	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	e r
0	P55197	AF10_HUMAN	Homo sapiens	Protein AF-10	31.79131	T266;T268;S269;T341;S343;S346;S352;S496;S790	S217;S252;S436;S532;S684;S686;S689	38665916;34019948;33214551;39302247;31492838;29237092;40307207;20068230;35132862;39534244;35254053	MVSSDRPVSLEDEVSHSMKEMIGG CCVCSDERGWAENPLVYCDGHGCS VAVHQACYGIVQVPTGPWFCRKCES QERAARVRCELCPHKDGALKRTDN GGWAHVVCALYIPEVQFANVSTMEP IVLQSVPHDRYNKTCYICDEQGRESK AATGACMTCNKHGCRQAFHVTCAQ FAGLLCEEEGNGADNVQYCGYCKY HFSKLKKSKRGSNRSYDQSLSDSSS HSQDKHHEKEKKKYKEKDKHKQKH KKQPEPSPALVPSLTVTTEKTYTSTS NNSISGSLKRLEDTTARFTNANFQE VSAHTSSGKDVSETRGSEGKGKKSS AHSSGQRGRKPGGGRNPGTTVSAAS PFPQGSFSGTPGSVKSSSGSSVQSPQ DFLSFTDSDLRNDSYSHSQQSSATK DVHKGESGSQEGGVNSFSTLIGLPS TSAVTSQPKSFENSPGDLGNSSLPT AGYKRAQTSGIEEETVKEKKRKGNK QSKHGPGRPKGNKNQENVSHLSVS SASPTSSVASAAGSITSSSLQKSPTLL RNGSLQSLSVGSSPVGSEISMQYRH DGACPTTTFSELLNAIHNGIYNSNDV AVSFPNVVSGSGSSTPVSSSHLPQQ SSGHLQQVGALSPSAVSSAAPAVATT QANTLSGSSLSQAPSHMYGNRSNS SMAALIAQSENNQTDQDLGDNSRN LVGRGSSPRGSLSPRSPVSSLQIRYD QPGNSSLENLPPVAASIEQLLERQW SEGQQFLLEQGTPSDILGMLKSLHQ LQVENRRLEEQIKNLTAKKERLQLL NAQLSVPFPTITANPSPSHQIHTFSA QTAPTTDSLNSSKSPHIGNSFLPDNS LPVLNQDLTSSGQSTSSSSALSTPPP AGQSPAQQGSGVSGVQQVNGVTVG ALASGMQPVTSTIPAVSAVGGIIGALP GNQLAINGIVGALNGVMQTPVTMSQ NPTPLTHTTVPPNATHPMPATLTNS ASGLGLLSDQQRQILIHQQQFQQLL NSQQLTPEQHQAFLYQLMQHHHQ QHHQPELQQLQIPGPTQIPINNLLAG TQAPPLHTATTNPFLTIHGDNASQK VARLSDKTGPVAQEKS	True	False	4.502	5.0	False	F