

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6P5E4	UGGG1_MOUSE	Mus musculus	UDP-glucose:glycoprotein glucosyltransferase 1	16.785659	S931	S1277	30059200;34678516;34887587;28135057	MCSRGDANTADAAAARRVTGLRYN MRLLIALLALPCLFSLAEANSKAITTSL TTKWFSAPLLLLLEASEFLAEDSQEKF WSFVEATQNISSDHHDTDHSYDA VLEAAFRFLSPLQONLLKFCLSLRSY SASIQAFQQIAVDEPPPEGCKSFLSV HGKQTCDLDTLESLLTAADRPKPL LFKGDHRYPPSSNPESPVVILYSEIGH EEFSNIHHQLISKSNEGKINYVFRHY ISNPSKEPVYLSGYGVLAIKSTEYKA KDDTQVKGTEVNATVIGESDPIDEV QGFLFGKLRELYPALEGQLKEFRKH LVESTNEMAPLKVWQLQDLSFQTAA RILAASGALSLVVMKDISQNFPTKAR AITKTAVSAQLRAEVEENQKYFKGTI GLQPGDSALFINGLHIDLDTQDIFSL FDTLRNEARVMEGLHRLGIEGLSLH NILKLNIPSETDYAVDIRSPAIWV NNLEVDSRYNSWPSSLQELLRPTFP GVIRQIRKLNLMVFIIDPVHETTAE LISIAEMFLSNHIPLRIGFIFVNDSE DVDGMQDAGVAVLRAINYVAQEVD GYHAFQTLTQIYNKVRTGETVKVEH VSVLEKKYPYVEVNSILGIDSAYDQ NRKEARGYYEQTGVGPLPVVLFNGM PFEKEQLDPDELETITMHKILETTTF FQRAVYLGELSHDQDVVEYIMNQPN VVPRINSRILTAKREYDLTASNNFY VDDFARFSALDSRGKTAAIANSMNY LTKKGMSSKEIYDDSFIRPVTFWIVG DFDPSGRQLLYDAIKHQKTSNNVR ISMINNPSQEISDSSTPIFRAIWAALQ TQASSAKNFITKMAKEETAALAA GVDIAEFSVGGMDVSLFKEVFESSR MDFILSHALYCRDVLKLLKKGQRVVIS NGRIIGPLEDNELFNQDDFHLENII LKTSGQKIKSHIQQLRVEEDVASDLV MKVDALLSAQPKGEARIEYQFFEDK HSAIKLKPKEGETYYDVAVVDPVTR EAQRLAPLLLVLTLINMNLRFVFN CQSKLSDMPLKSFYRYVLEPEISFTA DSSFAKGPIAKFLDMPQSPLFTLNL NTPESWMVESVRTPYDLNIIYLEEV DSIVAAEYELEYLLEGHICYDITGQ PPRGLQFTLGTSANPTIVDTIVMANL GYFQLKANPGAWILRLRKGSRDDIY RIYSHDGTDSPPDANDVVILNFK SKIIKVKVQKKADMANEDLLSDGTN ENESGFWDSFKWGFSGQKAEVVKQ DKDDIINIFSVASGHLYERFLRIMML SVLKNTKTPVKFWFLKNYLSPTFKE FIPYMAKKYNFYELVQYKWRWLH

								QQTEKQRIIWGYKILFLDVLFPVVV KFLFVDADQIVRTDLKELRDFNLDG APYGYTPFCDSRREMDGYRFBKSGY WASHLAGRKYHISALYVVDLKKFRKI AAGDRLRGYQGLSQDPNSLSNLD QDLPNNMIHQVPIKSLPQEWLWCE TWCDDASKKRAKTIDLCNNPMTKEP KLEAAVRIVPEWQDYDQEIKQLQTLF QEEKELGTLHTEETQEGSQKHEEL
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