

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q6PAR5	GAPD1_MOUSE	Mus musculus	GTPase-activating protein and VPS9 domain-containing protein 1	11.752367	S798;T800;S801	S227;T390;T458;S466;T470;S566;S569;S742;S746;S757;T762;S766;S876;S902;S903;S908;S964;S1017;S1044;S1076	34418053	MVKLDIHTLAHHLKQERLYVSSEKQ LIQRLNADVLTAEKLYRTAWIAKQQ RINLDRLIITSAEASPAECCQHAKILE DTQFVDGYKQLGFQETAYGEFLSRL RENPRLIASSLVAGEKLNQENTQSVI YTVFTSLYGNCIMQEDESULLQVRLY LIEFELKESDNPRLLRRGTCAFSIL FKLFSEGLFSAKFLTLATLHEPIMQL LVEDEDHLETDPNKLIERFSPAQQE KLFGEKGSDFRQKVQEMVDSNEA KLVALVNKFIGYLKQNTYCFPHSLR WIVSQMYKTLSCVDRLEVGEVRAM CTDLLLACFICPAVVNPEQYGIISDAP INEVARFNLMOVGRLLQQLAMTGT EEGDPRTKNSLGKFDKGCVA AFLDV VIGGRAVETPPMSSVNLEGLSRTVV YISYSQLITLVNFMKSVMSGDQLKE DRMALDNLLANLPQAKPGKSSSLD MTPYSTPQMSPATTPANKKNRPLIA TRSRSRSNMLMDLHMDHEGSSQET IQEVQPEEVLVISLGTGPQLTPGMM SENEVLNMQLSGGQGDVPVDENK LHGKPKTLRFSLCSDNLEGISEGPS NRSNSVSSLDLEGESVSELGAGPSG SNGVEALQLEHEQATTQDNLDDKL RKFEIRDMMGLTDDRDISETVSETW STDVLGSDFDPNVDEDRLQEIAGAA AENVLGSLCLPGSGSVLLDPCTGST ISETTSEAWSVEVLPSDSEAPDLKQE ERLQELESCSGLGSTSDDTDVREVS SRPSTPGLSVVSGISATSEIPNKIED LRSECSSDFGGKDSVTSPDMDDIAH GAHQLTSPPSQSESLAMFDPLSSH EGASAVVRPKVHYARPSHPPDPPIIL EGAVGGNEARLPNFGSHVLTAAEM EAFKQRHSYPERLVRSRSDIVSSVR RPMSDPSWNRPGNEELPPAAATG ATSLVAAPHSSSSSPSKDSSRGETEE RKDSDDERSDRSRPWWWRKRFVSAM PKAPIPFRKKEKQEKDKDDLGPDRF STLTDEPSPRLSAQQAQVAEDILDKYR NAIKRTSPSEGAMANDESAEVMGD GESAHDSPREEALQNISADDLPDSA SQAHPQDSAFSYRDVKKKLRRLALC SADSVAFPVLTHSTRNGLPDHTDPE DNEIVCFLKVQIAEAINLQDKSLMAQ LQETMRCVCRFDNRTCRLKLLASIAE DYRKRAPYIAYLTRCRQGLQTTQAH ERLLQRVLRDKEVANRYFTTVCVRL

LLESKEKKIREFIQDFQKLTAADDKT
AQVEDFLQFLYGVMAQDVIWQNAS
EEQLQDAQLAIERSVMNRIFKLAFFP
NQDGDILRDQVLHEHIQRLSKVVTA
NHRALQIPEVYLREAPWPSAQSEIRT
ISAYKTPRDKVQCILRMCSTIMNLLS
LANEDSVPGADDFVPVLVFLIKANP
PCLLSTVQYISSFYASCLSGEESYWW
MQFTAAVEFIKTIDDRK