

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P38167	ECM21_YEAST	Saccharomyces cerevisiae	Protein ECM21	9.365641	NaN	S18;S115;S140;S286;S527;S550;S775;S1035	33229814	MPFITSRPVAKNSSHSLSETDLNQS KGQPFQPSPTKKLGSMMQRRRSSTI RHALSSLLGGANVHSPAVLNNTTKG GNNNGNIRSSNTDAQLLGKKQNKQ PPPNARRHSTTAIQGSISDSATTTPR SSTSDTNRRTSGRLSVDQEPRISGGR YSQIEEDSTVLDFFDDHNSAVVSS DLSSTSLTRLANSKFNQFLIEYLT ARGLLGPKTVLSNEYLKISISTSGESV FLPTISSNDDEYLSRLNGLNDGTDD AEADFFMDGIDQQEGNTPSLATTAA ATESGGSINENRDTLLRENNSGDHP GSGSELNTRSVEIDSSMVSYIAVIV SVKKPTRFTDMQLELCRVKVFVWNT GVPPTKTFNEEFYNAASMKNLND ENFDLFLVPLSISPDDQMIENNSNDR QMRLFKNIPTEERLYLDKTKTKASLL NAIDVNKTHLYQPGDYVFLVPVVS NHIPETIYLP SARVSYRLRLATKAINR KGFYRQDSNSPQIVSPDSSSSLSST TSSLKLTETESAQAHRISNTLFSKV KNHLHMSSHQLKNEESGEEDIFAE YPIKVIRTPPPVAVSTANKPIYINRVW TDSLSEISFAQKYVSLNSEVPIKIKL APICKNVCVKRIHVSITERVTFVSKG YHEYDQTDPAKDPYNPYLDFAK RRKERSVSLFEIRTKKTRALREEI VENSFNDNLLSYSPFDDSDSKGNP KERLGITEPIIETKLFKPYEDLDR TAKIIPPYGIDAYTSIPNPEHAVANGP SHRRPSVIGFLSGHKGSKSHEENEK PVYDPKFHQTIKSNGLPVKTHTRL NTPKRGLYLDLHFSNVYCRHKLEI MLRISKPDPECPSKLRHYEVLIDTPIF LVSEQCNSGNMELPTYDMATMEGK GNQVPLSMNSDFFGNTCPPPTFEE AIVPASPISPMGSPNIMASYDPDL LSIQQLNLSRRTTSVSGPSGYSDDAGV PNVNRNSISNANAMNGSISNSAFVS GNSGQGVARARATSVNDRSRFNNL DKLLSTPSPVNRSHNSSPTNGLSQA NGTVRIPNATTENSKDKQNEFFKKG YTLANVKDDEEQEGIVSSSSADSLLS HGNEPPRYDEIVPLMSDEE