

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8C196	CPSM_MOUSE	Mus musculus	Carbamoyl-phosphate synthase [ammonia], mitochondrial	16.594134	T109;T110;S537;S872;T1078;S1331;T1332	S148;S189;S537;S540;S569;S896;S898;S1036;S1079;S1090;S1093;S1203;S1419;S1431	29562282;36064721;35285892	MTRILTACKVVKTLKSGFGFANVTTK RQWDFSRPGIRLLSVKAKTAHVLED GTKMKGYSGHPSSVAGEVVFNTGL GGYPEALTDPAYKGGILTMANPIGN GGAPDITARDELGLNKYMESDGIV AGLLVLYNSNDYNHWLTKSLGQW LQEEKVPAIYGVDTRMLTKIIRDKGT MLGKIEFEGQSVDFVDPNKQNLIAE VSTKDVKVFVGKGNPTKVVAVDCGIK NNVIRLLVKRGAEVHLPWNHDF QMEYDGLLIAGGPGNPALAQPLIQN VKKILES DRKEPLFGISTGNIITGLAA GAKSYKMSMANRGONQPVLNITNR QAFITAQNHGYALDNTLPAGWKPLF VNVNDQTNEGIMHESKPPFAVQFH PEVSPGPTDTEYLFDSFFSLIKKGGK TTITSVLPKALVASRVEVSKVLILGS GGLSIGQAGEFDYSGSQAVKAMKEE NVKTVLMNPNIASVQTNEVGLKQAD AVYFLPITPQFVTEVIKAERPDGLILG MGGQTALNCGVELFKRGVLKEYGV KVLGTSVESIMATEDRQLFSDKLNEI NEKIAPSF AVESMEDALKAADTIGYP VMIRSAYALGGLGSGICPNKETLIDL GTKAFAMTNQILVERSVTGWKEIEY EVVRDADDNCVTVCNMENVNDAMG VHTGDSVVVAPAQTL SNAEFQMLRR TSVNVVRHLGIVGECNIQFALHPTS MEYCIIEVNARLSRSSALASKATGYP LAFIAAKIALGIPLPEIKNVVSGKTS CFEPLDYMVTKIPRWDLDRFHGTS SRIGSSMKSVGEVMAIGRTFEESFQ KALRMCHPSVDGFTPRLP MNKEWP ANLDLKKELSEPSSTRIYAIKALEN NMSLDEIVRLTSIDKWFLYKMRDIL NMDKTLKGLNSDSVTEETLRKAKEI GFSKQISKCLGLTEAQTRELRLKK NIHPWVKQIDTLAAEYPSVTNYLYVT YNGQEHDIKFDEHGIMVLGCGPYHI GSSVEFDWCAVSSIRTLRQLGKKT VVNCNPETVSTDFDECDKLYFEELS LERILDYHQAECNGCIISVGGQIPN NLAVPLYKNGVKIMGTSPLOIDRAE DRSIFSAVLDELKVAQAPWKAVNTL NEALEFANSVGYPCLLRPSYVLSGSA MNVVFSSEDEMKRFLAEATRVSQEH PVVLTKFVEGAREVEMDAVGKEGRV ISHAISEHVEDAGVHSGDATLMLPT QTISQGAIEKVKDATRKIAKAF AISGP FNVQFLVKGNDVLVIECNLRASRSF PFVSKTLGVDFIDVATKVMIGESIDE KRLPTLEQPIIPSDYVAIKAPMFSWP RLRDADPILRCEMASTGEVACFGEGI HTAFLKAMLSTGFKIPQKGIIGIQQ SFRPRFLGVAEQLHNEGFKLFATEA TSDWLNANNVPATPVAWPSQEGQN PSLSSIRKLIRDGIDLVINLPNNNTK FVHDNYVIRRTAVDSGIALLTNFQVT KLFAEAVQKSRVDSKSLFHRYQYS AGKAA