

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P31327-3	CPSM_HUMAN	Homo sapiens	Isoform 3 of Carbamoyl-phosphate synthase [ammonia], mitochondrial	19.616108	NaN	S148;S537;S540;S569;S835;S896;S898;S1036;S1079;S1090;S1093;S1203;S1419;S1431	30379171;16408927;36064721;26374642;35083852;26853435;33465208;28411811;34019948	MPQIIKMRILTAFKVVRTLKTGFQFNVTAAHQKWKFSRPGIRLLSVKAQT AHIVLEDGTKMKGYSGHPSSVAGE VVFNTGLGGYPEAITDPAYKGQILTM ANPIIGNGGAPDTTALDELGLSKYLE SNGIKVSGLLVLDYSKDYNHWLWATK SLGQWLQEEKVPAIYGVDTRMLTKII RDKGTMLGKIEFEGQPVDVDPNKQ NLIAEVSTKDVVYKGNPTKVVAV DCGIKNNVIRLLVKRGAEVHLVLPWN HDFTKMEYDGIAGGPGNPALAEPLIQNVRKILESDRKEPLFGISTGNLIT GLAAGAKTYKMSMANRGQNQPVLN ITNKQAFITAQNHGYALDNTLPAGW KPLFVNVDQTNNEGIMHESKPPFAV QFHPEVTPGPIDTEYLFDSFFSLIKK GKATTTITSVLPKALVASRVEVSKVLI LGSGGLSIGQAGEFDYSGSQAVKAM KEENVKTVLMNPNIASVQTNEVGLK QADTVYFLPITPQFVTEVIKAEQPDG LILGMGGQTALNCGVELFKRGVLKE YGVKVLGTSVESIMATEDRQLFSDK LNEINEKIAPSFIVESIEDALKAADTI GYPVMIRSAYALGGLGSGICPNRETL MDLSTKAFAMTNQILVEKSVTGWKE IEYEVVRDADDNCVTVCNMENVDA MGVHTGDSVVVAPAQTLNSNAEFQM LRRTSINVVRHLGIVGECNIQFALHP TSMEYCHIEVNARLSRSSALASKATG YPLAFIAAKIALGIPLPEIKNVVSGKT SACFEPPLDYMVTKIPRWDLDRFHG TSSRIGSSMKSVGEVMAIGRTFEESF QKALRMCHPSIEGFTPRLPMNKEW PSNLDLRKELSEPSSTRIYAIKAIDD NMSLDEIEKLYIDKWFLYKMRDIL NMEKTLKGLNSESMTEETLKRAKEI GFSDKQISKCLGLTEAQTRELRLKK NIHPWVKQIDTLAAEYPSVTNYLYVT YNGQEHVDVFDHGMVVLGCGPY HIGSSVEFDWCAVSSIRTLRQLGKKT VVVNCPETVSTDFDECCKLYFEEL SLERILDIYHQEACGGCIISVGGQIPN NLAVPLYKNGVKIMGTSPLQIDRAE DRSIFSAVLDELKVAQAPWKAVNTL NEALEFAKSVDPCLLRPSYVLSGSA MNVVFSEDEMCKFLLEATRVSQEH PVVLTKFVEGAREVEMDAVGKDGRV ISHAISEHVEDAGVHSGDATLMLPT QTISQGAIEKVKDATRKAFAISGP FNVQFLVKGNDVLVIECNLRASRF PFVSKTLGVDFIDVATKVMIGENVDE KHLPTLDHPIIPADYVAIKAPMFSWP RLRDADPILRCEMASTGEVACFGEI HTAFLKAMLSGFKIPQKILIGIQQ SFRPRFLGVAEQLHNEGFKLFATEA

TSDWLNANNVPATPVAWPSQEGQN
PSLSSIRKLIRDGSIDLVINLPNNNTK
FVHDNYVIRRTAVDSGIPLLTNFQVT
KLFAEAVQKSRKVDSKSLFHRYQYS
AGKAA