

UniprotKB ID	Entry name	organism	full name	oglnacscore	oglnac sites	phosphorylation sites	PMIDS	sequence
Q8CHP6	PHC3_MOUSE	Mus musculus	Polyhomeotic-like protein 3	29.698267	T238	S231;S261;S269;S312;T607;T612;S614;S759;S760	19458039;22645316;21606357;30059200	MDSEPPSSGTSVSTTASSTTTTTITSS SRMQQPQISVYSGSDRHAVQVIQQA LHRPPSSAAQYLQOMYAAQQQHLM LHTAALQQQHLSSSQLSAAVQAS LSSGRPSTSPGTGSVTQQSSMSQTSIL SASPAPAQLMNRSTSSSTSGSITQ QTMLLGSTSPTLTASQAQMYLRAQ MLIFTPATTVAAVQSDIPVVSSSPSPS CQSAAAQVQNLTLRSQKLGVLSSSQ NGSPKSAGQTQSLTICHNKTTVTSS KISQRDPSPESKKGSGPGLSRSTAV TRTSSIHQLIAPASYSPIQPHSLIKHQ QIPLHSPPPKVSHHQLLLQQQQQI QPITLQSPSQDPPPSQHCIPLPNHGL SPAPSNAQPQHCSVPQSHPPPLTVS PNQAQSAQQSVVSPPPHSPSQSP TIIHPQALIQPHPLVSSALQTGPNLQ QAAADQVQSTAQLNLPShLPLPASP VVHIGPVQQSALVSPGQQMVSPTSH QQYSALQSSPIPIATPPQMSASPPAQ LPPLPLQSMQSLQVQPEILSQGQVL VQNALVSEELPAAEALVQLPFQTLF PPQTVAVNLQVQPPAPVDPPVVYQV EDVCEEEMPEESDECARMDRTPPP PTLSPAAVTVGRGEDLTSEHPLLEQV ELPAVASVSASVIKSPSDPTHASAPA PPLLIPAASTRSSSTSLASSTPSLENK PPQAIVKPQILTHVIEGFVIEGLEPF PVSRSLLIEQPVKKRPLLDNQVVNS VCVQPELQNNTKHADNSSDTEIED MMAEETLEEMDSELLKCEFCGKMG YPNEFLRSKRFTMSCAKRYNVSCS KKFALSRWNRKPDNQSLGHRGRRP SGPEGAAREHILRQLPITYPSAEEDV ASHEDPVPSAMTTRLRRQSERERER ELRDVRIKMPENSDLLPVAQTEPSI WTVDDVWAFIHS LPGCQDVADEFR AQEIDGQALLLLKEDHILMSAMNMK LGPALKICARINSLKDS

0