

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3UHK8	TNR6A_MOUSE	Mus musculus	Trinucleotide repeat-containing gene 6A protein	25.122525	S1622;T1623	S724;S863;S976;S1197;S1255;T1406;S1520;S1804;S1825	21606357;30059200;22517741	MRELEAKATKDVERNLSRDLVQEEE QLMEEKKKKKDDKKKKEAAQKKAT EQKIKVPEQIKPSVSOQPANSDNGT STATSTNNNAKRATASNOQPPPPQQ QQPQQEQQQQPQALPRYPREVPPR FRHQEHKQLLKRQHFVIAANLGS AVKVLNSQSESSAVTNQQPQNNGE VQNSKQSODINHNTSGSHYENCQR GPVSSTSDCSTSCKNAVNDLLEKEA WPSAPGSDPELAPECIDADASANSE SERNITVMASGNTGGEKDGLRNSTG LGSQSKFVVGSSNNVGHGSSTGP WGFPHGALISTCQVSDAPESKPES SNNRMNAWGTVSSSSNGGLNPSTL NSASNHGAWPVLENNGLALKGPVG SGSSGINIQCSTIGMPNNQINISK VSGSSTHGTWGSLOETCEPEVSGTQ KVSFSGQPQNITTETTGNNTTNFM TSSLPNSGSVQNNELPTSNPGAWRV STMNHPQIQAPVMNGTSLSHLSN GESKTGGSYGTTWAGYGSNYSGDKC AGPNGQANGDTVNATLMQPGINGP MGTNFQVNTNKGKGVWEPGTVNS QSSPWGSGNGANSRGGRRGWGSPA QNTGTGLSSVEWNLPSNQHSNDS ANGNGKLTNGWKSTEEDDQGSAT SQTNEQNSVWAKAGGTVESDGSAE STGRLEEKVTGESQSRDRRKIDQHT LLQSIVNRTDLDPVLSNSGWGQTPI KQNTAWDTETS PRGERKTDNGTEA WGSSATQTFNSGACTDKTSPNSND TSSVSGWGDPKPTLRWGDSKGSNC QGGWEDDSAATGMIKSNQWGGCK EDKSTWNDSQKSKQGWGDGQKSS QGWSISAGDNWGESSRSNHWGEA NKKSSSGSDSDRSISGWNELGKTS SFTWGNINPNNSGWDDESSKPNS SQGWGDPPKCNQSLGWGDSSKPVS SPDWNKQQDIVGSWGIPPATSKPPG TGWLGGPIPAKKEEPTGWEEPPS ESIRRKMEIDDGTSAWGDP SKYNYK NVNMWKNKIPEASGRSDQQAQMH RLLPAASAVSSKETSSSGWGEPWA EPSTPATTVDNGTSAWGKPIDSGPS WGEPITAASNASTWSSSVGPQSLS KSGPKSMQDGCWGDMDPLGSRPT GWEEEEDEIGMWNSNSSQELNSS LNWPPYTKMSSKGLSGKKRRRER GMMKGGNKQEDAWINPFVKQFSNI SFSRDSPEENVQSNKMDLSGGMLQ DKRMEIDKHS LNIGDYNRTVKGKGP SRPQISKESMERNPYFDKNGNPN MFGVGNTAAQPRGMQQPPAQLSS SQPNLRAQVPPLLSPQVPVSLKYA PNNGGLNPLFGPQQVAMLNQLSQL

								NQLSQISQLQRLLAQQRAQSQRSA PSANRQQQDQQGRPLSVQQMMQ QSRQLDPSLLVKQQTTPPSQQPLHQP AMKSFLDNVMPHTTPELQKGPSPV NAFSNFPIGLNSNLNVNMDMNSIK EPQSRLRKWTTVDSMSVNTSLDQN SSKHGAISSGFRLEESPFVPHYDFMN SSTSPASPPGSIGDGWPRAKSPNGSS SVNWPPEFRPGEPWKGYPNIDPETD PYVTPGSVINSLSTINTVREVDHLRDR NSGSSSSLNNTLPTSAWSSIRASNY NVPLSSTAQSTSARNSDSKLTWSPG SVTNTSLAHELWKVPLPPKNITAPSR PPPGLTGQKPPLSTWDNSPLRVGGG WGNSDARYTPGSSWGESSGRITN WLVLKNLTPQIDGSTLRITLCMQHGP LITFHLNLPHGALVRYSSKEEVKA QKSLHMCVLGNTTILAEFASEEEISR FFAQSQSLTPSPGWQSLGSSQSRLG SLDCSHSFSSRTDVNHWNAGLSG ANCGDLHGTSWGTPHYSTSLWGP PSSDPRGISSPINAFLSVDHLGGG GEM
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