

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q8TD19	NEK9_HUMAN	Homo sapiens	Serine/threonine-protein kinase Nek9	16.652558	T877	S2;S13;S16;S20;S29;T52;S76;T210;T254;S331;T333;S741;S801;S832;S868;S869;T886;S944;S978	34229054;28657654;23301498	MSVLGEYERHCDSINSDFGSESGGC GDSSPGPSASQGPRAGGGAAEQEEL HYIPRVLGRGAFGEATLYRRTEDDS LVVWKEVDLTRLSEKERRDALNEIVI LALLQHDNIIAYYNHFMNNTLLIEL EYCNGGNLYDKILRQDKLFEEMV VWYLFQIVSAVSCIHKAGILHRDIKT LNIFLTKANLIKLG DYGLAKKLNSEY SMAETLVGTPYYMSPELCQGVKYNF KSDIWAVGCVIFELLTKRTFDATNP LNLCKVIVQGIRAMEVDSSQYSLELI QMVHSCLDQDPEQRPTADELLDRP LLRKRRREMEEKVTLNAPT KRPRS STVTEAPIAVVTSRTSEVYVWGGGKS TPQKLDVIKSGCSARQVCAGNTHFA VVTVEKELYTWVNMQGGTKLHGQL GHGDKASYRQPKHVEKLGKPAIRQV SCGDDFTVCVTDEGQLYAFGSDYYG CMGVDKVAGPEVLEPMQLNFFLSN PVEQVSCGDNHVVVLRNKEVYSW GCGEYGRGLDSEEDYYTPQKVDVP KALIIVAVQCGCDGTFLLTQSGKVL CGLNEFNKLG L NQCMSGIINHEAY HEVPYTTSTFLAKQLSFYKIRTIAPGK THTA AIDERGRLLTFGCNKCGQLGV GNYKKRLGINLLGGPLGGKQVIRVS CGDEFTIAATDDNHIFAWGNGGNG RLAMTPTERPHGSDICTSWPRPIFGS LHHVPDLSCRGWHTILIVEKVLNSK TIRSNSSGLSIGTVFQSSSPGGGGG GGGEEEDSQESETPDPSGGFRGT MEADRGMEGLISPTTEAMGNSNGAS SSCPGWL RKELENAEFIPMPDSPSP LSAAFSESEKDTLPYEELQGLKVASE APLEHKPQVEASSPRLNPAVTCAGK GTPLTPPACACSSLQVEVERLQGLVL KCLAEQQKLQENLQIFTQLQKLNK KLEGGQQVGMHSGTQTAK EEME MDPKPDLSDSWCLLGTDSRPSL