

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q925H1	TRPS1_MOUSE	Mus musculus	Zinc finger transcription factor Trps1	15.818779	S803	S90;S127;S178;S216;S978;S1041;S1066;S1085	36852467	MVRKKHPPLRNVASEGEGQTLEPTA TESKVSQKNEKLSADQMSSENTDQS DVAELNSKEEHSTHGQEPSSSGKKD LQISGLSEKAGFNYESPSKGGSLVSF PHDEVTDNRNMLAFSSPAAGGVCEPL KSPQRAEADDPQDMACTPSGDSLET KEEHKMSPKATEETGPVQSGQANC QGLSPVSVASKNPQVPSDGGVRLSK PKGDLLVNDNPDPAPLSPELQDFKC NICGYGYGNDPTDLIKHFRKYHLG LHNRTRQDAELDSKILALHNMVQFS HSKDFQKVNRSVLSGVLQDISSSRP ALLNGTYDVQVTSGGTFIGIRKTPD CQGNTKYFRCKFCNFTYMGNSSTE LEQHFLQTHPNKIKVSLPSSEGVKPS EKNSNKSIPALRASDSGDVKGWQDK MTVKAGDDTPVGYSVPIKPLDSSRQ NGTEATSYYWCKFCFSCESSSLK LLEHYGKQHGAVQSGGLNPELNDK LPRGSVINQNDLAKSVEGEPLTKPE KGLSGAKKKDFPSKGAEDNMVTSY NCQFCDFRYSKSHGPDVIVVGPLL HYQQLHNIHKCTIKHCPFCPRGLCS PEKHLGEITYPFACRKSNCSHCALLL LHLSPGVAGSSRVKHQCHQCSFSTP DVDVLLFHYETVHESQASDVKQEAN HLLGSDGQQAVRDSKEHSCTKCDFI TQVEEEISRHYRAHSCYKCRQCSF TAADTQSLLEHFNTVHCQEITTA NGEEGGHAIPTIKEPKIDLKVYSL NPDSKMGETVPESIVKREKLDDKEG LKDKIWTESSTDDLGRVAWRGADIL RGSPSYTQASLGLLTPVSSSQEQTKT LRDSPNVEAAHLARPMYGLAVDTKG FLQGAPAGSEKSASLTQQYPAGES KTKDESQSLRRRRRGSGVFCANCLT TKTSLWRKNANGGYVCNACGLYQK LHSTPRPLNIIKQNNGEQIIRRRTRK RLNPEALQAEQLNKQQRGSQEEQV NGSPLERRSEDHLSSEHPREIPLPSL SKYEAQGS�TKSHSAQQPVLVSQAL DIHKRMQPLHIQIKSPQESTGDPGN SSSVSDGKGSSERGSPIEKYMRPAK HPNYSPPGSPIEKYQYPLFGVPPFVHN DFQSEADWLRFWFSKYKLSVPGNPH YLSHVPGLPNPCQNYVPYPTFNLPP HFSAVGSDNDIPLDLAIKHSRPGPTA NGASKEKTKAPPTVKNEPDLNVVKT EKVDRSTQDELSTKCVHCGIVFLDE

