	UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochndrion	endoplasmic reticulum	golgi apparatus	plasma membrane	extracellular region
0	Q8VID5	RECQ5_MOUSE	Mus musculus	ATP-dependent DNA helicase Q5	25.810548	NaN	S489;S492;T527;S728	29187734	MSARPFSTPFDRERRVRSTLKKVFG FDSFKTPLQESATMAVVKGAEDVFV CMPTGAGKSLCYQLPALLASGITIVV SPLIALIQDQVDHLLALKVQVSSLNS KLSVQERKELLSDLERDKPRTKLLYI TPEMAASASFQPTLNSLVSRNLLSYL VVDEAHCVSQWGHDFRPDYLRLGA LRSRLAHAPCVALTATATPQVQEDV FAALHLKQPVASFKTPCFRANLFYD VQFKELIPDVYGNLRDFCLKALGQK AENGSSSGCGIVYCRTREACEQLAIE LSSRGVNAKAYHAGLKASDRTQVQN EWMEEKVPVIVATISFGMGVDKANV RFVAHWNIAKSMAGYYQESGRAGR DGKPSWCRLYYSRNDRDQVSFLIRK ELAKLQEKRGNKPSDKATLLAFDAL VTFCEEVGCRHAAIAKYFGDAPPAC AKGCDYCQNPAAITKKLDALERSSS WSKTCIGPSQGNGFDPELYEGGRRG YGGFSRYDEGSGGSGDEGRDEAHK REWNLFYQKQMSLRKGKEPKIEEFT PPDEDCPLREASSRKIPKLTVKAREH CLRLLEEALISNHQAAGSTHGADLQ AKAVELEHETFRNAKMVNLYKASVL KKVAEIHKASKDGQLYDMESGTKSC GAAAEFSEPSDYDIPPTSHVYSLKPK RVGAGFSKGPCSFQTATELLGKSHS QKQAPEAMLEGGQEPPGWVCDLQD EDRSKPHPGYQEKALGSSVNCGDPS PEKKTKGSSQGSAKARASKKQQLLA TAARKDSQNITRFLCQRTESPPLPAS VPRSEDASPSCGDVPGKCTQEVGAQ GHLVAVFQTEGPRERPSTCSLRDQS FPEGQPSPLKETQAEKRPRPQQGNP ERRAQKRLRPSTKSSILAEAKDSTLA SDRSTENKVAQEPCQLSASGTSLRE AADIVVRHLTPFYKEGRFISKDLFKG FARHLSHLLAQQLSPGRSVKEEAQS LIKQFFHNRARCESEADWHSLRGPQ R	None	None	None	None	None	None	None	None	None