

| 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 | extrac region | |
|--------------|------------|------------|--------------|---------------------------|---------------|-----------------------|-------|---|--|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|---------------|-------|
| 0 | Q9D8N0 | EF1G_MOUSE | Mus musculus | Elongation factor 1-gamma | 42.943791 | NaN | NaN | 22645316;29187734;24788674;37453647;21540332;36288343;39627609;37507081 | MAAGTLYTYPENWRAFKALIAAQYS GAQVRVLSAPPHFHFGQTNRTPEFL RKFPAGKVPAPFEGDDGFCVFESNAI AAYVSNELRGSTPEAAAQVQWVS FADSDIVPPASTWVFPPTLGMHHNK QATENAKEEVKRI LGLLDTHLKRTRTF LVGERVTLADITVVC TLLWLYKQVLE PSFRQAFPNTNRWFLTCINQPQFRA ILGEVKLCEKMAQFDKFAESQPK KDTPRKEKGSREEKQKPAERKEEK KAAAPAPEEEMDECEQALAAEPKAK DPFAHLPKSTFVLDEFKRKYSNEDT LSVALPYFWEHFDKDGWSLWYAEY RFPEELTQTFMSCNLITGMFORL DK LRKNASFVILFGTNNSSSISGVWVF RGQELAFPLSPDWQVDYESY TWRKL DPGSEETQTLVREYFSWEGTFQHV G KAVNQGKIFK | False | False | 2.502 | 2.487 | 1.987 | 2.884 | 0.871 | 1.369 | 1.393 |