

| UniprotKB ID | Entry name | organism | full name | oglcnacscore | oglcnac sites | phosphorylation sites | PMIDS | sequence | intracellular | extracellular | cytosol | nucleus | mitochondrion | endoplasmic reticulum | golgi apparatus | plasma membrane | extracellular region |
|--------------|-------------|--------------|-----------------------------------|--------------|---------------|-----------------------|----------|---|---------------|---------------|---------|---------|---------------|-----------------------|-----------------|-----------------|----------------------|
| 0 Q6TGC4 | PADI6_HUMAN | Homo sapiens | Protein-arginine deiminase type-6 | 18.379595 | S3 | S10;S446 | 30379171 | MVSVEGRAMSFQSIHLSLDSPVHA VCVLGTEICLDLSCAPQKCCQCFTH GSGRVLIDVANTVISEKEDATIWWPL SDPTYATVKMTSPSPSVDADKVSVTY YGPNEAPVGTAVLYLTGIEVSLEVD IYRNGQVEMSSDKQAKKKWIWGPS GWGAILLVNCPADVGQQLDCKKT KKVIFSEEITNLSQMTLNVQGPSCIL KKYRLVLHTSKEESKKARVYWPQKD NSSTFELVLGPDQHAYTLALLGNHL KETFYVEAIAFPSAEFGLISYSVSLV EESQDPSIPETVLYKDTVVFRVAPCV FIPCTQVPLEVYLCRELQVGFVDTV TKLSEKSNQVASVYEDPNRLGRWL QDEMAFCYTQAPHKTTSLIDTPQA ADLDEFPMKYSLSPIGYMIQDTE HKVASMDSIGNLMVSPVVKVQKKEY PLGRVLIGSSFYPSAEGRAMSKTLRD FLYAQQVQAPVELYSDWLMTGHVD EFMCFIPTDDKNEGKGFLLLLASP SACYKLFREKQKEGYGDALLFDEL ADQLLSNGREAKTIDQLLADESLKK QNEYVEKCIHLNRDILKTELGLVEQ DIIEIPQLFCLEKLTNIPSDQQPKRSF ARPYFPDLLRMIVMGKNLGPVQKPF PQIKGTCCLEEKICCLEPLGFKCTFI NDFDCYLTEVGDICACANIRRVVFAF KWWKMVP | False | False | 1.315 | 3.748 | 1.518 | 1.05 | 0.795 | 1.294 | 1.748 |