

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9Y4A5	TRRAP_HUMAN	Homo sapiens	Transformation/transcription domain-associated protein	3.245925	S1139	S1628;S2051;S2077	30379171;34019948	MAFVATQGATVVDQTTLMKKYLQFV AALTDVNTPEDETKLKMMEVSENF ENVTSSPQYSTFLEHIIPRFLTFLQD GEVQFLQEKPAQQLRKLVEIIHRIP TNEHLRPHTKNVLSVMFRFLETENE ENVLICLRIIIHLHKQFRPPITQEIHH FLDFVKQIYKELPKVNVNRYFENPQVI PENTVPPPEMVGMITTIIVKVNPER EDSETRTHSIIPRGSLSLKVLAELPIIV VLMYQLYKLNHNVAEFVPLIMNTI AIQVSAQARQHKLYNKELYADFIAAQ IKTLSFLAYIIRIYQELVTKYSQQMVK GMLQLLSNCPAETAHLRKELLIAAK HILTTELRNQFPCMDKLFDESILIGS GYTARETLRPLAYSTLADLVHHVRQ HLPLSDLSLAVQLFAKNIDDESLPSS IQTMSCKLLLNLVDCIRSKSEQESG NGRDVLMRMLEVFLKFHTIARYQL SAIFKCKPQSELGAVEAALPGVPTA PAAPGPAPSPAPVPAPPPPPPPPPAT PVTAPVPPFEKQGEKDKEDKQTFQ VTDCRSLVKTLCVGVKTTITWGITSCK APGEAQFIPNKQLQPKETQIYIKLVK YAMQALDIYQVQIAGNGQTYIRVAN CQTVRMKEEKEVLEHFAGVFTMMN PLTFKEIFQTTVPYMERISKNYALQI VANSFLANPTTSALFATILVEYLLDR LPEMGSNVELSNLYLKLFLVFGSV SLFAAENEQMLKPHLHKIVNSSMEL AQTAKEPYNIFLLRALFRSIGGGSH DLLYQEFPLLPNLLQGLNMLQSGL HKQHMKDLFVELCLTVPVRLSSLLP YLPMLMDPLVSALNGSQTIVSQGLR TLELCVDNLQPDFLYDHIQPVRAEL MQALWRTLRLNPADSISHVAYRVLGK FGGSNRKMLKESQKLHYVVTEVQG PSITVEFSDCKASLQLPMEKAIETAL DCLKSANTEPYRRQAWSEVIKCFV AMMSLEDNKHALYQLLAHPNFTEK TIPNVIIISHRYKAQDTPARKTFEQALT GAFMSAVIKDLRPSALPFVVASLIRHY TMVAVAQCGPFLPCYQVGSQPST AMFHSEENGSKGMDPLVLIDAIAC MAYEEKELCKIGEVALAVIFDVASII GSKERACQLPLFSYIVERLCACCYEQ AWYAKLGGVVSIFLMLERLPLTWVL QNQQTFLKALLFVMMDLTGEVSNQ AVAMAKTTLEQLLMRCATPLKDEER AEEIVAAQEKSFHHVTHDLVREVT PNSTVRKQAMHSLQVLAQVTGKSVT

VIMEPHKEVLQDMVPPKKHLLRHQ  
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