

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P10636	TAU_HUMAN	Homo sapiens	Microtubule-associated protein tau	80.842688	T123 (P10636-8);S185 (P10636-8);S191 (P10636-8);S199 (P10636-8);S202 (P10636-8);T205 (P10636-8);S208 (P10636-8);T212 (P10636-8);S214 (P10636-8);T217 (P10636-8);S238 (P10636-8);S262 (P10636-8);T263 (P10636-8);S396 (P10636-8);S400;S404 (P10636-8);S412 (P10636-8);S413 (P10636-8);S422 (P10636-8);S525;S555;S717	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	12527113;19451179;30379171;27935279;28521765;26488919;20706749;26343633;28657654;26854602;23608187;34220449;24113653;21327254;24444746;15249677;28571750;22311971;32310828;30386294;27071766;22366723	MAEPRQEFVEMEDHAGTYGLGDRK DQGGYTMHQDQEGDTDAGLKESPL QTPTEDGSEEPGSETSDAKSTPTAE DVTAPLVDEGAPGKQAAAQPHTEIP EGTTAEEAGIGDTPSLEDEAAAGHVT QEPESGKVVQEGFLREPGPPGLSHQ LMSGMPGAPLLPEGPREATRQPSGT GPEDTEGGRHAPELLKHQLLGLDH QEGPPLKGAGGKERPGSKEEVEDDR DVDESSPQDSPPSKASPAQDGRFPQ TAAREATSIPGFPAEGAIPLPVDFLSK VSTEIPASEPDGPSVGRAKGQDAPLE FTFHVEITPNVQKEQAHSEEHLGRA AFPGAPGEGPEARPSLGEDTKEAD LPEPSEKQAAAAPRGKPVSRVPQLKA RMVSKSKDGTGSDDKKAKTSTRSSA KTLKNRCLSPKHPTPGSSDPLIQPS SPAVCPPEPSSPKYVSSVTSRTGSSG AKEMKLKGADGKTKIATPRGAAPP QKQANATRIPAKTPPAPKTPPSSGE PPKSGDRSGYSSPGSPGTPGSRSRTP SLPTPTREPKKVAVVRTPPKSPSSA KSRLQTAPVPMPLKLVKSKIGSTE NLKHQPGGGKVQIINKKLDLSNVQS KCGSKDNIKHVPGGGSVQIVYKQV LSKVTSKCGSLGNIHHKPGGGQVEV KSEKLDKDRVQSKIGSLDNITHVPG GGNKKIETHKLTFRNAAKAKTDHGA EIVYKSPVVSQDTSRHLNSVSSTGS IDMVDSPQLATLADEVASLAKQGL

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