

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|------------------------------------|--------------|----------------|---|---|---|
| P10637 | TAU_MOUSE | Mus musculus | Microtubule-associated protein tau | 39.465564 | 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 | 26192747;30059200;16630055;23807304;24326295;26816085 | MADPRQEFDTMEDHAGDYTLQLQDQEGDMDHGLKESPPQPPADDGAEPEGSETSDAKSTPTAEDVTAPLVDERAPDKQAAAQPHTEIPEGITAEAEAGIGDTPNQEDQAAGHVTOGRREGQAPDLGTSDWTRQQVSSMSGAPLLPQGLREATCQPSGTRPEDIEKSHPASELLRRGPPQKEGWGQDRLGSEEEVDEDLTVDSSQDSSPPSQASLTPGRAAPQAGSGSVCGETASVPLPTEGVSPLPADFFSKVSAETQASQPEGPGTGPMEEGHEAAPEFTFHVEIKASTPKEQDLEGATVVGVPGEEQKAQTQGPSVKGKTKASLQEPPGKQPAAGLGRPVSRVPQLKARVASKDRGTGNDEKKAKTSTPSCAKAPSHRPCLSPTRPTLGSSDPLIKPSSPAVSPEPATSPKHVSSVTPRNGSPGTKQMKLKGADGKTGAKIATPRGAASPAQKGTSNATRIPAKTTPSPKTPPGSGEPPKSGERSGYSSPGSPGTPGSRSRTPSLPTPPTREPKKVAVVRTPPKSPSASKSRLQTAPVPMPLKKNVRSKIGSTENLKHQPGGGKVQIINKLDLSNVQSKCGSKDNIKHVPGGGSVQIVYKVPVDSLKVTSKCGSLGNIHHKPGGGQVEVKSEKLDKDRVQSKIGSLDNITHVPGGGNKKIETHKLTFRENAKAKTDHGA EIVYKSPVVS GDTSPRHLSNV SSTGSIDMVDSPLATLADEV SASLAKQGL |