

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8BG87	TET3_MOUSE	Mus musculus	Methylcytosine dioxygenase TET3	34.48173	NaN	NaN	22645316;36852467;37903183;25568311	MSQFQVPLAVQPDLSGLYDFPQGQV MVGGFQGPGLPMAGSETQLRGGGD GRKKRKRRCGTCDPCRRENCGSCTS CTNRRTHQICKLRKCEVLKKAGLL KEVEINAREGTGPWAQGATVKTGSE LSPVDGPVPGQMDSGPVYHGDSRQ LSTSGAPVNGAREPAGPGLLGAAGP WRVDQKPDWEAASGPTHAAARLEDA HDLVAFSAVAEAVSSYGALSTRLYET FNREMSREAGSNGRGRPRPESCSEG SELDLTLQTALALARHGMKPPNCTC DGPECPDFLEWLEGKIKSMAMEGG QGRPRLPGALPPSEAGLPAPSTRPPL LSSEVPQVPPLEGLPLSQSALSIAKE KNISLQTAIAIEALTQLSSALPQPSHS TSQASCPLPEALSPSAPFRSPQSYLR APSWPVVPEEHPSFAPDSPAFPPAT PRPEFSEAWGTDTPPATPRNSWPVP RPSDPMAELEQLLGSASDYIQSVFK RPEALPTKPKVKVEAPSSSPAPVPSPI SQREAPLLSSEPDTHQKAQTALQOH LHHKRNLFLEQAQDASFPTSTEPQA PGWWAPPGSPAPRPPDKPPKEKKKK PPTPAGGPVGAEKTTTPIKTSVRKPI QIKKSRSRDMQPLFLPVROIVLEGLK PQASEGQAPLPAQLSVPPPASQGAA SQSCATPLTPEPSLALFAPSPSGDSL LPPTQEMRSPSPMVALQSGSTGGPL PPADDKLEELIRQFEAEFGDSFGLPG PPSVPIQEPENQSTCLPAPESPFATR SPKKIKIESSGAVTVLSTTCFHSEEG GQEATPTKAENPLTPTLSGFLESPLK YLDTPTKSLLDTPAKKAQSEFPTCDC VEQIVEKDEGPYYTHLGSPTVASIR ELMEDRYGEKGAIRIEKVIYTGKEG KSSRGCPIAKWIRRHTLEKLLCLV RHRAGHHCQNAVIVILILAWEGIPRS LGDPLYQELDTLTKYGNPTSRRCG LNDDRTCACQKDPNTCGASFSG CSWSMYFNGCKYARSKTPRKFRLTG DNPKEEEVLRNSFQDLATEVAPLYK RLAPQAYQNQVTNEDVAIDCRLGLK EGRPFSGVTACMDFCAHAHKDQHN LYNGCTVVCTLTKEFNRCVGOIPED EQLHVLPLYKMASTDEFGSEENQN AKVSSGAIQVLTAFPREVRRRLPEPAK SCRQRQLEARKAAAEKKLQKEKLS TPEKIKQEALAGVTTDPGLSLKGG LSQQSLKPSLKVEPQNHFFSSFYKSG

