

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	10.517576	NaN	S788;T1741;S2601;S2637	35083852;37217939;30379171	MKTKPVSHKTEITYRFLTFAERLGN VNIDIHRIDRTASYEEEVETVFFEGL LKWRELNLTEHFGKFYKEVIDKQCQ FNQLVYHQNEIVQSLKTHLQVKNSF AYQPLLDLVVQLARDLQMDFYPHFP EFFLTITSILETQDTELLEWAFTSLSY LYKYLWRLMVKDMSSIYSMYSTLLA HKKLHIRNFAAESFTFLMRKVSDKN ALFNLMFLLDLKHPKVEGQVGLLF EMCKGVRNMFHSCTGQAVKLILRK LGPVTETETQLPWMLIGETLKNMVK STVSYSISKEHFGTFFECLQESLLDLH TKVTKTNCCESSEQIKRLLLETYLILVK HGSGTKIPTPADVCKVLSQTLQVASL STSCWETLLDVISALILGENVSLPET LIKETIEKIFESRFEKRLIFSSEVMF AMKQFEQLFLPSFLSYIVNCFLLDDA VVKDEALAILAKLILNKAAPPTAGSM AIEKYPLVFSPQMVGFIKQKKTRSK GRNEQFPVLDHLLSIKLPNKDDTY LSQSWAALVVLPHIRPLEKEKVIPLV TGFIEALFMTVDKGSFGKGNLFVLC QAVNTLLSLEESSELLHLPVERVK NLVLTFFLEPSVLLLTDLYYQRLALC GCKGPLSQEALMELFPKLQANISTG VSKIRLLTIRILNHFDVQLPESMEDD GLSERQSVFAILRQAEVLPATVNDYR EKLLHLRKLRLHDVVQTAVPDGPLQE VPLRYLLGMLYINFSALWDPVIELIS SHAHEMENKQFWKVYYEHLEKAAT HAEKELQNDMTDEKSVGDESWEQT QEGDVGALYHEQLALKTDCQERLD HTNFRFLLWRALTKFPERVEPRSRE LSPLFLRFINNEYYPADLQVAPTQDL RRKGKGMVAEEIEEPAAGDDEELE EEAVPQDESSQKKKTRRAAAKQLIA HLQVFSKFSNPRALYLESKLYELYLQ LLLHQDQMVQKITLDCIMTYKHPHV LPYRENLQRLLEDRSFKEEIVHFSIS EDNAVVKTAHRADLFPILMRILYGR MKNKTGSKTQGSASGTRMAIVLRF LAGTQPEEIQIFLDLLFEPVRHFKN ECHSAVIQAVEDLDLSKVLPLGRQH GILNSLEIVLKNISHLISAYLPKILQIL LCMTATVSHILDQREKIQLRFINPLK NLRRLGIKMVTDIFLDWESYQFRTE EIDAVFHGAVWPQISRLGSESQYSPT PLLKLISIVSRNARYFPLAKQKPGH PECDILTNVFAILSAKNLSDATASIV MDIVDDLNLPDFEPTETVLNLLVT

GCVYPGIAENIGESITIGGRLILPHVP
AILQYLSKTTISAQVKKKKNRAQVS
KELGILSKISFKMKDKEQSSVLITLLL
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CVDPTSFLKPIAKLFSVIKKNLSRKLL
CTVFETLSDFESGLKYITDVVKLNAF
DQRHLDDINFDVRFETFQTITSYIKE
MQIVDVNYLIPVMHNCFYNLELGD
MSLSDNASMCLMSIICKLAALNVTE
KDYREIHRSLLEKLRKGLKSQTESI
QQDYTTILSCLIQTFPNQLEFKDLVQ
LTHYHDPEDFFENMKHIQIHRRA
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IIGAICKHLSWSAYMYLKHFIHVLO
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ELERVDEEEKEYTCKSLSDNGQPGT
PDPADSGGTSAKESECITKPVSFPLQ
NKEEIERTIKNIQGTITGDILPRLHKC
LASTTKREEEHKLVKSKVVNDEEVV
RVPLAFAMVKLMQSLPQEVMEANL
PSILLKVCALLKNRAQEIRDIAARSTLA
KIIEDLGVHFLLYVLKELQTTLVRGY
QVHVLTFTVHMLLQGLTNKLVQGD
LDSCLDIMIEIFNHELFGAVAEKEV
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LSYGLISENLPLLTEKEKNPVAPAPD
PRLPPQSCLLLPPTPVRRGGQKAVVSR
KTNMHIFIESGLRLLHLSLKTSKIKS
SGECVLEMLDPFVSLIDCLGSM DV
KVITGALQCLIWVLRFP LPSIETKAEQ
LTKHLFLLLKDYAKLGAARGQNFHL
VNCFKCVTILVKKVKSQITEKQLQ
VLLAYA EEDIYDTSRQATAFGLLKAIL
SRKLLVPEIDEVMRKVSKLAVSAQSE
PARVQCRQVFLKYILDYPLGDKLRPN
LEFM LAQLNYEHETGRESTLEMIAY
LFDTFPQGLLHENC GMFFIPLCLMT
INDDSATCKKMASMTIKSLLGKISLE
KKDWLFD MVTTWFGAKKRLNRQLA
ALICGLFVESEGVD FEKRLGT VLPVI
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SFLTITKLIKECNIIQFTKPAETLSKI
WSHVHSHLRHPHNWVWL TAAQIFG
LLFASCQPEELIQKWNTKKT KKHLP
EPVAIKFLASDL DQKMKSISLASCHQ
LH SKFLDQSLGEQVKNLLFAAKVL
YLLELYCEDKQSKIKEDLEEQALED
GVACADEKAESDGEEKEEVKEELGR
PATLLWLIQKLSRIAKLEAAYS PRNP

