| | 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** | Q5FFM4 | DNAK_EHRRG | Ehrlichia ruminantium | Chaperone protein DnaK | 25.830362 | NaN | T201 | 30930869 | MILCRGLFMAVIGIDLGTTNSCVAV MEGGDAKAIENSEGARTTPSIVAFTD SEVLVGDPAKRQATTNAKNTIYASKR LIGRRYQDTRDIKTSYDIVSAKNGDA WIKVRDKDYSPSQIGALILEKMKETA ERHLGCKVEKAVITVPAYFDDAQRQ ATKDAGKIAGLDVIRIINEPTAAALAY GLNKSDKQKVIAVYDLGGGTFDVSIL EIADGVFEVKSTNGDTMLGGEDFDH AIMEYLMDDFKKSTGIDLHSDAMA MQRIKEAAEKAKIELSSRMETDINLP FLSSDSTGPKHLSLKLTRATFENLVS DLVKRTIEPCKKALKDAGISADKIDE VVLVGGMTRVPKIIQTVKEFFGKEPH KGVNPDEVVAIGAAIQGGILAGDVRD VLLLDVTPLSLGIETLGGVFTPLIERN TTIPTKKSQVFSTAEDGQTAVTIKVF QGERKMANDNKLLGQFSLEGIPPAP RGMPQIEVTFDIDANGIVHVSAKDK ASGKEQAIRIQSSGGLTDDEIQNMIK EAESKAEEDEKRKKFVEVKNNAENL VHSTEKSLKEHGDKISNADKLDIEN AIRDLKDCISKDNIEDTDTMQNKLD HLMKVSMKLGEALYSNTNNATAGD NNTTDTGSSSNSDGSKVVDSDYQEI DKKDGK | None | None | None | None | None | None | None | None | None |