| | 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** | Q7XYY2 | MED25_ARATH | Arabidopsis thaliana | Mediator of RNA polymerase II transcription subunit 25 | 27.144759 | NaN | NaN | 28154133 | MSSEVKQLIVVAEGTAALGPYWQTIV<br>SDYLEKIIRSFCGSELNGERNPVSTV<br>ELSLVIFNSHGSYCACLVQRSGWTR<br>DVDIFLHWLSSIQFGGGGFNEVATA<br>EGLAEALMMFSPPSGQAQPSNDLK<br>RHCILITASNPHILPTPVYRPRLQNV<br>ERNENGDAQAESRLSDAETVASYFA<br>KCSVSLSVVCPKQLPTIRALYNAGKP<br>NQQSADLSIDTAKNTFYLVLISENFV<br>EACAALSHSATNLPQTQSPVKVDRA<br>TVAPSIPVTGQPPAPVSSANGPIQNR<br>QPVSVGPVPTATVKVEPSTVTSMAPV<br>PSFPHIPAVARPATQAIPSIQTSSASP<br>VSQDMVSNAENAPDIKPVVVSGMTP<br>PLRTGPPGGANVNLLNNLSQVRQV<br>MSSAALAGAASSVGQSAVAMHMSN<br>MISTGMATSLPPSQTVFSTGQQGITS<br>MAGSGALMGSAQTGQSPGPNNAFS<br>PQTTSNVASNLGVSQPMQGMNQGS<br>HSGAMMQGGISMNQNMMSGLGQ<br>GNVSSGTGGMMPTPGVGQQAQSGI<br>QQLGGSNSSAPNMQLSQPSSGAMQ<br>TSQSKYVKVWEGNLSGQRQGQPVLI<br>TRLEGYRSASASDSLAANWPPTMQI<br>VRLISQDHMNNKQYVGKADFLVFRA<br>MSQHGFLGQLQDKKLCAVIQLPSQT<br>LLLSVSDKACRLIGMLFPGDMVVFK<br>PQIPNQQQQQQQQLHQQQQQQQQI<br>QQQQQQQQHLQQQQMPQLQQQQQ<br>QHQQQQQQQHQLSQLQHHQQQQQ<br>QQQQQQQQHQLTQLQHHHQQQQQ<br>ASPLNQMQQQTSPLNQMQQQTSPL<br>NQMQQQQQPQQMVMGGQAFAQAP<br>GRSQQGGGGGQPNMPGAGFMG | None | None | None | None | None | None | None | None | None |