| | 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 | extr regi |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | P32921 | SYWC_MOUSE | Mus musculus | Tryptophan--tRNA ligase, cytoplasmic | 39.472249 | NaN | S355 | 33300544;22645316;28528544;24788674;25153642;39627609;37507081 | MADMPSGESCTSPLELFNSIATQGE<br>LVRSLKAGNAPKDEIDSAVKMLLSL<br>KMSYKAAMGEEYKAGCPPGNPTAG<br>RNCDSDATKASEDFVDPWTVRTSSA<br>KGIDYDKLIVQFGSSKIDKELINRIER<br>ATGQRPHRFLRRGIFFSHRDMNQIL<br>DAYENKKPFYLYTGRGPSSEAMHLG<br>HLVPFIFTKWLQDVFNVPLVIQMSD<br>DEKYLWKDLTLEQAYSYTVENAKDII<br>ACGFDINKTFIFSDLEYMGQSPGFYR<br>NVVKIQKHVTFNQVKGIFGFTDSDCI<br>GKISFPAVQAAPSFSNSFPKIFRDRT<br>DIQCLIPCAIDQDPYFRMTRDVAPRI<br>GHPKPALLHSTFFPALQGAQTKMSA<br>SDPNSSIFLTDTAKQIKSKVNKHAFS<br>GGRDTVEEHRQFGGNCEVDVSFMY<br>LTFFLEDDDRLEQIRKDYTSGAMLT<br>GELKKTLIDVLQPLIAEHQARRKAVT<br>EETVKEFMTPRQLSFHFQCFCFDT | False | False | 3.82 | 3.479 | 1.843 | 1.58 | 0.884 | 1.46 | 1.68 |