	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	extrac region
0	E6EK15	E6EK15_DROME	Drosophila melanogaster	Calcium-transporting ATPase	28.06426	S1114;S1116;T1140;S1143	NaN	40245499	MATIDGRPAQYGISLKQLRELMEHR GREGVMKIAENGGIHELCKKLYTSP NEGLSGSKADEEHRRETFGSNVIPP KPPKTFLTLVWEALQDVTLIILEVAAL VSLGLSFYKPADEDAPVLQEEEEHH GWIEGLAILISVIVVVIVTAFNDYSKE RQFRGLQNRIEGEHKFSVIRGGEVC QISVGDILVGDIAQVKYGDLLPADGC LIQSNDLKVDESSLTGESDHVKKGP DVDPMVLSGTHVMEGSGKMVVTAV GVNSQAGIIFTLLGAAVDEQEAEIKK MKKGENDGRSQIKGSQAPSQRETVT SEITKSESEGNHLPQSSSSGAAETG HKKEKSVLQAKLTKLAIQIGYAGSTIA VLTVIILIIQFCIKTFVIDEKPWKNTYA NNLVKHLIIGVTVLVVAVPEGLPLAV TLSLAYSVKKMMKDNNLVRHLDAC ETMGNATAICSDKTGTLTTNRMTVV QSYICEKLCKVLPTLSDIPQHVGNLIT MGISVNSAYTSNIMAGHNPGDLPIQ VGNKTECALLGFVQGLGVKYQSIRD EITEDKFTRVYTFNSVRKSMGTVIPR PNGGYRLYTKGASEIIMKKCAFIYGH EGTLEKFTRDMQERLIREVIEPMAC DGLRTISVAYRDFVPGKAAINEVHID GEPNWDDEENIMTNLTCLCVVGIED PVRPEVPDAIRKCQRAGITVRMVTG DNINTARSIASKCGILRPNDDFLILE GKEFNRRIRDSNGDIQQHLIDKVWP KLRVLARSSPTDKYTLVKGIIDSTVSE NREVVAVTGDGTNDGPALKKADVGF AMGIAGTDVAKEASDIILTDDNFSSI VKAVMWGRNVYDSIAKFLQFQLTVN VVAVIVAFIGACAVQDSPLKAVQML WVNLIMDTLASLALATEFPTPDLLLR KPYGRTKPLISRTMMKNILGQALYQ LIIIFGLLFVGDVILDIESGRGQELNA GPTQHFTIIFNTFVMMTLFNEINAR KIHGQRNVIEGLLTNPIFYTIWIFTMI SQVLIIQYGKMAFSTKALTLDQWLW CIFFGIGTLVWGQLITSVPTRKLPKIL SWGRGHPEEYTDGMNLGEERFDSI DSDKKPRAGQILWIRGLTRLQTQVIG GELQERLIPVPYSKSNTDQAIRVVNA FRQGLDARYGDHTNTSLAEVLRKQT SLSKRLSETSSIEYADNIPDELTIPEI DVERLSSHSHTETAV	NaN	NaN	NaN	NaN	NaN	NaN	NaN	NaN	None