| | 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** | O54890 | ITB3_MOUSE | Mus musculus | Integrin beta-3 | 25.685855 | NaN | T766;T772;T778;T784 | 39627609 | MRAQWPGQLWAALLALGALAGVVV<br>GESNICTTRGVNSCQQCLAVSPVCA<br>WCSDETLSQGSPRCNLKENLLKDN<br>CAPESIEFPVSEAQILEARPLSSKGSG<br>SSAQITQVSPQRIALRLRPDDSKIFSL<br>QVRQVEDYPVDIYYLMDLSFSMKDD<br>LSSIQTLGTKLASQMRKLTSNLRIGF<br>GAFVDKPVSPYMYISPPQAIKNPCYN<br>MKNACLPMFGYKHVLTLTDQVSRF<br>NEEVKKQSVSRNRDAPEGGFDAIM<br>QATVCDEKIGWRNDASHLLVFTTDA<br>KTHIALDGRLAGIVLPNDGHCHIGTD<br>NHYSASTTMDYPSLGLMTEKLSQK<br>NINLIFAVTENVVSLYQNYSELIPGTT<br>VGVLSDDSSNVLQLIVDAYGKIRSKV<br>ELEVRDLPEELSLSFNATCLNNEVIP<br>GLKSCVGLKIGDTVSFSIEAKVRGCP<br>QEKEQSFTIKPVGFKDSLTVQVTFDC<br>DCACQAFAQPSSPRCNNGNGTFEC<br>GVCRCDQGWLGSMCECSEEDYRPS<br>QQEECSPKEGQPICSQRGECLCGQC<br>VCHSSDFGKITGKYCECDDFSCVRY<br>KGEMCSGHGQCNCGDCVCDSDWT<br>GYYCNCTTRTDTCMSTNGLLCSGRG<br>NCECGSCVCVQPGSYGDTCEKCPTC<br>PDACSFKKECVECKKFNRGTLHEEN<br>TCSRYCRDDIEQVKELTDTGKNAVN<br>CTYKNEDDCVVRFQYYEDTSGRAVL<br>YVVEEPECPKGPDILVVLLSVMGAIL<br>LIGLATLLIWKLLITIHDRKEFAKFEE<br>ERARAKWDTANNPLYKEATSTFTNI<br>TYRGT | True | True | 2.421 | 4.539 | 2.15 | 2.002 | 1.508 | 5.0 | 3.634 |