

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
P52789	H XK2_HUMAN	Homo sapiens	Hexokinase-2	11.272169	T778;S792	NaN	23301498;30379171;28510447;35132862;30397120	MIASHLLAYFFTELNHDQVQKVDQY LYHMRLSDETLLEISKRFKEMEKGLGATTHPTAAVKMLPTFVVRSTPDGTEHGEFLALDLGGTNFRVLWVKVTDNGLQKVEMENQIYAIPEDIMRSGTQLFDHIAECLANFMDKLQIKDKKLP LGFTFSFPCHQTKLDESFLVSWTKGFKSSGVEGRDVVALIRKAIQRRGDFDIDIVAVVNDTVGTMMTTCGYDDHNC EIGLIVGTGSNACYMEEMRHIDMVEGDEGRMCINMEWGAFGDDGSLNDIRTEFDQEIDMGS LNPGKQLFEKMISGMYMGELVRLILVKMAKEELLFGGKLSPELLNTGRFETKDISDIEGEKDGI RKAREVLMRLGLDPTQEDCVATHRICQIVSTRSASLCAATLA AVLQRIKENKGEERLRSTIGVDGSVYKKHPHFAKRLHKT VRRRLVPGCDVRF LRSE DGSG KGAAMVTAVAYRLADQHRARQKTLEHLQLSHDQLLEVKRRMKVEMERGLSKETHASAPVKMLPTYVCATPDGTEKGDFLALDLGGTNFRVLLVVRVNGK WGGVEMHNKIYAIPQEV MHGTGDELFDHIVQCIADFLEYMGMGVSLPLGFTFSFPCQONS LDESILLKWT KGF KASGCEGEDVVTLLKEAIHRREEFDLDVAVVNDTVGTMMTTCGFEDPHCEVGLIVGTGSNACYMEEMRNVELVE GEEGRMCVNMEWGAFGDNGCLDDFRTEFDVAVDEL SLNPGKQRF EKMISSGMYLGEIVRNILIDFTKRGLLFRGRI SERLKTRGIFETKFLSQIESDCLALLQVRAILQHLGLESTCDDSIIVKEVCTVARRAAQLCGAGMAAVVDRIREN GLDALKVTVGVDGTLYKLHPHFAKVMHETVKDLAPKCDVSFLQSE DGSG KGAALITAVACRIREAGQR

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