

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P07756	CPSM_RAT	Rattus norvegicus	Carbamoyl-phosphate synthase [ammonia], mitochondrial	23.439218	NaN	S148;S189;S537;S540;S569;S896;S898;S1036;S1079;S1090;S1093;S1203;S1419;S1431	38843836	MTRILTACKVVKTLKSGFLANVTSK RQWDFSRPGIRLLSVKAQTAHVILED GTKMKGYSGFHPSSVAGEVVFNTGL GGYSEALDTPAYKQILTMANPIIGN GGAPDITARDELGLNKYMESDGIKV AGLLVLNYSHDYNHWLTKSLGQW LQEEKVPAIYVDTRMLTKIIRDKGT MLGKIEFEGQSVDFVDPNKONLIAE VSTKDVKVFSGKNPTKVVAVDCGIK NNVIRLLVKGAEVHLVFNHDFDFT QMDYDGLLIAGGPGNPAQAQLIQN VKKILESDRKEPLFGISTGNIITGLAA GAKSYKMSMANRGONQPVLNITNR QAFITAQNHGYALDNTLPAGWKPLF VNVNDQTNEGIMHESKPPFAVQFH PEVSPGPTDTEYLFDSFFSLIKKGGK TTITSVLPKPAVASRVEVSKVLILGS GGLSIGQAGEFDYSGSQAVKAMKEE NVKTVLMNPNIASVQTNEVGLKQAD AVYFLPITPOFVTEVIKAERPDGLILG MGGQTALNCGVELFKRGLKEYGV KVLGTSVESIMATEDRQLFSDKLEI NEKIAPSAVESMEDALKAADTIGYP VMIRSAYALGGLSGGICPNKETLMD LGTKAFAMTNQILVERSVTGWKEIE YEVVRDADDNCVTCNMENVDAM GVHTGDSVVVAPAQTLNSNAEFQMLR RTSINVVRHLGIVGECNIQFALHPTS MEYCIIEVNARLRSALASKATGYP LAFIAAKIALGIPLEIKNVVSGKTS CFEPSLDYMTKIPRWDLDRFHGTS SRIGSSMKSVGEVMAIGRTFEESFQ KALRMCHPSVDGFTPRLPNMKEWP ANLDRKELSEPSSTRYIAIAKALEN NMSLDEIVKLTSDIKWFLYKMRDIL NMDKTLKGLNSESVTEETLRQAKEI GFSDKQISKCLGLTEAQTRELRLKK NIHPVWKQIDTLAAEYPSVTNYLYVT YNGQEHDIKFDEHGIMVLGCGPYHI GSSVEFDWCAVSSIRTLRQLGKKT VVNCNPEVTVSTDFDECCKLYFEELS LERILDYHQAECNGCHISVGGQIPN NLAVPLYKNGVKIMGTSPLQIDRAE DRSIFSAVLDELKVAQAPWKAVNTL NEALEFANSVGYPCLLRPSVYLSGSA MNVVSEDEMKRFLEEATRVSQEH PVVLTKFIEGAREVEMDAVGKEGRVI SHAISEHVEDAGVHSGDATLMLPTQ TISOGAIEKVKDATTRIAKAFASGPF NVQFLVKGNDVLECNLRASRFP FVSKTLGVDFIDVATKVMIGESVDEK HLPTLEQPIPSDYVAIKAMPFSWPR LRDADPILRCEMASTGEVACFGEGI HTAFLKAMLSTGFKIPQKILIGIQQ SFRPRFLGVAEQLHNEGFKLFATEA TSDWLNANNVPATPVAWPSQEQGN PSLSSIRKLIRDGSIDLVINLPNNNTK FVHDNYVIRRTAVDSGIALLTNFQVT KLFAEAVQKARTVDSKSLFHRYQYS AGKAA	True	False	2.608	3.615	5.0	1.183	False	3.474	1.276