

UniprotKB ID	Entry name	organism	full name	oglcnaicscore	oglcnaicsites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
G3X982	AOXC_MOUSE	Mus musculus	Aldehyde oxidase 3	23.627151	NaN	S320	37453647	MSPSKESDELIFVNGKKVTERNAD PEVNLLFYLRKVIRLTGTKYGC GGDD CGACTVMISRYDPISKRISHFSATAC LVPICSLHGAAVTTVEGIGSTKTRHP VQERIAKGHGTQCGFCTPGMVMSIY TLLRNHPEPSTEQIMETLGGNLCRC TGYRPIVESAKSFCSSTCCQMNGE GKCCLDEEKNEPERKNSVCTKLYEK KEFQPLDPTQELIFPEL MRMAEES QNTVLTFRGERTTWIAPGTLNDLLE LKMKHPAPLVIGNTYLGLHMKFTD VSYPIIISPARILELFVVTNTKQGLTLG AGLSLTQVKNVLSDVVSRPKKTKOI YCALLKQLKTLAQQIRNVASLGGHI ISRLPTSDLNPILGIGNCILNVASTEG IQIPLNDHFLAGVDPDAILKPEQVLIS VFPVPRSSKWEFVSAFRQAPRQONAF ATV NAGMKVVKEDTNTITDLGILYG GIGATVISADKSCRQLIGRCWDEEM LDDAGKMICEEV SLLMAAPGGMEE YRKT LAISFLFMFYLDV LKQLKTRDP HKYPDISQKLLHILEDPLTMPYGM QSFQD VDFQQLQDPIGRPIMHQSG IKHATGEAVFCDDMSVLP GELFLAV VTSSKSHAKIISLDASEALASLGVVD VVTARDVPGDNGREESLYAQDEVI CVGQIVCAVAADSYAHAQQA AKKVK IVYQDIEPMIVTVQDALQYESFIGPER KLEQGNVEEAFQCADQILEGEVHLG GQEHFYMETQSVRVVPKGEDKEMD IYVSSQDAAFTQEMVARTLGIPKNRI NCHVKRVGGAFGGKASKPGLLASVA AVAAQKTGRPIRIFILERRDDMLITGG RHPLLGKYGIFMNNNGKKAADIQLY INGGCTPDDSELVIEYALLKLENAYKI PNLRVRGRVCKTNLPSNTAFRGFGF POGAFVTETCMSAVA AKCRLPPEKV RELNMYRTIDRTIHNQEFDPNLLQ CWEACVENS SYNRKKADEFNQQ RFWKKRGIAIIPMKFSVGF PKTFYYQ AAALVQIYTDG SVLV AHGGVELGQGI NTKMIQVASRELKIPMSYIHLDEMS TVTVPNTVTTGASTGADVNGRAVQN ACQILMKRLEPIIKQNPSGTWEEVW KEAFVQSISLATSATGYFRGYQADM DW EKGEGDIFPFYFVGAACSEVEIDCLT GAHKNIRTDIVMDGSFSINPAVDIGQ IEGAFVQGLGLYTLLEELKYSPEGLY TRGPHQYKIASVTDIPEEFHV SLLTP TPNPKAIYSSKGLGEAGTFLGCSVFF AIAAAVAAAREERGLSPIWAINSPAT AEVIRMACEDQFTNLVPQTDSKCK PWSIPVA	None	None	None	None	None	None	None		