

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	go ap
P10637	TAU_MOUSE	Mus musculus	Microtubule-associated protein tau	60.932047	S400;S529;S692	T18;S35;S50;T58;T60;T100;S188;S369;T445;S451;T461;T467;T468;S470;T473;S477;S483;S487;T489;S490;S491;S494;T497;T504;S506;T509;T523;S527;S529;S531;S554;S577;S581;S585;S597;S616;S644;S648;T686;S688;S692;T695;S696;S701;S708;S714;T719	35357812;35879403;36288343;24326295;26192747;30059200;16630055;23807304;26816085	MADPRQEFDTMEDHAGDYTLLODQEGDMDHGLKESPPQPPADDDGAEPEGSETSDAKSTPTAEDVTAPLVDERAPDKQAAAQPHTEIPEGITAEAGIGDTPNQEQQAAGHVTVQGRREGQAPDLGTSDWTRQQVSSMSGAPLLPQGLREATCQPSGTRPEDIEKSHPASELLRRGPQKEGEGWGDRLGSEEEVDEDLTVDESSQDSPPSQASLTPGRAAPQAGSGSVCGETASVPLPTEGVSPLPADFFSKVSAETQASQPEPGTGPMEEGHEAAPEFTFHVEIKASTPKEQDLEGATVVGVPGEQKAQTQGPSVKGKTKASLQEPGKQPAAGLPGRPVSRVPLKARVASKDRTGNDEKKAKTSTPSCAKAPSHRCLSPTRPTLGSSDPLIKPSSPAVSPEPATSPKHVSVTPRNGSPGTKQMKLKGADGKTGAKIATPRGAASPAQKGTSNATRIPAKTTPSPKTPPGSGEPPKSGERSGYSSPGSPGTPGSRSRTPSLPTPPTREPKKVAVVRTPPKSPSASKSRLOTPVPMPLDKNVRSKIGSTENLKHQPGGGKVQIINKKLDLSNVQSKCGSKDNKIHVPGGGSVQIVYKPVDSLKVTSKCGLGNIIHKPGGGQVEVKSEKLDKDRVQSKIGSLDNITHVPGGNKKIETHKLTFRENAKAKTDHGAIVYKSPVSGDTSRHLNSVSTSGSIDMVDSPQLATLADEVASLAKQGL	True	True	4.621	4.701	2.707	2.226	2.3