| | 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** | P08018 | PBS2_YEAST | Saccharomyces cerevisiae | MAP kinase kinase PBS2 | 24.475439 | NaN | S68;S269;S514;T518 | 33229814 | MEDKFANLSLHEKTGKSSIQLNEQT<br>GSDNGSAVKRTSSTSSHYNNINADL<br>HARVKAFQEQRALKRSASVGSNQSE<br>QDKGSSQSPKHIQQIVNKPLPPLPVA<br>GSSKVSQRMSSQVVQASSKSTLKNV<br>LDNQETQNITDVNINIDTTKITATTIG<br>VNTGLPATDITPSVSNTASATHKAQL<br>LNPNRRAPRRPLSTQHPTRPNVAPH<br>KAPAIINTPKQSLSARRGLKLPPGGM<br>SLKMPTKTAQQPQQFAPSPSNKKHI<br>ETLSNSKVVEGKRSNPGSLINGVQS<br>TSTSSSTEGPHDTVGTTPRTGNSNN<br>SSNSGSSGGGGLFANFSKYVDIKSG<br>SLNFAGKLSLSSKGIDFSNGSSSRIT<br>LDELEFLDELGHGNYGNVSKVLHKP<br>TNVIMATKEVRLELDEAKFRQILME<br>LEVLHKCNSPYIVDFYGAFFIEGAVY<br>MCMEYMDGGSLDKIYDESSEIGGID<br>EPQLAFIANAVIHGLKELKEQHNIIH<br>RDVKPTNILCSANQGTVKLCDFGVS<br>GNLVASLAKTNIGCQSYMAPERIKSL<br>NPDRATYTVQSDIWSLGLSILEMAL<br>GRYPYPPETYDNIFSQLSAIVDGPPP<br>RLPSDKFSSDAQDFVSLCLQKIPERR<br>PTYAALTEHPWLVKYRNQDVHMSE<br>YITERLERRNKILRERGENGLSKNVP<br>ALHMGGL | None | None | None | None | None | None | None | None | None |