

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O75691	UTP20_HUMAN	Homo sapiens	Small subunit processome component 20 homolog	10.197167	NaN	S788;T1741;S2601;S2637	35083852;30379171	MKTKPVSHKTENTYRFLTFAERLGN VNIDIIHRIDRTASYYYYEVEYFFEGL LKWRELNLTEHFHGKFKYKEVIDKCS FNQLVYHQNEIVQSLKTHLQVKNSF AYQPLLDLVVQLARDLQMDFYPHFP EFFLTITSILETQDTELLEWAFTSLSY LYKYLWRLMVKDMSSISMYSTLLA HKKLHIRNFAAESFTFLMRKVSDKN ALFNLMFLDLDKHPEKVEGVGQLLF EMCKGVRNMFHSCTGQAVKLILRK LGPVTETETQLPWMLIGETLKNMVK STVSYISKEHFGTFFECLQESLLDLH TKVTKTNCCESSEQIKRLLLETYLILVK HGSGTKIPTPADVCKVLSQTLQVASL STSCWETLLDVISALILGENVSLPET LIKETIEKIFESRFKRLIFSFSSEVMF AMKQFEQLFLPSFLSYIVNCFLLDDA VVKDEALAILAKLILNKAAPPTAGSM AIEKYPLVFSQPMVGFYIKQKKTRSK GRNEQFPVLDHLLSIKLPNKNDDTY LSQSWAALVVLPHIRPLEKEKVIPLV TGFIEALFMTVDKGSFGKGNLFVLC QAVNTLLSLEESSELLHLPVERVK NLVLTFFLEPSVLLLTDLYYQRLALC GCKGPLSQEALMELFPKLQANISTG VSKIRLLTIRILNHFDVQLPESMEDD GLSERQSVFAILRQAEVLPATVNDYR EKLHLRKLRLHDVVQTAVPDGPLQE VPLRYLLGMLYINFSALWDPVIELIS SHAHEMENKQFWKVYEHLEKAAT HAEKELQNDMTDEKSVGDESWEQT QEGDVGALYHEQLALKTDCQERLD HTNFRFLLWRALTKFPERVEPRSRE LSPLFLRFINNEYYPADLQVAPTQDL RRKGKGMVAEEIEEPAAGDDEELE EEAVPQDESSQKKKTRRAAAKQLIA HLQVFSKFSNPRALYLESKLYELYLQ LLLHQDQMVQKITLDCIMTYKHPHV LPYRENLQRLEDRSFKEEIVHFSIS EDNAVVKTAHRADLFPILMRILYGR MKNKTGSKTQGSASGTRMAIVLRF LAGTQPEEIQIFLDLLFEPVRHFKNG ECHSAVIQAVEDLDLSKVLPLGRQH GILNSLEIVLKNISHLISAYLPKILQIL LCMTATVSHILDQREKIQLRFINPLK NLRRLGIKMTDIFLDWESYQFRTE EIDAVFHGAVWPQISRLGSESQYSPT PLLKLISIWSRNARYFPLAKQKPGH PECDILTNVFAILSAKNLSDATASIV

MDIVDDLNLPDFEPTETVLNLLVT  
GCVYPGAENIGESITIGGRLLPHVP  
AILQYLSKTTISAEKVKKKKNRAQVS  
KELGILSKISKFMKDKEQSSVLITLLL  
PFLHRGNIAEDTEVDILVTQNLKHL  
CVDPTSFLKPIAKLFSVIKNKLSRKL  
CTVFETLSDFESGLKYITDVVKLNAF  
DQRHLDDINFDFVFETFTTITSYIKE  
MQIVDVNYLIPVMHNCFYNLELGD  
MSLSDNASMCLMSIICKLAALNVTE  
KDYREIHRSLLEKLRKGLKSQTESI  
QQDYTTILSCLIQTFFPNQLEFKDLVQ  
LTHYHDPMDFFENMKHIQIHRRRA  
RALKKLAKQLMEGKVVLSKSLQNY  
IMPYAMTPIFDEKMLKHENITTAATE  
IIGAICKHLSWSAYMYLKHFIHVLO  
TGQINQKLGVSLLVIVLEAFHFDHKT  
LEEQMGKIENEENAIEAIELPEPEAM  
ELERVDEEEKEYTCKSLSDNGQPGT  
PDPADSGGTSAKESECITKPVSFPLQ  
NKEEIERTIKNIQGTITGDILPRLHKC  
LASTTKREEEHKLVKSKVVNDEEVV  
RVPLAFAMVKLMQSLPQEVMEANL  
PSILLKVCALLKNRAQEIRDIARSTLA  
KIIEDLGVHFLLYVLKELQTTLVRGY  
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KQILSKVMEARRSKSYDSYEILGKFV  
GKDQVTKLILPLKEILQNTTSLKLAR  
KVHETLRRITVGLIVNQEMTAESILL  
LSYGLISENPLLTEKEKNPVAPAPD  
PRLPPQSCLLLPPTPVRRGGQKAVVSR  
KTNMHIFIESGLRLLHLSLKTSTIKS  
SGECVLEMLDPFVSLIDCLGSM DV  
KVITGALQCLIWVLRFP LPSIETKAEQ  
LTKHLFLLLKDYAKLGAARGQNFHL  
VVNCFKCVTILVKKVKSQITEKQLQ  
VLLAYAEEDIYDTSRQATAFGLLKAIL  
SRKLLVPEIDEVMRKVSKLAVSAQSE  
PARVQCRQVFLKYILDYPLGDKLRPN  
LEFMLAQLNYEHETGRESTLEMIAY  
LFDTFPQGLLHENC GMFFIPLCLMT  
INDDSATCKKMASMTIKSLLGKISLE  
KKDWLFD MVTTWFGAKRRLNRQLA  
ALICGLFVESEGVDFEKRLGTVLPVI  
EKEIDPENFKDIMEETEEKAADRLLF  
SFLTITKLIKECNIIQFTKPAETLSKI  
WSHVHSHLRHPHNWVWL TAAQIFG  
LLFASCQPEELIQKWNTKKT KKHLP  
EPVAIKFLASDL DQKMKSISLASCHQ  
LHSKFLDQSLGEQVVKNLLFAAKVL  
YLLELYCEDKQSKIKEDLEE QEALED

