| | 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** | D3Z158 | D3Z158_MOUSE | Mus musculus | glutamine--tRNA ligase | 25.385059 | NaN | NaN | 28528544 | MATPDSLALFTGLGLSENKARETLK<br>NEALSTQLREAATQAHQILGSTIDKA<br>TGVLLYDLVSRLRDTRRRSFLVSYIA<br>NKKIHTGLQLSAALEYVRSHPQDPID<br>TKDFEQECGVGVVVTPEQIEEAVEST<br>INKHQLQLLAERYRFNMGLLMGEA<br>RAALRWADGKMIKNEVDMQVLHLL<br>GPKMEADLVKKPKVAKARLEETDRK<br>TAKDVVEKGENYKTPGYVITPYTMDL<br>LKQHLEITGGQVRTRFPPEPNGILHI<br>GHAKAINFNFGYAKANNGICFLRFD<br>DTNPEKEEAKFFTAIYDMVTWLGYT<br>PYKVTYASDYFDQLYAWAVELIHGGL<br>AYVCHQRVEELKGHNPLPSPWRDR<br>PKEESLLLFEAMRKGKFAEGEATLR<br>MKLVMEDGKMDPVAYRVKYTPHHR<br>TGDKWCIYPTYDYTHCLCDSIEHITH<br>SLCTKEFQARRSSYFWLCNALKVYC<br>PVQWEYGRLNLHYAVVSKRKILQLV<br>AAGAVRDWDDPRLFTLTALRRRGFP<br>PEAINNFCARVGVTVAQTTMEPHLL<br>EACVRDVLNDAAPRAMAVLEPLQVV<br>ITNFPAPKPLDIRVPNFPADETKGFH<br>QVPFASTVFIERSDFKEESEPGYKRL<br>ASGQPVGLRHTGYVIELQNIVRGSSG<br>CVERLEVTCRRADAGEKPKAFIHWV<br>SQPLVCEIRLYECLFQHKNPEDPVEV<br>PGGFLSDLNPASLQVVEGALVDCSV<br>ALAKPFDKFQFERLGYFSVDPDSHQ<br>GQIVFNRTVTLKEDPGKI | None | None | None | None | None | None | None | None | None |