

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|------------|--------------|-------------------|--------------|--------------------------|------------------------------------------------------------------------------|----------------------------|---------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Q80U78 | PUM1_MOUSE | Mus musculus | Pumilio homolog 1 | 31.870043 | S212;S220;S787;S798;S801 | S19;S75;S98;S106;T112;S124;S159;S197;S209;S229;S305;T515;S710;S715;S807;S823 | 22645316;22517741;36852467 | MSVACVLKRKAVLWQDSFSPHLKH HPQEPANPNMPVVLTSGTGSQAQP QPAANQALAAAGTHSSPVPGSIQVAG RSQDDAMVDYFFQRQHGEQLGGGG SGGGGYNTSKHRWPTGDNIIHAEHQ VRSMDELNHDFQALALEGRAMGEQ LLPGKKFWETDESSKDGPKGIFLGD QWRDSAWGTSDHSVSQPIMVQRRP GQSFHVNSEVNSVLSRSESGGLGV SMVEYVLS SSPGDSCLRKGGFGPRD ADSDENDKGEKKNKGTDFDGLGD LKEEGDVMDKTNGLPVQNGIDADV KDFSRTPGNCQNSANEVDLLGPNQ NGSEGLAQLTSTNGAKPVEDFSNM ESQSVPLDPMEHVGMELQFDYSG TQVPVDSAAATVGLFDYNSQQQLFQ RPNALAVQQLTAAQQQYALAAAHQ PHIAGLAPAAFVNPYIISAAPP GTDP YTAGLAAAATLGPVVPVPHQYYGVTP WGVYPASLFQQQAAAAAATNSATQ QSAPQAQQGQQQVLRGGASQRPLTP NQNQQQQTDPLVAAA AVNSALAF GQGLAAGMPGYPVLAPAAAYDQTGA LVVNAGARNGLGAPVRLVAPAPVIIS SSAAQA AVAAAAASANGAAGGLAGT TNGPFRPLGTQQPQPQQQPSNN LASSFYGNNSLSSNSQSSSLFSQG SAQPANTSLGFGSSSLGATLGSALG GFGTAVANSNTGSGSRRDSL TGSSD LYKRTSSSLAPIGHSFYSSLSYSSSPG PVGMP LPSQGP GHSQT PPPSLSSHG SSSSLNLGGLTNGSGRYISAAPGAEA KYRSASSASSLFSPTSSTLFS S SRLRY GMSDVMP SGRSRLLEDFRNNRYPN LQLREIAGHIMEFSQDQHGSRFIQL KLERATAAERQLVFNEILQAAYQLM VDVFGNYVIQKFFFEFGSHEQKLALA ERIRGHVLSLALQMYGCRVIQKALEF IPSDQQVINEMVRELDGHVLCVKD QNGNHVVQKCIQCVQPSLQFIIDA FKGQVFALSTHPYGCRVIQRILEHCL PDQTLPILEELHQHTEQLVQDQYGN YVIQHVLEHGRPEDKSKIVAEIRGNV LVLSQHKFASNVEKCVTHASRTER AVLIDEVCTMNDGPHSALYTM MKD QYANYVVQK MIDVAEPGQRKIVMHK IRPHIATLRKYTYGKHILAKLEKYIM KNGVDLGPICGPPNGII |