

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi appa
Q9Y6A5	TACC3_HUMAN	Homo sapiens	Transforming acidic coiled-coil-containing protein 3	28.004703	T17;S424	S25;S39;S71;S175;S177;S250;S317;S402;S434;S558	23301498;40307207;34019948;31373491	MSLQVLNDKNVSNKNTENCDFLF SPPEVTGRSSVLRVSKENVPPKNL AKAMKVTFQTPLRDPQTHRLSPSM ASKLEAPFTQDDTLGLENSHPVWQ KENQQLIKEVDAKTTHGILQKPV DTDLLGDASPAFGSGSSSESGPGAL ADLDCSSSSQSPGSSSENMVSPGKV SGSPEQAVEENLSSYSLDRRVTASE TLEDPCRTEAQHKAETPHGAEIECK AETPHGAEIECRHGGVCAPAAVATS PPGAIKPEACGGAPLQGLPGEALGCP AGVGTVPADGTQTLTCAHTSAPES TAPTINHLVAGRAMTLPQEEVAAGQ MASSRSRSGPVKLEFDVSDGATSKRA PPRRRLGERSGLKPPLRKA AVRQK APQEV EDDGRSGAGEDPPMPASR GSYHLDWDKMDDPNFIFFGGDTKS GCSEAAQPPESPETRLGQPAEQLHA GPATEEPGPCLSQLHSASAEPTV VQLAAETPTAESKERALNSASTSLPT SCPGSEVPVTHQQGQPALELKEESF RDPAEVLGTGAEV DYLEQFGTSSFK ESALRKQSLYLKFDPLLRDSPGRPVP VATETSSMHGANETPSGRPREAKLV EFDFLGALDIPVPGPPPVPAPGGPP LSTGPIVDLLQYSQKDLDAVVKATQE ENRELRSRCEELHGKNLELGKIMDR FEEVVYQAMEEVQKQKLSKAEIQK VLKEKDLITDLNSMEKSFSDLFKR FEKQKEVIEGYRKNEESLKKCVEDY LARITQEQRYQALKAHAEKQLQA NEEIAQVRSKAQAEALALQASLRKE QMRIQSLEKTVEQTKENEELTRIC DDLISKMEKI	True	False	4.718	1.641	False	False	4.0