

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9QXS1	PLEC_MOUSE	Mus musculus	Plectin	19.567819	T2762	S728;T823;S1055;S1443;S1729;S2639;S2781;T2788;S2809;T2893;T3040;T3369;T3792;T3797;T4037;S4061;S4389;S4391;S4392;S4393;S4396;S4397;S4398;S4399;T4400;S4403;S4413;T4418;T4546;S4614;S4620;T4622;S4623;S4625;S4629;T4630;S4633;S4649;S4682;S21;T26	33300544;22517741	MVAGMLMPLDRLRAIYEVLFREGV MVAKKDRRPRSLHPHVPGVNLTQV MRAMASDLKARGLVRETFAWCHFVW YLTNEGIDHLRQYLHLPPEIVPASLQ RVRRPVAMVIPARRRSPHVQTMQGP LGCPPKRGPLPAEDPAREERQVYRR KEREEGAPETPVVSATTVGTLARPGP EPAPATDERDRVQKKTFTKWNKHL IKHWRAEAQRHISDLYEDLRDGHNL ISLLEVLSGDSLPREKGRMRFHKLQ NVQIALDYLRHRQVKLVNIRNDDIA DGNPKLTLGLIWTIILHFQISDIQVSG QSEDMTAKEKLLWSQRMVEGYQG LRCDNFTTSWRDGRLFNAIHRHKP MLIDMNKVYRQTNLENLDQAFSVA ERDLGVTRLLDPEDVDVQPDEKSII TYVSSLYDAMPVPGAQDGVRADEL QLRWQEYRELVLQLLQWIRHHTAAF EERKFSSFEIEILWCQFLKFKETE LPAKEADKNRSKVIYQSLEGAVQAG QLKIPPGYHPLDVEKEWGKLVHVAILE REKQLRSEFERLECLQRIVSKLQME AGLCEEQLNQADALLQSDIRLLASG KVAQRAGEVERDLKADGMIRLLFN DVQTLKDGRHPQGEQMYRRVYRLH ERLVAIRTEYNLRLKAGVGGAPVTQVT LQSTQRRPELEDSTLRYLQDLLAWV EENQRRIDSAEWGVDLPSVEAQLGS HRGMHQSIEEFRAKIERARNDESQLE SPATRGAYRDCLGRLDLQYAKLLNS SKARLRSLESLHGFVAAATKELMWL NEKEEEEVGFWDSDRNTNMAAKKE SYSALMRELEMKEKKIKEIQNTGDR LLREDHPARPTVESFQAALQTQWS WMLQLCCCIEAHLKENTAYFQFFSD VREAEELQKLQETLRRKYSCDRTIT VTRLEDLLQDAQDEKEQLNEYKGH SGLAKRAKAIVQLKPRNPAHPVRGH VPLIACDYKQVEVTVHKGDQCQLV GPAQPSHWKVLSSGSSEAAVPSVCF LVPPPNEAQEAARLEAQHQALVT LWHQLHVDMMKSLLAWSLSDIQLI RSWSLVTFRTLKPEEQRQALRNLEL HYQAFLRDSQDAGGFGPEDRLVAER EYGSCSRHYQQLLSLEQGEQEESR CQRCISELKDIRELQLEACETRTVHRL RLPLDKDPARECAQRIAEQKKAQAE VEGLGKGVARLSAEAEKVLALPEPSP AAPTLRSELETLGKLEQVRSLSAIYL

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