	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum
0	P24928	RPB1_HUMAN	Homo sapiens	DNA-directed RNA polymerase II subunit RPB1	24.396123	S1534;S1619;S1621	S27;S217;T1840;S1843;S1845;S1847;S1849;S1850;T1854;S1857;T1860;S1861;T1863;S1864;T1867;S1868;T1870;T1874;S1875;T1877;S1878;T1881;S1882;T1885;T1894;S1896;S1899;S1906;T1909;T1912;S1913;T1915;T1916;S1917;T1919;S1920;T1923;T1926;S1927;T1929;T1930;S1931;T1933;S1934	26807597;38665916;34019948;26853435;22371499;37689115;34725712;22605332;37217939;29351928	MHGGGPPSGDSACPLRTIKRVQFGV LSPDELKRMSVTEGGIKYPETTEGG RPKLGGLMDPRQGVIERTGRCQTCA GNMTECPGHFGHIELAKPVFHVGFL VKTMKVLRCVCFFCSKLLVDSNNPK IKDILAKSKGQPKKRLTHVYDLCKGK NICEGGEEMDNKFGVEQPEGDEDL TKEKGHGGCGRYQPRIRRSGLELYA EWKHVNEDSQEKKILLSPERVHEIF KRISDEECFVLGMEPRYARPEWMIV TVLPVPPLSVRPAVVMQGSARNQDD LTHKLADIVKINNQLRRNEQNGAAA HVIAEDVKLLQFHVATMVDNELPGL PRAMQKSGRPLKSLKQRLKGKEGRV RGNLMGKRVDFSARTVITPDPNLSI DQVGVPRSIAANMTFAEIVTPFNIDR LQELVRRGNSQYPGAKYIIRDNGDRI DLRFHPKPSDLHLQTGYKVERHMC DGDIVIFNRQPTLHKMSMMGHRVRI LPWSTFRLNLSVTTPYNADFDGDEM NLHLPQSLETRAEIQELAMVPRMIV TPQSNRPVMGIVQDTLTAVRKFTKR DVFLERGEVMNLLMFLSTWDGKVP QPAILKPRPLWTGKQIFSLIIPGHINC IRTHSTHPDDEDSGPYKHISPGDTKV VVENGELIMGILCKKSLGTSAGSLVH ISYLEMGHDITRLFYSNIQTVINNWL LIEGHTIGIGDSIADSKTYQDIQNTIK KAKQDVIEVIEKAHNNELEPTPGNT LRQTFENQVNRILNDARDKTGSSAQ KSLSEYNNFKSMVVSGAKGSKINIS QVIAVVGQQNVEGKRIPFGFKHRTL PHFIKDDYGPESRGFVENSYLAGLTP TEFFFHAMGGREGLIDTAVKTAETG YIQRRLIKSMESVMVKYDATVRNSIN QVVQLRYGEDGLAGESVEFQNLATL KPSNKAFEKKFRFDYTNERALRRTL QEDLVKDVLSNAHIQNELEREFERM REDREVLRVIFPTGDSKVVLPCNLLR MIWNAQKIFHINPRLPSDLHPIKVVE GVKELSKKLVIVNGDDPLSRQAQEN ATLLFNIHLRSTLCSRRMAEEFRLSG EAFDWLLGEIESKFNQAIAHPGEMV GALAAQSLGEPATQMTLNTFHYAGV SAKNVTLGVPRLKELINISKKPKTPS LTVFLLGQSARDAERAKDILCRLEHT TLRKVTANTAIYYDPNPQSTVVAEDQ EWVNVYYEMPDFDVARISPWLLRVE LDRKHMTDRKLTMEQIAEKINAGFG DDLNCIFNDDNAEKLVLRIRIMNSD ENKMQEEEEVVDKMDDDVFLRCIE SNMLTDMTLQGIEQISKVYMHLPQT DNKKKIIITEDGEFKALQEWILETDG VSLMRVLSEKDVDPVRTTSNDIVEIF TVLGIEAVRKALERELYHVISFDGSY VNYRHLALLCDTMTCRGHLMAITR HGVNRQDTGPLMKCSFEETVDVLM EAAAHGESDPMKGVSENIMLGQLA PAGTGCFDLLLDAEKCKYGMEIPTNI PGLGAAGPTGMFFGSAPSPMGGISP AMTPWNQGATPAYGAWSPSVGSGM TPGAAGFSPSAASDASGFSPGYSPA WSPTPGSPGSPGPSSPYIPSPGGAMS PSYSPTSPAYEPRSPGGYTPQSPSYS PTSPSYSPTSPSYSPTSPNYSPTSPSY SPTSPSYSPTSPSYSPTSPSYSPTSPS YSPTSPSYSPTSPSYSPTSPSYSPTSP SYSPTSPSYSPTSPSYSPTSPSYSPTS PSYSPTSPSYSPTSPSYSPTSPNYSPT SPNYTPTSPSYSPTSPSYSPTSPNYTP TSPNYSPTSPSYSPTSPSYSPTSPSYS PSSPRYTPQSPTYTPSSPSYSPSSPSY SPASPKYTPTSPSYSPSSPEYTPTSPK YSPTSPKYSPTSPKYSPTSPTYSPTTP KYSPTSPTYSPTSPVYTPTSPKYSPTS PTYSPTSPKYSPTSPTYSPTSPKGSTY SPTSPGYSPTSPTYSLTSPAISPDDSD EEN	None	None	None	None	None	None