| | 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** | P40505 | SDS3_YEAST | Saccharomyces cerevisiae | Transcriptional regulatory protein SDS3 | 24.475439 | NaN | S166;S211 | 33229814 | MAIQKVSNKDLSRKDKRRFNIESKV NKIYQNFYSERDNQYKDRLTALQTD LTSLHQGDNGQYARQVRDLEEERD LELVRLRLFEEYRVSRSGIEFQEDIE KAKAEHEKLIKLCKERLYSSIEQKIK KLQEERLLMDVANVHSYAMNYSRP QYQKNTRSHTVSGWDSSSNEYGRD TANESATDTGAGNDRRTLRRRNASK DTRGNNNNQDESDFQTGNGSGSN GHGSRQGSQFPHFNNLTYKSGMNS DSDFLQGINEGTDLYAFLFGEKNPK DNANGNEKKKNRGAQRYSTKTAPP LQSLKPDEVTEDISLIRELTGQPPAPF RLRSD | None | None | None | None | None | None | None | None | None |