	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellula region
0	P10636-8	TAU_HUMAN	Homo sapiens	Microtubule-associated protein tau	25.916349	NaN	T18;T29;S46;S61;T69;T71;T111;S214;S396;T470;T486;T492;T498;S502;S508;S512;T514;S515;S516;S519;T522;T529;S531;T534;T548;S552;S554;S579;S602;S610;S622;S641;S669;S673;T711;S713;S717;T720;S721;S726;S733;S739;T744	36888912	MAEPRQEFEVMEDHAGTYGLGDRK DQGGYTMHQDQEGDTDAGLKESPL QTPTEDGSEEPGSETSDAKSTPTAE DVTAPLVDEGAPGKQAAAQPHTEIP EGTTAEEAGIGDTPSLEDEAAGHVT QEPESGKVVQEGFLREPGPPGLSHQ LMSGMPGAPLLPEGPREATRQPSGT GPEDTEGGRHAPELLKHQLLGDLH QEGPPLKGAGGKERPGSKEEVDEDR DVDESSPQDSPPSKASPAQDGRPPQ TAAREATSIPGFPAEGAIPLPVDFLSK VSTEIPASEPDGPSVGRAKGQDAPLE FTFHVEITPNVQKEQAHSEEHLGRA AFPGAPGEGPEARGPSLGEDTKEAD LPEPSEKQPAAAPRGKPVSRVPQLKA RMVSKSKDGTGSDDKKAKTSTRSSA KTLKNRPCLSPKHPTPGSSDPLIQPS SPAVCPEPPSSPKYVSSVTSRTGSSG AKEMKLKGADGKTKIATPRGAAPPG QKGQANATRIPAKTPPAPKTPPSSGE PPKSGDRSGYSSPGSPGTPGSRSRTP SLPTPPTREPKKVAVVRTPPKSPSSA KSRLQTAPVPMPDLKNVKSKIGSTE NLKHQPGGGKVQIINKKLDLSNVQS KCGSKDNIKHVPGGGSVQIVYKPVD LSKVTSKCGSLGNIHHKPGGGQVEV KSEKLDFKDRVQSKIGSLDNITHVPG GGNKKIETHKLTFRENAKAKTDHGA EIVYKSPVVSGDTSPRHLSNVSSTGS IDMVDSPQLATLADEVSASLAKQGL	None	None	None	None	None	None	None	None	None