	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	O75808	CAN15_HUMAN	Homo sapiens	Calpain-15	24.917859	NaN	S296;S335;S338;S1070	28411811	MATVGEWSCVRCTFLNPAGQRQCSI CEAPRHKPDLNHILRLSVEEQKWPC ARCTFRNFLGKEACEVCGFTPEPAP GAAFLPVLNGVLPKPPAILGEPKGSC QEEAGPVRTAGLVATEPARGQCEDK DEEEKEEQEEEEGAAEPRGGWACP RCTLHNTPVASSCSVCGGPRRLSLP RIPPEALVVPEVVAPAGFHVVPAAPP PGLPGEGAEANPPATSQGPAAEPEP PRVPPFSPFSSTLQNNPVPRSRREVP PQLQPPVPEAAQPSPSAGCRGAPQG SGWAGASRLAELLSGKRLSVLEEEA TEGGTSRVEAGSSTSGSDIIDLAGDT VRYTPASPSSPDFTTWSCAKCTLRN PTVAPRCSACGCSKLHGFQEHGEPP THCPDCGADKPSPCGRSCGRVSSAQ KAARVLPERPGQWACPACTLLNALR AKHCAACHTPQLLVAQRRGAAPLRR RESMHVEQRRQTDEGEAKALWENI VAFCRENNVSFVDDSFPPGPESVGF PAGDSVQQRVRQWLRPQEINCSVFR DHRATWSVFHTLRPSDILQGLLGNC WFLSALAVLAERPDLVERVMVTRSL CAEGAYQVRLCKDGTWTTVLVDDM LPCDEAGCLLFSQAQRKQLWVALIE KALAKLHGSYFALQAGRAIEGLATLT GAPCESLALQLSSTNPREEPVDTDLI WAKMLSSKEAGFLMGASCGGGNM KVDDSAYESLGLRPRHAYSILDVRDV QGTRLLRLRNPWGRFSWNGSWSDE WPHWPGHLRGELMPHGSSEGVFW MEYGDFVRYFDSVDICKVHSDWQE ARVQGCFPSSASAPVGVTALTVLERA SLEFALFQEGSRRSDAVDSHLLDLCI LVFRATFGSGGHLSLGRLLAHSKRA VKKFVSCDVMLEPGEYAVVCCAFNH WGPPLPGTPAPQASSPSAGVPRASP EPPGHVLAVYSSRLVMVEPVEAQPT TLADAIILLTESRGERHEGREGMTCY YLTHGWAGLIVVVENRHPKAYLHVQ CDCTDSFNVVSTRGSLRTQDSVPPL HRQVLVILSQLEGNAGFSITHRLAH RKAAQAFLSDWTASKGTHSPPLTPE VAGLHGPRPL	None	None	None	None	None	None	None	None	None