| | UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochndrion | endoplasmic reticulum | golgi apparatus | plasma membr |
|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|---|
| **0** | A0A0B4KFJ8 | A0A0B4KFJ8_DROME | Drosophila melanogaster | [histone H3]-trimethyl-L-lysine(9) demethylase | 33.728526 | S535;S543;T551;T553 | NaN | 28604694;40245499 | MKMSEVPRIKVFRPTWEEFKDFPKY<br>VAYMESQGAHKAGLAKVVPPPEWV<br>PRRSGYADLDALNVTIPAPICQVVTG<br>KQGYYQQINIQKKPLTVKQFSELAST<br>ERYATPKHFDFEDLERKYWKNITYV<br>APIYGADVSGSITDTDQDSWNINRLG<br>TILDYVNKDYNIQIDGVNTAYLYFGM<br>WKTTFAWHTEDMDLYSINYLHFGA<br>PKTWYVVPPECGRKLEKVANQYFPA<br>SYKNCNAYLRHKMTLISPQILKQHD<br>VPVSKITQEAGEIMITFPFGYHAGFN<br>HGFNCAESTNFAMERWIEYGKRAV<br>QCTCSNDMVKISMDTFVKRFQSDR<br>YDLWMEGRDVGRHPEDPPNAVLSA<br>APLPPHLDVLLCDKKMKKQCNPTKA<br>KSFKERNPDLDLDEIQQNPNVPDDV<br>KAMLKESVLTLDTGDLATDEADFPN<br>EDAMSLQSPANLKTKQELLEYIDDG<br>TEDDDEEEDFKRRKQKRRYDADYD<br>DDWLASKRKTNSRNNRGRSPRTKD<br>DRSISPASSTSSTSRGARRGKASGTP<br>RKTPARRKKDSITTSPAVSSAATAVK<br>TPTSAVVAGTTSIATTTTPPADGGGD<br>AKKERQSLQFMQQSRKFEGKIPKLS<br>QTSAATGGATAAAEASTSKSSQGQQ<br>QQIVYVNMLPAANTLNGIPQQQQQQ<br>QYASTDGNVYQLQNEILCDANGHAV<br>TAATASYQTTASSPQQQQQQQQQQ<br>QNVATSVAATIYQRQCY | NaN | NaN | NaN | NaN | NaN | NaN | NaN | NaN |