

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							
E9PUL3	E9PUL3_MOUSE	Mus musculus	NaN	25.775171	NaN	NaN	34678516	MVPGQLVLLFLTLHLLQNTESMVB LNSNGYEGVVIAINPSVPEDERLIPSI KEMVTQASTYLFTEATERRFYFRNVSI LVPITWKSKEYLTPKQESYDQADVI VADPHLQHGDDPYTLQYGCQCDRG QYIHFTPNFLTDNLGIYGRGRVFN HEWAHLRWGVFDEYNMDRPFYMS RKNTVEATRCSTDITGTSVVRECOG GSCVSRRCRRDAKTGMQEAKCTFIP NKSQTARGSIMFMQSLDSVVEFCTE KTHNVEAPNLQNKMCNLRSTWDVI KASADFQNASPMTGTEAPPLPTFSL LKSQRVVCLVLDKSGSMRLGSPIT RLTLMNQAAELYLIQIEKESLVGLV TFDSTATIQTNLIRIINDSSYLAISTKL PQYPNGGTSICNGLKKGFEAITSSDQ STSGSEIVLLTDGEDNRISSCFQEVK HSGAIHHTIALGPSAARELETLSDMT GGLRFYAKEDVNGLIDAFSGISSKSG SISQQALQLESKAFNVGAGAWINST VPVDSTVGNDDFFVITWTVRKPEIIL QDPKGNKNTTSDFFQEDKLNIFSVRL RIPGIAETGTWYSLLNKGATSQLLT VTVITRARSPTTLPIVATAHMSQSTA QYPSRMIVYARVSGFLPVLGANVT AVIEAESGNQVTLELWDNGAGADTL KNDGIYSRYFTDFHNGGRYSLKVN QARKNMAKLNKQKNKSLYIPGYVE NDQIVLNPPRPEIPEATEATVEDFSR LTSGGSFVSGAPPDGDHARVFPSP KVTDLAEAFIGDHIHLTWAPGKVL DKGRAYRYVIRMSGHSLALQEDFSN STLVNTSSVMPKEAGSKETFKFKPE TFKIENGTQVYIAIQADNEARLSSEV SNIAQAVKFIPPQVYLPSTPPGLSTP STPPGLSTPSTPPGLSTPSTPPGLSTP STPPGLSTPSTPPGLSTPSTPPGLSTP STPPGLSTPSTPPGLSTPSTPPGLSTP STPPGLSTPSTPPGLSTPSTPPGLSTP STPPGLSTPSTPPGLSTPSTPPGLSTP VSVPSLTVFVLVATLFI	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None	None

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