

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
043151	TET3_HUMAN	Homo sapiens	Methylcytosine dioxygenase TET3	24.568079	S381;T431;T439;T441;T445;S1379	NaN	35254053;35289036;37340703;24394411;30620550;35132862;24304661;30379171	MSQFQVPLAVQPDLPGLYDFPQRQVMVGSFPGSGLSMAGSESQLRGGGDGRKKRKRKCGTCEPCRRLENCGACTSCTNRRRTHQICKLRKCEVLKKKVGLLKEVEIKAGEGAGPWGQGAAVKTGSELSPVDGPVPGQMDSGPVYHGDSRQLSASGVPVNGAREPAGPSLLGTGGPWRVDQKPDWEAAPGPAHTARLEDAHDLVAFSAVAEAVSSYGALSTRLYETFNREMSREAGNNSRGRPRGPEGCSAGSEDLDTLQTALALARHGMKPPNCCDGPCEPDYLEWLEGGKIKSVMEGGEERPLPGPLPPGEAGLPAPSTRPLLSSEVPQISPOEGLPLSQSALSIAKEKNISLQTAIAIEALTQLSSALPQPSHSTPQASCPLPEALSPPAPFRSPQSYLRAPSWPVVPPEEHSSFAPDSSAFPPATPRTEFPEAWGTDTPPATPRSSWPMPRPSDPMAELEQLLGSADYIQSVFKRPEALPTKPKVKVEAPSSSPAPAPSPVLQREAPTSPSEPETHQKAQTALQQHLHHKRSLFLEQVHDTSFAPSEPSAPGWPPPSSPVRLPDRPPKEKKKKLPTPAGGPVGTAKAAPGIKPSVRKPIQIKKSRPREAQPLFPPVRQIVLEGLRSPASQEVQAHPPAPLPASQGSVAVPLPPEPSLALFAPSPSRDSSLPTQEMRSPSPMTALQPGSTGPLPPADDKLEELIRQFEAEFGDSFGLPGPPSVPIQDPENQQTCLPAPESPFATRSPKIQIESSGAVTVLSTTCFHSEEGQEATPTKAENPLTPTLSGFLESPLKYLDTPTKSLDTPAKRAQAEFPTCDCVEQIVEKDEGPYYTHLGSQPTVASIRELMEERYGEKGAIRIEKVIYTGKEGKSSRGCPIAKWVIRRHTLEEKLLCLVRHRAGHHCQNAVIVILILAWEGIPRSLGDTLYQELTDLRKYGNPSTRRCGLNDDRTCACQGKDPNTCGASFSGCSWSMYFNGCKYARSKTPRKFRLAGDNPKEEEVLKRSFQDLATEVAPLYKRLAPQAYQNQVTNEEIAIDCRLGLKEGRPFAGVTCMDFCAHAHKDQHNLYNGCTVVCTLTKEDNRCVGKIPEDQLHVLPLYKMANTDEFGSEENQNAKVGSGAIQVLTAFPREVRRLPEPAKSCRQRQLEARKAAAEEKKIQKEKLTPEKIKQEALELAGITSDPGLSLKGGLSQQGLKPSLKVEPQNHFSFKYSGNAVVESYSVLGNCRPSDPYSMNSVYSYHSYQAQPSLTSVNGFHSKYALPSFSYGFSSNPVFPSQFLGPGAWGHSGSSGSFEKKPDLHALHNSLSPAYGGAFAELPSQAVPTDAHHTPHHQPPAYPGPKEYLLPKAPLLHSVSRDPSFPAQSSNCYNRSIKQEPVDPLTQAEPVPRDAGKMGKTPLSEVSQNGGPSHLWGQYSGGSP

MSPKRTNGVGGSWGVFSSGESPAIV
PDKLSSFCASCLAPSHFTDGGWGLF
PGEGQQAASHSGGRLRGKPWSPCK
FGNSTSALAGPSLTEKPWALGAGDF
NSALKGSPGFQDKLWNPMKGEEGR
IPAAGASQLDRAWQSFGLPLGSSEK
LFGALKSEEKLWDPFSLEEGPAEEP
PSKGAVKEEKGGGGAEIEEEELWS
DSEHNFLDENIGGVAVAPAHGSILIE
CARRELHATTPLKPNRCHPTRISLV
FYQHKNLNQPNHGLALWEAKMKQ
LAERARARQEEAARLGLGQQEAKLY
GKKRKWGGTVVAEPQQKEKKGVP
TRQALAVPTDSAVTVSSYAYTKVTGP
YSRWI