

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|-------------|--------------|---------------------|---|---|---|
| Q86TC9 | MYPN_HUMAN | Homo sapiens | Myopalladin | 7.305184 | T390;T704;S714;S715 | S101;S131;T251;S644;S759;S813;S818;S867;S907;S928 | 30379171;33214551;31492838;31373491;34019948;30620550 | MQDDSI EASTSISQLLRESYLAETRH RGN NERSRAEPSSNPCHFGSPSGAA EGGGGQDDLPDLSAFLSQEELDES NLARLAINYDPLEKADETQARKRLSP DQMKHSPNLSFEPNFCQDNPRSP SSKESPOEAKRPQYCSETQSKKVFL NKAADFIEELSSLFKSHSSKRIRPRA CKNHKSKLESQNKVMQENSSSFSD LSERRERSVPIPIPADTRDNEVNHA LEQQEAKRREAQAASEAAGGDTP GSSPSSLYEEPLGQPPRFTQKLRSR EVPEGTRVQLDCIVVGIPPPQVRWYC EGKELENSPDIHIVQAGNLHSLTIAE AFEEDTGRYSCFASNIGTDSTSAEI YIEGVSSDSEGDPNKEEMNRIQKP NEVSSPPTTSAVIPPVPAQAHLVAQ PRVATIOQCQSPTNYLQGLDGKPIIA APVFTKMLQNLASSEGQLVVFECRV KGAPSPKVEWYREGTLIEDSPDFRIL QKKPRMAEPPEICTLVIAEVFAEDS GCFTCTASNKYGTVSSIAQLHVRGN EDLSNNGSLHSANSTTNLAAIEPOP SPPHSEPPSVEQPPKPLEGLVNVH NEPRSSRIGLRVHFNLPEDDKGSE ASSEAGVVTTTRQTRPDSFQERFNGQ ATKTPEPSSPVKEPPPVLAKPKLDST QLQLHNQVLEQHQLQNPPSP KEFPFSMTVLNSNAPPAVTTSSKQV KAPSSQTFSLARPKYFFPSTNTTAAAT VAPSSSPVFTLSSTPQTIQRTVSKESL LVSHPSVQTKSPGGLSIQNEPLPPGP TEPTPPPFTFSIPSGNQFQPRCVSIP VSPTSRIQNPVAFSSVLPVSLPAIPPT NAMGLPRSAPSMPSQGLAKKNTKS PQPVNDDNIRETKNAVIRDLGKKITF SDVRPNQQEYKISSFEQRLMNEIEF RLERTPVDESDEIQHDEIPTGKCIA PIFDKRLKHFVTEGSPVFTTCIVGI PVPKVYWFKDGKQISKRNEHCKMR REGDGTCSLHIESTTSDDDGNYTIM AANPQGRISCSGHLMVQSLPIRSRLT SAGQSHRGRSRVQERDKEPLQERFF RPHFLQAPGDMVAHEGRLCRLDCK VSGLPPELTWLLNGQPVLDPASHK MLVRETGVHSLLDPLTQRDAGTYK CIATNKTGQNSFSLELSVVAKEVKKA PVILEKLQNCGVPEGHPVRLPCRIG MPPPVFYWKDNETIPCTRERISMH QDTTGACLLIQPAKKS DAGWYTLS AKNEAGIVSCTARLDIYAQWHHQIPP PMSVRPSGSRYGSLTSKGLDIFSAFS SMESTMVYSCSSRSVVESEDEL |