

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|-------------------------|--------------|----------------|--|---|---|
| Q9ULL5 | PRR12_HUMAN | Homo sapiens | Proline-rich protein 12 | 17.61578 | S201;S202;T363 | S332;S340;S651;T738;S865;S1077;S1135;T1304;S1308;S1381;S1382;S1387;T1561;S1568;T1705;S1925 | 31492838;29237092;34019948;30620550;35254053;35289036 | MDRNYPSAGFGDPLGAGAGWSYER SAKASLVYSSRTSHPETDILHRQAY AAPHPLOQSYATNHHHPAGLSGLFDTG LHHAGSAGPDASVMNLI SALESRGP QPGPSASSLLSQFRSPSWQTAMHTP GPTLFIISGALPGSSTFPSSALSAYQ HPASFGSRFPVPSLSLQDPPFSPP ANGLLSPHDVLHLKPSQAPTVPSSL GFERLAGGGV LGPAGLGP AQTPPYR PGPPDPPPPRHLP TQFNLLASSAA AAAAEQSSPQLYNFSGAAPGPPPE RALPRQDTVIKHYQR PASAQPPPPP PAHALQH YLSCGGSYPSMGHRANL ACSP LGGGEPSPGAGEPSKAGPSGA TAGASGRATGPEAAGGGGAGGGGG GYRPIIQSPGYKTGKGGYGAAAGGAT RPPPPRSTATPKCQSLGGPAAAYATG KASGAGGAGGQAYSPGQPQGLLGPO AYGQGFGGGQAQDLSKAPSYSGGPP QPPSGPPPPGLATCQSYSPDQLQGG LYGVQGEPPYGPAAHSQGLPTASPSL SYSTGHSPALSGHGGWGPPSSLGG GGEASPSHIIRPLQSPPATGRPPGVG SPGAPGKYLSSVLASAPFLAPPGAGS YAAGAGGYKKG DGSELLAGPGGPP AERTEDEEF LIQHLLQAPSPRTSGA DGLVGEDGAADASKGLGGSGGAGG PPGTPYELAKEDPQRYHLQSVIRTS SLDEGATAALELGLGRLKEKKKGPE RGGETPEGLATSVVHYGAGAKELGA FLQKSPPPPPPTAQSTQPTPHGLLLE AGGPDLP LVLPPPPQLLPSVLSHAP SPSPSASKVGVHLLLEPATRDGAPQPP PPPPPPPPMPLQLEAHLRSHGLEP AAPSPRLRPEESLDPPGAMQELLGA LEPLPPAPGDTGVGPPNSEGKDPAG AYRSPSPQGTKAPRFVPLTSICFPDS LLQDEERSFFPTMEEMFGGGAADD YGKAGPPEDEGDPKAGAGPPPGPPA YDPYGPYCPGRASGAGPETPGLGLD PNKPPPELPSTVNAEPLGLIQSGPHQA APPPPPPPPPAPASEPKGGLTSPIF CSTKPKKLLKTSSFHLLRRRDPPFQT PKKLYAQEYEFEADEDKADV PADIRL NPRRLPDLVSSCRSRPALSPLGDIDF CPPNPGPDGPRRRGRKPTKAKRDGP PRPRGRPRIRPLEVPTTAGPASASTP TDGAKKPRGRGRGRGRKAE EAGGT RLEPLKPLKIKLSVPKAGEGLGTSSG DAISGTDHNSLDSSLTREKIEAKIKE VEEKQPEMKSGFMASFLDFLKS GK RHPPLYQAGLTPPLSPPKSVPPS VPA RGLQPQPATPAVPHPPPSGAFGLG GALEAAESEGLGLGCPSPCKRLDEE LKRNL ETLPSFSSDEEDSVAKNRDL QESISSAISALDDPPLAGPKDTSTPD |

