

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P11586	C1TC_HUMAN	Homo sapiens	C-1-tetrahydrofolate synthase, cytoplasmic	18.555225	NaN	S318;S413;S490	18369606;34019948;16408927;34725712;26853435;28510447;32119511;38665916;30397120;37217939;3214551;35138101	MAPAEILNGKEISAQIRARLNQVQTQ LKEQVPGFTPRLAILQVGNRDDS NL YINVKLKAAEEIGIKATHIKLPR TTTE SEVMKYITSLNEDSTVHGFLVQL PL DSENSINTEEVINAI APEKDVDG LTSI NAGKLARGDLNDCFIPCTPKGC LELI KETGVPIAGRHAVVGRSKIVGAP M HDL LLWNNATVTTCHSKTAHLDE E VNKGDILVVATGQPEMVKGWIKP G AIVIDCGINYVPDDKKPNGRKVV GDV AYDEAKERASFITVPPGGVGPMT VA MLMQSTVESAKRFLEKFKPGKWM I QYNNLNLKTPVPSDIDISRCKPKP I GKLAREIGLLSEEVELYGETKAKV LL SALERLKHRPDGKYVVVTGITPTPL G EGKSTTTIGLVQALGAHLYQNVFAC VRQPSQGPTFGIKGGAAGGGYSQV IP MEEFNHLTGDIHAITAANNLVAAAI DARIFHELTQTDKALFNRLVPSVNG VRRFSDIQIRRLKRLGIEKTDPTLTD EEINRFARLDIDPETITWQRVLDTND RFLRKITIGQAPTEKGHTRTAQFDIS VASEIMAVLALTT SLEDMRERLGKM VVASSKKGEPVSAEDLGVSGALT V MKDAIKPNLMQTLEGTPV FVHAGPF ANIAHGNSIIADRIALKLVGPEGFV V TEAGFGADIGMEKFFNIKCRYSGLC PHVVVLVATVRALKMHGGGPTV TAG LPLPKAYIQENLELVEKGF SNLKKQI ENARMF GIPVVAVNAFKTDTESEL DLISRLSREHGAFDAVKCTHWAEGG KGALALAAQAVQRAAQAPSSFQLLYD LKL PVEDKIRIIAQKIYGADDIELL PEA QHKAEVYTKQGFGNLPICMAKTHLS LSHNPEQKGVPTGFILPIRDIRASV G AGFLYPLVGT MSTMPGLPTRPCFYDI DLDPETE QVNGLF

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