

UniprotKB ID	Entry name	organism	full name	oglcnaScore	oglcna sites	phosphorylation sites	PMIDS	sequence
Q15648	MED1_HUMAN	Homo sapiens	Mediator of RNA polymerase II transcription subunit 1	6.527807	S596;S671;S673;S682;S1021;S1026;S1027;T1032;T1035;S1036;T1215;T1259;S1263;S1265;T1267;S1289;S1290;S1427	S588;S664;S795;T805;S887;S953;T1032;T1051;T1057;S1156;S1207;T1215;S1223;S1302;S1347;S1403;S1433;T1440;T1457;S1463;S1465;S1479;S1481;S1482	33214551;35254053;30379171;29351928;34725712;38253038;35132862;31492838;35138101;37217939;34019948;26960196;23301498;35289036	MKAQGETEESKLSKMSSLLERLHA KFNQNRPWSETIKLVRQVMEKRVV MSSGGHQHLVSCLETLQKALKVTSL PAMTDRLESARQNLGSHLSASGT ECYITSDMFYVEVQLDPAGQLCDVK VAHHGENPVSCEPELVQQLREKNFD EFSKHLKGLVNLNLPDGNLKLTK MYLALQSLEQDLSKMAIMYWKATN AGPLDKILHGSVGYLTPRSGGHLMN LKYYVSPSDLDDKTASPIILHENV SRSLGMNASVTIEGTSAVYKLPAPLI MGSHFPVDNKWTPSFSSITSANSVDL PACFFLKFPQIPVSRFAVQKLQNC GIPLFETQPTYAPLYELITQFELSKDP DPIPLNHNMRFYAALPGQQHCHYFLN KDAPLPDGRSLQGTLVSKITFOHPG RVPLILNLRHQVAYNTLIGSCVKRTI LKEDSPGLLQFEVCPSESFRFSVSFQ HPVNDSLVCVMDVQDSTHVSKL YKGLSDALICTDDFIKVVQRCMSIP VTMRAIRRKAEITQADTPALSIAETV EDMVKKNLPPASSPGYGMTGNNP MSGTTTPTNTFFPGGPITTLFNMSMS IKDRHESVGHGEDFSKVSQNPILTS LQITGNGGSTIGSSPTPPHHTPPV SMAGNTKNHPMLMNLKDNPAQD FSTLYGSSPLERQNSSSGSPRMEICS GSNKTKKKSSRLPPEKPKHQTEDD FQRELFSDMDVDSQNPFDVNMAD TLDTPHITPAPSQCSTPPTYPQVP HPQPSIQRMVRLSSSDSIGPDVTDIL SDIAEEASKLPSTSDDCPAIGTPLRD SSSSGHSQSTLFDSDVFQTNNNENP YTDPADLIADAAGSPSSDPTNHFFH DGVDFNPDLNSQSQSGFGEEYFDE SSQSGDNDDFKGFASQALNTLGVP MLGGDNGETKFKGNNQADTVDFSII SVAGKALAPADLMEHHSQSGPLLT TGDGKEKTQKRVKEGNGTNSSTLS GPGLDSKPGKRSRTPSNDGKSKDKP PKRKKADTEGKSPSHSSSNRPFTPP TSTGGSKSPGSAGRSQTPPGVATPPI PKITIQIPKGTVMVGKPSHSHSQTSS GSVSSSGSKSHSHSSSSSSSASTS GKMKSSKSEGSSSSKLSSSMYSSQG SSGSSQSKNSSQSGGKPGSSPITKH GLSSGSSSTKMKPQGKPSLMLNPSL SKPNISPSHSRPPGSDKCLASPMKP VPGTTPSSKAKSPISSGSGGSHMSGT SSSSGMKSSSGLSSGSLSQKTPPS SNSCTASSSSFFSSGSSMSSSQNH GSSKSKSPSRNKKPSLTAVIDKLKH GVVTSKPGGEDPLDGQMGVSTNSS SHPMSSKHNMSGGEFQKREKSDK DKSKVSTSGSSVDSKKTSESKNVG STGVAKIIISKHDGGSPSIKAKVTLQK PGESSGGLRPMASAKNYGSLIS GSTPKHERGSPSHSKPAYTPQNL SESESGSSIAEKSYQNSPSSDDGIRP LPEYSTEKHKHKHKKKVKDKDRD RDRDKDRDKKSHSIKESWSKSPI SSDQSLSMTSNTILSADRPSRLSPDF MIGEEDDDLMVALIGN

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