

UniprotKB ID	Entry name	organism	full name	oglcnacscore	oglcnac sites	phosphorylation sites	PMIDS	sequence
Q01097	NMDE2_MOUSE	Mus musculus	Glutamate receptor ionotropic, NMDA 2B	18.754664	NaN	S882;S886;S917;S920;T962;T1039;S1058;S1061;S1064;T1109;T1133;S1143;T1155;S1255;S1259;S1303;T1472	22645316;33300544	MKPSAECCSPKFWLVLAVLAVSGSK ARSQKSAPSIGIAVILVGTSDVEAIKD AHEKDDFHHLSSVPRVELVAMNET DPKSIITRICDLMSTRDKIQGVVLA TDQEAIAQILDFISAQTLTPILGIHGG SSMIMADKDESSMFFQFGPSIEQQA SVMLNIMEEYDWYIFSIVTTYFPGYQ DFVNKIRSTIENSFVGVWELEEVLLLD MSLDDGDSKIQNQLKQLQSPIILLYC TKEEATYIFEVANSVGLTGYGYTWIV PSLVAGDTDTPSEFPTGLISVSYDE WDYGLPARVRDGAIIITTAASDMLSE HSFIPEPKSSCYNTHEKRIYQSNML NRYLINVTFEGRNLSFSEDGYQMHP KLVILLNKERKWERVGVKWKDKSLQ MKYYVWPRMCPETEEQEDDHLISIVT LEEAPFVIVESVDPLSGTCMRNTVPC QKRIISENKTDEEPGYIKKCKGFCI DILKKISKSVKFTYDLYLVTNGKHGK KINGTWNGMIGEVVMKRAYMAVGS LTINEERSEVVDFSVPIETGISVMV SRNGTVSPSAFLEPFSADVWVMM FVMLLIVSAVAVFVFEYFSPVGYNRC LADGREPGGPSFTIGKAIWLLWGLV FNNSVPVQNPKGTTSKIMVSVWAFF AVIFLASYTANLAAFMIQEEYVDQVS GLSDKKFQRPNDFSPPFRFGTVPNG STERNIRNNYAEMHAYMGKFNQRG VDDALLSLKTGKLDIFIYDAAVLNY MAGRDEGCKLVTIGSGKVFASSTGYG IAIQKDSGWKRQVDLAILQLFGDGE MEELEALWLTGICHNEKNEVMSSQ LDIDNMAGVFYMLGAAMALSLITFI CEHLFYWQFRHCFMGVCSGKPGM VFSISRGIYSCIHGVAIEERQSVMNS PTATMNNTHSNILRLLRTAKNMAN LSGVNGSPQSALDFIRRESSVYDISE HRRSFTHSDCKSYNNPPCEENLFS YISEVERTFGNLQLKDSNVYQDHYH HHHRPHSIGSTSSIDGLYDCDNPPF TTQPRSISKKPLDIGLPSSKHSQSD LYGKFSFKSDRYSGHDDLIRSDVSDI STHTVTYGNIEGNAAKRRKQYKDS LKKRPASAKSRREFDEIELAYRRRPP RSPDHKRYFRDKEGLRDFYLDQFRT KENSHPHEHVDLTDIYKERSDDFKR DSVSGGGPCTNRSHLKHGTGDKHG VVGVPAPWEKNLTNVDWEDRSGG NFCRSCPSKLHNYSSTVAGQNSGRQ ACIRCEACKAGNLYDISEDNSLQEL

