

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q3TUF7	YETS2_MOUSE	Mus musculus	YEATS domain-containing protein 2	36.92943	S371;S526;S600;T601;T602;S668;S701;T718;T939;T973;T998;S1004	S118;S120;S157;T406;S446;S462;S464;S470;S472;T477;S534;S573;S625;T1204	22645316;36852467;21606357;34678516;35822049;22517741	MSGIKRTIKETDPDYEDVSVALPNKR HKAIESSARDAAVQKIETIIEQFALE MKNKEHEIDVIDQRLIEARRMMDKL RACIVANYASAGLLKVSEGLKTFDP MAFNHPAIKKFLESPSRSSSPTNQR SETPSANHSESDSLSQHNDFLSDKD NNSNVDVEERPSTGEQRPSRKAGR DTSSISGSHKRELRNADLTGDETSRL FVKKTIVVGNVSKYIPDPKREENDQS THKWMVYVGRSRREPSINHFVKKV WFFLHPSYKPNDLVEVREPPFHLTR RGWGEFPVRVQVHFKDSQNKRIDII HNLKLDRTYTGLQTLGAETVVDVEL HRHSLGEDSVYPQSSSDVCDAPPP TLTLPAAVKASAVAQSPAAAAAPVG EGFPETTEAERHSTFYSLPSSLERTP TKVTTAQKVTFSSHGNSAFQPIASSC KIVPQSQVNPESPGKSFQIPITMSCK IVSGSPISTPPSPPLRPTPTSTPVHLK QGTASSGVSNPHVIVDKPGQVIGAST PSTGSPTSKLPVASQASQGTGSPIPKI HGSSFLTSTVKQEESELFASMPPLCPI GSHPKVQSPKAVTGGGGAFTKVIKQ EPGEAPHVSTTGAASQSAFPQYVTV KGGHMIAVSPQKQVISAGEGTTQSP KIAPSKVVGVPVGSALPSTVKQAVAI SSGQILVAKASSSVTKAVGPKQVVTQ GVAKAIVSGGGGTIVAQPVQTLTKTQ VTAAGPQKSGSQGSVMATLQLPATN LANLANLPPGTCLYLTTNSKNPSGK GKLLLIPOGAILRATNNANLQSGSAA AGGSGSSGAGGGSGGGGGSGAGGT PSTSGPGGGPQHLTYTSYILKQTPQG TFLVGQPSQTPGKQLTTASVVQGTL GVSSSAQGGQTLKVISGQKTLTFTQ AATAGQASLLKLPDNTLKSVPAAAPQL AKPGTTMLRVAGGVITAAPSPAVAFS ANGAVHQSEGSTPVSSSVGSIKTPG QPQVCVSQATMATCKGPAAVAGTAA SLVSAPSSISGKATVSGLLKVHSAQS SPQQAVLTIPSQLPLSINTSGGVQT VLMPVNKVVQSFSTSKLPTTVLPISV PNQAAPSSAPVAIAKVTEPETPGPN CISQENQVAVKTEESSELSNYVIKVD HLETIQLLTAVVKKIPLITAKGDDAS CFSAKSLEQYYGWNIGKRRAAEWQR AMTVRKVLQEILEKNPRFHHLTPLK TKHIAHWCRCCHGYTPDPESLRHDG DSIEDVLTQIDSEPECLSSFSTADDL CRKLEDLQQFQKREPENEEVDILS LSEPLKTNIKKEQEEKQEEMRFYLP PTPGSGFVGDITQKIGITLQPVALHR NMYASVVEDMILKATEQLVSDILRQ ALAVGYQTASPNRIPKEITVSNIHQAI CNIPFLDFLTNKHMGRNLNEDQ