

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q9HC35	EMAL4_HUMAN	Homo sapiens	Echinoderm microtubule-associated protein-like 4	13.68832	S869;T882	S7;S13;S16;S61;T96;S134;S144;S146;S171;S200;T201;T226;T237;T490;T609;S891;S895;T897;T899;S903;S978;S981	23301498;35254053;28657654;33214551;28510447;34725712	MDGFAGSLDDSSISAASTSDVQDRLS ALESRVQQQEDEITVLKAALADVLRRLA LAISEDHVASVKKSVSSKQGPSRAV IPMSCITNGSGANRKPSTSAVSIAG KETLSSAAKSGTEKKKEKPOGQREK KEESHNSNDQSPQIRASPSQPSSQPL QIHRQTPESKNATPTKSIKRPSPAEK SHNSWENSDDSRNKLSKIPSTPKLI PKVTKTADKHKDVIINQEGEYIKMF MRGRPITMFI PSDVDNYDDIRTELPP EKLKLEWAYGYRGKDCRANVYLLPT GKIVYFIASVVVLFNVEERTQRHYLG HTDCVKCLAIHPDKIRIATGQIAGVD KDGRPLQPHVRVWDSVTLSTLQIIGL GTFERGVGCLDFSKADSGVHLCIIDD SNEHMLTVWDWQKKAKGAEIKTTN EVLAVEFHPTDANTIITCGKSHIFF WTWGSNSLTRKQGFYKYEKPKFVQ CLAF LGNDVLTGDSGGVMLIWSKT TVEPTPGKGPVYQISKQIKAHDGS VFTLCQMRNGMLLTGGGKDRKIILW DHDLNPEREIEVPDQYGTIRAVAEG KADQFLVGT SRNFILRGTFNDGFQIE VQHTDELWGLATHPFKDLLLTCAQ DRQVCLWNSMEHRLEWTRLVDEPG HCADFHPSGTVVAIGTHSGRWVFLD AETRDLVSIHTDGNEQLSVMRYSID GTFLAVGSHDNFIYLYVSENGRKYS RYGRCTGHSSYITHLDWSPDNKYIM SNSGDYEILYWDIPNGCKLIRNRSDC KDIDWTYTCVLGFQVFGVWPEGSD GTDINALVRSHNRKVIAVADDFCKV HLFYPCSKAKAPSHKYSAHSSHVT NVSFTHNDSHLISTGGKDMSHIQWK LVEKLSLPQNETVADTTLT KAPVSST ESVIQSNTPTPPSQPLNETAEESR ISSPTLLENSLEQTVEPSEDHSEEE SEEGSGDLGEPLYEPCNEISKEQA KATLLEDQQDPSPSS