

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
A2AMY5	A2AMY5_MOUSE	Mus musculus	NaN	18.691598	S392;S439;S468;T658;S876;S1017;S1019;S1020	NaN	30016717;28528544;25153642;34887587	<p>MTSVSNDRCRGAREKPMPTAHAA  QSQKQVVQATAEQMRLAQVIFDKN  DSDFEAKVKQLMEVTGKNQDECIVA  LHDCNGDVNKAINILLEGNSDTTSW  ETVGGKKKNFGRESSENKENREKR  TEREASRGRGTNNRKGRGGNRVRE  FKGEENGIDCSQGDKPAERGKRARG  RGFGRGRGRGTGRFSAQSMGTFNP  ADYSESMSTDGCGTKLAVWEAAQN  GTDEGPEGLAKSHSMSQEPSPKSSY  GLKGAWKNSVEEWTTEDWTEDLSE  TKVFTASSAPAENHVTPGHSIDLVAL  LHKPAPPTQATEVNSFETSQQQGF  QALVFTNSQHNNQMAPGTANSTSA  SSYSPQSLSSVLGSGFGELPQSNMV  NISNSQILDKLPGLSPFPAASSAQ  QNDTASPPATTAAWDLKPSAPQPSV  LSRLDFKSQPEPSPVLSQLSQRQQH  QTQAVSVPPGLESFSSLAKPRESTA  GDGPSTVSRLLQLPNMTVENIVSAH  QPQPKHIKPKRRVPPASKVPVSAVE  MPGSSDVTGLNVQFGALEFGSEPSL  SEFGSAASASENSNQIPISLYPKSLS  EPLNASFPMTSAVQSSTYTTSVVTSS  TLTSSALSSTSPVTSSSYDQSSVHT  RIAYQSSASPPDSAPGSVANGHGGG  RSQHTVDTTSSVPAPKKTDPALPSV  STLPGPASCTALLPSSAQHTATLPSL  TPAAAELSSSPLSQLSSSLSGHQNS  MTSAHATRSTSTPHTHASVESTASS  AAFSAAATSAPSAPSSGVVLPGSMST  VSSLCLGGTTVSVSSSTRATALVTS  GKAPPNLPQGVPPLLHNQYLVGPGG  LLPAYPIYGYDELQMLQSRPMDYY  GIPFAAPTALASRDGNLANNPYSGD  VTKFGRGDSASPAPPTTPAQAAQSQ  SQTHHTAQQPFLNPGLPPGYSTGL  PYVTGVPSAFQYGPTMFPPTSAPKQ  HGVALSTPPTPFQASGYGQHAYST  GYDDLTQGTAAAGDYTKGGYGGSSQA  PNKSTGSGPGKGVSVSSGTGLPDMT  GSVYNKTQTFDKQGFHAGTPPPFSL  PSALGSTGPLAPAAAPGYAPAPFLHI  MPAHQQPHSQLLHHHLQQDAPSGS  GQRSQPSSLQPKSQASKPTYGSAPY  WTN</p>