	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	P53135	SLD3_YEAST	Saccharomyces cerevisiae	DNA replication regulator SLD3	24.475439	NaN	NaN	33229814	METWEVIASVKEATKGLDLSLDHPLI IKSEDVPSNILQLLQQKNRRQLKHIC MKSRKEYFLLEEYGPGFWVKWPYN YFNGYSLPERRTEVVTTVERERAKR ETLKTWDELKFKELLHLWSEEPKGS CKLEKDKDLKLDMNPPDMKGESKI NDYYSDPKEYIESKYYDALFSIHTPLA YFVKSNLVRLKNTCRTKYGSDSYKIA YQAMLQKFLLSIVQFKDRHDNRLLL EPFSSPIADEKRKNCLTKFVIQDENK NSSTIADLCVVLKSREIKLQILLLLEII GLNDLDWNFRDFEKKYKLKLKKRS LNLTKKGLVRRRSKKKTSEKDKGIE RITTSLDYCEQLDLYLDRACILDILLS SETPNPDAIEASNGTIQEHKKNILDK SKEASLVGFINYVLIPYFNKKVPHAV EFIIQKLKGPSMRPKRALKKVNDST NVSSPNTVETYNRLSTSQRASRSSII NSVPSSPALRRVDANLFSRKSIASPT PELLNSRTNSNLNEFLESETRSLKR PSQLGRTKSDLTMNHLQKRQFSVS DLSTTRVPNSSTITLKTPFSHSTINAY KTMNNSFRRVGKRKDINETIRLHER VDSEENVQVQATPAVKKRTVTPNKK AQLQSIIESPLNFKDDDTHEGRKNT SNITSTPTNKPPENSSKRRVRRRLFA PEST	None	None	None	None	None	None	None	None	None