

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
F6TFN2	F6TFN2_MOUSE	Mus musculus	NaN	25.161187	NaN	NaN	34887587	XDSGYGDIWCPDRGEFPAPPGSHRR EDSFESLDSLGSRLTSCSSDFTLRV GKEGCESDIDWEFTFKMQDYNKDD MSYRRISAIEPKSALPFNRFLPNKSK QPSYVPAPLRKKRPDKHEDNRRSWA SPVYETDGTGTFSRKSMDSVSAEDV QNLROVRYEEMQKIKSQLKEQDOK WQDDLAKWKNRRKSYTSDLQKKKE EREIEKQALEKSDRSSKTFREMLQ DRESRGSTVTSWRRTYSSDDLDDG VLPPTVLTSETSYQSERLEDEARHP AEMPKESTTFANREDSVVAETQLA SHSPEEQRLAPSPSEEPQASGSSA EPHQAPLLSTEPRRTPLPSEEPHQAS LSSTLQRPSSLSSQSWSTRAESTRI SASLPRSYQRTDTARLTSVVTARPFQ TSSRGISLPRSMTDDGWKYNGDV EVVKREQLDLGRATGPKPDSSHFI GSASEKEVIATEDVASLSPTLPSSSL SHDRAVSSKATFSSMSGLDSVSDSG EGRGSPLREVSRLDQFSDMRVSIN QTPGNKPVFGFTIKWDISGIFVASVE QGSPAEFSQLQVDDEILAINNTKFSY KDTKKWEEVMANAQETGNLVMDV RRYGKSGSSETKWIDTTSGIYNSDKS SSLSVTDFSESLSQSPYTESKEVNGI HEESNPFDSKASESILKLNKRRSQ FFEQSSDSVAPDLPVPTLSAPSRW AWDQEEERRRQERWQKEQDRLLQE KYQREQEKLREEWORAQQAEREN SKYLDEELMVLNSNSISLTSREPVA TWEATWSEGSKSLDSEGTRAGEED RGQLEDDAVYEDQSQKLELEQEKK RKEQEAQEEERRKQEAQAWAEAE AKAWAEAQARAEAEAKARAEQARA EAEAQKRAEAQKLAERERETSVKIY QYRRPIDSYDLPKREDDTSGMLPTD RSKSRSTTELNDPLIEKNGSSKYSER IGATSFSHRSKKDQAPSEAEALERQ QILQEMRKRTSLHNDNSWIRQRSAS VNKEPICLPGIMRRGESLDNLSQR PGSWRQSPWHSQPAGVYASSSVQD FSRPPPQLLSTSNRAYMRNPSSAVP PPAGSVKTSVPGSPSRSHSPSMSQS GSQLRNRVSGKRVCSYCNILGKG AAMIIETGLSYHLHCFKCVSCGCDL GGSSSGAEVRIRNHQLYCNDCYLRF KSGRPTAM	None	None	None	None	None	None	None		