| | 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** | A6X935 | ITIH4_MOUSE | Mus musculus | Inter alpha-trypsin inhibitor, heavy chain 4 | 25.685855 | NaN | NaN | 39627609 | MKSPAPAHMWNLVLFLPSLLAVLPT<br>TTAEKNGIDIYSLTVDSRVSSRFAHT<br>VVTSRVVNRADAVQEATFQVELPRK<br>AFITNFSMIIDGVTYPGVVKEKAEAQ<br>KQYSAAVGRGESAGIVKTTGRQTEK<br>FEVSVNVAPGSKITFELIYQELLQRR<br>LGMYELLLKVRPQQLVKHLQMDIYI<br>FEPQGISILETESTFMTPELANALTT<br>SQNKTKAHIRFKPTLSQQQKSQSEQ<br>DTVLNGDFIVRYDVNRSDSGGSIQIE<br>EGYFVHHFAPENLPTMSKNVIFVID<br>KSGSMSGKKIQQTREALVKILKDLSP<br>QDQFNLIEFSGEANQWKQSLVQATE<br>ENLNKAVNYASRIRAHGGTNINNAV<br>LLAVELLDRSNQAELLPSKSVSLIILL<br>TDGDPTVGETNPTIIQNNVREAINGQ<br>YSLFCLGFGFDVNYPFLEKMALDNG<br>GLARRIYEDSDSALQLQDFYHEVAN<br>PLLSSVAFEYPSDAVEEVTRYKFQH<br>HFKGSEMVVAGKLQDQGPDVLLAK<br>VSGQMHMQNITFQTEASVAQQEKE<br>FKSPKYIFHNFMERLWALLTIQQQL<br>EQRISASGAELEALEAQVLNLSLKYN<br>FVTPLTHMVVTKPEGQEQFQVAEKP<br>VEVGDGMQRLPLAAQAHPFRPPVR<br>GSKLMTVLKGSRSQIPRLGDAVRAS<br>RQYIPPGFPGPPGPPGFPAPPGPPGF<br>PAPPGPPLASGSDFSLQPSYERMLSL<br>PSVAAQYPADPHLVVTEKSKESTIPE<br>ESPNPDHPQVPTITLPLPGSSVDQLC<br>VDILHSEKPMKLFVDPSQGLEVTGK<br>YENTGFSWLEVTIQKPHLQVHATPE<br>RLVVTRGRKNTEYKWKKTLFSVLPG<br>LKMTMNMMGLLQLSGPDKVTIGLL<br>SLDDPQRGLMLLLNDTQHFSNNVK<br>GELGQFYRDIVWEPPVEPDNTKRTV<br>KVQGVDYLATRELKLSYQEGFPGAEI<br>SCWTVEI | False | True | 1.303 | 1.324 | 1.344 | 0.974 | False | 5.0 | 4.264 |