| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | O14715 | RGPD8_HUMAN | Homo sapiens | RANBP2-like and GRIP domain-containing protein 8 | 29.035765 | S918 | T19;S21 | 30620550;31492838 | MRRSKADVERYVASVLGLTPSPRQK<br>SMKGFYFAKLYYEAKEYDLAKKYICT<br>YINVQERDPKAHRFLGLLYELEENT<br>EKAVECYRRSVELNPTQKDLVLKIAE<br>LLCKNDVTDGRAKYWVERAAKLFPG<br>SPAIYKLKEQLLDCEGEDGWNKLFD<br>LIQSELYVRPDDVHVNIRLVELYRST<br>KRLKDAVAHCHEAERNIALRSSLEW<br>NSCVVQTLKEYLESLQCLESDKSDW<br>RATNTDLLLAYANLMLLTLSTRDVQ<br>ENRELLESFDSALQSAKSSLGGNDE<br>LSATFLEMKGHFYMYAGSLLLKMG<br>QHGNNVQWRALSELAALCYLIAFQV<br>PRPKIKLREGKAGQNLLEMMACDRL<br>SQSGHMLLSLSRGKQDFLKEVVETF<br>ANKIGQSALYDALFSSQSPKDTSFLG<br>SDDIGKIDVQEPELEDLARYDVGAIR<br>AHNGSLQHLTWLGLQWNSLPALPG<br>IRKWLKQLFHRLPHETSRLETNAPE<br>SICILDLEVFLLGVVYTSHLQLKEKC<br>NSHHSSYQPLCLPFPVCKQLCTERQ<br>KSWWDAVCTLIHRKAVPGNLAKLRL<br>LVQHEINTLRAQEKHGLQPALLVHW<br>AKYLQKTGSGLNSFYGQLEYIGRSV<br>HYWKKVLPLLKIIKKNSIPEPIDPLFK<br>HFHSVDIQASEIVEYEEDAHITFAILD<br>AVNGNIEDAVTAFESIKSVVSYWNL<br>ALIFHRKAEDIENDALSPEEQEECR<br>NYLTKTRDYLIKIIDDGDSNLSVVKK<br>LPVPLESVKQMLNSVMQELEDYSEG<br>GPLYKNGSLRNADSEIKHSTPSPTKY<br>SLSPSKSYKYSPETPPRWTEDRNSLL<br>NMICQQVEAIKKEMQELKLNSSKSA<br>SRHRWPTENYGPDSVPDGYQGSQT<br>FHGAPLTVATTGPSVYYSQSPAYNSQ<br>YLLRPAANVTPTKGSSNTEFKSTKEG<br>FSIPVSADGFKFGISEPGNQEKKREK<br>PLENDTGLQAQDIRGRKKGRGVIFG<br>QTSSTFTFADVAKSTSGEGFQFGKK<br>DLNFKGFSGAGEKLFSSRYGKMAN<br>KANTSGDFEKDDDAYKTEDSDDIHF<br>EPVVQMPEKVELVTGEEGEKVLYSQ<br>GVKLFRFDAEVRQWKERGLGNLKIL<br>KNEVNGKLRMLMRREQVLKVCAN<br>HWITTTMNLKPLSGSDRAWMWSAS<br>DFSDGDAKLERLAAKFKTPELAEEF<br>KQKFEECQRLLLDIPLQTPHKLVDT<br>GRAAKLIQRAEEMKSGLKDFKTFLT<br>NDQTKVTEEENKGSGTGVAGASDTT<br>IKPNAENTGPTLEWDNYDLREDALD<br>DSVSSSSVHASPLASSPVRKNLFRFD<br>ESTTGSNFSFKSALSLSKSPAKLNQS<br>GTSVGTDEESVVTQEEERDGQYFEP<br>VVPLPDLVEVSSGEENEQVVFSHRA<br>EIYRYDKDVGQWKERGIGDIKILQNY<br>DNKQVRIVMRRDQVLKLCANHRITP<br>DMSLQNMKGTERVWVWTACDFAD<br>GERKVEHLAVRFKLQDVADSFKKIF<br>DEAKTAQEKDSLITPHVSRSSTPRES<br>PCGKIAVAVLEEITRERTDVIQGDDV<br>ADAASEVEVSSTSETTTKAVVSPPKF<br>VFVSESVKRIFSSEKSKPFAFGNSSA<br>TGSLFGFSFNAPLKSNNSETSSVAQ<br>SGSESKVEPKKCELSKNSDIEQSSDS<br>KVKNLSASFPTEESSINYTFKTPEKE<br>PPLWHAEFTKEELVQKLRSTTKSAD<br>HLNGLLREIEATNAVLMEQIKLLKSE<br>IRRLERNQEREKSAANLEYLKNVLL<br>QFIFLKPGSERERLLPVINTMLQLSP<br>EEKGKLAAVAQDEEENPSRSSG | None | None | None | None | None | None | None | None | None |