLNVTTSIHKILINRVVRKVDVDSIVV PLPFSSHELFIEDSGTMYVITTPAGLI IKWAHLTGIIHIFGPFNLSSYTEG LCGICNDNPDDDLRMQNGTIITNME DIEFIGSWEIEKSFVTRRRPVRNC TEYDCSHCIELLNREGFIPCHDKVSP RDFCEKMWINYTYFWSYECDAISAY VALCNKFDICIQWRTPDYCLSCPEG KEYQPCVRPCEARTCLNKWFYGHSS CLNLREDCVCKNGTILHRPKTLCLIP EQECVCTDREEHPRSAGEIWNNGGID ECTLYKCLELDGNIPIEIVCEEPSPI CERTAEVWIGIVDKLTCCSKKVCVCGCD	None	None	None	None	None	None	None		

MSLCDRTIPPCTNSQKLIVGYSALSC
CPQYECECDTVRCPDISTPVCRRDDQ
FVLQVRQGEPCCFYPSCVCKTCTEP
TPQCTDGEFLTVNINTTHLCCPOYY
CVCEPDLCPPPSLECAKDMNLVKEN
VSGQCCPNWRCECNCE TLVMPTCD
VGEFAAIDQNFQTD CGCVQYLCEKD
DVCVFQEVSVLNPQQSLIKYLEEFC
YIIECLDEKDN YDFHTLNVTMVNC
SKDCDAHQIYIPSSDYDCCGTCKNI
SCKFIMENGT SVIYEEGSTWHYNCS
TYECVNTEEGATILNYSMVCPPFNE
TECKLNEGIVKLYNEGCKICKREER
ICQKVIIKSHKKQDCVSSISVASCD
GKCP SATIYNIN VESH LRFCKCCRE
NGVRNVTVPLHCSGNGTEVMYTLQ
EPIDCTCQWN