

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochnd
Q501J7	PHAR4_MOUSE	Mus musculus	Phosphatase and actin regulator 4	60.959274	S190;S192;S193;T194;T199;S202;T206;T231	S116;S118;S129;S145;S264;S285;S335;S337;S420;T425;S436;S446;S457;S503;S505;S549;S582;S620	22517741;22645316;40885482;36852467;36084651;35822049;40021952;34678516;30059200	MEDPSEEAEPSPGDPGMGMSVEA GDTTPPTKRKSKFSALGKIFPWKW RKKKSSDKFKETSEVLERKISMRKP REELVKRGVLEDPEDQDGEDSGKLS HAALKNGHTTPIGSARSSSPVLVEEE PERSLRNLTPEEESKRLGSGTSQP NSEAEPGPEHAPKQPLLPKRPSSS CEAKEVPAGSTARSVSSTSGSTTVTS AATTAATDMTKTVKSFVGPPTAPAPA PRTLPAAPASANTAATTTAPAKQPI PPPKPAQRNSNPIAELSQAMNSGTV LSKPSPLPPKRGIPSTSIPLPAAS FTTKTANDQREKTIVSLCLEPPLIIPPS SPSPPLPTHIPPEPRSPLVPAKTFQI VPEVEFSSSDLFQDISQQEDQKTEV PKKIQQSFGESHPSRLPPLPLHIRI QQALTSPLVTPPLEGTHRAHSLLE NSDSFSEDTGTLGRTRSLPITIEMLK VPDDEEEEQTCPFVEDVTVTSATPSL PLCLREEEEKESDSEGPICYRDEEE DDDDDESHQSALANRVKRDITLAM KLSSRFSEPETNLNSWPRKSKEEW NEIRHQIGNTLIRRLSQRPTAEELEQ RNILQPKNADRQAEKREIKRRLTR KLSQRPTVAELLARKILRFNEYVEVT DAHDYDRRADKPWTKLTPADKAAIR KELNEFKSSEMVEHVDSKHFTRYH RP	False	False	2.036	2.93	0.633