

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
D1FNM8	D1FNM8_MOUSE	Mus musculus	Calcium-transporting ATPase	13.680566	NaN	NaN	22645316	MTNPPGQSVSANTVAESHEGEFGC TLMDLRKLMELRGADAVAQISAHYG GVQEICTRLKTSPIEGLSGNPADLEK RRLVFGKNVIPPKRPKTFLELVWEAL QDVTLLIILEIAAVISLVLSFYRPPGGD NEICGHIASSPEEEEEGETGWIEGAA ILASVIIIVLVTA FNDSKEKQFRGL QSRIELEQKFSIIRNGQLIQLPVAEIV VGDIAQIKYGDLLPADGILIQGNDLKI DESSLTGESDHVKKTLDKDPMLLSG THVMEGSGRMVVTAVGVNSQTGIIF TLLGASEEEDDDDKKKKGKKQGAPE NRNKAKTQDGVALEIQPLNSQEGLD SEDKKIKIARIPKKEKSVLQGLTRL AVQIGKAGLIMSVLTVVILILYFVVDN FVIQRREWLPECTPVYIQYFVKFFIIG VTVLVVAVPEGLPLAVTISLAYSVKK MMKDNNLVRHLDACETMGNATAIC SDKTGTLTMNRMTVVQAYIGGTHYR QIPQPDVFPKVLIVNGISINCAYT SKIQPPEKEGGLPRQVGNKTECGLL GFVTDLKQDYQAVRNEVPKLFKV YTFNSVRKSMSTVIRKPEGGFRMFS KGASEIMLRRCRILNKEGEIKSFRS KDRDNMVRNVIEPMASEGLRTICLA YRDFDGTPEPSWDIEGEILTSLICIAVV GIEDPVRPEVPDAIAKCKRAGITVRM VTGDNVNTARAIATKCGILTPKDDFL CLEGKEFNLSLIRNEKGEVEQEKLDKI WPKLRVLARSSPTDKHTLVKGIIDST AGEQRQVVAVTGDGTNDGPALKKA DVGFMAGIAGTDVAKEASDIILTDDN FTSIVKAVMWGRNVYDSISKFLQFQ LTVNVVAVIVAFTGACITQDSPLKAV QMLWVNLIMDTFASLALATEPPTES LLRRRPYGRNKPLISRTMMKNILGH AVYQLLIVFLLVFA GDTLFDIDSGRK APLNSPPSQHYTIVFNTFVLMQLFN EINARKIHGEKNVFA GYRNIIIFCTV VLGTFFCQIMIVELGGKPFSCSTSLTM EQWMWCLFIGIGELLWGQVISA IPT KSLKFLKEAGHGSDKEDISRDT EGM DEIDLAEMELRRGILWVRGLNRIQ TQIDVINKFQTEAPLKRVRNMTQH LDVKLVPSSYSAAVASLR TCPSISSAI SSAVTSPVVGSE