	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	extracellular region
0	P19332	TAU_RAT	Rattus norvegicus	Microtubule-associated protein tau	29.427024	S711	T18;S35;S50;T58;T60;T100;S191;S204;S388;T464;T480;T486;T487;S489;T492;S496;S502;S506;T508;S509;S510;S513;T516;T523;S525;T528;T542;S546;S548;S573;S596;S600;S604;S616;S635;S663;S667;T705;S707;S711;T714;S715;S720;S727;S733;T738	19692427	MAEPRQEFDTMEDQAGDYTMLQDQEGDMDHGLKESPPQPPADDGSEEPGSETSDAKSTPTAEDVTAPLVEERAPDKQATAQSHTEIPEGTTAEEAGIGDTPNMEDQAAGHVTQEPQKVEIFSQSLLVEPGRREGQAPDSGISDWTHQQVPSMSGAPLPPQGLREATHQPLGTRPEDVERSHPASELLWQESPQKEAWGKDRLGSEEEVDEDITMDESSQESPPSQASLAPGTATPQARSVSASGVSGETTSIPGFPAEGSIPLPADFFSKVSAETQASPPEGPGTGPSEEGHEAAPEFTFHVEIKASAPKEQDLEGATVVGAPAEEQKARGPSVGKGTKEASLLEPTDKQPAAGLPGRPVSRVPQLKARVAGVSKDRTGNDEKKAKTSTPSCAKTPSNRPCLSPTRPTPGSSDPLIKPSSPAVCPEPATSPKYVSSVTPRNGSPGTKQMKLKGADGKTGAKIATPRGAATPGQKGTSNATRIPAKTTPSPKTPPGSGEPPKSGERSGYSSPGSPGTPGSRSRTPSLPTPPTREPKKVAVVRTPPKSPSASKSRLQTAPVPMPDLKNVRSKIGSTENLKHQPGGGKVQIINKKLDLSNVQSKCGSKDNIKHVPGGGSVHIVYKPVDLSKVTSKCGSLGNIHHKPGGGQVEVKSEKLDFKDRVQSKIGSLDNITHVPGGGNKKIETHKLTFRENAKAKTDHGAEIVYKSPVVSGDTSPRHLSNVSSTGSIDMVDSPQLATLADEVSASLAKQGL	None	None	None	None	None	None	None	None	None