

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extra regic
P53836	CA120_YEAST	Saccharomyces cerevisiae	CCR4-NOT transcriptional complex subunit CAF120	24.475439	NaN	S491;S510;S518;S538;S556;S871;S885	33229814	MRIFSGDNKVVDSLASNPLMLSPSN FGGDFGSRLKVNVTSKKKLNDSST SPMESSPVSPPELVLPILTLNAHTRR YHEGVFLILQDLNNGTHAARKWK DVYGVLLGTQLALWDAKELAEFTDP SCPVSEKKLKEVASKPTYINLTDATL RTLDNNDNIVMECGKNTNALVVST TLKNRYFLQFGNKESFNAWNNSAIRL CLYECSSLOEAYTGAFISSRGAKLGD IRILLTNRKYDYKDWVSVRFAGAMP WKRKYAVISQSSSKKGFGEINLY ENDKVKKNHAMATIVEAKALYAVY PSSPKLIDSSSTIIKVVGSVKFEKESA QEKDVFIMPEKHQAVPSYDITIRFLIP AMDTFKLYGRPEKLLSSKNDPHSLL FGLPVLPHIYYLEVEDLLPLTNSVSS LHWSNNEWKEHISDILQRKIAQGYC GCNSTSNITSPLPSPFLGSADLFERA DGVLSPKLSYGSKSSNNSSKNSLP KRERVKLSSSEODLNNSDPSIKR KSPPLVISESPHKVHTPTDASFRTRV TEGSPYAKQRHPKPFASSVNDSPSD RAKSRTVPYNNNDRKATTPEKFERG ETSCGKNVDESLEKVRNMKLEIPES NFDKFMTDKNLLSVDSKCSNEKKL SVESDLSAIYEKYSNGPFGHTEGLN GSSDETYLRFQRASVHSESNYNSRK SFTPSDFSNGNEEHAVLQELNSLT QRINELGMESINSNSDSDRINGSYS QVDFGNNDEDDMNLFDPDFMAQ DQLRAEERDYNKDDRTPLAKVPAAF QSTGLGITPDDDIERQYITEHRSRHE VPKRSPEKPSNPLEIGNPYAKPGTRL NTTHTHSKTDRSITPQRGQVPVPSGQ QISSYVQPANINSPNKMYGANNNSAM GSPRNPKTRAPPGPYNOGWNNRPS PSNIYQRPHPSDTQPQAYHLPGNPY STGNRPNMQAQYHPQQVPMPILOQ PNRPYQPYAMNTHMGSPGGYAGAA PPFQANVNYNTRPQPWPPTNSPS AHYRPPNLLNQPNQNSAGYYRPPAP QLQNSQARPKKDGFSQFMPSATT KNPYAQ	None	None	None	None	None	None	None	None	

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