	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	extracellula region
0	P0CG38	POTEI_HUMAN	Homo sapiens	POTE ankyrin domain family member I	18.01798	NaN	NaN	30397120;37217939;38665916	MVAEVDSMPAASSVKKPFVLRSKM GKWCRHCFPCCRGSGKSNVGTSGD QDDSTMKTLRSKMGKWCCHCFPCC RGSGKSNVGTSGDHDDSAMKTLRS KMGKWCCHCFPCCRGSGKSNVGA WGDYDDSAFVEPRYHVRREDLDKL HRAAWWGKVARKDLIVMLRDTDVN KQDKQKRTALHLASANGNSGVVKLL LDRRCQLNVLDNKKRTALTKAVQCQ EDECALMLLEHGTDPNIPDEYGNTT LHYAIYNEDKLMAKALLLYGADIESK NKHGLTPLLLGVHEQKQQVVKFLIK KKANLNALDRYGRTALILAVCCGSA SIVSLLLEQNIDVSSQDLSGQTAREY AVSSHHHVICQLLSDYKEKQMLKIS SENSNPEQDLKLTSEEESQRFKGSE NSQPEKMSQEPEINKDGDREVEEE MKKHESNNVGLLENLSNGVTAGNG DDGLIPQRKSRTPENQQFPDNESEE YHRICELVSDYKEKQMPKYSSENSN PEQDLKLTSEEESQRLKGSENGQPE KRSQEPEINKDGDRELENFMAIEEM KKHGSTHVGFPENLTNGATAGNGD DGLIPPRKSRTPESQQFPDTENEEY HSDEQNDTQKQFCEEQNTGILHDEI LIHEEKQIEVVEKMNSELSLSCKKEK DFLHENSTLREEIAMLRLELDTMKH QSQLRKKKYLEDIESVKKKNDNLLK ALQLNELTMDDDTAVLVIDNGSGM CKAGFAGDDAPRAVFPSIVGRPRQQ GMMGGMHQKESYVGKEAQSKRGIL TLKYPMEHGIITNWDDMEKIWHHT FYNELRVAPEEHPILLTEAPLNPKAN REKMTQIMFETFNTPAMYVAIQAML SLYTSGRTTGIVMDSGDGVTHTVPIY DGNALPHATLRLDLAGRELTDYLMK ILTERGYRFTTMAEREIVRDIKEKLC YVALDFEQEMAMAASSSSLEKSYEL PDGQVITIGNEWFRCPEALFQPCFL GMESCGIHETTFNSIMKSDVDIRKD LYTNTVLSGGTTMYPGMAHRMQKE IAALAPSMLKIRIIAPPKRKYSVWVGG SILASLSTFQQMWISKQEYDESGPSI VHRKCF	None	None	None	None	None	None	None	None	None