

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q24368	ISWI_DROME	Drosophila melanogaster	Chromatin-remodeling complex ATPase chain Iswi	13.191182	NaN	S67	28604694	MSKTDTAAVEATEENSNETTSDAAT SSSGEKEAEFDNKIEADRSRRFDLL KQTEIFTHFMTNSAKSPTKPKGRPK KIKDKDKEKDVADHRHRKTEQEED ELLAEDSATKEIFRFDASPAYIKSGE MRDYQIRGLNWMISLYENGLGILA DEMGLGKTLQTISLLGYLKHFNQA GPHIVVPKSTLQNWVNEFKKWCP LRAVCLIGDQDTRNTFIRDVLMPE WDVCVTSYEMCIREKSVFKKFNWR YLVIDEAHRIKNEKSKLSEILREFKTA NRLITGTPLQNNLHELWALLNLL PDVFNSSDFDEWFNTNTCLGDDA LITRLHAVLKPFLRLKAEVEKRLK PKKEMKIFVGLSKMQRDWYTKVLLK DIDVNGAGKVEKMRLQNILMQLR KCTNHPYLFDAEPGPPYTTDTHLV YNSGKMAILDKLLPKLQEQSRVLIF SQMTRMLDILEDYCHWRNYNYCRL DGQTPHEDRNRQIQEFNMDNSAKF LFMLSTRAGGLGINLATADVVIYDS DWNPQMDLQAMDRAHRIGQKKQV RVFRLITESTVEEKIVERAEVKLRD KMVIQGGRLVDNRSNQLNKDEMLN IIRFGANQVFSSKETDITDEDIDVILE RGEAKTAEQKAALDSLGESSLRTFT MDTNAGEAGTSSVYQFEGEDWREKQ KLNALGNWIEPPKREKANYAVDAY FREALRVSEPKAPKAPRPPKQPIVQD FQFFPRLFELLDQEIYFRKTVGYK VPKNTLGSATKQVQREEQRKIDEA EPLTEEEIQEKENLLSQGFTAWTKR DFNQFIKANEKYGRDDIDNIAKDVE GKTPEEVIEYNAVFWERCTELQDIER IMGQIERGEGKIQRRLSIKKALDQKM SRYRAPFHQLRLQYGNKKGKNYTEI EDRFLVCMLHKLGFDKENVYEELRA AIRASPQFRFDWFIKSRTALELQRR NTLITLIERENIELEEKERAEEKKKAP KGSVSAGSGSASSNTPAPAPQPKAS QKRKSEVVATSSNSKSKKKK