

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi appa |
|--------------|-------------|--------------|-----------|--------------|----------------|---|---|---|---------------|---------------|---------|---------|---------------|-----------------------|------------|
| Q60865 | CAPR1_MOUSE | Mus musculus | Caprin-1 | 45.841649 | T265;S418;T428 | S10;S113;S333;S341;T623;T634;T637;T649;T660;T663;T668 | 34887587;40885482;25153642;37453647;28528544;28135057;34678516;29187734;39950214;40997131 | MPSATSHSGSGSKSSGPPPPSGSSG SEAAAGAAAPASQHPATGTGAVQTE AMKQILGVIDKCLRNLKKGKGLDD YQERMNKGERLNQQLDAVSKYQE VTNNLEFAKELQRSFMALSQDIQKT IKKTARREQLMREEAEQKRLKTVLE LQYVLDKLGDDVVRTDLKQGLSGVP ILSEELSLLEDFYKLVDFERDMSLR LNEQYEHASIHLDLLEGKEKPVCG TTYKALKEIVERVFQSNYFDSTHNN QNGLCSEEEAASAPTVEDQVAEAEAP EPAEYYTEQSEVESTEYVNRQFMAE TQFSSGEKEQVDEWTVETVEVNSL QQQPQAASPSVPEPHSLTPVAQSDP LVRORVQDLMAQMGGPYNFIQDS MLDFENQTLDPAIVSAQPMNPTQN MDMPQLVCPQVHSESRLAQSNQVP VQPEATQVPLVSSSTSEGYTASQPLYQ PSHATEQRPQKEPMDQIQATISLNT DOTTASSSLPAASQPQVFQAGTSKPL HSSGINVNAAPFQSMQTVFNMNAP VPPANEPETLKQSQSOYQATYNQSF SQPHQVEQTELQQDQLQTVVGTYH GSQDQPHQVPGNHQPPQQTGFP RSSQPYNSRGVSRGGSRGARGLM NGYRGPANGFRGGYDGYRPSFNT NSGYSQSQFTAPRDYSGYQRDGYQQ NFKRGSQSGPRGAPRGRGGPPRP NRGMPQMNTQQVN | False | False | 3.796 | 3.4 | 1.498 | 1.919 | 1.845 |