

| UniprotKB ID | Entry name | organism | full name | oglnacscore | oglnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane |
|--------------|-------------|--------------|---------------------------------|-------------|----------------|-----------------------|--|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|
| Q86T24 | KAISO_HUMAN | Homo sapiens | Transcriptional regulator Kaiso | 16.737824 | T251;T257;S502 | T251 | 34019948;33214551;39302247;30379171;39531497;40307207;30620550;35138101;35254053 | MESRKLISATDIQYSGSLLNSLNEQR GHGLFCDVTVIVEDRKFRAHKNILS ASSTYFHQLFSVAGQVVELSFIRAEI FAEILNYIYSSKIVRVRSDLLDELIKS GQLLGVKFIAELGVPLSQVKSISGTA QDGNTEPLPPDSGDKNLVIQKSKDE AQDNGATIMPIITESFSLSAEDYEMK KIIVTDSDDDDVIFCSEILPTKETL PSNNTVAQVQSNPGVAISDVAPSA SNNSPPLTNITPTQKLPTPVNQATLS QTQGSEKLLVSSAPHTLTPNIILLNQ TPLSTPPNVSSSLPNHMPSSINLLVQ NQQTPNSAILTGNKANEEEEEEIIDD DDDTISSSPDSAVSNTSLVQADTSQ NTSFDGSLIQKMQIPTLLQEPLSNL KISDIIRNTNDPGVGSKHLMEGQKI ITLDTATEIEGLSTGCKVYANIGEDTY DIVIPVKDDPDEGEARLENEIPKTSG SEMANKRMRKVKHDDHYELIVDGRV YYICIVCKRSYVCLTSLRRHFNIHSW EKKYPCRYCEKVFPLAEYRTKHEIHH TGERRYOCLACGKSFINYQFMSSHI KSVHSQDPGSKLYRLHPCRSLQI RQYAYLSDRSSTIPAMKDDGIGYKVD TGKEPPVGTSTTSTQNKPMTWEDIFI QQENDSIFKQNVTDGSTEFEFIIPE Y | True | False | 4.373 | 4.743 | False | False | False | 4.373 |