

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus
Q9Y2J2	E41L3_HUMAN	Homo sapiens	Band 4.1-like protein 3	16.361899	T953;T980	S88;S420;S443;S460;T469;T492;T706;S708;S960;S962;T1081	28657654;33214551;35138101;34019948;34725712;32119511;38665916	MTTESGSDSESKPDQEAEPQEAAGA QGRACAPVPEPPKEEQQALEQFAA AAAHSTPVRREVTDKQEFAARA AAK QLEYQQLEDDKLSQKSSSKLSRSP LKIVKKPKSMQCKVILLDGSSEYTC DVEKRSRQVLFDFKVCHEHLNLEKDYF GLTYRDAENQKNWLDPAKEIKKQVR SGAWHFSFNKVFYPPDPAQLSE DIT RYYLCLQLRDDIVSGRLP CSFVTLAL LGSYTVQSELGDYDPDECGSDYISEF RFAPNHTKELEDKVI LHKSHRGM T PAEAEMHFLEN AKKLSMYGVDLHH AKDSEGVEIMLGV CASGLLIYRDLR L INRF AWPKVLKISYKRN NFYKIRPG EFEQFESTIGFKLP NHRAAKRLWKV C VEHHTFFRLLPE APPKFLTLGSK FRYSGRTQAQTR RASALIDRP PAPYFE RSSSKRYTMSRSLD GVEVGTQYATT KGISQTNLITVTPE KKAEEERDEEE DKRRKGEEVTP S AIRHEGKSPGLGT DSCPLSPSTH CAPTSPTELRRR CKE NDCKLPGYEPS RAEHLPGEPALDS D GPGRPYLGDDQD VAFSYRQQTGKGT T LFSFSLQPE SFP SLDDDDGYLSF PN LSETNLLPQSLQ HYLPIRSPSLV PCF LFIFFFLLS AFSVPYALTL SFPLALC LCYLEPKAASLS ASLDNDPDS SSEEE TDSERTDTA ADGETTATE SDQEEDA ELKAQELEKT QDDL MKHQ T NISEL K RTFLETSTDT AVTNEW EKRLSTSPV RLAARQED APMIEPLV PEETKQSSG EK LMDGSEIF S LLESARK PTEFIGGV TSTSQSWVQ KMETTESS GIEPT VHHLPLST EKVVQET VLVEERR VVH ASGDASY SAGDSG DAAAQ PAFTG IK GKEGSALTE GAKEEG GEEVAK AVLE QEETAAAS RERQEE QSA AIHIS ETLE QKPHFES STVKTET ISFGSV SPGGVK LEISTKE VPVHT ETKT ITYESS QVDP GTDLEPG VMSAQ TITSET TTTT HITKTVK GGISE TRIEK RIVIT GDADI DHDQALA QAIKE AKEQ HPDMS VTKV VVHKETE ITPEDGED	True	True	4.612	2.287	1.866	0.842	0.79