

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
F4J8G7	VIR_ARATH	Arabidopsis thaliana	Protein virilizer homolog	27.144759	NaN	NaN	28154133	MVRSEPCVLFQAQTFVHPQLDEYVDE VIFAEPVIIACEFLEQNASSSSQAVS LVGATSPPSFALEVFVRCGEGSKFKR LCNPFLYTPSAPYPLEVEAVVTNHLV VRGSYRSLSLIVYGNIVKDLGQYNIIL EGRSVTDIVSSTEGNLEDLPLVLHSV NRTIEECLSSLDIVSLPLAAVDLPVEV KRLLQLLLKIFDKLATNDVVKFVDT VVGVSYSYVDNDVDFLKNKNC SAV TSSLD SGLFHDIVDRVKEDILD LNEI QESDVALGLFSFLESETYLATSQQLV VMLSPYIQFERDSLCTVLPKLSKGKA TLLGLSLAFLLCSGREGCLOFVNSG GMDQLVYLFHGDGQNSTITITLLLLG VVEQATRHSVGCEGFLGWWPREDG SIPSGKSEGYCLLLKLLMQKPCHEIA SLAIYILRRLRIYEVISRYEFAVLSALE GLSN SHGAATHNLNMLS DAKS QLQ KLQNLMKSLGSVEDPSPSAYAERSL VSDHSEGWLSYKATSKLTSSWTC PF YSSGIDSHILALLKERGFPLSAALLS MPELH SKVGDIMDVFTDIAMFIGNII LSFMFSRTGLSFLHHP ELTATHIQS LKGSDVLDNKEECVPLHYASILSKGF TCSLLEIGINLEMHLRVVSAVDRLLK SIQQTEEF LWILWELRDVRSRDCGR EALLTLGVFPEALAVLIEALHSAKDM EPAVENSGISPLNLAICHSAAEIFEVI VSDSTASCLHAWIEHAPVLH KALHT LSPGGSNRNDAPSRLKWKIDAGVVY HKHGVGGLLR YA AVLASGGDAQLSS SSILALDLTPAENGAGESTNVSEMN VLDNLGKVIFEKSFEGVNLSDSSISQ LTTALRILALISDNSTVAAALYDEGAV TVVYAILVNC SFMFERSSNIYDYLVD DDHGCSSISDFLSERNREQSLVDLLI PSLVLLISVLQRLOGTKEQYRNTKLM KALLRLHREVSPKLAACAADLSSHYP DSALGFGAVCHLIVSALVCWPVYGW IPGLFHTLLSGVQTSSVPALGPKETC SFLCILSDILPEEGVWFWKSGMPLLS GLRKLAVGTLMGQPKEKQINWYLEP GPLEKLNHLTPNLDKIAKHQH HAV SALVVIQDMLRVFIVRIACQRVEHAS ILLRPIFSSIRDGILDQSSTRDTEAYM VYRYLNFLASLLEHHPHAKGLLLEEGI VQLLVEVLERCYDATYPSENRVLEY GIVSASSVIQWCIPAFRSISLLCDSQV PLLCFQKELLASLSAKDCALIPPFV LKF CQVLPVGNELLSCLGAFKDLSS CGEGQDGLVSLFLHFLFSGTEESVSE RWC DTNSLSLDQLDMKKNPPFLSC WIKLLNSINSKDG LSSLAMKAVNVL SVGSIRLCLDGKSLDSKKVAALKSLF GLPSEFSGTDTFREENIGLIEQMVTL LSSMTSGSDSSATAEMKPYLHEASQ SLLSLLK DGNIDDIISCKGVFVSPGN LDMDDL VSRNIEDDLYQRGLEDKF WWECPETLPERLPQSSLP AKRKLPT LESSRRAKGENSSVDIPTQNSIQRG MGSVSLPPAPTRRDAFRQRKPNTSR PPSMHVDDYVARERSVDTAGNSNAI TISRAGSSGRPPSIHVDEFMARQRE RGQNPSTIVGEAVVQVKNP TPARD TEKVAGKPKQFKADPDDLQGIDIVF DGECEGPDDKLPFLQPDENLMQP APVMVEQNSPHSIVEETESDANGSS QFSHMGTPVASNV DENAQSEFSSRI SVSRPEMSLIREPSISSDRKFVEQAD EAKKMAPLKSAGISESGFIPAYHMP GSSGQNSIDPRVGPQGFYKSGQQH TGHIHGGFSGRGVYEQKVMNPQPL PLVPPPSVSPVIPHSSDLSLNQSSPFI SHGTQSSGGPTRLMPPLPSAIPQYSS	None	None	None	None	None	None	None		

