

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q92576	PHF3_HUMAN	Homo sapiens	PHD finger protein 3	18.790935	S488;S527;S868;S1722;S1724;S1758;T1770;S1773;S1786	S97;S125;S283;S299;S680;S1014;S1133;S1148;S1178;S1614;S1642;S1898;S1925	35254053;35132862;34019948;31492838;23301498;35289036;27114449;38665916;29351928;28411811;30379171	MDIVDTFNHLIPTEHLDDALFLGSN LENEVCEDFSASQNVLEDSLKNML SDKDPMLGASNQFCLPVLDSNDP NFQMPNSTVVGGLDDIMDEGVVKES GNDTIDEEELILPNRNLRDKVEENS VRSPRKSPLMAQEQVRSRLRQSTIA KRSNAAPLSNTKKASGKTVSTAKAG VKQPERSQVKEEVCMSLKPEYHKE NRRCSRNSGQIEVVPEVSVSSSHSS VSSCLEMKDEDGLDSKHKCNPNPGE IDVPSHELNCSSLSETCVTIGEKKNE ALMECKAKPVGSPFLFKFSDKEEHEQ NDSISGKTGETVVEEMIATRKEQD SKETVKLSHEDDHILEDAGSSDISSD AACTNPNTENSLVGLPSCVDEVTE CNLELKDTMGIADKTENTLERNKIE PLGYCEDAESNRQLESTEFNKSLE VVDSTSTFGPESNILENAICDVPDQNS KQLNAIESTKIESHETANLQDDRNS QSSSVSYLESKSVKSKHTKPVHISKQ NMTTDAPKKIVAAYEVVHVKTKVN VKSVKRNTDVPESQQNFHRPVKVRK KQIDKEPKIQSCNSGVKSVKNOAHS VLKKTLDQDTLVQIFKPLTHSLSDKS HAHPGCLKEPHHPAQTGHVSHSSQ KQCHKPQQAPAMKTNVSHVKEELE HPGVEHFKEEDKLLKPKPEKNLQPR QRRSSKSFSLDEPPLFIPDNIAIRRE GSDHSSSFESKYMWTPSKQCGFCK KPHGNRFMVGCGRCDDWFHGDCV GLSLSQAQQMGEEDKEYVCVCCA EEDKTEILDPTLENQATVEFHSG DKTMECEKLGSKHTTNDRTKYIDD TVKHVKILKRESGEGRNSSDCRDN EIKKWQLAPLRKMGQPVLPRRSSEE KSEKIPKESTTVTCTGEKASKPGTHE KQEMKVKVEKGVNLVHPAASASK PSADQIRQSVRHSLKDILMKRLTDS NLKVPEEKAQKATKIEKELFSFFRD TDAKYKNKYRSLMFNLKDPKNNILF KKVLKGEVTPDHLIRMSPEELASKE LAAWRRRENRTIEMIEKEQREVER RPITKITHKGEIEIESDAPMKEQEA MEIQEPAANKSLEKPEGSEKQKEEV DSMSKDTTSQHRQHLFDLNCIKICIG RMAPPVDDLSPKKVKKVGVGVARHKS DNEAESIADALSSTSNILASEFFEEE KQESPKSTFSPAPRPEMPGTVEVES TFLARLNFIWKGFINMPSVAKFVTK AYPVSGSPEYLTEDLPDSIQVGGRISP QTVWDYVEKIKASGTKEICVVRFTPV TEEDQISYTLFAYFSSRKRYGVAAN NMKQVKDMLYPLGATDKIPHPLVP FDGPGLELHRPNLLGLIIRQKLRQ HSACASTSHIAETPESAPPALPPDK KSKIEVSTEEAPEEENDFNFSFTTVL HKQRNKPQQLQEDLPTAVEPLME VTKQEPKPLRFLPGVLIGWENQPTT LELANKPLPVDLILQSLGTTGQVYD QAQSVMEQNTVKEIPFLNEQTNISKI EKTDNVEVTDGENKEIKVKVDNISE

STDKSAEIETSVVGSSSISAGSLTSL
LRGKPPDVSTEAFLTNLSIQSKQEET
VESKEKTLKRQLQEDQENNLQDNQ
TSNSSPCRSNVGKGNIDGNVSCSEN
LVANTARSPQFINLKRDPQAAGRS
QPVTSESKDGDSCRNGEKHMLPG
LSHNKEHLTEQINVEEKLSAEKNS
CVQQSDNLKVAQNSPSVENIQTSA
EQAKPLQEDILMQNIETVHPFRRGS
AVATSHFEVGNTPSEFPSKSITFTS
RSTSPRTSTNFSMPRPQPNLQHLLK
SSPPGFPPGPPNFPPQSMFGFPPH
LPPPLLPPPGFGFAQNPMVPWPPVV
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FYQVKDIRRPERRHSDPWGRQDQQ
QLDRPFNRGKGDRQRFYSDSHHLK
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DRDRKSREEGHKDKERARLSHGDR
GTDGKASRDSRNVDKPKPKSEDY
EKDKEREKSKHREGEKDRDRYHKD
RDHTDRTKSKR