

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence	intracellular	extracellular	cytosol	nucleus	mitochondrion	endoplasmic reticulum	golgi apparatus	plasma membrane	es
P35438	NMDZ1_MOUSE	Mus musculus	Glutamate receptor ionotropic, NMDA 1	37.975063	S278;S282;T302;T437	S889;S890;S896;S897	33300544;40885482	MSTMHLLTFALLFSCSFARAACDPK IVNIGAVLSTRKHEQMREAVNQAN KRHGSWKIQLNATSVTHKPNAIQMA LSVCEDLISSQVYAILVSHPTPNNDH FTPTPVSYTAGFYRIPVLGLTTRMSIY SDKSIHLSFLRTVPPYSHQSSVWFE MMRVYNWNHILLVSDDHGEGRAAQ KRLETLLEERESKAEEKVLQFDPGTK NVTALLMEARDLEARVILSASEDDA ATVYRAAAMLNMTGSGYVWLVGER EISGNALRYAPDGIIGLQLINGKNES AHISDAVGVVAQAVHELLEKENITDP PRGCVGNTNIWKTGPLFKRVLMSKK YADGVTGRVEFNEDGDRKFANYSIM NLQNRKLVQVGIYNGTHVIPNDRKII WPGGETEKPRGYQMSTRLKIVTIHQ EPFVYVKPTMSDGTCKEFTVNGDP VKKVICTGPNDTSPGSPRHVTPQCC YGFCDLLIKLARTMNFYEVHLVA DGKFGTQERVNNSNKKEWNGMMG ELLSGQADMIVAPLTINNERAQYIEF SKPFKYQGLTILVKKIIPRSTLDSFM QPFQSTLWLLVGLSVHVVAVMLYLL DRFSPFGRFKVNSEEEEEALTLSS AMWFSWGVLLNSGIGEGAPRSFSA RILGMVWAGFAMIIVASYTANLAAFL VLDRPEERITGINDPRLRNPDKFIY ATVKQSSVDIYFRRQVELSTMRYHM EKHNYESAAEAIQAVRDNKLHAFIW DSAVLEFEASQKCDLVTTGELFFRS GFGIGMRKDSPWKQNVSLSILKSHE NGFMEDLDKTWVRYQECDSRSNAP ATLTFENMAGVFMLVAGGIVAGFLI FIEIAYKRHKDARRKQMLAFAAVN VWRKNLQDRKSGRAEPDPKKTATF RAITSTLASSFKRRRSSKDTSTGGGR GALQNQKDTVLPRAIEREEGQLQL CSRHRES	False	True	2.629	2.846	2.737	5.0	2.01	5.0	3.