

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
F7A4A7	OTOGL_MOUSE	Mus musculus	Otogelin-like protein	9.108089	NaN	NaN	33300544	MVPWRALSLPILLVSLRGYVCASSVL SETSESEFYENEQRRALLAVQFEATS PRYFFHEAINWGESKIKGSCPHECL NGAFCSKTGTCDQCQIFQALGTRCQI VPNMGSGRDGICKTWGQYHFETFD GIYYFPGSCSYIFAKDCGNLEPQYT VWVHNSPKCLGSVYSCYRSISLFFS NQEEIRIYGHEIRKNGISLSLPQTLG QVHLEKVADYILVKTTFGFLAWDGI SGIYLLKSEEHRGKSCGLCANYNGIQ SDDFVILQEDYTEDIAMFANSWLVL TPDDTKCVPTSPDFPNPCSSGMPAF EAIFFKCQILLQFPFLSCHEYIDPYLY IASCVNDLCKTDDDETYCRAATEYA RACSHAGFPIQDWRDDFPACTDKCD DSFVHRDCISCCPPSCTFEKQCLGS NLHCLDGCYCADGLIMDNGTCISLE SCPCSFHGLAYSVGSKIEQECTECVC VGGVWNCTEHDCPVQCSVVGDSHF TTFDGRHYSFIGLCQYILVKGTKDR FTITLQKAHCEQNLGLVCLQSITLILE DDFNKQVTLRGGQIVTSPNQGFTL NGIVEIQTLSSLFILLRTTFGLKILFAI DGERIYIQLSSAWKRRTLGLCGTFN GNIRDDFLSPSGMIEGTPQLHAHAW RVSSTCFAPVHVPMPVDPCNINQQNI GYAAHCDVIHQELFAPCHVYVSPGL YYQLCRHDACKCGSPCLCNALAHYA YLCGQRGVPIDFRAHISFCAVVCQKG MLYHHCSSLCLRCTLSLSSPEQCKD DCAEGCNCPEGKFYEETLNFCVPIY HCRCHYRGSYIYQPGELIPTPSGLCQC SNGTVKCDELATPSTVHACPEGKEY FDCRFPDPALPAGGINCETTANLA MNFTCAPSSPCISGCVCAAGMAEHK GKCYVPESCPCIWKDWEYSSGEVIST PCYTCVCRGMFNCTYYPCPAVCTV YGDRHYHSFDGLEDYISDCQVFLIK STDDSDISVISQNKCFDNDIVCSKS VLISIGDTEIYLNDAFYKQKRSGFFLE SRPEYQLWKAGFYIYIYFPEEDITILW DEKTTIHIKVGQPQWKNKLAGLCGNF DKCTSNDMTTSNNIEVRNAQVFGD SWALGQCEDLMEALKPCEAHQNKF PYAKRECSILYSDVFAPCRNVIDVTS FAKNCHEDTCNCNLGGDCECLCTS VAAYAYKCCQEGVPVHWSPTVCAL DCEYYNQGLGEGPYMLASYGQSGLV LGANMTSRSVFLPRSNRGNLFFI

FMITPGLFKEKTSSLALVSLESAERP
NYFLYVHDNDTSLKLRANSEFH
QRATFFHHQGLWIPGYSAFELYSKK
GYFIVFMGSSVKASKYDDSEEFKQS
SSFSIEEIQAVVPYRRMCEWRMPECA
TPCFKTCSDPEALACTFLPPVEGCLP
YCPKNMILDETTLKCVHPEDCIPLFP
TEPALPPDITPSDMTPTPGLECEPQQ
FDPVYNCSQYICLNMEWTFYNWSL
NCPKDLEMPDCGFRGWVQVNTDI
CCPEWECPCRCMSLSELSIITFDGN
SAALSSMASYLVRVPGEIVVHIDK
CSMNQNGHALKKPASFGRISGLCFK
KLNVTTSIHKILINRVVRKVDVDSIV
PLPFSSHELFIEDSGTMYVITTPAGLI
IKWAHLTGIIHFQFNLSSYTEG
LCGICNDNPDDDLRMQNGTIITNME
DIELFIGSWEIEKSFVETMRRPVRNC
TEYDCSHCIELLNREGFIPCHDKVSP
RDFCEKMWINYTYFWSYECDAISAY
VALCNKFDICIQWRTPDYCPLSCEG
KEYQPCVRPCEARTCLNKWFYGHSS
CLNLREDCVCKNGTILHRPDKTLCIP
EQECVCTDREEHPRSAGEIWNNGID
ECTLYKLEDGNIPIEPVCEEEPSPI
CERTAEVVIGIVDKLTCCSKKVCDCD
MSLCDRTIPPCTNSQKLIVGYSALSC
CPQYECECDTVRCPDISTPVCRRDDQ
FVLQVRQGEPCCFYPSCVCKTCTEP
TPQCTDGEFLT VNINTTHLCCPQYY
CVCEPDLCPPPSLECAKDMNLVKEN
VSGQCCPNWRCECNCETLVMPTCD
VGEFAAIDQNFQTDGCGVQYLCEKD
DVCVFQEVSVLNPGQSLIKYLEEEFC
YIIECLDEKDNYTDFHTLNVMTVNC
SKDCDAHQIYIPSSSDYDCCGTCKNI
SCKFIMENGTSVIYEEGSTWHYNCS
TYECVNTEEGATILNYSMVCPPFNE
TECKLNEGIVKLYNEGCKICKREER
ICQKVIKSIKKQDCVSQSSISVASCD
GKCPSATIYNINVESHLRFCKCCRE
NGVRNVTVPLHCSGNGTEVMTLQ
EPIDCTCQWN