

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
P01132	EGF_MOUSE	Mus musculus	Pro-epidermal growth factor	25.655977	NaN	NaN	38359610	MPWGRRPTWLLAFLLVFLKISILSV TAWQTGNCPGPLERSERSGTCAGP APFLVFSQGSISRDPDGTNHQQLV VDAGISADMDIHYKKERLYWVDVER QVLLRVFLNGTGLEKVCNVERKVS LAIDWIDDEVLWVDQNGVITVTD TGKNSRVLLSSLKHPNIAVDPIERL MFWSSSEVTGSLHRAHLKGVVKT LETGGISVLTLDVLDKRLFVWQDSG EGSHAYIHSCDYEGGSVRLIRHQAR HLSLSSMAFFGDRIFYSVLKSKAIWIA NKHTGKDTVRINLHPSFVTPGKLMV VHPRAQPRTEDAAKDPPELLKQRG RPCRFGLCERDPKSHSSACAEGYTL SRDRKYCEDVNECATQNHGCTLGC ENTPGSYHCTCPTGFVLLPDGKQCH ELVSCPGNVSKCSHGCVLTS DGPRC ICPAGSVLGRDGKTCTGCSSPDNGG CSQICLPLRPGSWECDPFGYDLQS DRKSCAASGPQPLLLFANSQDIRHM HFDGTDYKVLLSRQMGMVFALDYD PVESKIYFAQTALKWIERANMDGSO RERLITEGVDLEGLALDWIGRRIYW TDSGKSVVGGSDLSGKHHRIIIQERI SRPRGIAVHPRARRLFWTDVGMSPR IESASLQGS DRVLIASSNLEPSGITI DYLTDTLYWCDTKRSVIEMANLDGS KRRRLIQNDVGHPPFLAVFEDHLWV SDWAIPSVIRVNRKTGQNRVRLQGS MLKPSLVVVHPLAKPGADPCLYRN GGCEHICQESLGTARCLCREGFVKA WDGKMCLPQDYPILSGENADLSKEV TSLSNSTQAEVPDDDGTESSTLVAEI MVSGMNYEDDCGPGGCGSHARCVS DGETAECQCLKGFARDGNLCSDDIDE CVLARSDCPSTSSRCINTEGGYVCRC SEGYEGDGISCFDIDECQRGAHNCG ENAACTNTEGGYNCTCAGRPS SPGL SCPDSTAPSLLGEDGHHLDRNSYPG CPSSYDGYCLNGGVCMHIESLDSYT CNCVIGYSGDRCOTRDLRWELRH AGYGQKHDIMVAVCMVALVLLVL GMWGTYYYRTRKQLSNPPKNPCDE PSGSVSSSGPNSSSGAAVASCQPW FVLEKHQDPKNGSLPADGTNGAVV DAGLSPSLQLGSVHLTSWRQKPHID GMGTGQSCWIPSSDRGPQIEGNS HLPYSYRVPVGEKLSLQSANGSCH RAPDLPRQTEPVQ	False	True	3.219	3.71	3.157	3.169	2.882	5.0	5.0