| | 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** | P31977 | EZRI_RAT | Rattus norvegicus | Ezrin | 26.1995 | NaN | T146;S366;T478;S535;T567 | 38843836;34502162 | MPKPINVRVTTMDAELEFAIQPNTT<br>GKQLFDQVVKTIGLREVWYFGLQYV<br>DNKGFPTWLKLDKKVSAQEVRKEN<br>PVQFKFRAKFYPEDVADELIQDITQK<br>LFFLQVKEGILSDEIYCPPETAVLLGS<br>YAVQAKFGDYNKEMHKSGYLSSERL<br>IPQRVMDQHKLSRDQWEDRIQVWH<br>AEHRGMLKDSAMLEYLKIAQDLEM<br>YGINYFEIKNKKGTDLWLGVDALGL<br>NIYEKDDKLTPKIGFPWSEIRNISFN<br>DKKFVIKPIDKKAPDFVFYAPRLRIN<br>KRILQLCMGNHELYMRRRKPDTIEV<br>QQMKAQAREEKHQKQLERQQLETE<br>KKRRETVEREKEQMLREKEELMLR<br>LQDFEQKTKRAEKELSEQIEKALQLE<br>EERRRAQEEAERLEADRMAALRAKE<br>ELERQAQDQIKSQEQLAAELAEYTA<br>KIALLEEARRRKEDEVEEWQHRAKE<br>AQDDLVKTKEELHLVMTAPPPPPPP<br>VYEPVNYHVQEGLQDEGAEPMGYS<br>AELSSEGILDDRNEEKRITEAEKNER<br>VQRQLLTLSNELSQARDENKRTHN<br>DIIHNENMRQGRDKYKTLRQIRQGN<br>TKQRIDEFEAM | True | True | 4.741 | 2.817 | 1.903 | 1.945 | 1.859 | 5.0 | 2.352 |