| | 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** | Q6SZW1 | SARM1_HUMAN | Homo sapiens | NAD(+) hydrolase SARM1 | 18.379595 | NaN | S548;S558 | 30379171 | MVLTLLLSAYKLCRFFAMSGPRPGA<br>ERLAVPGPDGGGGTGPWWAAGGRG<br>PREVSPGAGTEVQDALERALPELQQ<br>ALSALKQAGGARAVGAGLAEVFQLV<br>EEAWLLPAVGREVAQGLCDAIRLDG<br>GLDLLLRLLQAPELETRVQAARLLEQ<br>ILVAENRDRVARIGLGVILNLAKERE<br>PVELARSVAGILEHMFKHSEETCQR<br>LVAAGGLDAVLYWCRRTDPALLRHC<br>ALALGNCALHGGQAVQRRMVEKRA<br>AEWLFPLAFSKEDELLRLHACLAVA<br>VLATNKEVEREVERSGTLALVEPLVA<br>SLDPGRFARCLVDASDTSQGRGPDD<br>LQRLVPLLDSNRLEAQCIGAFYLCAE<br>AAIKSLQGKTKVFSDIGAIQSLKRLVS<br>YSTNGTKSALAKRALRLLGEEVPRPI<br>LPSVPSWKEAEVQTWLQQIGFSKYC<br>ESFREQQVDGDLLLRLTEEELQTDL<br>GMKSGITRKRFFRELTELKTFANYST<br>CDRSNLADWLGSLDPRFRQYTYGLV<br>SCGLDRSLLHRVSEQQLLEDCGIHL<br>GVHRARILTAAREMLHSPLPCTGGK<br>PSGDTPDVFISYRRNSGSQLASLLKV<br>HLQLHGFSVFIDVEKLEAGKFEDKLI<br>QSVMGARNFVLVLSPGALDKCMQD<br>HDCKDWVHKEIVTALSCGKNIVPIID<br>GFEWPEPQVLPEDMQAVLTFNGIK<br>WSHEYQEATIEKIIRFLQGRSSRDSS<br>AGSDTSLEGAAPMGPT | True | False | 4.583 | 2.443 | 4.708 | 1.682 | 1.2 | 2.266 | 2.184 |