| | 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** | Q8CAA7 | PGM2L_MOUSE | Mus musculus | Glucose 1,6-bisphosphate synthase | 24.474581 | NaN | S175 | 33300544 | MAENADDDLNSNLLHAPYLTGDPQ<br>LDTAIGQWLRWDKNPKTKEQIENLL<br>RNGMNKELRDRLCCRMTFGTAGLR<br>SAMGAGFCYINDLTVIQSTQGMYKY<br>LERCFSDFKQRGFVVGYDTRGQVTS<br>SCSSQRLAKLTAAVLLAKDIPVYLFS<br>RYVPTPFVPYAVQELKAVAGVMITAS<br>HNRKEDNGYKVYWETGAQITSPHD<br>KEILKCIEECVEPWNDSWNDNLVDT<br>SPLKKDPLQDICKKYMEDLKKICFYR<br>DLNSKTTLKFVHTSFHGVGHDYVQL<br>AFQVFGFKPPIPVPEQKDPDPDFSTV<br>KCPNPEEGESVLELSLRLAEKENARI<br>VLATDPDADRLAVAELQENGRWKVF<br>TGNELAALFGWWMFDCWKKNKPN<br>ADVKNVYMLATTVSSKILKAIALKEG<br>FHFEETLPGFKWIGSRIKDLLGNGK<br>EVLFAFEESIGFLCGTSVLDKDGVSA<br>AAVVAEMASFLDTRKVTLMEQLTKV<br>YEIYGYHMSKTSYFLCYDPPTIKTIFE<br>RIRNFESPKEYPKFCGAFAILHVRDI<br>TTGYDSSQPNKKSVLPVSKNSQMIT<br>FTFQNGCVATLRTSGTEPKIKYYAEM<br>CASPGQSDTTFLEEELKKLIDALIEN<br>FLEPSKNALVWRSV | None | None | None | None | None | None | None | None | None |