

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
O60391	NMD3B_HUMAN	Homo sapiens	Glutamate receptor ionotropic, NMDA 3B	13.589521	NaN	NaN	37217939	MEFVRALWGLALALGPGSAGGHP QPCGVLARLGGSVRLGALLPRAPLA RARARAALARAAALAPRLPHNLSLEL VVAAPPARDPASLTRGLCQALVPPGV AALLAFPEARPELLQLHFLAAATETP VLSLLRREARAPLGAPNPFHLQLHW ASPLETLLDVLVAVLQAHAWEDVGL ALCRTQDPGGLVALWTSRAGRPPQL VLDLSRRDTGDAGLRARLAPMAAPV GGEAPVPAAVLLGCDIARARRVLEAV PPGPHWLLGTPLPPKALPTAGLPPGL LALGEVARPPLEAAIHDIVQLVARAL GSAAQVQPKRALLPAPVNCGLDQPA GPESPGRFLARFLANTSFQGRTPGV WVTGSSQVHMSRHFVWVSLRRDPR GAPAWATVGSWRDGLDLEPGGAS ARPPPQGAQVWPCLRVTLLLEHPF VFARDPDEDGQCPAGQLCLDPGTN DSATLDALFAALANGSAPRALRKCC YGYCIDLLERLAEDTPDFELYLVGD GKYGALRDGRWTGLVGDLLAGRAH MAVTSFSINSARSQVVDFTSPFFSTS LGIMVRARDTASPIGAFMWPLHWST WLGVFAALHLTALFLTVEYWRSPYG LTPRGRNRSTVFSYSSALNLCYAILF RRTVSSKTPKCPTGRLLMNLWAIFC LLVLSSYTANLAAMVGDKTFEELS GIHDPKLHHPAQGFRFGTVWESSAE AYIKKSFPMHAHMRRHSAPTTPRG VAMLTSDPPKLNAFIMDKSLLDYEV SIDADCKLLTVGKPFIEGYGIGLPQ NSPLTSNLSEFISRYKSSGFIDLLHD KWYKMVPCGKRFAVTETLQMSIYH FAGLFVLLCLGLGSALLSSLGEHAFF RLALPRIRKGSRLQYWLHTSQKIHRA LNTEPPEGSKEETAEAEPGPEVEQ QQQQDQPTAPEGWKRARRAVDKE RRVRFLLEPAVVVAPEADAEAEAAPR EGPVWLCYGRPPAARPTGAPQPGE LQELERRIEVARELRQALVRRGQLL AQLGDSARHRPRLLQARAAPAEAP PHSGRPGSQE