

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P33438	GLT_DROME	Drosophila melanogaster	Glutactin	10.212237	NaN	NaN	33925313	MKPLLLVLALCGAQVHAHSVGLRPD YNDYSEDEDTRRDWLPEPLKVPVWQ SETRYAQPQEAVVQAPEVGOILGISG HKTIANRPNVNAFLGIRYGTVGGGLAR FQAAQPIGYQGRVNATVQSPNCAQF PELDRLRLSESRGENVDDCLTLDIYA PEGANQLPVLVVFVHGEMLFDDGGSE EAQPDYVLEKDVLLVSINYRLAPFGF LSALTDELPGNVALSDLQLALEWLQ RNVVHFGGNAGQVTLVGQAGGATL AHALSLSGRAGNLFQQLILQSGTAL NPYLIDNQPLDTLSTFARLARCPPPS INPSAQGLKPLYDCLARLPTSQLVAA FEQLLLQNEHLGLTQLGGFKLVVGD PLGFLPSHPASLATNSSLALPMIIGA TKDASAFIVSRIYDQLARLQSRNVSD YIDVLRHTAPPSEHRLWKQWALRE IFTPIQEQTASLQTVAPGLLELSNYIL YRAPVINSISQSYRSVPAYLYTFDYRG EHHRFGHLSNPLPFGVDASLSDDSV YLFYPPEASRLNPLDRSLSRALVTM WVNFATTGVPNPSSGVWPQATSEY GPFLRFTNNQQSPLELDPHFGEIY LPNYRVYKPTTNFSPITTTTTTTTTT TTTTSRPYAYNPYANWQNRPSQQHP NWHPADPEYVRAQEARQQEFIRERE QRRREQQLRDQQRYPQQEPREQQD ERIRQQREQEERLRQQREQEERLRQ QRELEERIRQQQEREQYEREQQERE QREREELERQQREREQQQPEQQPE YNPEPVNPWGYPVQEPQPDDNPED GRLPYPSYEQYGPEGNENLPETDAN RNFSEEDREQQQQEQLRREQQEQQ EREYQLQLEREQQEREQQERGOQE PGPEEYPSYEEYSRALQEKNAERDRI YAEERERERQQQETLLQENQQHPE QSLPEEQPTHPNYEAYDGRSYAEE QEREQQRRDQVEQEREEQPDEDQG EEYERSPDEEEAAEQDVLKVEDFPS YEAYLEAATKLREEQEEQEKLEERY RAQQEEEDRIQAERERNSRN