	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	Q5XGY5	Q5XGY5_XENLA	Xenopus laevis	Pyruvate dehydrogenase E1 component subunit beta	26.754248	NaN	NaN	18617508	MAALRSLFCGGKSSVSVFLQRSFHR SGPAALQVTVRDALNQAMDEEIERD ERVFLLGEEVAQYDGAYKISRGLWK KYGDKRVMDTPITEMGFAGIAVGAA MAGLRPICEFMTFNFSMQAIDQVIN SAAKTHYMSGGLVSVPIVFRGPNGA SAGVAAQHSQCFAAWYGHCPGLKV VSPWNAEDAKGLLKASIRDDNPVVF LENELMYGVPFELSEEVQSKDFVVPI GKAKIERPGSQITLVSHSRSVGHCLE AASVLEKEGIDCEVINMRTIRPMDTE TIEASVVKTNHLVTVEGGWPQFGVG AEICANIMEGPAFNYLDAPVVRVTGT DVPMPYAKILEENCVPQVKDIIFAVK KTLNVL	None	None	None	None	None	None	None	None	None