

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q5DTT3	TASO2_MOUSE	Mus musculus	Protein TASOR 2	15.818779	T880;S1202;S1203	S19;S219;S1021;S1082;S1430;S1540;S1962;S1990;S2012;S2015;S2019	36852467	MAAPTSKIILELNKNALISPWKGQFII QGCLLCDITLWSTYGTVVPLQLPREL DFKYVMDVSSLKESLPEAAFRRRSY LEQKVCCQDLCFDLYEVELTNKQGE NIDKLMKEYVKNKELALIKCLEDKSFF ILFTSSALTPPEPGFGAEQMGLHGLH LFHAPQTAGAKDLKVEDGISLKVIPIL PALSVALLEAKKSLSEEGIPPNILVKH SFQELYKVDKSLSLMAPPQDGVEDT ASTGKLSHAFDLPPPLETCPSESLTH LKCYSFSDPAGYTLDLAALDLLAHP QFPCIADGVCDAGFSLVMTDPPEFL DSEMEIRKTETAEKSGRKLKVKKKA VTPSSNQRVQPKRKASTTAVTLPSKR VSLGRPTSKRTVPRTDNRSCNPTLK LVKGQFPQKRKRGAEVLAQIVQKT RLERKKQEASVSKDAPVPTNTRAK KQEKSPGRIASQPKPPMKKSPQKRK VNVARGRRNTRKQLQPAEKEIALHL QSEISSDGQKDGLNLSTSQQESISMI PKGPPENSVISCDSQLNMLADLAL SSAAASIPSCKPRNLPCVSDLPRNNV LLTKENPLLGASDHEYHKGVKSQKA VLLPKPYSDEKISSEDLTRSQEENL VPCAQPLPIAQAPHSEARELSDASQ NSVVVEHSYALLLAEQSQKHLHQRK LPSPAFVKNIGKPEAGTPVGKVMVP FRHLQNTSPLQKHSEDSLMKHKSL FVSSTLKEFFCSHTVLKCDGFSKITF KCEGEYVFLDSKYTSNPLEKTVLRA LHGPWNTDLPENVDDVKLLLHIWV ALFYSKNNKVIKSPRKVVEHKNPAK YVCINRSLESLDHSEIEAFSNERPS VEGSVDPLETKETHIGHATNMTFP GPNRVLPFINPPTTRDLELCVQNDQ KEVFTGECHLDTSGNQNFYSCNTE VTGGKTKQELSNKLETSNVVLGCVV SAQSHGTCIPSEDKTCQSTKMVSYN DSVTQATLTTAYDEASSELMCQKSV FDNLENKVDSFHPSPLIKTDAVQDVI QHSSHINNECQPSVEKREDNVECM MVNLDPVTLAFEKNASVPIHTEVHT TDKPTGFNIELVKRVSPASSVQYPMS ALEEVQTQSSRDVPSLAMSEYKDSK CLSASSVKKETPPESLCLLQKEIPPLT SSADEGLIMEALSLVSSSYSLASDK TKCPQDDSLQTQNGLSMSLDEVLEP SKVNVVSSTSVTLREQSPNCIPAMS DVAGASTVIMNSGSSSLNQEKILQTF

SLVFPKQTDLSLKREEVSMELSGEE
ADINLTLTISPPTSPSEEIAAGEIEQF
QKTPVSNVGGQQRSEEMVEPEEEER
LIRNREINSASCMSVYPVESRELLKN
CTPEVTEKVNVTLDFFPGPLIEVSPA
SSPDPIVQPGDRPSSPCCLKLHSSQS
EKTNKFSQIKSGEVTIPEKESLFLGP
ESPKGQDKLAEVQVQISAEMLOITT
NAEVEGRVNM PGKVTKVSVPSEHS
ENLSFLEKVQCNTLNLNLSLPAKYG
GNFKPLEKSGNSLEAGCMENRNVD
VKHLALESSVPSCSPRKVVENKSLT
DTLVSITTS GIVNMSLKQTSSKNIKK
NVCDS DVKTDSDVKTEADSNMQTE
AVNSALIDKTDVQAYSHPEVSKFVSS
SDSAQCTYHAKPVSVEPGFQTQEIPV
VRMASLLKNIGVELHEEKMDLSATG
LQSNMSAKDEQKTMHVLQDTICE
VKEFLNGDVFSQNAHSCQNTVDFS
KSISEEPSASFVPEFVDAICGVYKEH
TFNESP NMVHETKADAETLSRNT
SVNASMFCGPRSGAYVQDSHDCKS
CKFDVENLRGNHESQKDAVKDSCD
SFTSLNNSDDTWACSSKISTLETHIP
PRDQETEPRLISPNCIPRYIQIPDSH
GIPKTYANFTITKEFKDTTRRLHSLK
RHRNLSANCNLLSSWTSTWHVTDD
LTQHTLDLEYLRFDHKLKQIKKGGG
QQSSFPKESLVQISSGTSPSTQTSEA
SGLHLPPE SRSPILVTVVRADTRQOS
HHRRGCSPSSLDGSSSFWKKCSQ
SRNLKNSERSQTVPFHLNKLKYNST
LKE SRNDISLILNEYAEFNKIMMNS
KQIVSQDEELNVAEAVFQEAYQP
RQPVS YEDVITDL CATLHVKLKGVVR
EACKSPFWFYL VETEDKS FLLR TKSI
LKKGGHIEIELLDFCQAFHRENETLL
VIKNEDIISHLHQIPSLK LKHFP SV
VFAGVDSPEDIVNETYQELFRSGGF
VVSDDVILESLTLVQLKEILKILEKLN
EN GKWKWLLHYRESKNL KEDVRVD
SIAHKKNLILKSYQSVNIIDLLHYHN
CDSPSSTKAEIFKCLLN LQIQHISAR
FAVFLTDKPTVSTEIFENNGILVTDV
NYFTENIQIAAPFRSSYW