

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|--------------|--------------|--------------|-----------|--------------|---------------|-----------------------|----------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| D3YUW8 | D3YUW8_MOUSE | Mus musculus | NaN | 25.385059 | NaN | NaN | 28528544 | MADTDLFMECEEEEELEPWQKISDVI EDSVVEDYNSVDKTTTSVSVSQQPVS APVPIAAHASVAGHLSTSTTVNSNGA QNSDSTKKTTLVTLIANNNAPGTQFV KPTVGVPPQVFSQMTVPVRPGSTMPVR PTTNTFTTVIPATLLTIRSTVPQSQSQQ TKSTPSTSTTPTATQPTSLGQLAGQP PGQSNQTSNPKLAPSPFPSPAVSIAS FVTVKRPVGTGENSNEVAKLVNTLN TVPSLGQSPGPVVVSNSSAQRSTSG PESSVKVTSSIPVFDLQDGGKICPR CNAQFRVTEALRGHMCYCCPEMVE YOKKGSLSDAEPSVPSAAKPSPEKT APVTSTPSSTIPALSPPTKVPENEN AGDAVQTKLIMLVDDFYGRDGGKA AQLTSFPKVATSFRCPHCTKRLKNNI RFMNHMKHHVELDQQNGEVDGHT ICQHCRYQFSTPFQLQCHLENVHSP YESTTKCKICEWAFESEPLFLQHMK DTHKPGEMPYVCQVCQYRSSLYSEV DVHFRMIHEDTRHLLCPYCLKVFKN GNAFQQHYMRHQKRNVYHCNKCR LQFLFAKDKEHKLQHHKTRKPKQ LEGLKPGTKVTIRASRGQPRTPVSS NDAPSGTLQEAALSTDPVFLYP PVQRNIQRAVRKMSVMGROTCLE CSFEIPDFPNHFPTYVHCCLCRYSTC CSRAYANHMNNHVPRKSPKYLALF KNSVSGIKLACTSCTFATSVGDAMA KHLVFNPSHRSSNILPRGLSWMSHL RPGQASERVFDWSMKNYLPPLVP NKAATVKPVGVTPEPQELAGPVLQ ALPSPASTATPPATPTHQPSALPPS ATEGTECLNVSEQEEGSPVTQDPEP ASGGGGGSGVGKKEQLSVKCLRVL FALCCNTEQAAEHFRNPQRIRRWL RRFQASQGENLEGKYLSEAEKLA EWWLIQREQQLPVNEETLFOKATKI GRSLEGGFKISYEWAVRFMLRHHLT PHARRAVAHTLPKHVAENAGLFIEF VORQIHNQDLPLSMIVAIDEISLFLD TEVLSDDDRKENALQTVGTGEPWCD VVLAILADGTVLPTLVVFRGQANRFA NVPDSILLEAKDSGYSDEIMELWS TRVWKKHTACQHSKSLMVMDCCHR THLSEEVALLSASSTLPAVVPAGCS SKIQPLDVCIKRTVKNFLHKKWKEQ AREMADAACSDVLLQLVLVWLGE VLGVIGDSPELVQRSFLVASVLPQPD GNVNSPTRNADMQEELIASLEEQLK LNGEQSEHSASAPRPRSSPEETVE PESLHQLFEGESETESFYGFEEADL DLMEI | None | None | None | None | None | None | None | | |

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