

| UniprotKB ID | Entry name | organism | full name | oglcnaacscore | oglcnaac sites | phosphorylation sites | PMIDS | sequence |
|--------------|-------------|--------------|--------------------------------|---------------|--|-----------------------|--|--|
| Q5T6F2 | UBAP2_HUMAN | Homo sapiens | Ubiquitin-associated protein 2 | 45.727653 | S20;S303;S306;S310;S314;T372;S384;T389;T390;T391;S393;T394;S398;T399;S400;T404;T405;S407;T414;S415;S417;S418;S421;T486;T487;S492;S494;S552;S565;S633;S634;S635;S637;S737;S776;S987;S988;S994;S1004;S1006;S1008;S1095;S1104 | S432;S439;S473;S630 | 30379171;33214551;31492838;30620550;30397120;30059200;28314751;21740066;29249144;28657654;29351928;31373491;29237092;34019948;22121020;27114449;28604694;20305658;22661428;32574038;32870666;23301498;28510447 | MMTSVSSSDHCRGAREKPKQISAAQST QPQKQVVQATAEQMRLAQVIFDKN DSDFEAKVKQLMEVTKGNQDECIVA LHDCNGDVNKAINILLEGNSDITTSW ETVGCCKKNFAKENSENKENREKK SEKESRGRGNNNRKGKRGNGRGRE FRGEENGIDCNQVDKPSDRGKRAR GRGFRGRGRGAGRFSTQGMGTFN PADYSDSTSTDVCGTKLVVWEAAQN GADEGTELASNTHNIAQDLSNKSSY GLKGAWKNSVEEWTTEDWTEDLSE TKVFTASSAPAENHILPGQSIDLVAL LQKPVPHSQASEANSFETSQQQFG QALVFTNSQHNNQMAPGTGSSSTAV NSCSPQSLSSVLGSGFGELAPPKMA NITSSQILDQLKAPSLGQFTTTPSTQ QNSTSHPTTTTSDWDLKPPTSQSSVL SHLDFKSQPEPSPVLSQLSQRQQHQ SQAVTVPPPGLESFPSQAKLRESTPG DSPSTVNKLLQLPSTTIENISVSVHQ PQPKHIKLAARRIPPASKIPASAVEM PGSADVTGLNVQFGALEFGSEPSLS EFGSAPSENSNQIPISLYSKLSEPL NTLSMSTSAVQNSTYTTSVITSCSLT SSSLNSASPVAMSSSYDQSSVHNRI PYQSPVSSSESAPGTIMNGHGGGRS QOTLDPKTTGPPSALPSVSSLPSTT SCTALLPSTSQHTGDLTSSPLSQLSS SLSSHQSSLSAHAALSSSTSHTHAS VESASSHQSSATFSTAATSVSSSASS GASLSSSMNTANSLCLGGTPASASS SSSRAAPLVTSGKAPPNLPQGVPLL HNQYLVGPGGLLPAYPIYGYDELQM LQSRLPVDYYGIPFAAPTALASRDGS LANNPYPGDVTKFGRGDSASPAPAT TPAQPQSQSQTHHTAQPPFVNPAL PPGYSYTGPLYTGMPSAFQYGPTM FVPPASAKQHGVNLSPTPPFQAS GYGQHGYSTGYDDLTQGTAAGDYSK GGYAGSSQAPNKSAGSGPGKGVSVS SSTTGLPDMTGSVYNKTQTFDKQGF HAGTPPPFSLPSVLGSGTGPLASGAAP GYAPPPFLHILPAHQPPHSQLLHHH LPQDAQSGSQRSQPSSLQPKSQAS KPAYGNSPYWTN |

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