| | 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** | O76074 | PDE5A_HUMAN | Homo sapiens | cGMP-specific 3',5'-cyclic phosphodiesterase | 17.826004 | NaN | S102 | 38665916 | MERAGPSFGQQRQQQQPQQQKQQ QRDQDSVEAWLDDHWDFTFSYFVR KATREMVNAWFAERVHTIPVCKEGI RGHTESCSCPLQQSPRADNSAPGTP TRKISASEFDRPLRPIVVKDSEGTVS FLSDSEKKEQMPLTPPRFDHDEGDQ CSRLLELVKDISSHLDVTALCHKIFL HIHGLISADRYSLFLVCEDSSNDKFL ISRLFDVAEGSTLEEVSNNCIRLEW NKGIVGHVAALGEPLNIKDAYEDPR FNAEVDQITGYKTQSILCMPIKNHRE EVVGVAQAINKKSGNGGTFTEKDEK DFAAYLAFCGIVLHNAQLYETSLLEN KRNQVLLDLASLIFEEQQSLEVILKKI AATIISFMQVQKCTIFIVDEDCSDSFS SVFHMECEELEKSSDTLTREHDAN KINYMYAQYVKNTMEPLNIPDVSKD KRFPWTTENTGNVNQQCIRSLLCTP IKNGKKNKVIGVCQLVNKMEENTGK VKPFNRNDEQFLEAFVIFCGLGIQN TQMYEAVERAMAKQMVTLEVLSYH ASAAEEETRELQSLAAAVVPSAQTLK ITDFSFSDFELSDLETALCTIRMFTD LNLVQNFQMKHEVLCRWILSVKKN YRKNVAYHNWRHAFNTAQCMFAAL KAGKIQNKLTDLEILALLIAALSHDL DHRGVNNSYIQRSEHPLAQLYCHSI MEHHHFDQCLMILNSPGNQILSGL SIEEYKTTLKIIKQAILATDLALYIKRR GEFFELIRKNQFNLEDPHQKELFLA MLMTACDLSAITKPWPIQQRIAELVA TEFFDQGDRERKELNIEPTDLMNRE KKNKIPSMQVGFIDAICLQLYEALTH VSEDCFPLLDGCRKNRQKWQALAE QQEKMLINGESGQAKRN | None | None | None | None | None | None | None | None | None |