	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	Q9GZU2	PEG3_HUMAN	Homo sapiens	Paternally-expressed gene 3 protein	21.01871	S1009	NaN	37217939;37340703	MLPPKHLSATKPKKSWAPNLYELDS DLTKEPDVIIGEGPTDSEFFHQRFRN LIYVEFVGPRKTLIKLRNLCLDWLQP ETRTKEEIIELLVLEQYLTIIPEKLKP WVRAKKPENCEKLVTLLENYKEMY QPEDDNNSDVTSDDDMTRNRRESS PPHSVHSFSDRDWDRRGRSRDMEP RDRWSHTRNPRSRMPPRDLSLPVV AKTSFEMDREDDRDSRAYESRSQDA ESYQNVVDLAEDRKPHNTIQDNME NYRKLLSLVQLAEDDGHSHMTQGH SSRSKRSAYPSTSRGLKTMPEAKKST HRRGICEDESSHGVIMEKFIKDVSRS SKSGRARESSDRSQRFPRMSDDNW KDISLNKRESVIQQRVYEGNAFRGG FRFNSTLVSRKRVLERKRRYHFDTD GKGSIHDQKGCPRKKPFECGSEMRK AMSVSSLSSLSSPSFTESQPIDFGAM PYVCDECGRSFSVISEFVEHQIMHT RENLYEYGESFIHSVAVSEVQKSQV GGKRFECKDCGETFNKSAALAEHRK IHARGYLVECKNQECEEAFMPSPTF SELQKIYGKDKFYECRVCKETFLHSS ALIEHQKIHFGDDKDNEREHERERE RERGETFRPSPALNEFQKMYGKEK MYECKVCGETFLHSSSLKEHQKIHT RGNPFENKGKVCEETFIPGQSLKRR QKTYNKEKLCDFTDGRDAFMQSSE LSEHQKIHSRKNLFEGRGYEKSVIH SGPFTESQKSHTITRPLESDEDEKAF TISSNPYENQKIPTKENVYEAKSYER SVIHSLASVEAQKSHSVAGPSKPKV MAESTIQSFDAINHQRVRAGGNTSE GREYSRSVIHSLVASKPPRSHNGNE LVESNEKGESSIYISDLNDKRQKIPA RENPCEGGSKNRNYEDSVIQSVFRA KPQKSVPGEGSGEFKKDGEFSVPSS NVREYQKARAKKKYIEHRSNETSVI HSLPFGEQTFRPRGMLYECQECGEC FAHSSDLTEHQKIHDREKPSGSRNY EWSVIRSLAPTDPQTSYAQEQYAKE QARNKCKDFRQFFATSEDLNTNQKI YDQEKSHGEESQGENTDGEETHSE ETHGQETIEDPVIQGSDMEDPQKDD PDDKIYECEDCGLGFVDLTDLTDHQ KVHSRKCLVDSREYTHSVIHTHSISE YQRDYTGEQLYECPKCGESFIHSSFL FEHQRIHEQDQLYSMKGCDDGFIAL LPMKPRRNRAAERNPALAGSAIRCL LCGQGFIHSSALNEHMRLHREDDL LEQSQMAEEAIIPGLALTEFQRSQTE ERLFECAVCGESFVNPAELADHVTV HKNEPYEYGSSYTHTSFLTEPLKGAI PFYECKDCGKSFIHSTVLTKHKELHL EEEEEDEAAAAAAAAAQEVEANVHV PQVVLRIQGLNVEAAEPEVEAAEPEV EAAEPEVEAAEPNGEAEGPDGEAAE PIGEAGQPNGEAEQPNGDADEPDGA GIEDPEERAEEPEGKAEEPEGDADE PDGVGIEDPEEGEDQEIQVEEPYYDC HECTETFTSSTAFSEHLKTHASMIIF EPANAFGECSGYIERASTSTGGANQ ADEKYFKCDVCGQLFNDRLSLARHQ NTHTG	True	False	2.684	4.865	1.842	1.392	0.997	1.944	1.877