ABSTRACT OF THESIS

FURTHERING PHARMACOLOGICAL AND PHYSIOLOGICAL ASSESSMENT OF THE GLUTAMATERGIC RECEPTORS AT THE DROSOPHILA NEUROMUSCULAR JUNCTION

*Drosophila* larval neuromuscular junctions (NMJs) serve as a model for synaptic physiology. The molecular sequence of the postsynaptic glutamate receptors has been described; however, the pharmacological profile has not been fully elucidated. Despite the postsynaptic molecular sequence used to classify the receptors as a kainate subtype, they do not respond pharmacologically as such. Kainate does not depolarize the muscle, but dampens evoked EPSP amplitudes. Quantal responses show a decreased amplitude and area under the voltage curve indicative of reduced postsynaptic receptor sensitivity to glutamate transmission. ATPA, a kainate receptor agonist, did not mimic kainate’s action. The metabotropic glutamate receptor agonist t-ACPD had no effect. Domoic acid, a quisqualate receptor antagonist, blocks the postsynaptic receptors without depolarizing the muscle, which supports the presence of quisqualate subtype receptors. The results suggest a direct postsynaptic action of kainate due to partial antagonist action on the quisqualate receptors. There does not appear to be presynaptic auto-regulation via a kainate receptor subtype or a metabotropic
auto-receptor. A complete pharmacological profiling of the known receptor subtypes at this NMJ has not yet occurred; however, this study aids in furthering the ongoing investigations to provide a clearer picture of pharmokinetic profile and specificity of the receptor subtypes.

KEY WORDS: Synaptic facilitation, NCX, calcium, *Drosophila* larval heart

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THESIS

A thesis submitted in partial fulfillment of the requirements for the degree of Master in the College of Arts and Sciences at the University of Kentucky
By
Junyoung Lee

Lexington, Kentucky

Director: Dr. Robin Lewis Cooper, Associate Professor

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I came to the University of Kentucky from South Korea two years ago. I am in active duty as a captain in the Korean Army. I was sent here by the Korean Army to learn Biology and experimental methods to address how the nervous system functions. When I began pursuing this masters program, I had little knowledge of the biological field. When I was a student at the Korean Military Academy I focused on Chemistry and Military Science. The time I spent in the biology department was stressful and compounded with frustration, as the language was new to me and the courses moved along very rapidly. So I feel extremely happy and lucky to have made it through the program. It was not a miracle that I survived in this foreign country, as I had very helpful people around me. It is a great pleasure to have a chance to be aquatinted with them.

The first person I appreciate is Dr. Robin L Cooper who is my mentor. I emphatically state that he is one of the best professors I have experienced. He has helped me a lot to feel comfortable both inside and outside the lab. His patience with me was nothing short of miraculous and he even never lost smile on his face. 설레임과 두려움을 안고 미국으로 온 저에게 항상 웃는 얼굴로 학업 및 생활하는데 불편함이 없도록 직접 보살펴 주시고, 많이 바쁜 와중에도 저의 영동한 질문도 그냥 넘기는 것 없이 자상하게 대해주시는 모습을 보면서 남몰래 눈물을 흘린 적도 있었습니다. In fact, it is impossible for me to relay of all gratitude and respect in this letter. 이곳 렉싱턴에서의 교수님과 함께했던 시간은 소중한 추억으로 간직되어 평생 잊지 못할 것입니다. 쿠퍼 교수님 그리고 사모님 감사합니다.

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마지막으로, 제 인생에서 가장 중요하고 많은 부분을 차지하는 부모님께 감사드립니다. 자식들 뒷바라지 하느라 많은 것을 희생하신 우리 부모님. 아버지, 어머니의 존재 자체만으로도 저에게는 그 무엇보다 큰 힘이 되었습니다. 이제는 제가 아버지, 어머니의 든든한 기둥이 되어 얼굴에 항상 미소가 끊이지 않도록 노력하겠습니다. 사랑해요~♡.

p.s. 본문에 구체적으로 적지는 않았지만, 우리 사랑하는 누님, 이윤환 중대장님, 군 선배님 및 후배, 백두산 부대와 승리부대에서의 동생들, 그리고 켄터키 대학에서 물심양면으로 많은 도움을 주신 형님, 누님들께도 감사드립니다.
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Chapter One

Background of the *Drosophila* neuromuscular junction and research goals

The nervous system in animals is used to convey information of the surrounding external and internal environment, as well as to react to it. In the majority of cases, the electrical signals neurons use are relayed to target cells by chemical communication. When this occurs for close targets, the site of communication is referred to as a synapse. The chemical released from the presynaptic neuron is referred to as transmitter. This transmitter enters into the synaptic cleft and then binds to receptors on a target cell. The target cell can be a neuron, a muscle or an endocrine cell or even a combination of cell types. The post-synaptic receptors determine if the target cell will be excited or inhibited. These determine actions by the receptor itself, being a ligand-gated ion channel or by initiation of a cellular cascade to influence ion channels (Berridge, 1998; Miyazaki, 1995). Receptors that mediate secondary cascades are referred to as metabotropic receptors, whereas receptors that act as a receptor mediated ion channel are termed ionotropic receptors.

The neurotransmitter molecules are stored in vesicles within the presynaptic nerve terminal. These vesicles fuse to the presynaptic membrane with evoked stimulation because of the rise in $[\text{Ca}^{2+}]_{i}$ (Sudhof, 2004). The $[\text{Ca}^{2+}]_{i}$ is generally very low compared to that outside the neuron ($[\text{Ca}^{2+}]_{o}$ is 10 mM). The entry of $\text{Ca}^{2+}$ into the nerve terminal causes the mobilization of both the reserve and the readily releasable pools of vesicles (Rosenmund and Stevens, 1996; Kuromi and Kidokoro, 1998).

The whole process of vesicle fusion must be finely regulated as not to over or under excite the target cells or else the information will not be correctly conveyed. Thus, there are various levels of excitation and inhibition to regulate nerve terminals, such as ion pumps, ion exchanges and auto-receptors for negative feedback. Because of the difficulty in accessibility of postsynaptic dendrites and the wide complexity of the central nervous system in general, most of our understanding in synaptic properties at a vesicle level (i.e., quantal level) has been gained by studying accessible and relatively simple preparations, such as neuromuscular junctions (NMJ) in invertebrates and amphibians (Bennett and Pettigrew, 1975; Nudell and Grinnell, 1983; Wilkinson and Lunin, 1994; Wilkinson et al., 1992). The frog, locust and crayfish neuromuscular junctions were initially used because of the relatively ease for accessibility and viability in relatively minimal physiological salines (Fatt and Katz, 1953a, b; Dudel, 1965, 1981; Dudel et al., 1983).
More recently the *Drosophila melanogaster* has become a model for synaptic transmission at the larval NMJ. This is because a known genome that can be manipulated for altered protein expression to examine mechanistic cellular functions of synapses (Atwood et al., 1993; Betz et al. 1993; Jan and Jan 1976; Kurdyak et al., 1994; Li et al., 2002; Li and Cooper, 2001; Pawlu et al., 2004; Ruffner et al., 1999; Sigrist et al., 2002, 2003; Stewart et al., 1994, 1996). A recent review in Ca²⁺ regulation which influences synaptic transmission in the preparation has appeared (Desai-Shah and Cooper, 2009).

Since the larval *Drosophila* NMJ is becoming a synaptic model of choice recently for many reasons, it is important to understand the pharmacological and physiological profiles of this synapse. The *Drosophila* NMJ consists of excitatory motor neurons and muscle. Recent studies described which postsynaptic receptor subtypes are expressed in muscle. There are five different subunits, which fall into kainate/AMPA type glutamate receptors based on molecular sequence homology (Betz et al., 1993; Völker et al., 2000; Marrus et al., 2004; Qin et al., 2005; Guerrero et al., 2005). However, in an experiment investigating a pharmacological sensitivity on the postsynaptic receptors for intact 3rd instar *Drosophila* NMJ, kainate did not depolarize the muscle but reduced the EPSP amplitude. Also, AMPA did not alter the EPSP amplitude nor alter the resting membrane potential (Bhatt and Cooper, 2005). Thus, the postsynaptic receptors at the *Drosophila* NMJ should not be classified based on a pharmacological profiling nomenclature as a kainate/AMPA subtype of glutamate receptors. Based on my investigations and the previous study (Bhatt and Cooper, 2005), kainate appears to be blocking the glutamate activities at the larval *Drosophila* NMJ.

To date, the pharmacological assessment of this NMJ has not been tackled carefully. No one has even addressed if the motor nerve terminals may possess glutamate auto-receptors as noted to occur in other arthropod species (Schramm and Dudel, 1997). Thus, I choose to address these issues in my studies.

In this research project, I first support the idea that the postsynaptic receptors at the *Drosophila* NMJ are not of a kainate subtype of glutamate receptors and that application of kainate reduces the evoked EPSP amplitude. I approached these studies by use of pharmacological and physiological investigations. Specifically I addressed the spontaneous vesicular postsynaptic responses at the 3rd instar NMJ using domoic acid and glutamate. Domoic acid is a potent agonist of the kainate subtype of glutamate receptors in vertebrates. I also addressed how exogenous application of kainate could be causing the reduction of the EPSP amplitudes.

The hypotheses I am testing is based on preliminary findings at this NMJ and combined reports on other glutamatergic synapses in various animal
models. The hypotheses addresses why kainate produces a reduction in the EPSP amplitude. My hypotheses are as following;

1) Kainate has a direct postsynaptic action as an antagonist.
2) Kainate has a direct presynaptic action reducing glutamate release, possibly through activating glutamatergic autoreceptors on the nerve terminals.

Addressing these aims is very important in understanding processes in synaptic regulation at the neuromuscular junction of this model preparation. Several disorders are known for altered synaptic function in the vertebrate brain. In order to understand these disorders better, one can use invertebrate models such as Drosophila, which can serve as good electrophysiological and genetic models to study synaptic regulation on a fundamental level.

In order to elucidate my hypotheses about kainate’s action at the Drosophila NMJ, I conducted several experiments as described in detail within Chapter Two. In short, I examined the action of kainate on spontaneous fusion of vesicles with exposure to kainate. This is diagrammatically shown in Figure 1-1 with kainate either having an action on the presynaptic terminal or having an action on the postsynaptic receptors as to alter receptivity to glutamate. Since $[\text{Ca}^{2+}]_i$ is tightly regulated within the nerve terminal for rapid transmission, if kainate had a presynaptic action via alteration in the $[\text{Ca}^{2+}]_i$, I would predict that short-term facilitation or depression, known to occur at this synaptic preparation, would be greatly affected. Thus, I used a four-pulse stimulation train and measured the facilitation index before and during application of kainate at various concentrations of kainate as well as $[\text{Ca}^{2+}]_o$. Since transmission is so sensitive to $[\text{Ca}^{2+}]_o$, altering the concentration with various concentration of kainate exposure might unmask any subtle effects of kainate. This general model to address this experimental paradigm is shown in Figure 1-2. In addition to the action of kainate other pharmacological agents were examined to help in describing the pharmacological profiles of the putative presynaptic and known postsynaptic glutamatergic receptor subtypes. The working models of action for ATPA (Figure 1-3) and t-ACPD (Figure 1-4) are shown for action on evoked transmission.

In addition, the reports which characterized the molecular sequence of the glutamate receptors at the Drosophila NMJ to the human’s kainate receptor subunits, did not provide the statistical similarities between them. The paper published in 2005 by Qin et al., designated some amino acid sequences as similar even though they appear to be far from being similar. Besides, in this earlier study only five amino acid sequences of the postsynaptic receptor subunits at the Drosophila NMJ were examined although there are around 30 different glutamate receptor genes in Drosophila genome (Littleton and Ganetzky,
2000). Thus, I show statistical similarities of the aligned sequences (Figure 1-5a,b) and provide a phylogeny tree with the known 36 predicted amino acid sequences of the glutamate receptors in Drosophila genome (Table 1-1).
Figure 1-1. Schematic model of predicted kainate action at the *Drosophila* neuromuscular junction, which is the site of communication from the axon terminal of motorneuron to the muscle. Kainate is added into the synaptic cleft without neural stimulation to measure any possible direct effect on either the presynaptic nerve terminal or on postsynaptic sensitivity. Spontaneous responses are involved in one vesicle fusion events. If kainate has an effect presynaptically then one would expect an alteration in the frequency of spontaneous response. If there was an affect on the postsynaptic receptors then one would expect a change in the amplitudes or shape of the spontaneous events. The frequency and amplitude of spontaneous responses before and during exposure to kainate are compared.
Figure 1-2. Schematic model of facilitation effect on presynaptic nerve terminal. Ca$^{2+}$ plays a critical role in vesicle fusion in the presynaptic nerve terminal. [Ca$^{2+}$] in the presynaptic nerve terminal can be changed at each pulse of induced facilitation. Four-pulses with 20Hz is induced in the presynaptic nerve terminal, which may alter the dynamics affecting several factors involved in vesicle fusion inside the cell. This presynaptic activity at each pulse will have a slightly different condition. If there is a kainate effect on Ca$^{2+}$ dynamics it would likely alter the vesicle fusion events during facilitation. Facilitation Index (FI) ($4^{th}$ pulse amplitude / $1^{st}$ pulse amplitude -1) is measured before and during kainate exposure to determine if there is a presynaptic action of kainate. (RP: reserve pool, RRP: readily releasable pool, ER: intracellular Ca$^{2+}$ storage, ES: endosome, VGCC: voltage-gated Ca$^{2+}$ channel, KAR: kainate receptor)
Figure 1-3. Schematic model of predicted ATPA action at *Drosophila* NMJ. (RS)-2-Amino-3-(3-hydroxy-5-tert-butylisoxazol-4-yl)propanoic acid (ATPA) is a kainate receptor agonist. If ATPA has an action either presynaptically or postsynaptically it would alter the frequency of amplitude of spontaneous events and the size of evoked responses. Excitatory postsynaptic potential (EPSP) amplitudes and membrane potential in the muscle are measured with the bath application of ATPA to compare if they are affected.
trans-(±)-1-Amino-1,3-cyclopentanedicarboxylic acid monohydrate (t-ACPD) is a well-known metabotropic glutamate receptor agonist. If t-ACPD has an action either presynaptically or postsynaptically it would alter the amplitude of evoked response. Excitatory postsynaptic potential (EPSP) amplitudes and membrane potential in the evoked response are measured with the bath application of t-ACPD to see if there is a t-ACPD action through the metabotropic glutamate receptors. In the figure, narrow arrow is described to show the action of t-ACPD on the postsynaptic receptor because of a small possibility of a postsynaptic action on receptors at the Drosophila NMJ. The receptors are known as ionotropic glutamate receptors.
Figure 1-5a. Sequence alignment with predicted amino acid sequences of the postsynaptic receptor subunits at Drosophila NMJ and human’s kainate receptor subunits. Compared subunits are shown on the left of the first row in aligned sequences. Identical and similar amino acids matched are colored by red and putative extracellular domains are shaded by yellow, respectively. ClustalW2 is used for sequence analysis.
(Figure 1-5a cont.)

EIYKPTVNEFMVWTPDN---GKKRLLN---LELESAGTTQDFSEQVRKVTVTHYEFPYFMKEDE---NFREGRKmeyeAVDLIKSELMEFDFYEMIVNGN
EYNNPIIDRVTWIKNKEFQLVDFKLENSTQLKQRLQNKFEDSFQKPIRTVATRVGKPYSWREEPEGVYHEGERNSFEGYAVDLIYNLAECKFDNFEPRVDRN
EYDPLDIGIAIDWPGR---QITQLNVQAVKQKMYRATQKSNPQSYNEGMYLTLIDTAMHEARMSGFYFVVFAPQ
IELOQSGWTVSEUG------DFEFQR--PPQAVNFDIDDGSLVNKTFLIVSIVATPKYASLVESID--TLGNNQFFGYGVDMLEKADLKLGVNFTFVNGND
ILOMQGIVKVTWDEDR------QYRFDQALPDDAQFQSDNV--SLANKFIIILLSVNPQYAQLVETAYQY--QLGNSQEGYGVDMLEKADLKLGNFTFVNGND
ISLKEEGTEKNSNSNS---GLNMND---SNKDSSITDSLALRLTIVT-TLEETEVYMKYSDK--PLYQNDGFEGYCLDLKELSINLGFYIDVKLVDPG

TM1

-+-GKYNPETKVQWDGIIKRLIDHHAQIGVCDLTITQAR-RSVVDTFVPFQMLGISEILHYKSPPEPKNQFAFLFVAVYWIYIMFAQLMTAFVFIAKLSYRELPFEN
KYGYDANTDEWDGIIKRLIDNNASIGICDLTITQAR-RSVVDTFVPFQMLGISILSYKEPPPKAIDYAFNLQNKVPYLVMIAAMMIFAALIFTRGDIDQYEDQPVQ
YQGKLDKETKQWNGIIIGEIIINNDAHMGICDLTITQAR-KTAVFDTFVPFQMLGVSILAKSSHVEKTLDAYLAFPGGSWILISYFVMFPLKTIVARISKMDWENPH
-YGSPNKTNTSUGMLKEIEVGRAIAITDLTSEREEVIDSPFINNLGIAILYVHPQKAPLFSFAIDDFSSVWLYLGIAELGVLSCFIIGGSLPIEWDQPYY
-YGSPNKTNNSTGMLREIMTGRADIATDLTSEREOALDFIPMNLIAlIYLKPQATPELFPTMDFFSESYWFLGFELGVSLSFFILGRLSPEWDPNY
-KYGAQNOKGEWNGMKELEDHRALAVPLPITVYVEKIVDPSKFMTMLGISEILYRKNPNGTNGFVFSFNLPSQIMYVLLACELGVSLVFLVIARTFYEWYNF

TM2

PAIQDPDELENIWVNNSTWLMGVSIMQCGDILPRGMRILRITGWMWFFALMLSLSTYATLANFLLSNWWQSSIKSLQDIEQD-KVPEGMRGGSGLISFFSESDND
ENVNREMERQGINHSLNAWLVLGSLNQCGDLPRLPRLULATAFWTFALLISQTYIAKLAAPFSSKIAQIGGSLHLVDQN-KVQFTPITRSASVSYFSSND
PCNRDFPEVLENQWRIHNTGWLTVASIMTAGCIDLPRSPQVRMEATWIIAFAIVANNSYATLANFLLSNWWQSSIKSLQDIEQD-KVPEGMRGGSGLISFFSESDND
PCIEEPHELQQFTINNLSWFTTAGQGESEIAKPSALTITISAIWETFLLIMWSSYTLANEFITNETPSIINSVKLDADNNKVDDQYAGAKRGTSTRNFFSSE
PCIEEPHELQQFTLGNISWFTTAGQGESEIAKPSALTITISAIWETFLLIMWSSYTLANEFITNETPSIINSVKLDADNNKVDDQYAGAKRGTSTRNFFSSE
PCNPDSDKVENFTLNSWFWVFGALMQCGDILPRGMRILRYVTFWFFFTLLIIIVTSYATLANFLLSNWWQSSIKSLQDIEQD-KVPEGMRGGSGLISFFSESDND

TM3

TDYQRAWNQMKDFNPSAFSTSNKEGVARVRKEGGAYFLMETTSLTYNIERNCDLTQIKEQIHEKHYGLAVFLSDYRTNLVSILQSEREGLQKMKWKNKNHNN
TDNRMWANMKLSFPFDPATKNNевGVRKSLKSYTFLMMETTSLQYVQRCCELTQIGESFGEKHYALVNAFDRFSNLNSGIRSLSFLKLNKWNFSNES
TVYRRAFLINNNDPSAYTSDKNGEVDRVVRKRNQDRMFMLMETTTLHYHERQNCRLDRSVKGEKHYALVFFGAYRSLNVSLAIKLSELEYDLQKWWKPNPNA
PIYXKMNENNA-HPEMLMNNOQVQDKVKS-TYKAFMESTSIEINFTVRECNLTKGDPEAKDEMGIAYMKWNYPYDFKNKALNELQCEGVIAKLNKWNNEVGA
ERYKMNKFSK-NEQYLTEDNMEGNVRKVTK-THAFMESTSIEYNTKRECNLKIKDAGDEKDGYAMKWDPHRGKFVNNALELQCEGVLEKMKWNWNEVGT
STYEMKWAFMSRQQTALVRNSEDQGIRVLT---TDYALLIMESIYEYVTQRNCNLTIQGLDSSKYGTVGFSGPYRDKITIAIIQLOEQLKHMKEKWWGRNG-
(Figure 1-5a cont.)

TM4

-TCDS-YHEVDGDELSIIERLGVFLVLAGGVILGVIDLGEFLWNVQNVAEEVTPWQAFKAILIFALKFWVRKKPMRIS--SSDKSSSRSS-----GSRRSSK
-TCDSNVPTIDDQFDMDSVGLLFLVVLIVGVVGLVIGVAFELWVQVRSKEKIPPMHALKAEFYFVIRFWLTPKPLHTYR--QSRDSTSTGYSSLEQITSASSAK
-SCFEEDPDAATPDTFEEGIFYTLYAGIALALIGITElizabethFVLYVQVQALEERLTFKDAFKKEIFVLCVWNMKPIVAGTPISSVRTPRSLDKLDRTPKSSR
GVCSAKSDDDPSEILGVDLVLSIYVVLVIGSIISIIISIIILCWYFVYKAKNYEVFPFCDALAEFFIVIFVIFSENERPLKSAQSIYRSRRNNSQSIESLKTDSSENMP
GICATKEDAPDATPLMNNLEGVFFVLLVGCALLYGISWVLFVMKHHRVVLRLDALKFFQFVIDFNNVYRVLKNASAYRSLRQSSMSVASVQESQ-----
---CPEEDNKEASALGVENGIFIVLAAAGLVLVSFVAINGEFYKSRKNNDIEQCLSNAEMELGISLKNQKKIK--KKSRTKGGSSFTSILTCQRTQRTKETVA

EKSRSKTIVS-------------------------
KKKKTTRIEK------------------------
RVIVGRSEEMREAMQGSQSSGSSNAGREKEARV
VED-----------------------------
--------------------------------
(Figure 1-5a cont.)

GluR-IIA

GluR-IIB

GluR-IIC

GluR-IID

GluR-IIE

GluR-6

------------------MRLCPVYIAFIIIIFLEGIIALGGDDRNEITVG----AIFYENEKIEIELSFDQAFREVN--NMKFSELRFVTIKRYMTNDSTFLQQ
------------------MHGLQFLVLLAALAFILAFGNEDTLVIG----AIFDDTMKLADAFSAALEEV---AIN-PALKLDAIKRYYVSTIDVLQD
MKKELSNGSSARWFMPGNSREAGMPWQIILLGCMWAFPMCRSRRQQING----AFFYDELELEKFMVNAVINGPESQTMRFYPPKLLKPEDEGVTMQE
------------------MHFCWIS----IIILSLSQVAQYFGGNYAESASSQSIIRG----LITDDATDRIQTFHELVSQV---NELGVPVLGETQEVAYQNSVQAFA
------------------MFNFHVFILWLSFISIHISNVQAGYENFG---YDNQVESLEEQVIG----LIDTDQTEQNNIVFDHAIDVAN----QEVGSTLTSKLEEVENYGDGAQYSYQ
------------------MKIIFPILSNPVFVRRTVKLLLCLLWIG----YSQGTTHVLRGQIFYEVESGPMGAEELAFLRAFTVNTRNLTRLFPNTLYTDQKINLYDSFEASK

ITCELISNQVAIAFPGSSHSSAVSDIVAQIANATGIFHIYEYDLKLEATRFPQEQQLNHQMSNVAPFLSLVSLRAYSIEIKSNGYVEWRYFTTLIYETPEG---LGQDJILMQQA
ISCDLIGSGVAIAFPGSSHSSAVSNIDVLCNMTIGIFQLFDHPQCNSSRNRNHQLTVNAVFPTLMFSAFDILASKTDFKSTFTIAERSSSHLHQLHIWAKQ
HACCLIDIGNQVAIAFPGSSHSSAVSDIVALCSTNGIFHIEPIDSGQAEGQAEKPNHQMTLNYPQAPQAILSKKAYADIVQNF---GWRKFTIVYDADDARARALLQDLQRLR
QLCRQMQGSQVGAVFPGPAARHTASHTLNLNASDIKIFIPYF---HLSWGSNPDGFLHPSFEDANLYDIVQNF---KSRVICFYESAEY---LKLHDLMTRYQ
KLCRMLETGIAGVFPGSSHRTAVHMLSCDAMDIPHIYSS-----YMS---ENAEGFNLHPHDAPLAKALTSLTEFP----WTRFIIYFYESAEY---LNLHNLTTMLQG
KAQDQLSGQVVAIFPGSSHTSSANAVQSCNALGPHIQTRWK----HQSNDKDSFYVSVYLPDFSSLRSAIDLDLVQFQ---KWKTCTTVYYDSTG---LIRQELIKAPS

LNSDYVKLRLNDAYYDRILWLKTEDFTEQRIILDCEPKTLKELLLKSIFDKLQGFPRNFWFLHLHDTHNSGLRDIY---EDFANITVSRKLVDANDPFEFKKRT
LHAKIGIKQEEFERGDDLYRLKNKANREKKEVILDCPSDINVAINASIGYMNMTSPGNNVTLNLDTHLSGQFYS---RDFPTVAVAARITRYTPFVPHVDEIDV
VHNDVYVRFKFEKDDDDFVMWKSISRG---ERRVLDCEFNMLVEILNSTFEFLTGQYNHIFLNLTYTDHLELAAEDETTAVNITARALLWNFDFPPILSYQG
IKGFVKMNYRLNLNLNGYKIFPELVLAGQLRVEQALQLQVILQMNEDTYTQILNLMHLDTFLELEY秆---EANITGRMIFPSDQEVRLDMEEK
KGTSTVITYRDMQLNGYKQVLRLVRKESDNVYIVVGSSETMPEFLNRQAOQVIYRNDKYYKGNLDFHDFSLFEDDEYLS----EANITGLRPSFENKNAVELLMK
RNYNLKLQIFPLAEDTAKDFPLEKMKRIGKEFQHPFIVFEDCSHEMAAGILKQALAMGTYEYYYIFITTLDLFDALDFEPYRS----GNNMTGFRILNTENTQVSSIEK

LT-KVDQILGN-----QTMLPIIYDAVVFAPASSRNAIVIAMQPFHPP-----NRHCQSSSPWMGAFIVNEMKTIISEDDEVEPHFKENMKLDEYQGHRFNLEYK
FDNSVRTFRS----SLQSLQVYDSLVLFNYE-LALEISQRPFGFYRTYFNSSCG-RGFQWQPGRPLVEQMQKHTPKMVPPKTQRLQINADQRGREDNFNLQYN
YVTDQRDNYEVDSDPRTLHDILHDLALQLFQAQWLNASFYDPDMVRPRITCDFAASGGRTWANGRLARLMKATSG-VNVNFTPDSLQFEDQORITFINELBYD
LHQRELGESEP-----VNSGSTMFITEMEALYDVARVIAETTKHLIPYQF-----QLMNQSCREDHVDQPGDSTFNYMRSLIEKETTIGRIYFEGNVRKGFTDVIE
LGQYTDPQDE-----FRNQSCPENVYMTALYDVARVLQLKFLFFPQ-----MFQPSCQRTESQTLQFGDSDSNQKNMRTLRTDLRTGPIYFEGNVRKVHLQV
WSMERILQAPP-----PDSALLGDGFNMTDAALMYDAVHVVSIVVAVQPFOQMT-----VSSLCNRKHPWR-FGTRFMSLIKEIAHEWGLTGRTFINKNTGDLRFDLQVIS

PTVNEPMWVTDPN-----GIIKRLN-----LELESAGTTQDFSEQKVRVYTVPHTHEEYPFMMKEDH---NFRGREKKEYGAYDLIKSLKSEMFDYEFMVNGN---GK
PIIDRTVTHWKEFQLDVEFLKRENSTQALKRLQKNEDFSQKPQRYTAVTVGKYFWSREREPEGQHEGYNERFRAYDLIVALYAEQCKDFNDEFVRDKYGS
PLDQIGIAWDPRG-----QITQQLNDVFAQKAHKMRVATRIPPYFYSNEXARELNLGNQALQGYAVDLIARLVQGMYEWVPVADQQYCK
LQTSGVLKVTGWEED----GKDFEFQRPQPAVNFDMDDGDLSLHNTKLYIVLSVATKPDLYSNLSD----TLISQNQFOQGYVGLIKEAKALKGLFNTDREGGAN-DYCS
LQPSGVIKVTGWEDE-----DRQYPQRLAPTAQFDSVLDN-LASKTFLILLSVPNQAYQVLQTEY-----QEGSNSQEGYVGLIKEAKALKGLFNTDREGGAN-DYCS
LKEEGLEIKGWDP-----ASGLNMTE-----SQKGPANIANTDSLSNRSNLIVTTLQEPHEVLFSKDKS-----PLEYGNDFEYGIDDQLRELSTILGFYEEIRLVEDKYGAG
(Figure 1-5a cont.)

SKTVS-----------------------------
TRRIEK-----------------------------
GRSSEEMREMAQSGSSSGSNNAGRGEKEARV
--------------------------------
--------------------------------
(Figure 1-5a cont.)

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<td>GluR-IIIE</td>
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| GluR-IIA     | ITCELSNNGVAIFFPGSSKAAASIIDLAVQAIAITGIFHIEYDLDKLEATRFRQEQNLHQMINSVIPSLSVLSRAYEIIKSN---YEWTFTTLIYETPEG---LARIQDIQMIQCA |
| GluR-IIB    | ISCDLIGSGVAAIFFPGSSKTAISDIDIVLCTIGFHPQQPQNSRNRENHQQLTVNVALPELFSLAASFDLASKTFDSEQKSTIAERSSH---LIRQHLAILAQWQ |
| GluR-IIC    | HACDLDNVGAIFFPGSSKAAASDIVALVCNSTIGHIFHEDIQDIEGAEKIPNHQMTRNLYPQPAQIALSKAYADIVQNF---GWRKFTIVYDADDRANAAILOQDQLQLE |
| GluR-IID    | QLCLLMQSGVGAIFGFPARAHTASLNLNCDSSKDPFYPQ---HLWGSNPDGFNLHPSEDIANALYDIVQNF---KEKRFICFQSYAESAY---LIIHLTMTRYQK |
| GluR-IIIE   | KLCRMLETGIAVGFGPSRSSHTAVHLMSCIDAMDPHIY---YMS---ENAEGFMNLHPHAPDALKALYSLITF---WTRFIFIYSEAY---LNIHELTLMLQG |
| GluR-6      | KACDQLSLGVAAIFFPGSSHSSANAVSICNALGVPHIQTRWK---HQVSDLNSDFYSVLPDFSSLRISDFLLQFFFF--KWKTIVTVYDSTG---LIRQELVIKASM |

------------------MRLCPVVIYAFIIIIGFLEGIIALGGDDRNEITVG----AIFYENEKIELSFDQAFREVN--NMKFSERLRFVTIKRYMPTNDSFLLQQ |
| GluR-IIA     | LNSDNYVKLRNLADYADDRYILWRKETDTEHRQIIILDCEPKTLKELKLKVSIDFKLQGFPPRFNPWFLTHLDTHNSGLDIYIN---EDFKANITVSLRJKVDAMPFPERK(TM |
| GluR-IIB    | LHKAGIKQEEFIRGDDYRILKINN---AREKTVILPCDSLIVDRVNAISGYNMTGSFPIHLNTNLSSLPLGDFYS---RDPFTVAVAARITYPVPEVHVEIDV |
| GluR-IIC    | VHNDDVRVRKFH-KDDDPFRVMKWSIRG---ERRVVDCEFNMWLLELLNSTFEFGTQGHYHFLINLETYDHELEALADNETFAVNTAALLVNFDPFDPSLYLQH |
| GluR-IID    | IKGFPVKMRDYLNLINNSNLY-effectsILQVAGTHAQYUGEECYNATNQNDLTFELDEEYKS---EANITGIRMFSPDOEVRDLMEK |
| GluR-IIIE   | KGSQYVRILYDQLNLNGNYQKVLRVKSVDNRRVVVSQETMEQPLAAQOAVGQINIEXECYKIANGLDFSDSFLEDEEYKS---EANITGLRFLPSEKNAVREKQMK |
| GluR-6      | RYNLRKIRQPLQDPTDAKFLKLKEMKRGKEFHIFVIFCSCHEAMAGILKZALAMGEEPEHYFITALLLDFALDLVQP---GVMNTGRFILNTENTQVSSIEK |

------------------MRLCPVVIYAFIIIIGFLEGIIALGGDDRNEITVG----AIFYENEKIELSFDQAFREVN--NMKFSERLRFVTIKRYMPTNDSFLLQQ |
| GluR-IIA     | LT-KVQDIQLGN-----QTMLPILYDAVVLFFASSARNVIAAMQPFHFP---NRHCGSSSPWMLGAFIVNEMKTISEDDEEPHFKTENMKLDEYQRHINEFLNIEYK |
| GluR-IIB    | FDNSVDTRFS--------SLGLQYDSILFYN--ALLEISQRPF-FYYP---NSFCG-RGFQWQPGLRLEVELQKIQITPKMVKKPFQTKRQLINADQREDPFNLVYN |
| GluR-IIC    | YVTQRDNVYESSDDPRTLHLDHLCDLALFAQFSSWNASPFYDPDMVRPRTICDFAASAGGRTWANGYLRALMKGTSV-VNTNFRSELQFDFEQR0ITFINUYD |
| GluR-IID    | LHIQEGSEFP------VINSGSTFIMEMALTYDAVRVIAETKHIPYQP---QMLNCSERHDNQVDGSTFRNYMRSLIEKЕКТГГГТРИГ ГРУЕВРКГКФТДГГС |
| GluR-IIIE   | LGYPTDQDE--------FRNGSNCFPVMATTEYAQVTPVLKLFK----MPFQNCQRTESILKELQFSSKNMTKRLMTDTLCTIPYEGFNGVKYLGHLVDEE |
| GluR-6      | WSMERQAPP----PDSGGLDLGNTTDDAIALMYDASHVHSVSSAVQQFPQMT---VSSLQCNRHKPRW-FGTRFMSLIKEAHWEGLTGRITFNKTNGLRTDFDLVIS |

------------------MRLCPVVIYAFIIIIGFLEGIIALGGDDRNEITVG----AIFYENEKIELSFDQAFREVN--NMKFSERLRFVTIKRYMPTNDSFLLQQ |
| GluR-IIA     | PTVNEMPWVTPDN-------GKIKRLLN-------LELESAGTTQDFEQRKVTYVTYHEEPFYMKEDEHE---NFRGKEKYEYAVDLISKSELMEFDEYMFIYNQG---GK |
| GluR-IIB    | PIIDRTVHIIWNEKFQLVDEPKLRSTQALKQLKRLQNKEDFQSPIFYRTAVTRVKGPYFISREPEEGVHLEYNFRGKLPEAVAILYMLAQECFDFNFPFVRDNKYG |
| GluR-IIC    | PLQDGIAWDPRG--------QTITQNLVDVQAQMMYYCTRIPAYPSYNEUTARELNLTGNALYQYGAVDIALIARHGVEYFVVFPAVADQYCK |
| GluR-IID    | LQTSGLKVGTRWEE------GKDFEQR-PPQAQVNFNINDGDLDSNKNKFTILSVATKYASLVESEID---TLLQNCNQFOGGYVGLKDIAKDGFKNTRFDGGN--DYG |
| GluR-IIIE   | LQPSGIKVTGTDWELRQYQRQPLATTAQFDSDVLN---SLANTIFILLVSIFPNKYQAQYTVK---QELENSQGYEVGAVGKDIAKDFNFTVNGN--DYG |
| GluR-6      | LKEEGLEKIGTDPLES-------AQGLNMTE---SQKGPANITDLSNRSRLILITVILEPEYLVFSDKS---PŁYGNDRFEGCYCIDLRELSTILTGEYETIRLVEDKYGA |
(Figure 1-5a cont.)

YPETKQWDGIIKRKLIDHHAQIGVDLTLTIIQMSRVSVDFTVPFMQLGISILHYKSPFEPKQFAPFAPPFVAVWYMIYAQLIYMTAFVFIALRSLYREWLPNPNAIQ
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FNNKTNSTSGLMKEVEGRADIAATLTDIIITSEEEVDFSPFNMGAKLAILVYKQAKPEAPFLSFMDPSSEVWLYLGAYLGVSLCFIIIGLSPIEWDNYPYCIE
YNKSTENSTGLMREMTGRADIAATLTDIIITSEEEVDFSPFNMGAKLAILYKQAPKQATFLSFMDPSSEVWVFLGSFLGVSLSFILGSLPSFDEDNPYCIE
QDDANGQWNGMVRELIDKDIALAVAPALITYVREKVIDFSKPFMLIGISILYKVNGTNPQFVFSGFLNPFLSPFIWMYILLAYLGVSCVLFVIARFSYPYEYNYPHCNP

TYM2

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REMERQNWHLNSLWLWLVGSMLNQGCDLLFRLPMRGLLTAFWIFWALLISQTYIAKLAFFITSSKLADGGSLGDHDLVDQNKQOQFTRGASTVYFSEISNNTDN
DPEVLENQWRIHTNLWTASIMTAGCIDLFRSPQVRMEATWIFIAIIIIANSTYANLAAFLTSKKMGSALTNLDDLASKC-KVKFITYGGSTYNLIALDSNETVR
EPEELENQFTINWSLWFTIAGLQQSEIAFKALSTRTISAIWFFWFTYLWSSYTANLAAFLTIENPTSPIVSKDLADKDVQGAKTGRSTNFSESEPI
EPEELENQFTIGNSWFTTIGALQQSEIGFKALSTRTFASFWFVWFSTLWSSYTANLAAFLTIEKPQSLINTSDALNDGVGAYAKKGTGRSTNFMTSAEERYK
DSDDVVENFTLNNSFIFGVLALQGQSELMFKALSTRTVGGIGWFSTLIIISSYTANLAAFLTVERMESIPDSADLAKQTEKYYAVEGDGAAYMTFFKKSTYD

RAWNQMKDFNPASATSNKEGVARKKEQGGAFLMETTSLLTNYIERCNDLTLQIEQIGEKHYGIALVPGLGSDYTNLSVISILQSERLQLMKNKWKNNHV-TCD
MAWNKMLSFKPDASKNEEGDVRLKLSQTYLAFLMETTNQNYVQKNCETLQQGSEFEQHEGAIAVFLHPLRFRSNLSVGILRSELELKLNKWSNS-TCD
LAPNLMNNDDSAYSINLQLEGVRLKRNQGRFLMETTENLYHEQCNQLDRSGVEFKQFGEAAYFSLNVAKLSEELYLDLQKWKNPNA-SCF
KMNEYMLANPE-MLLENQGQGVKKGSKTQAGAFLMETTNNQNYVQKNCETLQQGSEFEQHEGAIAVFLHPLRFRSNLSVGILRSELELKLNKWSNS-TCD
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TM3

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SNFTOIDQGFDMSVGLGFVFLVQVVVGLIVGVAEFHWQYQIVSVKEKIPMLALAEFYFVIRFWLTRKPLHTYR=QRSDSTSTGYSSELIQTSASSAABKKK
EEPDDPDTFOMFTEELRGIYFLAYAGIALAFPLIFELFVYQQVSLFLELRTFDFKAEKIFVLCWVNNRKPVIAVGTFISSVRTPRSLDKLDSRLTPSRV
AKDSDDGSPGSLGDNLSGTYLWVGLIGISIIISLLCWCYFVYKKAKEYVVFCDALAEFIFVIRFSENERPLKASQYISYRRSNSSQSIESLRLKTLSEENMEVED
TKEADPATLPDMNIELGVEFVFLGVLSSCALLGYIISWLVFVMMCBAHHYRPLDRAKLEEFPQVVDNYYNYVRLNKASIALYRSRSQSMSSVAVAQESQ-----
PEEESKEASALGVQINGIFIVLAILAGVLSVFVAVGFELEYKSKNAQ---------------------------LEKESSIWLVPPYHPDTV------------------
(Figure 1-5a cont.)

SKTVS---------------------------
TRRIEK--------------------------
GRSSEEMREMAQGSSGSNGNAGRGEKEARV
(Figure 1-5a cont.)

GluR-IIA

-------------MRLCPVVIYAFIIIGFLEGIIALGGDDRNEITVGAIFYE---NEKEIELSDFQAFREV---NMRKSELRFVITKRYMPTNSDLLQQT

GluR-IIB

-------------MMLQFLVLLALAIASGANEDTLVIKIGAFFD---TEMLKADSALEELVN---AIN-PALKLDAIKRYVTVDSDIVLQDI

GluR-IIC

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GluR-IID

-------------MHFCWIS---IIILSLSTQVAFQYGGNAYEEAASQGSIRLGLTDD---AETRDIITFEDAIQSVRNAGGNYQFQAFO

GluR-IIE

-------------MFNNHFILWLSFIELSIIHNVNAQYENFFG-YNDSQLEVSPIGLITUDEQ---NTEQMNIVFDHAIADVAN---QEVGTSLSLTKEEVEYGDAYSQF

KA1

---------------------------------------------------------------------------------------------------------------------------------

PMRPSAPVLLVPAWLVMACSPHSLRIAAILDPMCECSERGLSITLAKRNISRAPERLGKAVKEVDIFELLRDSEYETAT

TCELISNGVAIAFGESSK-ASDIVALVAIQANATIGPHIYEYDILKLEATRQEQNLHQSMISNAVPSLVSLSRAYEFLIKSN-YEWRTFTLIIYETPEG---LARQDQIMNQA

SCDLIGSVAIAFGESSK-TNDIVELVLNMTGIGHQLQDFWILQQPSNRERANHQLVTNAVPMELFASFDIILASKTFDWSFTTAYERSSH---LIRQDILAWKQ

ACDDILGVAIAIFGEESK-ASDIVALVAIQCNSTGPHIYEFDISDEGIAEQKPNQMTLNLYPESAQILSAKAYDIVQNF---GWRKFTIVYDDADDARAARQLQDLQRE

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LCRMLGETGIAQVFGESSR-HTAVEHLMS_IDCAMDIHPHYSMS-ENAEG-----------FNLHPHPDAKALYSSTLETF---NWTRFIPFLSEAY---NLINLENTMLMG

MCQILPKGVVAVLGPSSPSASSISSINICGEKVEFHVKEAEEVFKQFQFQQR---FTTLMLHSNTDISVAVAGILNFF---NCTACILICAKAE---LLNLEKLLRQFL

LNSDYVKLRLNLD-YADDDYLWKEDETETHEQRIILDCEPILTLLKVSDFKQFPRNFWFLTHQTHHNSGRLDIYN---EDFKNATVSVKLKVDDANPFERKKT

LHKGAIKQFEF---RGGDYILVETIKN-AREKHVLDDDPSILYDVNASIQYNMGTSQFQWNLNTHSLDQOSP---BDFTAVANISVRRTVYDDFVHEID

VHNDDVVRKRFH---KDDDFRVMWKSIRG---ERRVVLDCPMLNLVFLNSSTEGFQYNYHPRMLTNYTDAI1ELAADADETPAFVNIATALLVNPDPPPSYLFY

IKGFVQIVQMDLNLNGNYSVWLIRK-SREDSSIPSHGSLGRLAK--UNSYGRQVQEPMDQUNLDEYETIIYGGLNNLHTDFLEDEY----EANITGRIIFSPDIDQEEVRLME

KSTGVTILRYDMLQNLNGNQYLKRVRKV-SDVRNLFLVQGSETMPEFNQAOVQVLQINDXYIIYGLDFHSDFDDEEY----EANITGLRSFPEKMAKELLM

ISKTDSLVRMLDD---TRDPTPLKKEID-RDKTATIHANASMSHTILLAELGTVSAYYYTFITLNLFSQLRQMSDVLDDL---RNVILGFSIFNQSSHAFQEEFA

RLT-KVDQIILGN-----QTMLPLIYIDAVLFAWASSARNVIAAMQPMHFP-----NRHCGSSSPWLMGAFIVNEMKTISDEDDEVPHFKCTENMKLDEYQGRINFLEI

VDFNSVDRFTS---SLGQLVYDSTLFEYNYALLEISQRPQ-FYIP-----NSFCG-RGFQWQPGFRLVEQMKQITPKMVKPFQTLQRLNADQGQERE لدي

GYVTQRDNYESED-PPRTLHDSLHLHDALQLAQFSWQHNSAFYPYDMRMVPRVINTCDFADASGGRTWANGYRALMKTGSG-VNNTNFRSTSLQFEQDGQRITFENI

KLHQLGEQEPVNSGSTFTTEMALTYDAVRVIAETTKHLYPQFOM-----LNCSEHERHDQVQDPSTFRNMYR---SMLEIKEITITGRYFEGNVRKQSTFDVIE

KLGYPFTQDE-FRNSCIPTLQEMALYDFQALTPKMLKP---QNCSTQRTSVSRSQDSKFSKKM--------TQLRDLTDLTIPYFEGNVRKVHLVDYE

SRLQSWQENCVDHPFTG-PAILSSALFDAYAVYDVAQAEQINRSQEQIE----VTKPLSCGSAIQWQHSTLMNLVR---MVEGLETGHIENFSQGQSRNYALKILQ

YKPTVNEPMMVWTDP---GIKKRLNL-----LEELASATGQDFESQKRYTVTVTHE-EPYFMKKEDHE---NFRGREKYEYAVDLISKSELMEDFYEFMIVVNGN-

YNPIIDRTHVINKEFQVLVDEKLRENSTQALKRQLNKEDFSQKPIRYTAVR--KPYFSWREEPEGVYHEGNEREGVAVDILYMLAECFQFNFQEPVRDNK

YDPLDGGIPADRF----------------QITQLNVLVAKQKMKYVATRIQ-PYFSYNETARENLHGLNAGYYAVIDALARHGFEEYVFVPVADAQO

LQTGSGKLVGTWEKGR---DEPQQR-PFQAVNEDIDDGLNQSVK7TVLSAVKPTAALVESID---TGLIGNQFOQYGVDIIKELADKMEKFQ-FR-DGGND

LQPSIGKVVTWTDRED----------------QYRQPRLAPTTAQFQDSVND-LSANLKFTILLSVNPKVYLKSHYATQ---QLEQNSQYEGTVKELLADKMEKFQ-FV-NGGND

FRMNGFROIQGWHLAEG-------------LSMDHLYSANISDTLFNTTLVVTITELPMLKGNHQ---EMEGNDREYFCVDMLKELAELFRNYKIRLVGDGV
(Figure 1-5a cont.)

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YGYSYNKTSTNSGLREIMGRADLAAFTLITSMEEVDIFSIPFMNMGIALYVQPKQPAFALSFPMDFPSSEVWFLGFSFLVGSLSFFILGIESEDWNYPY
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TM2
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NVN-REMERQNIWHLNSALWLVGLSGMLQNCGCTIMLPIILMALTAFWIFWIFALLSQTGYIAKLAIAFSTSGIAHDLVD-QNCKQFSTIRGGATSVYESND
CNR-DFFELENQWRHNTGLTBASMTAGCDLPRSPQFVMFEATWVWFAIIANANTANLAEFLTSKMEGSTEANLKLAS-QCKKPKQTIYGGSTYNLLADNE
CIE-EFELEENQFINTNLSWFTQALLQGEIAIPKALSTRTSIAIWVFVTLMVSSYTAANYLAFTIENPTSPINSVDLADNIDQVYAGKTGTSRNFSTSEE
CIE-EFELEENQFINTNSFTQALLQGEIAIPKALSTRTSVASFWWIFVTLSQTYIAKLAIAFTSSKMEGSLATNLDV-AQKQVQFSTIRGGATISVYESND
CAQGRCNNLVNYLSGLSQVSSGFSLQGSSTAPRASLRTCVVSIGWAFATLIIISSYTAANYLAEFTQIWPDESVESVJDAD-CTAIEYTIHGSSMTFQNSR

TM3
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SSTVVLAFNLMNNDFSAITKDNLEGVDVRVRKNGPYDFMLTTTLEEHRQNCILSRSVEGEFKEHYAIAPFFGAEBRNSLSVAILKSLERGELYLQWKWPNPA
PIYKMNENLAH-FEMLMEMNQGGQVGGKSTYKAYFLMESTSTIFNTVRECNLTVGDPLLEKDYGIAVMKNWYRDKFKNALLEOQCQVLARKWNNHEVA
ERYKKNMEFSS-EQFOYLEDMNEGVYNVKTNTYHAFLMESTSTIEYNTKRCNLTKIGPALDEKGYIAMKDWPMRKFFNNALEEOQCQVLARKWNNHEVG
TQTQRWNMYKMSFKESVFVKEEGIARVLNS--NAYLAEFTQIWPQMDPSESVESVVDADQDEEDHRAKGMGGSEMEGKKLVSSARLQNVAAAIPPPAMLEF

TM4
-TCDSYHEVDGDELSSI1LGVLFVLVLAGVLGLVILGIFEFILPNVQNVAAEVEFTPQAFKAELIIFALKFWVRKPMRISS--SSDKSSRSRSLC-GSRRRSK
-TCDSNVPTIDDOQDMDSVGLFVVLVIVGVVGVIGVEAEFWHIVQRISVEKIFPMLAKAEFYVIFWFLTRKPLHTVR--QSRDSTSTGSYSSLVQITASSAK
-TCEFEPDFPDTDMTEELRGIYFYLGILAILFLIIEFLFLQYQVALEELIIFKDAFEKIFIRYFVLCWVKNKRPIVAGTPISVVRPTSLSDKTLPSSR
GVCASAKSDDDPGPSLGDVNLSGYIVVVLGISIISISSILCWFNTYVYKAKNYVEVFCDALAEERFIRVFIRFSNEPERKLSAQSIYSSRNSQSSSTESLKTSE
GICATKEDAPATPLMDRLLEGVFVLLVSGLSCALLYGISWVLFSMKAHHRVFPLRLDEKKEFQVFDNNVYVVLFKNSAYISYSSRQSGMSVSAYQESQ--
----PEEEDRHKGLMELOGIFVVFILCGLVAMAMLEFVTLSR-HSEATEVSVCQEMVTELRIS1IIQCDSISHPRRRAAAVPVPPRIPPIPEERRRGATLNSGK
(Figure 1-5a cont.)

EKSRSKTVS--------------------------------------------------------------
KKKKTRRIEK----------------------------------------------------------
RVVIGRSSEEMREMAQGSGSSGSNNAGRGEKEARV-------------------------------
VED---------------------------------------------------------------
-----------------------------------------------------------------
LCGAGEPDQLAQRLEAALVARGCTHIRVCPECRRFQGLRARFSPARSEESLEWEKTNSSEPE
(Figure 1-5a cont.)

SKEKRSKTVS---------------------------------------------------------------
AKKKKTRRIEK---------------------------------------------------------------
SRRVVRIRSEEMREMAQGGSNGAGNAGRGKEARV------------------------------------
MPVED--------------------------------------------------------------------------------
-----------------------------------------------------------------------------------
LYSAGAGGDAGSAHGGPQLDPGPPSARPAAPTPCQCHVRVQECRRIQALRASGAGGPRPGLGVPAEATSPPRPRPGAGPRELAEHE
Figure 1-5b. Sequence alignment with predicted amino acid sequences of the postsynaptic receptor subunits at Drosophila NMJ (GluRIIA, GluRIIB, GluRIIC, GluRIID and GluRIIE) and human's kainate receptor subunits (GluR5, GluR6, GluR7, KA1 and KA2). Sequence comparisons are achieved to investigate the similarities of each subunit of Drosophila NMJ and human. Since homologous sequences are different lengths, gaps are inserted to gain the best match. Clustal W2 is used for sequence analysis.
(Figure 1-5b cont.)

GluR-IIB

MHGLQFLVLLALAIASG--------ANEDTLVIKGAFFFTEMELAD-----AFSAALEEVEN---AINPALKLDAIKRYVTVDSSITLQDISCDLGSGVAATT
MEHTLLAQPGPLWTRDTSWALLYFLCYLPTAPQVLIRGIFETVENEPVNVEELAFKAVTSINRNRTLMPFTTLYTIDRINLFLDSFEASRRADCDQLALGVAAL

GluR-5

FGPSSKTNSDIVEVLCNMTGIPHLQFDWHPQSNRERMNHQ1TVNAVPEMELFLSAAFSDLASKTFDWKSFTIAAYERSSSHLRLQHILAWKQLHKAGIKMQEFERGD
FGPSHSSSVSAVQISICNALEVPHIQTRKHPSDVNI----DLFYINLYPDYAASIRALDLVLY----YNKVTKVTVVYEDSTGRLQELIKAPSRYNIKIKIRQLPSGN

-STQALKQRLQNKEDFSQPKIRYTVATRVGKYFWSRREEPESGVHYENGNFEGYAVDLILYMLAQECKDFDFNFEVPRLKNGYSDANTDEWDIGILRQIDNNAIQIGC
MTDSNKKSSHNTIDSLANK-----TLVTTTILEEPYMYRERSDK-----FLYGNREDREGCFCLELNSNLGFIYDLCVLPDGKYGQAQND-KGEnwGmveLKEdLHrAdLAVA

-TM1

DLTTIQARRSVVVDVTVPFMQLGISILSYKEPPKADIAYFLNPYNAEVLWLFMVIAAMMTAFALIFTRIDQYEWDPQVENVNRMEQRN1WHLNSNALWVLGSLMNLQ
PLTITVYREKVIFDSKPFMTLGISILYRKPNGTNPGVFSPLNLSPDILWYMLLACLGLSVCLFVIARPFTPYEWYNPHCPNDSDVVENNFTLNSFWFGVGLMQQ

-TM2

GCDLLFPRGLPMRLLTAFWIFALLISQTYIAKLAAPFITSSKIAGDidGSLHDLVDQNKVQGTRGGATSYYFSESNDTDRMAWKNMLSPKPDAMFTKNEEGVDRVKG
GSELMPKALSTRVGGWFFTLLISSSYTANLAAFLVTREREMPSIDADLAKCQTKIEYGAVRDSMTFIFKSKISTYEMWAFMSRSSQQALTVRNSDEGIQRVL

-TM3

LSKGTYAPLMETTNLQYYVQRNCETIQGESFGKHYGIAYPLNADFRSNLSVGILRLSERELFKLKNWKSNSNECTSDNVPTIDDQFDMDVSVGGLFVVHLGV
TID----YALLMESTIEYVTQRNCNLTIQGGLIDSKGYYVGTPFIGPSYRDKTIAILQIQEGKILHMMEKWKWRNGCGPEEEDN----EASALGVEIGGIFIYVLAGL

-TM4

VVGLVIGVAEFLWHVQRIVSKEKIPPMALKAEBYFVIRFWLTRKPLHTYRQSRDSTSTGYSLEIQITSASSAKKKKTRRIEK
VLSVFVAIEFYIKSSNKNDIEQCLSNAIMEELGISL---------QKKIKKSKRTGKSSFTSILCHQRRTORKETVA--
GluR-IIC
MKKELSGNSARWQPOMFGGSNREAGMWRILLLGCMWSAFFMCRSRGGQIN-IGAFPYDEDELe----------LEKFMTTVNAINGPSEQTMRFYPLIKRLLKEDG5VMTQ
------------------------MEHGTLLAQGPLWTRDSTWALLYFLCYIPFTPATQVIPRIGGFETVENENPVNEELAFKAVTSINRNRMT1MNTTLYD1QIINLFDSFEAS

GluR-5
EHACDLDNGVAAIFGPSSKAASDIVALVCNSTGHPHIEFDISDEGIQAEKPNHQMTLNYPAAILSKAYADIVQNFWRKFTIVYDADDARAAARLQDLIQRELREV
RRACDQLAIAGVAALFGPSXSSVSAQVSICNALEVPHIQTRWKHPSVVDKQ---LYINLYPDYAIASRAILDLYYNYWKTIVVYEDSTG--ILRQLQELIKAPSR

HNDVVRVRKFKH-DDDFRVWMSKISRGER---RVLVCPEPNMLVELLNCOSEFTGQTNHYHFLTNELTYDHDEEELIAADENETFAVNTIAARLLVNVDFPPPYSLPYGY
YNIKIKIRQLPSGNKDAPLLEKEMMKGKEYFYVIFDCSHETAEEILKQILFMBGMTEYHYHFFTTLLDFALLDLEYRYS----GVNMTGFRLLNIDNHFVSSIEKKS

TQRDNIVYESSDPRTLHDLIHDACLQLFAQSWRNASFFYPDHRMVVFRTCPFAASGGRPAWAMGRLARLMKGTSGVNNNTFRTSILOFEQDEMQRIFMIEVYDPLD
MER---IQAQPRPEARLGLDMMTTEAAAMLYDVMVIAAS--HRASQLTVSSLOCHRHPWRLGPRMFLIKEAR---WDGLTGHITITTGLRKRDFDDLSILKKE

G--IGIAWDPQGITQLNVDVKAQKMKMYRAT---------IGPPSYNETARELSLNTIGNYLQYQYGAVDLIDIAIAHVGFEYFYVFTVADQQYKGLKDETKQWNGI
EGTEKI1HNSNSNNLMTSNDKKSSNITDSDLSRTTVITILEPEVMVRKGDKL---YGNDRFCYCLDLKELSNIDGIYDVLVLEPKYQAGCMBRGEWNGM

(Figure 1-5b cont.)
(Figure 1-5b cont.)

GluR-II
---MHFWLSLILLSLRVTQAQFYGGAAYEASSQSLRLG---LITDADTRIQTTEHAISVNN---ELGVPLVGEITEQVAYGSVQAFAQLQRMQSGVGA

GluR-5
MEHGTLAQPGLWTRDTSWALLYLYLCPQTVPLRIGGIFETVNEPVNEELAAPFKAVTSINRNRTPMFNLTYDIRQFRNSFESARACDQLALGVAAL

FGPAARHTASHLNDCAKDIPIYPHLSWSGSPQG----FNLLHPSFEDIANYDIVNQFEWRSFICYESAEYKLIDLMTRYGKPIVKYMRDNLNLGNYK
FGPSHSSVCVQQSINALEVPHIQTRWKLPSVDNDFLDYFLNLYPDYAIISRALLDVLYVNWKVTVPPVESTGLILQRKELIKAPRNYIKIKIRAPRPS-GNKDAK

SVLRIRKSEDRIIVVVGSTTGAELLRLQAQQGVMNEDTYYIIIGNILNHTDFLEHRYASEANITGIRMRSPDQEEDLMELKLHQELGEFVNGSS---TFIIIME
PLLKEMMKGKPEFYVIFDCSHEATIKQILFMGMMTEYHHYFTTLDLFALDLEGYRSGNVMTGFRLLINIDNHPVSSIEKWSMERLQAPFRPETGLDGMMLTE

MALTYADAQVIAETTHTLPIPYQOPSMLNCSEHIDNVQPDDSTFRVNYRSSEIKEKMITIGRIFEGYNREKSFVDTLQETSGLVKGWTEEGKDFEFQRPPQAVNFNDI
AAILMYDAVYMSIAKASHPRASQLTQVSSLQCHRHKPR-WLGPRFMNLKEARWDDLTHGITFNKTNLRLKDFDLIDISLKEEGTEKIGIINWSNLNMNTDSNKDKSSNIT

DDGSLVNKTFIVLISVATKPYASLVEISIDTILGNQFQGYVMDLKEALAKLGFNPFTFRDDGNYDFSNKTNSTSGLMLEKEVGRADLAITDLTITSEEREDIFS
D---SLANRPLIV-T--LIPEEFYMVRSDKPLFYNDGREFCGCIILKEHINSNLGIFYTVKDDPGKVGACNKEWNGMVKEHIDRDLAVAPLTTIVYREKVIDFS

TM1
IPFMNLGIAILYVKFQKAPFALFSMDPFSFSWLYLDIALGVSCLFPITGRLSPIEDNYPFCEIEPEELNQFTINSLWFTTGGALQQGSEIAPKALSTRITIS
KPFMILGISLYKRPNGTNPGVFSNLPPLPSIWMYLYLACLGVSCVLFVIRFTPFEWYNPHPCNPSDVVQENFTLLNSFWFGVAGMQGQGSELMPKALSTRIVG

TM2

TM3
AIWFFTLIMVSSYIANTLAAFLTIEFSNPSPINSVKLADNKKDQVYKAKRTGSTRNFFSTSEPIYIKMNEN-AHIPMLMENNOQGVDKVSGYAFLMESTSI
GIWFFTLIIISSYIANTLAAFLTVERMESPIDADDLAKQTIEGYAVIRGDGSTMFFKSKISTYEKMWFMSRRQQTALVRNSDEGIRVLT-ТИДАМЕСТSI

TM4
EFNIVKRCNTKGCPLDELKGYGAIMVKNWPFYRKFNFKNALLELQEQVVLARLKNKWWVNGAGVCASKDDDGGPSLGVNLGSIYVLLVGISSISISILCWF
EFVTRFFCNCNLTQICIGLDSKGYGVPFGPSYRDKITIAILQLQQEELKHKKEKKEWRRGMC---PEEINKEASALGVENIQSIFIVAAGLVLVSFVALIPEFITYK

VYKAKNYEVFPFCDALAEAFRIVFRSNERPLKSAQIYSRSRNSSQSIESLKTDSIENMPVED
SRKNNDEEQCLSFNAIMNELGISLKNQKIK---KKSRTKGKSSFTSILTCQRTQRTKETVARED---
(Figure 1-5b cont.)

**GluR-IIE**

MFFNHFVILWSLFSIHSVNAAQYENFGGYDNQSLESMPVGGGATLTDQNTEQMNIVFDHAIDVAN----QEVGTSTLSLKEEYNYGDAYQSYGKLCRMLEGITA

---MEHGTLLAQPGWTDRTDSWALLYFCYLPQTAPQVLRIGGFETVENDINGPVNEESFAKPAVTISNRRNLMPNNTLYDQRINLFSFEASRACDQLALGVA

**GluR-5**

GVFGPSRHTAVHMSIDAMIPHYSYMSENAD----FNLPHPHDALAKALYSLITEFNWTRFIFLYYESAEAYVNINLONELTTMLGSGTGVITVRYDMLQNGN

ALFGPSHSSVSAVQSCINALEVPHPITQRTWHSVVDNKDFYINLYPDYAISRAIDLVLVNYKWTVTVYDESTDGLIRLQELIHKASRYNIKIKIRQLPSSNK-D

YKQVLRVRKSVDRNIVVGVSETMPPEPLNQAQVGVINEDEKYYIIGNLDHFSDFDEFSEYKESEANITGLRLFSPERMAKVELMKLGYPTQDEFNGS----CPIT

AKPLLEMKFGKEFYVINIFDCSHEVAAEILQILPMGMTEYXYHFSTTLDLFDALDELYRSVGNMTGFRLNIDNPHVSSIEKWSMERLQAPPRPETGFLDLGGMT

VEMENTYDAVQLAQPQKMPQNCQRETEVRDGCASSFKNYWRTLALTDRLLTGPIYFEGN---VRKGYHLDFIELQPSIGKVECTWDIDRQYRFQRLAPTTA

TEAADMYAVYMRVAIASHRASQQTSLQCHRHPWR-LGQFMPNFLKEAVWG----LTGHTIFNTNGLARKDFDLDDIISLKEEGTEKIGIWNNSNSGNN--MTDSNKD

QFDSVDNSSLKFIILLSVPNFQYAQVLETVYEQLENSQYEGYVLIDKELDGLGFNFNVGNGDGNYSNKTSTNETOMLEIETMDATLDLITTSREQA

KSSNITDSLSANRTTVIT-ILEEYPYMRKSDPYMGKFGFCYSLDLKELSNILGIFYDVKLVDGYKAGQNDKEEWNMQVELIHDRRAVLAVPLITYVVEKV

**TM1**

LDFTFPNFLGIAILYKQFQKAFELFMPDPSSEEVWFGLGSGPSLHRLSSEPDNPYPCIEEPEELENQFSTLGNSWFTTGALLQQGSEIGPKALST

IDFKPFFMTGLISILYRFQGNTPGFVSFLNPSSDIWMVYLLACLVGCVLFLVARETPYEYNYMPCNPSDVVENFLLNFSFWGFVMQQGSPELMPKALST

**TM2**

RTVASHWFWFVFTLVWVSSYTANLAAAFTEIPEFOSILSGVDDLADNKGQVYAGKGTSTRNFMTSAEERYKKKNKFMSENFCQYLDENMLMEGNVRVKTNYAFLMSE

RIVGGIWWFVFTILLISSYTANLAAAFTEIPEFOSILSGVDDLADNKGQVYAGKGTSTRNFMTSAEERYKKKNKFMSENFCQYLDENMLMEGNVRVKTNYAFLMSE

**TM3**

RTVASHWFWFVFTLVWVSSYTANLAAAFTEIPEFOSILSGVDDLADNKGQVYAGKGTSTRNFMTSAEERYKKKNKFMSENFCQYLDENMLMEGNVRVKTNYAFLMSE

RIVGGIWWFVFTILLISSYTANLAAAFTEIPEFOSILSGVDDLADNKGQVYAGKGTSTRNFMTSAEERYKKKNKFMSENFCQYLDENMLMEGNVRVKTNYAFLMSE

**TM4**

TSIEYNRKRECLNLKIGDALKDEKGYIAMRDKPHRGKPFNNALLELSEQQVGLEKMKNNKWNVGTGICATKEDAPDATPLDMNNLEGVFVVLGVSQSCALLYGISSW

TSIEYTVQRNCLNTIGGCLISGKYGVTPIGSPYRDKIITAIAQQLRCEKGLHMMKKEKRNQG-----PCEENKEASALGVENIGGIFIVLALAGLVLSVFVAIGEF

TVLFMICAHYVRPLRDALKEEFQFVIDFNNYVRVLKNSASIYESRSQSMSVSAVAQESQ---IYKSRKNDIEQCSSFNAIMEELGISLKNQQKIKKSKRTKGGSFSTILTCQRTQRRTKETFVA
(Figure 1-5b cont.)

**GluR-IIA**

----MRLCPVVIYAPIIIGLEGETALGDDGRNEITVGAFYENE----

**GluR-6**

MKIIIFPILSNPVPFRRTKLLLCLLWIGSYGQTTHVLFRFGIFEGYEVEGSPGMAEELAFRFAVNTINRRIKTTLPPNTLTDDTQKINLYDSFEASKKACDQLASLGAVAF

YRILKWEKDTEDEPFVIQRIILDCEPKTLKELIVSDFKQLQGPFRNWFLTHLDTHNSSLGRLDIYNEDEFPKAN-ITSVRLKVVDAPFERTKTRLTKVDQILGNQTMFLIL

AKPLKEMKR-GKEFHVIFDCSHEAAVILQALGMQMTYEHYIPTTLDFALDVEPYRSGVNTMTGFIIRI1NTTQVSSIEKWSMERLQAPPKPDSGGLLDGFM

YDAVVFPLASSAFNVIAAMQFHPFPN---HCQGSSPWMLGAFIVNEEMKTEDEDVEPHFKTENMKLDEYGRQIEHFNLIEYKPTVNEPMMWVTDFQNGIKKLRLNNLELE

TTDAALMYDAVHVVSVAQFPQMTVSSLQCNRHKPWRFGTRFMSLIKEAHWEGITGRIFN---KTNGLRTDFDLQVSLKKEEGLIKEGTWDPASGLNMES-QKGM

SAGTTQDFSEEQRKVYTVTHTBPEYFMMEKDEHFRGKEKEGYAVDLSKLSELMEDYEYIMVNGN---GKNPETKQWDGIIIRKLIDHHAQSVGCDLTITQMRSSHF

PANITDSLK---RSILVTILLEEPVFKSVFDLYCSNDFEGYICDILRRELSTLTIYEVIRLVEDGKYQAEDANGQWNMVREILDHKADIIVAPAITYVREK

**TM1**

VVDFTPFMQLGCSILHLKYSPPEPKQAFLEKWAFQVENMYMIFAQLMIMLFVFTARLYREMLPNFAIQDPELENIWNVNNSTWILMGSMOQCGDILRPGP

VIDFSPKMFTLGISILYRPIPNGTNGVFSFLINPLSPIYMNLLAYGVSCVLFVIAFHYRYNHEPFCNPDSTTVENNFTLLNSFVCQGALMOQGSELMKALS

**TM2**

MRILCTQMNFEAIKMILTSTYTANLAAFLTSNWDQSSIKLQDLIEQDKVKHFGSMRGSTSLFFSESNDTDYQRAWQNMKDFNSAFSTNKEGVARVRKEKGGYAFLM

TRIVGSIWWMFTLIIISSYTYANLAFIVERMESPISSADLAOKQTKEYGAVEDGATMTFFKSKISTYDKM@AFMSSRQSLVKSNEEGIQQVLTSDYFAFLM

**TM3**

ETTSLYNIERNCDLTQIGEQIEKHAYGLAVPLGSYRTNLNVSILQSEREGLQMRRNKWKKMNIVTCDDSYHEVGDGDELIIIEdwardVeLVAGVLLGIVMIGEF

ESTTEFVTQRNCNLTQICGGLIDSKGYGTVFMSGYPRDAAITAQLQIEGKLLHMMEKWKWRGN--GCPEESEKEASALGVQNNIGFIVLAAALVLSSVFAVGF

**TM4**

FWNQVIVEEREVTWFQAFKAEILFALFKFVVRKPKMRISSSDKSSSRRSGSRSSKESKRSKTVS

LYKSKKNAQLE------------------------KESSILWLVPPYHPDTV------------------------
(Figure 1-5b cont.)

GluR-IIB

________________________MHGLQFLVLLALAIASGANEDTLVIKGAFFDTMEKLADAFSAALEELNS------AINFALKLDAIKRYTVDSIVLQDISCDLIGSQAATF
MKIIIFPILSNPVRRTVKLLLCLLWIGYSESQGTTHVLRFGFFIYEVESGPMGAEALAFRAVFANTIRNRTDLFPNTTLYTDQKINLYDSFEASKKACDQLSLGVAATF

GluR-6

________________________GPSKTNSDIVEVLNQMTGPHLQFDWHPQSNRERMNNQLTVNVAPMEELFLSAAPSDDLASKTFD3WKSTTAYERSSSHLRILQHILAWKQLHKGAGKQEFERG-D
GPSHSSSANAVCSIGNALGVPHIQTRWKHQVSDD----NKDFYVSYLPSFDSSLSRAILDLDVFFKWKRTVTUVDYDDSTGSLIRQLKIEPSRNYRLIRKIRQLPADTF

________________________DYRILWKSNAREKFLVLCDPSDLIDVNVASIGYNMTGSKNLFLTLTLDTHSGKDFYSRDFTVAAAVVRR---IRTVFPPPVDEIDVDFFNDVTRFSSLGSLQDV
DAKPLLEMKRGKEFHVFDCSHEMAAGILQALAMGMTEYHYIFTTLDLFAVLDFEPYRYSQVMTGFRILNENTQVSSIIEKWSMERLQAPPKPSGILQDMFM

________________________YDSIVLFYNALLEIS----QRPGFYIPNFSGEGF-FWQPCPLVEQMNQFITKMPKVFPKTQRLQINADQOREDFNLEVYNIIRVTMINKEFQLVDFEKKRENS
TTDAALMYDAHVHSVAVQOFPQMVTSSLQCNHKPWFROGRFSMIIKAEWAGLTLGRITFNEKTNGLRTDFDLDSLKLEEGKLEKIG-------TWDPSGLNMM

________________________TQALKQKLQNKEDFSQPKIRYTYAARRVKGFKIFSNRREEPGYEHGEGEFEGAVDLVMMLQECFEDNFEVRONKYGSYNDANTDEWDGIIQLIDRNQPAGCID
TESGKKGKPTANITDSLSNR---SLIVVTEELEYFLFVFKSDKD--PLYGNDREFGYIDCDILRELSTILGITYEIRLVEQDKYGQDDANGQWNNMVEHLIDHDKADLVAF

________________________LTITQARVSDVDFTPVFMQLGILSYKEPPKADIYAFLNPYNAFVWLFVMMITAFALFITGGIDQYEWDQVENVNREMERQNIWHESNALWLVMLQSMNQG
LAIITYVERKIDFSKPFMTLGISILYRKPNGTVPGPFSFNLQSPIDMYILLAYLGVSCILFVEYAESPYESYNEWHPCNFDSDVVEENFTLNSFQFGVALMQDG

________________________CDLLEFGPMLLTAFWDNALLISSOTYIAKLALAFITSSKIAGIDSFDLDVQNKVQGTIRRGATSVYFSESNDTNRMANNKMLSFKDPAFTKNNKEEGVVDK
SELMEKASTRIGGIQWFTLISSYITTANLAFTVRMMEMPSADDLAKQTKIEYGAEDGAMATFPPKKSKISTYDKAMFSSRFQSVLVSNEEGQIVRKL

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(Figure 1-5b cont.)

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(Figure 1-5b cont.)

GluR-II D  
---MHFCWISLIIISLSRVQAQFYGGNAYEASGSQSIQLG----LTDDATDRIRQTPEHAISVNN----ELGVPLVGETEQVAYGNSVQAFACLRLMQSGVGAFFMKIIIPILSNFPVFRRVTKLCLLCLLWIGYSQTGTHVLRFGIFYEYVESPMGMEALEAFRAAVNTINRNLTPNMTTTLTDTQINLYDSFEASKACQDQLGVAEAF

GluR-6  
GPAARHTASALNAACDSDKIDPFPIYP+LSWGSNPDG----FNHLHFSPEDIANALYDVNQFEWSPFIFCYESAEYKLILDHMTYRIGKIPVKVMRDLNINLYKSGSGLSSSANAVQISIGNALGVPHTIQTWFKHQSVDNKKDSFYVSLYDFSSLSSAILDLQFFKKWKTQTVYVDDSTGQLRQELKAPFRLNRLKIRLQQPADTK-DAKP

VLIRRNKSEDSDRIVGVTGGSTTGAVELLROAQQVGIMNEFYTYIIGNLHLHFDLEYSEYKSEAMINTGRMFSPOQEEVEVDMEKLMHQLQGESEPVSNGL--STIFMEMLLKEMKKGKEFHVFIDCSHEMAAGILQALAMGMMTEYYHYIFFTTLDFALDVEPYSVQGISNMGTFRILNTENTQVSSIEKWSMERLOQAPFKPDSGLDDGFTLDA

ALTYDARVIAETTKFLPYQPSLNCESRHIDNVFDPGSTRNYMRSLEIKEKTITGIYFEGN--VRKGTITDVIELQTSGLVLQTKWEEGKDFEQRPPQAQQND

ALMYDANVHSVSVAVVQQFP----QMTSSSLQNRKHFWRGTRFMSLIEAHWELTGRIFNKTNGLRDTDFLDVDISLKEELEGKIGTWDPASGLNMTESSQGKAPNI

IDDGSIVNKTFLIVLSFGAVLSVEIDTLIGNNSQFQGVYDVILEKADLKGFTFRDGNN--DYSFENKTNSTSGMLKEIVEGRADLAILDTLTITSEREEVIDTDS---LSNRSL1TITILEEPYVLFKSKDFLYGNDFRFGYFMDINSLSTILTGLEYTIEVLDGGYQADANGQWNGMLVRELDHKGKADVLAPLATTYVREKVID

FSIFPNMLGIALYKVQKAPPAFLSFMDPPSSSVDLGYLAIYLGVSLCFLFLLGLSIHEWDNPYFCIEPEELENQFTINNSLWFTTGALLQQGSEIAPIKARSTRTFSKPFMTLGIISLYRKPNGTNFVGFSFLNPSSDIHMYILLAYLGVSCVFVIAEFSPFYWYNPHCPNPSDVVENNFITLNSFWFGVAGMLQGQGSELMPKALSTRI

TM3

ISAIWFFETLIISSYMTANLAAFLLNTPTSPINSVKDLADNKKDQVYQAKRGTSTRNFSTSEPIYIKMNEYLN-AHPEMLMENQQGVDKVSQKTYAFLMEST

VGGIWWFFETLIIISSYMTANLAAFLTVERMESPSADDLAKOT-KIYGEAVEDGATMTFKKSKISTYDKMWWAFMSSRQSVLKSNEEGIQRVL-T-SDAYAFLMEST

TM4

SIFITVRECNLTQDKGPLDEKGYGIVMVKNSPYRDKFNKALLELQEQVILARLKNKWNEVAGVCSASASKDDGPSLGTVDNLSTTVLVLWSISIISILCWC

TIEFVIQRNCNLTIQGGLIDSKGYGVTQ PGMSPYRDKTIAILQEOEQLGKHMNKEKMMWNRGNCPEEEGKE---ASALGVCNIIGFITLAAEFLVLSSVVFVAVGEFL
Figure 1-5b cont.

GluR-IIE

-------------MFFNHFVILWSLFSIHISVNWAQYENFGGYDNYQSLESVPGLLTDMQTEQNNIVFDHAIĐVAQVEGTSLLTSKKEEVNVYGDAYQSYGKLCRMLET

MKIIIFILSNPFRRTVKLLLCLLLWIGYSQGTTVLHVRFGG--IFEYVESGPMQAEELAFRAFVANTINRNTLLPN-----TTLYTDQINYLDSFEASKACDQSL

GIAGVGFPRSSRHTAVHLSICDANDIPIHYSYMSENENAE-----FNLHPHDALAKALYSLITEFNMTRFIYFLYESAEYINILNLETMLMGSMTGTVTVRLYDMQL

GVAALIFGPSSANAVQISICNALGVPHIQTRWKHQVSDKFSYVSLYPDFSLRSLAUDLQFPPKKTIVTVDYYTDSSTFLGRLQLQKAPSRYNLRLIKRQLPAUT

NGNYKQLRRRVRKSDNVRIVVGSSESTMPEFLNQAQVQGIIHEDKITYIIGNLDFHSDFLDEEYKISEANITGLRLFSPEKMAVKELLMKGYPTDQDEFNGLS-----C

K-DAKPLLKEKRGKKEFHVIFDCSHEMAAGILKQLAMGM4TEHYHYIETFLLDFALDVPEYRSGVMTGFR1LNTENTQVSSIEKWSMERIQAPPKPSGGLDG

PITVEMALTĐAVQLFAQTLKNLFFKPMPPQNCQSRTESVVGDGSGSFKNYMTTLRLTDTRLATLPTIYFEGN--VRKGYHLDVIELPSG1KVGTVDEDBQYRPQRLAP

FMTTDAALMYADVHVSFAVQQFPMQTVSSLQCNHPFIFR-FGTRFMS LIKEAHWEG-LLTGRTFNKTNGLRTFDDLVISILKEELEKIGTWDPASGLN-—MTES

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QKGKPAINTDSLNRSKSLIVTT-ILEEFPVLFFKSKDPLGYNDRFYGCYICDNLRELSLTIYGEYI1RLVEGDGYQAQDMANGNWGMYRELDHKLADLAVPLAITYV

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TM3

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FLMESTSIEYNTRCENLKKIGDALDEKGYIIAMRKDWPHRGKFNNALLELQEQCVMKMKWNEVCTGICATKEDAPDATPLDMNNEGFVFVILVCSCALLY

FLMESTTIEFVTQRNNCNLTQIGGLIDSKGYGVTFMGSPYRDKITITAILQLEEGKLHM4KEKWWRGNC-----CPEEESKEASALGQIVNIGIFIVLAAG-----

GTTSWLFFMVKKAHHLRVPLRDAAKEI{BQFVIDFNYRVRVLKNASIIYSRQRSSMSVAVQESO

--ILSVEFVAVGEFLYSSKNTA{KESSWLVPPYHPDTV-----------------------------
(Figure 1-5b cont.)

**GluR-IIA**

---MRCPVVIYAIFIIGLEGIALGDDDRNEITVGAIFYNE---KEIEELEFDQAFREVNNMKFSELRFVTIRKRYMTNDSFLLQQTTCELISNGVAMTAWPRRLRSLWYAGQVCAIIPDSRGMHPVRIGGGFYADGPNAQMVAEEHAFRFSANTINRNTLPLNNTTLTQRIHFDSFEATKACDSSLQALGVV

**GluR-7**

AIFGQSSKLANDAIAIQANATGPHIEYDLKCLEATRQEQLHNQMSINVAPSLSLSRALYFEIKISNYEWTFTLVIETPGLRKLQDLIMQALNSDVKLRNLDAYAIFGQSSQCTNAVQSCYANEVPHIQLRKLHPPLDNK---DTFYVNLYPDAALSLEASLHAIDLQVY-LKWRSATYVVDSTGLRILQELIMAPSRYRNIRLIRQQLPIDEADDYRIWKEIDETTFEPQRDELDCEPRTLKELKVLSIFQRLQGPRNNWFLHTLDTHSNGLRDYIENEDFKAN---ITSVRKLVDANPFERKKTERTKVDQILNGQNTMSDSSIPJLDEMRK-GREPRIIFDSCHTMAAQILQAMAMGMMTEYHYFIFIITLVIDLPEPRYSGLNLVFRDILQVEAEWKEKERSQRESEGGMELPILYDAVVLFASSARNVIAAMQPFPHPN-RHCGSSSPWMLGAFLVNEKMTISEDDEVDVEPHFEKTNMMLDEYQRHFNLEIYKPTN---EPSMVTDPDNGIKKRLLGVMMDAALLLYDAVHIVSVCYQRAQPMTVSNLQCHRKAQRFGGRFMNFKEEAQWELTGRIVFN----KTSGLRTDFLDIIISLKEDGLEKVGVWSPADGLN---I

**TM1**

NLELESAAGTTQDFSEQMKVTFVTVVYEEFYFMKMKHEDHENFREGREYKEGYAVLDISLKSELMEFDYEFMPVGGNGKYNP---ETKQWDGIIRKLIDHHAQIGVCDLTTVEYAKGRGNVTDSTLNRSLIVTVLEEPMVFMRKSDRTLYGRDRECFEGKILDEKLLEAHIGFSEIRLIVE-DEKYGAQDLCQWNGMKVKEIDHKAIDLAVAPLTT

**TM2**

QMRSVYDFTVPFMQLGISILHyKSSPPEPKNQFAFLEPFAVEWVYIMFQALIMTLAFLVIRLSYREWLPNNPAIDQDPDELEN1WNVNNSTWLMGGSIMQGCIDLHVREKAIDFSKPMTLGVSLYRKPNGTNSPFVSNFLSPFMYVYLLAYLGVSCVLFIARFSPYEWYDAHPCNPGSEVVENFTLLLNSFWFGMSLMQGGSEM

**TM3**

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**TM4**

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NLELESAAGTTQDFSEQMKVTFVTVVYEEFYFMKMKHEDHENFREGREYKEGYAVLDISLKSELMEFDYEFMPVGGNGKYNP---ETKQWDGIIRKLIDHHAQIGVCDLTTVEYAKGRGNVTDSTLNRSLIVTVLEEPMVFMRKSDRTLYGRDRECFEGKILDEKLLEAHIGFSEIRLIVE-DEKYGAQDLCQWNGMKVKEIDHKAIDLAVAPLTT

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AVGEFVYKLRKTAEREQVRS-----------------------------LRAWSLHRMGNGDSR----
(Figure 1-5b cont.)

GluR-IIB

---------MHGLQFLVLLALIASGANE----TTLVIKIGAFFDTMKDAFSAALNTEVAIN--------PAKLDAIKRKYTVDSLSVLQDSCDLIGGVA

GluR-7

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AIFGPSQGSCTNQVSICNALEVPHIQLRWHHLPLDK---MTFVYNLPDYASLSHALDLLW--YKLWRSATVYDSTGLRLQELIMAPSRYRNLKIRQPLI

-GDDYRIILWKRIRNNAKEKFVLWDCPSDILDVLVNASIGYNMTGSHNLFLTLDTHLSGDGFRSDFRDVAVAAAVRIRTYVPPFVHEDIDVDNS----VDTRFSSL

DSDDSRPLKLKEMKRGRELRIIFDCSHTMAAQILKQAMAMGAMMMTEYYHFIHTLDLYALDLEPHYRYS---VNLGTGRIILVIDIAHVAIYAEKEKSMERLQAAPRSGGL

GSQLYDVISVLFYNALLIES----QRFGFYIPNFSGGRG-FWQPGRPLVEQMKQITKMWKPFKTQRLQINADQREDFNPVEVYNNPILIDVTRTVHIpNKEFQYDFEK

LDGVMTADAAALYDAVHIVSVCYQAPQMTVNSLCHRHCAIRFGGRMFNIKEAQWEGLTGRIVFV----KTSLGLRTFDLDIISLKEDEGKEVV------VWSAPA

IGICDLTQCARFSVBDFTVPMGQMGSLYSYKEEPKKQADIYALNPYAEFLWFMIMQAMITAFAALITGRIDQYEWQVFENVNREMERQNIWHNSALWVLWLGSL

LAVAPLTTIITHVREKAIDFSKPFMTLGVSILYRPPNGTNPSVFSFVFLSPQIWMYVLLAYLGVSCVLFVIARFSPYEDHAPCNPGSVVENNFTLLNSFEGMGS

MLNQGCDLPLMRLTAFWWIFALLISQTYIKAALATPKKIQAGDIQSLHLIDVQNKVQFGTIRGGAQTSVYFESNDTQDNRAWAKLMSPFPAFTKNNNEDGV

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QRALTAD--YALLMESTTIETVQRNCNLTQIGGLDSKGYGIGTMGPMSGNYDITIALQEDKLIHMKEWVWQGSGCEPENK---EASALGIQKIGGIFIVL

TM1

TM2

TM3

TM4

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AAGLVLSVLAAGEVYKLRTAEREQVS-----------------------LRAWLSHRMGNGDSR------------------
(Figure 1-5b cont.)

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<th>GluR-7</th>
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| TM1 |
| TM2 |
| TM3 |
| LTVAISMATGCDILRSPQVRMEATWIFAIITANSTANLAAFLTSKMEGSITANLKLDSAQKKVPFTGTYIGGSTYNLADSNETVYRLAFNLNNDDPSAYKTKD-
| FGMGSLMQQGSESMPKALSTRIIGGWFFTLIIIISYANTLANAFLTVERMESPISDADLAKQTKIEYAVKDGATMTFFKKSIKFEMWFAMSSK-PSALVKE |
| NLEGVDRVRKNDRGYDYMFLMETTTELYHERQCNCDLRSLVGEFKEKHYAILVPFAEYRSRLNTAIALKLSERELYDLKQKNPNAASCREEPDADTPMTFEELR-|
| TM4 |
| IFYTLAGILIALIGITEFLVYQQVAALEERLTFKDAFKKEIRFVLCVWNRRKPIVAGTPPISSVTTRPRSLKSDRTFKSSRVRVIGRSSEEMREAMAQSGSSS |
| FTIVLAAGLVLSSVVAQVEFYKLRKTAEREQVSLR-------------------------------------AWSLHRMFNGDSSR---------------------GSNAGRGEKEARV |
(Figure 1-5b cont.)

GluR-IID

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GluR-7

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TM1

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TM2

IGGIWWFPTLIIISSYTANLAAFLTVERMESPIDADLAKQT-KIEYGAIVDKGATMFKKKSKSTFEKMWFAFSKPSALVKNNEEGIQBAALT-ADYALLMESTT

TM3

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YEYTVQRNCNLTIQGLLIDSKSYGIGTFMGSPYRIDKTIAILQLQEDIKLHIMKMWRRGS-----PEENKEASALGIIQKIGIFIVLAAGLVLSVLAVGFEVY

TM4

FVYKKAKEYEVPFCDALAEFEFRIVFSENERPLKSAQSIYSYRSMSSQSIESTFLTDSEENMPVED

KLRTAE---------------------------------------------------------------------REQVSLRAWSLHRMGNGDSR----------------------------------------------------------------
(Figure 1-5b cont.)

GluR-IIE

---MFNFHVILWLSFISHSVNWAGYNYQ---LESVPILGLTDQNETEQMNIVFMTDHAIDVANQEVG---TSLTSLKEEAVVDAYQSYGKLCRMLETGIA

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GluR-7

GVFGPSRRTAVHSMICDMIDPHIYSYSEAE----FNLHPHPADLAKLAVSLITEFKNWTRFIFLYESEAELINLNLTTMLGGTGGTTV/LR/LQAGQLNGN

AIGFGPSQGSCTNAVQStunicalevPHQLRLWKLHPDPNKTDFYTVNLGYDAPYASHLWALQWSATVVDGSTQILRQELIMAPSRYNIRLKR ICQLPI S-D

YKQVLVRKSRVDNIVVGVSETMPPEFLNQAQQVQIGINEDEYKIIYINGLDFHSHDFLEEEYKYSANITGLRLFSPFMAKAVKELMKLYGPTQDEFRNGS----CPIT

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TDAALYYDAVHIVSCYQVARQMTVNSLQCHRHKAWFQG-RFMNFIKEAQWEG--LTGRIVFNKTSGLRTDFDLDDIISSLKPDGILKEKVGVWSPADGLNITEVAK--G

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TM1

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TM2

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TM3

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DIVAVTIANATGPHIEYDSDLKAEATQIEQNLWNSMVLSRAFYEIKSYNWEirtFtllFytStGRLQLDQMMQNIAINSVDVYLKLRLNADYYRIDYRIKGET
SIISNICGKEKVPHFPKVAP--EEFVKFQFQFRTTLNHPNSNDISVAAGILN-FFNCTTACLIACAETCMLNLEKLRLQFLISKDTSVRMLDDTRDPPTPLKEIR

DETFHEQRTILDCPKTLKELKVKSIDFKLQGQRWNFWLHTLTHSNGLRDIYNEFDFAN---ITSVRKLVVPANFKEERKTTRLTKVQDQLGNQTLMLPIYDAVV
LDD---KTATIIIHANASMSHITLLKAELGMVASYYYTIFTNSTLSHFSLQRMDSLVDRVINLGFISIFNQSHAFFQEEAFQSLQSWQENCDFHFPTGAPAILSSLIFDA
Y

FASSARNVIAAMQPFPHRNCHRCSSSLSPWMLGAFIVNEMKTISSEDDVVPHEKTENMKLEDEYCRQHRNFLEIYKPTVNN--EPNVMVTPDNGIIKLRNLLELSEAGTTQ
D AVVTAVQELNRSQIEGVPKLSCGQAQIWQHTSRLMYRMVELEGLTGHIEFN-----SKGQRSNYALKILQFRNLGFRQIGWHAEGLS---MDSHLYASINTD

FSEQRKVVRVTYVTHYEFPYMFMEDEHNFRCREKYEYAVDLISKSLEMDFYMIVMNVNGKYNPETK-QWDGIIKRLIDHHAAIQGVCMLTITQMMRVSVDFTVFULL
LFN---TTLVVTIIILENYMLKQNGHEMQENDRJYFECGKLMKELAEILRFNYKILRLVGDGYVGEPEANGTWTGMVGEILAKCLAVAGLTTIAERKEVIDFKEF

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M2

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EGVARVRKKEGGYAFMETTSLTYWWAFTLIIISSYTANLAFLTVQVMVPESVDDLADQATIAEYGTIHGGSMTFFQNSRYQTVQMRWNYMSKQPSEVSTEGIARVLNSN--YAF
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NIERNCDLTQIEGQIGEKHYGLAVPLGSYDRTNLSVISLQLSERGELQKMKKNWWKHNHTEDSYEVDGDELSLQELGGVFLVISAGVLIAGVIGIFELFVWAV
QNYQCRNCNLQIECSILLDCFGYIGMNPVSVFRDFSLAUDLQLEQNRLIELRKWEWEGKCPKEEHRAG--GEMWGNIGIVFLVLCGLIVAIFMALEFLWTLRHS

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EATEVSVQEMVTELSRIILCQDSIHPRARRAAVPPRFPPIPEERRFRGTATLSNGKLCGAGEPDQLAQRLAQEAALVARCGTHIRVCPECRRFQLRARPSPAR
ESLEWEEKTTNSEEPE
Figure 1-5b cont.

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NAINSNICGEKEVF--HFKVAPEEFVFPQFQFRFTLNLHPNTNDSVAVAGIIN--FFNCTTAACLICAECALLNLEKLIRQFLISKDTLSVRLDLDTRTDPTLLEKIRUD

REKVFVLLDCPSDILVDVNASIGYNGTMSFNLHFLTNLTDNLHSIGDFYSSR-----DRTVAVAAVIRRTYVFPVPVHDIEVDNFDVSDFRSSLGQLVYSDFLYN
ETATIIIAHNASMSHTILKLAAELGMVSAAYTYIFTNLEFSIQRMDLVDVDRVNIQFGISIFQNSHAFFQEFQSLENQSWQENCDDVPFTGPEFALLFDAVYAVVT

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AVQELNRSQIEGIVKPLSCGSAIQIWHTGTSMLNYLR-----MVELEGLTGHIENSKGQRSNYALKILQFRGNFQIGGQWVHVAELSMDSLHYASNISDTLFN--

DFSQKPIRYVTARVGGFYSWREEPEGEGVHYEGNERFEGAYVDDLILYMAECKFDFNEVEVRDNKYQSGYDANTDENDIQLRQDDNNAQIGCIDLTITQAARSVDVF
---------TLTVVTITLLEPNYMLKQ--NHQMEMEGNDREYGFCEGFLAKELAEIFNLRYKIRLVGEQVGGVESANE-GTNGMTVGELAIRAKDVAAGLTTAERKEVDF

TM1

TVFFWQLGLSILSYKEPPKADYIAFLNPNYNAEWWFVMIAMMTAFALIFTGRIDQYEQDQPVENVN-REMERQNIWHLSNALWVLGLSMNLQGCDLLPRGLPMRL
SKFMTLGLSILSYHVMRKPGYFSPFSPFGVWLFMLLAVLSCVFLVALRTPYEQSHPCAGRCNLLNVYQLGNLSLWFPVGGFMQQGSTIAFALSTRC

TM2

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VSGVWWAFTLLIISSYTANLAAPITTVQMDVPESVDDAQDQAITEYGTIHHGSMSTTTFOSSRQSTQRQMWNYMSKQPSVFKSTEEGIAVNLN--YAFLESST

TM3

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SVGLFVVLIVGVVGLVIGEAELW
MNYRYRCRNCLNTQIGGLDLTKYGIGMPVGVSFRDEFDLAILQLENNRLELIKRWWEAGKCPKEDHAKGLC---MNGFIIVLICGLIVAEIFMAMLEFLW

TM4

HVQR-----ISVKEKIPMLALKAEFYFIRFWLTKPLHTYRQ-----SRDSTSTGYSSEIQITSASSSAKKKTTRRIEK
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--------------
PARSEESLEWEKKTNSSEPE
(Figure 1-5b cont.)

GluR-IIC

KA1

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TM1

TM2

TM3

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TM4

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MAQGSQSSSSGNSNAGREKEARV----------

CPECRRFQGLRARPSPARSESLEWKEKTSSEPE
(Figure 1-5b cont.)

GluR-IID
KA1

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ARHTASHLLN-ACDSKDIFFIYHLSWGS---NPDGFLNLHPSPEDIANALYDIVNXWEWSRFIFCYESAENYKILDHLMTRYIGKGPVIMVRYDLNINQNYKVS

SSPASSISISASICRICEKVEFHCLKVAPEEFVFQFQFRPTNLNLHPSNDTISVAGVILNFNCTTACLICAKACAELNLEKLQLRFLISKDLTSLVRMLDDTRP---TPL

LRRIRKSDSRRVVGVSTGVAELLRSQAVQGVIMNEDYTYIIIGNNLHNFDEELEYKSEANITGIWIMPSDCVQVQVRLQEMKLBQTELG---SEFVNSGSTFTMENA

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AVYAVTVAVQELNRSQEGIGPLSCSGS---AQIWHQGTSLNYLPRMELEG---LTGHIENSKQGQNSYALKIQFTPQNRQIQQWHAVAGILMSMDSHLAYSNIDT-

DGSLVKNKQFIVLISVATPVASLVEISILTILGNQQFGQYGVDILKELAKLGFLNTFEDGNDYGSFKNNTNSTMGLKEIVELAGRAIDLACTLSTSEREEVIDSI

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ETYQRKNCLLTIQGLLDTKGYIGPMGVSVFREDFLAILQLOEHNREILRKWWEGS----KCPKEEHRAKGLMENIQGIVVLICGLIVAFPAMMLELW

VYKAKARYEVFPCDALEEERFIVRSFNERPLKSAQISY---------------------------------------------RSRNS

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-SQIESLKTDSENNMPVE

PARSESELEWEKKDNSSEPE
(Figure 1-5b cont.)

GluR-II

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| YQRNCNITQICGLDTKGYGIQMPVGSDFDFTLAILQLENRELIFIRKWECCQ---KCPKEDHRAIDKLMENMIGGFVVLICGLIVAFMAML-EFLWTL |

**TM4**

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**RSEESLEWEKTNSSEPE**
(Figure 1-5b cont.)

**GluR-IIA**

**KA2**

- MRLCPVIYAFTIIIGFLEGIIALGDDDRNEITVGAIFYENKEKIERSFDQAFREVNNMKFSELRFVTIKRYMPTNDSLFLQQITCELISNGVAAIFGPSSK-ASAD
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- IVAQIANTGPHIEYDLEATRQECENHQMSINVIPSLSLGSRAYFEIKSANYEWFTFLTYETPGLARLDLMQMLNASQDVYKLRNLADYADDYRLKETD
  TVSHICGKEIPIHFGP---EETPRLQYLRFAVSLYPSNEDVSLAVSIKL-FNYPSASICAKAECLLRLEELVRGFLISKETLSSLMDDRDTPLLKEIRD

- ETPHEQRIILDCPEKTLKELLKVSIDFKLPQNRWFNLTHLDTHNSGLRDYIYNEE-----FKANTISVRLKVDANDFPERKKLTLKVDQILGQNMTMLPLILYDAVY
  D---KVSTIIIDANASISHLILRKASELGMTASFYKYILTMDFPILHLDIGVEDSSNILGFSMFNTSHPFYEFVRSLNMSWRENEASTYLGL-PALSAALMFDAVH

- IFASARNVIAAMQPFHPPNRHCGSSPMMLGAFIVNEIMKTISEDDVEPHFKTENMKLDEYGQRIHNFLEIYKPTVN-EPMMVWTPONGIKKPLINLELESAGTTQ
  VVSAVR-EINRSEQIVGKPLACTSANIWHGTSLMNLYRMVEYDGLT-----GRVEFNSKQRTNYLRIKESRQGRELGVYWSN------RTLAMNATTLDINL

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AVRELNRSEQIGVKPLACTSANIWPHGTSMLNYNL------MVEYDGLTGVEFNSSKQGRNTYTLIRILEKSRQGHREIG----------VWYSNRTLAMNATTLDIHEL

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(Figure 1-5b cont.)

**GluR-II (KA2)**

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ILPKGVSVLGSPSSPASASTVSHICGEKIEPHIKVGP---EETPRLQYLRFASVSLPHPNSDEVSASLSRILSKFNSYSPLASLCAK--AECLLRLEELVRGFLISKET

VRVKFHKDDDFRVNWKSIRGERR---VVLDECPNMLVELLLNSTEMFLGLTGQYNHFLTLENLTEEDDDLALLADNEIFVANVTIAARLLNVDPFFYLPYFYGVTQDRDN

LSVRMDDSRDPTPLLKEIRDDKVIITIIDANASISHLIRKASELGMTSAFKYILITMDFPILHLDGIVED------SNILGFSMFTSHFYPEFVRLNMSWRE

IVYESSDPRTLTHDILHIHDALQLFAQSWKNSFFEFPDRMVRPRITCDFAASGGRTWAMRYARLMLKGTSGVNNTHFRSTSLQFLEDGQRTFNIEVYPLDG---IG

NCEASTYLGPALSAALMFDAVHVVVSAR---ELNRSEIGVPLKACTSAN-----WPHGTSLNNYR-----MVEYDGLTGRVEFNKQORNTNYTLRILEKSRQGHRE

IAIWDFRQGITQNLVDVKAQCKKMIYRVATRIGPYYFSYNETARLEN---LTGNALQGYAVDLIAARHVGFEYVFVFPADQQYKGLDKEKTQWNGIGIEINNDA

IGVWNRYNRALTNNATTLLDINLSQTLANKTVITPITPLLMYFRENPQALSGNEEFEGCVEMRELAYFLFRYLRLVEDGLGAFEPN-GSTWGMVGEILNKRA

HMIGCDLTTQARKTAVDFTTVMQQLGVILAYKSHVEKTLDAYLFAPGGEVWIMILSVFVMTFLKIIVARISKMDWENPHPCNRD-PEVLENQWRHNTGWLTVDLAVAAFTIATAREKVIDDFSFPMTLGTISILYRHVMGRKPGYFSFGLDFSPAVWLFMLLAYAVCFLALAARSYPNWPNFPLRARPHERNENQTLGNSLWFPV

---TM1---TM2---

ASIMTAGCDILRSPQVFQRMFEATWWMIFAILTANSTYANLAFLTSSKMGSIANLKLSDAQKVKVGTGYGSTYNLLADNETVYRLAFNLMNDPSAYTKDNLE

GGFMQQGSGEIMPRALTCSRVCVGVWAVTFLLIIISSYTANLAFLTQV8MEVPVESADDLADQTNIEYTIHGASTMTFPQNSRTQYTRQWMNMQSKPSVFKSKE

---TM3---

CDVRKFRNGDVYFMLMENTTLYHREQNCNDRSVGKEFKGHEYIAAVPFQAGEYRNSVLAVKLSERGEYDLKQKKWPNASCSEEPDPDAFTDPMTFEELRGIY

GIAVRILSNR-----AFLLESTMNEYHRRLNCNLTQIGCGLDKTGKYGIGMPLGSFREDTIALQITQENRLEILKRWMEG--GRCPEEHDHAKGLMENIGGIFI

---TM4---

TLYAGILIALFAGITEFLYVQQVALEERLTLFKDAFKKEIRFVLCWINNRRK-------------------------------IVAGTPISSVRTFRRLDKS---

VLICGIIIAYVFAVMEFIFWSTRRAEAVEVSVCQEMLQELHRHAVSCRKTSRSRRRPPGSPSALLSLRAVREMRLSNGBKLYSAAGGDAAGAHSFGQRLDDPGPP

LDRTFKSSRRVIGRSSEMREMAQQGSSSSNNAGRGEKEARV-------------------------------SGARPAAAPTPCTHRVCCQECRRIQALRAGSAGAPRGLGVPAEATFPFRPGPAGPRELAEHE
(Figure 1-5b cont.)

**GluR-IID**

MHFCWISLTLSSLRVQAFYGGNAYEASSGQSIMRLGLITDDATDRIRTFEHASUNITELGVLGETEQVAYGSVQAFALCRIMQSGTVGAVFQPAH-THTSGAHPAEELLLLLVAFASPSCVQLSLRMAILODDQVTCGRGERLALALAREQINGIVIEPAPAR---VDEVDFELQDSDQETYTDTCQILPKGVVSVLGPOSSPASAS

HLLNACSDSDPFIYYPHLSWGS-----NPDFGFLHPSIPEDIANAALYDIVNFESRFSPICYESAFLKILDHLMTRYIKGVIKVMYRDNLNGNYKSVLRRIRKSTVSHNICEKEHIPIKGPEEPTPRLQYLFRASVSLVPNEDVSLAVSRLIKSFSNYPSASLICAKECLLLREELVRGFLIKETLSVRMLDSSDP---TPLLKEIRDD

EDSRIIVVVGSTTGAELLRQAQQVGMNEDTYTIIIGNLNLHTFDDEEYEKSEANITGIRRIFSPDQEEVRDLMLKHLQEQLEGSEPVNSGTFITMEMALTYDARVIAKVTIIIDANASISHLIILRASKELGMTSAYIKYILTTMDFPIHLHDGIVEDBNSNILGVMSFNTSHFYPEPFRSLENMWSREN-CEAASYGPAASAILMFDAVHVVP

ETTENLYPQP-OQMLCNSRHERDNVPDPGRFSTRNMYRSLSEIKEKTITGRIYFEGN-VRKGFTDIEIAQSGTGLKVGTDEGKDFEIFPOQAPVFNDDGSLVNTKSADVRENRSQEIGVPLACTSANWPHGTSLMLNYRMLVYEDG--LTVRVEFNSQGRRTNYTTRILEKSRQG_HREIGVWSNATLAMNATTLDINLSQT---LANK

FIVLSVATKFYASLVESIDTLIGNQQFQGYGVDDLKEILADKGLNFTRDDGDNYDSGNKTNTS3GMLKEIFEGRAVADLILDITSETREEVIDSFIPFMMNLGIA

**TM1**

ILVYKPFQKAPFALFSMDFFSEWLQLGIAYGLSLCFFIIGRLSPIEDWNPYFCIEE-PTEELEGFQTTNINLWFPTIGALLQQGSEIAFKALSTRTISAIWFFTL

ILVRHVHMRKFGYFSFLDFSPPAVWLFMLLAYALCVSLPLAARLSPEWYNPHRPCLRARPENLOYQTLLGNLFPVGGFMQGQGEIMPRALSTRCVSGVWWAFFT

**TM2**

**TM3**

IMVSSYTANLAALFTIENPTSPINSVKDLADNKDDVQYGAKRTGSTRNFFSTSEEPIYIKMNEYLFNAHPELMENNQQGVDKVSKGTYAFLMESTSIEFNTVREC

I11SSYTAANLAALFTVQRMEVFESADDLADQTN-IYEYTIHAGSTMTFFQNSRQYQTQRMWNYMQSKQPSVFKSTEESIA8VIN-SRYAFFLESTMNEYHRRLNC

**TM4**

NLTKVGDPLDCEGKGIAMVENKPNYMPTKALLELQEQGVLAFLNKWNNEVAGVCASAKSDGGDPSLEGDNLSIYVVVLIGISISILCWYFTVKAKNY

NLTIQIGGLDOKGFIIMFLGSPFRDEITAILQLQENNRLEILKWKWEGG----RCPCEDDRKALGMENIKEFIVLIGC11IAFVAVMFIWFRSRAESE

EVPFCDALAEFFRIVIRFSENE---DSE------ENMPVED----------

SGAGAPPRLAVPAEATSPPRPPGPGPRELAEHE
(Figure 1-5b cont.)

GluR-II

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<td><strong>MPAELLLLIVAFASPCQVLSSLRMAAILDDQVCTGQRGELALALAREQINGIEEVPAKAR----VEVDIFELQRDS-----QYETTDTCQPILPKGVVSVLGFSS</strong></td>
<td><strong>RVRKSDNRIVVGVGSETMPFELNQAQVGYINEQYIIIGNLFSDFLDEEYIKSEANITGLRLFPEKMAVKEELIMGLKGYPTDOQERNGSCPITVEMALTDAY</strong></td>
<td><strong>ANNFIILSVPNKPYAQTVETYKOLENQSYEGYGDVLKIELADVKNFTTVNQQDYGSYKSNESTGMLREIMTGADLAI TDITSSREQALDFIPMN</strong></td>
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<td><strong>ANSTTVSHICKEIPKIVGPEETRPQYLRFFASVSLQYLFPSDNLAVSRILKSISSTAGLACAEELRLEERLVGFLISKE</strong></td>
<td><strong>HTVVAFAVREINRSQEGVIVPLACTSANIVPHGTSIMNLYRMVEYDGL</strong></td>
<td><strong>GPPQKRFVQKATFELFTMDFPSSEVWVFLGGSFGLVGLSFFILGLPLSEDNYPYCPIEE--PEELENQFTLGSNWI TTTGALLQQGSEQPKALSTRTVASFWW</strong></td>
<td><strong>AFTLIIISSYTANLAAFLTVQRMEVPHSEADLDADQTN--IEYGTIHAGSTMTTFQNSRYQTYQMWNYPMSQKQSFVFST</strong></td>
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<td><strong>LGIALLYLKVQATFELFMTMDFPSSEVWVFLGGSFGLVGLSFFILGLPLSEDNYPYCPIEE--PEELENQFTLGSNWI TTTGALLQQGSEQPKALSTRTVASFWW</strong></td>
<td><strong>LGISILYRVHMGKGPYSFLDFDPSSAVWLFMLLAYLAVSVLFLAAPLSYETYNPHPCLRARPHILENQTYLGNSWLPFG</strong></td>
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<td><strong>FPTTIISSYTANLAAFLTVQRMEVPHSEADLDADQTN--IEYGTIHAGSTMTTFQNSRYQTYQMWNYPMSQKQSFVFST</strong></td>
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<td><strong>ALRASAAGAPPRGLGVPAETSPRPRPGPAGPRELAEHE</strong></td>
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--- **GluR-II KA2** ---
Table 1-1. Glutamate receptor genes in *Drosophila* genome. 36 different glutamate receptors genes are found in NCBI database (National Center for Biotechnology Information).

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<tr>
<th>Subunit name</th>
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<td>GluR-IIC (GluRIII)</td>
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<td>33275</td>
<td>CG 2902</td>
<td>NP 730940</td>
<td>40665</td>
<td>CG15732</td>
<td>NP 572795</td>
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<td>GluR-IIID (=kaiRIA)</td>
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<td>44483</td>
<td>CG 30379</td>
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<td>CG 6185</td>
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<td>GluR I</td>
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Table 1-1. Glutamate receptor genes in *Drosophila* genome. 36 different glutamate receptors genes are found in NCBI database (National Center for Biotechnology Information).
Chapter Two

Furthering Pharmacological and Physiological Assessment of the Glutamatergic Receptors at the *Drosophila* Neuromuscular Junction

INTRODUCTION

The function and pharmacology of glutamate receptors in general are of prime interest since they are the major excitatory receptor type used within the mammalian central nervous system (CNS). There are extensive investigations into the pharmacology, physiology, molecular structure and regulation in expression as well as turnover in mammalian preparations of glutamate receptors (Dingledine et al., 1999; Hollmann and Hainemann, 1994; Lipsky and Goldman, 2003; Mayer, 2005; Pauly et al., 2005; Seeburg, 1993; Thiagarajan et al., 2005). Earlier studies on glutamate receptors at the neuromuscular junction of insects and crustaceans paved the way to dissecting the functional aspects of these receptors within vertebrates (Anderson et al. 1976; Cull-Candy and Parker, 1983; Gration et al., 1981; Jan and Jan, 1976; Patlak et al., 1979; Shinozaki and Ishida, 1981; Shinozaki and Shibuya, 1974). Still today, invertebrate models serve a vital role in describing regulation and developmental aspects of glutamate receptor function (Featherstone et al., 2005; Guerrero et al., 2005; Littleton and Ganetzky, 2000; Logsdon et al., 2006; Pawlu et al., 2004; Rasse et al., 2005). Considering that *Drosophila melanogaster* is a true model organism with a known genome and used for successful rapid induction of mutations in various studies of development and synaptic mechanisms, it is important to understand the pharmacological and physiological function within this model synaptic neuromuscular preparation (Atwood et al., 1993; Betz et al. 1993; Jan and Jan 1976; Kurdyak et al., 1994; Li et al., 2002; Li and Cooper, 2001; Pawlu et al., 2004; Ruffner et al., 1999; Sigrist et al., 2002, 2003; Stewart et al., 1994, 1996).

The molecular sequence of the glutamate receptors on the *Drosophila* skeletal muscle has been described; however, to date there are few detailed investigations on the pharmacological profile of the intact larval *Drosophila* NMJ (Chang and Kidokoro, 1996; Chang et al., 1994; Delgado et al., 1989; Schuster et al., 1991; Zhang et al., 1999). Glutamate receptors are defined as NMDA (N-methyl-D-aspartic acid) and non-NMDA subtypes (i.e. AMPA, kainate and δ) based on their pharmacological sensitivity activating the receptors. The molecular sequences of the glutamate receptors for *Drosophila* NMJs were performed on the body wall muscles in larvae by isolating the skeletal muscle and the body wall (skeletal muscles attach to the cuticle body wall). The total mRNA was extracted from this muscle tissue and probed against an "affymetrix *Drosophila* genechip". A number of *Drosophila* glutamate receptor subunits were found to be present. The sequences isolated were identified by comparison to whole larval DNA known sequences for glutamate receptors. The ones identified
are GluRIIA, GluRIIB, GluRIIC and GluRIID. GluRIIE was later isolated when real-time PCR was used to follow up on the amplified mRNA of the four receptor subtypes in quantification of expression levels. The expression profile of the five glutamate receptor subunits has been conducted in body wall muscle by in situ hybridization for embryos and larvae (Guerrero et al., 2005; Marrus et al., 2004).

With the RT-PCR and the genomic profiling the sequence indicates that there are 5 glutamate receptor subunits (DGluR-IIA, DGluR-IIB, DGluR-IIC, DGluR-IID, DGluR-IIE) expressed in skeletal muscle and all of them are ionotropic receptors. In comparing the molecular sequence of these 5 muscle-expressed glutamate receptors with vertebrates, they showed the most similarity to the vertebrate kainate/AMPA receptor subtypes; therefore, they have been defined as a kainate receptor subtype without pharmacological examination (Betz et al., 1993; Guerrero et al., 2005; Marrus et al., 2004; Qin et al., 2005; Völkner et al., 2000). However, a recent physiological and pharmacological study demonstrates that the responses are primarily of a quisqualate subtype and not a kainate or an AMPA subtype since the muscle is not depolarized by kainate or AMPA at concentrations as high as 1mM (Bhatt and Cooper, 2005). In fact, it appears that kainate might block the glutamate receptors since the excitatory postsynaptic potential (EPSP) is reduced in the presence of kainate (on average by 40% for 1mM and 500μM but about 20% for 100μM). The reduction in the EPSP amplitude occurred without any significant change in resting membrane potential; thus, kainate did not depolarize the muscle as observed for application of glutamate or quisqualate (Bhatt and Cooper, 2005).

In addition, the procedures used to identify the molecular sequence of the glutamate receptors at the NMJ, by homogenizing the skeletal muscle and body wall, did not allow for assessment for potential presynaptic glutamate receptors. In the past procedure, the CNS was removed and thus the cell bodies of the motor neurons were not incorporated in the assay. Considering that there are presynaptic glutamatergic receptors for neurons in the vertebrate CNS (Park et al., 2006) and at crayfish NMJs (Schramm and Dudel, 1997), there may well be pre-synaptic autoreceptors present at the Drosophila NMJ. Crayfish NMJs show many similarities to the larval Drosophila NMJs (Atwood and Cooper, 1995, 1996) which is not so surprising since they are both arthropods. However, mixed results for the glutamatergic autoreceptors were obtained at the crayfish NMJ in that the responses depended on the depolarized state of the nerve terminal for the facilitatory action of exogenously applied glutamate and for the metabotropic agonist t-ACPD (Schramm and Dudel, 1997). In addition, some preparations showed a depression of presynaptic release while yet other preparations exhibited excitation at the same concentration of t-ACPD. Regardless of the response at the crayfish NMJ, there appears to be one or more types of presynaptic, glutamatergic receptors for the crayfish preparation (Schramm and Dudel, 1997). The earlier observation of the rapid reduction in the EPSP by kainate at the Drosophila NMJ is most likely due to one or both of the following factors: (1) antagonist action on the postsynaptic glutamate receptors; (2) reducing the number of presynaptic vesicular fusion events. It is possible both mechanisms might occur. Direct blocking action on postsynaptic receptors is a
possibility for kainate as many known agonists can be antagonists for subtypes within a receptor family (Chang and Weiss, 2002; Eberle et al., 2001; Hruby et al., 2003). This is likely as a part of the extracellular receptor domains for glutamate receptors must be similar in order to bind glutamate. However, at the 3rd instar Drosophila NMJ the binding of kainate does not transduce activation of the glutamate ligand gated ion channels (Bhatt and Cooper, 2005; Dudel et al. 1992; Heckmann and Dudel, 1997). There is also precedence for kainate, a potent agonist for some vertebrate metabotropic receptors, to act presynaptically in reducing evoked transmitter release via autoreceptors on the presynaptic terminals (Chittajallu et al., 1996; Frerking et al., 2001; Kamiya and Ozawa, 1998; McLamon and Quastel, 1988; Park et al., 2006). Kainate is known to bind metabotropic glutamate receptors (Zhang et al., 1999) and have a presynaptic action in altering transmitter release from presynaptic terminals in vertebrates (Kamiya and Ozawa, 2000; Kamiya et al., 2002; Park et al., 2006). In rats, kainate and the kainate receptor agonist, ATPA, act presynaptically in reducing glutamate release from primary sensory neurons (Kerchner et al., 2001). However, in the rat neocortex, kainate receptors have been demonstrated to have both facilitatory as well as inhibitory actions due to dose dependent effects. The mechanism proposed for the biphasic effect is that low doses excite the presynaptic terminal while high doses have an inhibitory effect on the postsynaptic target (Campbell et al., 2007). Thus, depending on the type of preparation and dosage, kainate might excite or inhibit the presynaptic terminal. The evidence suggests a mechanism of action via kainate-sensitive glutamatergic autoreceptors through a metabotropic action within the presynaptic terminals or by direct action on the presynaptic voltage-gated Ca2+ channels (see reviews Lerma, 2003 and Nicoll et al., 2000). In addition, the actions of glutamate at newly hatched Drosophila larvae revealed presynaptic actions in enhancing the occurrences of spontaneous events without any alteration in the amplitude of the quantal currents (Zhang et al., 1999). The mechanisms suggested for this phenomenon are that presynaptic autoreceptors are present which enhances synaptic transmission. In addition, the pharmacological agonist (1S, 3S)-ACPD for mGluR mimicked the effect of glutamate (Zhang et al., 1999). This helps to establish that autoreceptors at this NMJ are likely in the 3rd instar and that there maybe developmental differentiation in presynaptic receptor subtypes to account for the depressing action of kainate in the 3rd instar.

The purpose of this study is to test for these possibilities. I have approached this by accessing the characteristics of single quanta (i.e., peak amplitude) and frequency of spontaneous events with intracellular measures of miniature EPSPs (mEPSPs) from the whole muscle. I also examined further pharmacological profiles of the Drosophila NMJ to build knowledge of this model synaptic preparation.

**METHODS**

Electrophysiology and staging of flies
All the electrophysiology and staging of larvae are routine procedures and have been described previously. The dissection technique was previously reported (Ball et al., 2003; Campos-Ortega and Hartenstein, 1985; Li et al., 2002). The physiological saline is HL3 (in mM): 1.0 CaCl2 2H2O, 20 MgCl2, 70 NaCl, 5 KCl, 10NaHCO3, 5 trehalose, 115 sucrose, 5 BES (N,N-bis[2-hydroxy-ethyl]-2-aminoethanesulfonic acid) and adjusted to a pH of 7.2 (Stewart et al., 1994). All experiments were performed at room temperature (20-21°C). The recording techniques have been previously described for intracellular EPSP measures (Dasari and Cooper, 2004; Sparks et al., 2004; Stewart et al., 1994) and extracellular focal quantal recordings (Cooper et al., 1995a; Harrison and Cooper, 2003). The compound amplitude of the excitatory postsynaptic potentials (EPSP) elicited by Is and Ib motor nerve terminals in segment 3 of muscle m6 were monitored as in Kurdyak et al., (1994) and Ruffner et al., (1999). The identified m6 muscle was used in each preparation. Only preparations with a resting membrane potential of -50mV or greater were used and the HL3 dissection medium was completely replaced by HL3 media containing the pharmacological compounds.

**Quantal analysis**
All quantal analysis of intracellular whole muscle recordings were as described previously (Cooper et al., 1995b). The quantal analysis of measuring spontaneous frequency and characteristics of the quantal shape are standard approaches.

**Pharmacology**
In an earlier report (Bhatt and Cooper, 2005), actions of NMDA, AMPA, L-glutamate, kainate, quisqualic acid, NBQX, AP5, and DNQX, with regard to synaptic transmission and direct effects on the muscle fibers, was examined. This study showed that kainate reduced evoked EPSPs. The same procedures for exchanging the bathing medium with pharmacological agents were used. The compound trans-(±)-1-Amino-1,3-cyclopentanedicarboxylic acid monohydrate (t-ACPD) is a known mGluR agonist (Holscher et al., 1997) and has been shown to have some effect on newly hatched *Drosophila* larvae (Zhang et al. 1999). I examined the actions of ATPA, a known agonist of the kainate receptor in vertebrates, and 1,3-ACPD on synaptic responses. The ATPA, domoic acid and the trans-(±)-1-Amino-1,3-cyclopentanedicarboxylic acid monohydrate (t-ACPD) were obtained from Sigma

**Statistical analysis**
A non-parametric analysis (Wilcoxon rank sum) or a parametric Student’s t-test were used for assigning a significance at a p<0.05 depending on the appropriate experimental design as explained in the RESULTS section.

**Sequence Comparisons**
All of the glutamate receptor amino acid sequences used are annotated in National Center for Biotechnology Information (NCBI) database. There are 36
different genes encoding 36 different kinds of glutamate receptor subunits in *Drosophila melanogaster*. Among them, five subunits consisting of postsynaptic receptors were compared to five different kainate type glutamate receptor subunits in *Homo sapiens*. Both whole protein and extracellular domain sequences of each subunit were aligned to show the similarity and calculate the statistical matches between two species, respectively. The bl2seq in NCBI and Clustal W were used to align two different protein sequences. In Clustal W the default settings were used except ITERATION to alignment and NUMERITER were set to 3.

**RESULTS**

In measuring the evoked EPSPs and spontaneous quantal events a 3rd instar larvae is filleted so that each segment is observed. By stimulating a selected segmental nerve, only one nerve root is recruited which is important since the muscle fibers are electrically coupled between segments. Various identified muscles with the rather simplistic innervation profiles are shown in Figure 2-1A. In these studies, I utilized muscle 6 (m6) because of the well characterized innervation as depicted in Figure 2-1B (Atwood et al., 1993; Kurdyak et al., 1994; Li et al., 2002). The synaptic properties of the Is and Ib motor nerve terminals have been previously described (Atwood et al., 1993; Kurdyak et al., 1994; Li et al., 2002). I obtained similar measures as earlier reported. I did ensure to observe all three sized EPSP events (Ib, Is, and Ib+Is) in order to subsequently monitor the compound Ib+Is EPSP. The resting membrane potential was also monitored and only preparations which did not appear damaged or with a resting membrane potential more negative than -50mV, in normal physiological saline, were used for assessing the effects of various pharmacological agents.

Application of a saline containing kainate (1mM) rapidly reduces the EPSP amplitude and the effect can be partially reversed by extensive exchanging the bathing media back to normal saline (Figure 2-2A). The extent in the reduction of the EPSP amplitude is dose-dependent, with an almost 50% reduction at 1mM (Figure 2-2B). All three concentrations produced a significant reduction in the EPSP amplitudes (n=6 for each concentration; p<0.05 Wilcoxon rank sum non-parametric). There is no significant effect on the resting membrane potential on m6 for any of the three concentrations tested (Figure 2-2C). However when considering sham control experiments of exchanging the bathing medium with saline and providing the same time period in which EPSP measure were made there is run down in the EPSP amplitude by 13.5% (Figure 2-2D). The individual values are shown next to the mean value. The sham procedures produced a significant effect in reducing the EPSP responses (n=6; p<0.05 Wilcoxon rank sum non-parametric). The mean value was not subtracted from experimental conditions with pharmacological agents in the remaining results, but this effect should be considered. Thus, exposure to 500µM and 1mM kainate produced a significant difference from sham control, whereas 100µM did not (p<0.05 Student’s t-test).
In examining the effects of the AMPA and kainate receptor agonist, domoic acid (10μm and 1mM), the postsynaptic receptor sensitivity to spontaneous quantal responses and the frequency of spontaneous events were assessed. It was readily apparent that domoic acid at 1mM rapidly reduced the amplitude of spontaneous mEPSP quantal events (Figure 2-3A) as well as the frequency of occurrences (Figure 2-3B). Domoic acid at 10μM or 1 mM did not produce an effect on the resting membrane potential (Figure 2-3C). Glutamate at 10mM produced similar effects in reducing the mEPSP amplitude and frequency but this was compounded by the rapid depolarization that glutamate produced (p<0.05; Figure 2-3C). The differences in the AMPA and kainate receptor agonist and glutamate are that, domoic acid did not significantly depolarize the muscle; however, the mEPSPs were still affected. The combination of domoic acid (1mM) and glutamate (10mM) produced a reduced depolarization of the muscle and as expected, a reduced mEPSP amplitude and frequency occurred (Figure 2-3B, C). These results suggest domoic acid is an antagonist to the postsynaptic glutamate receptors. The effect on the reduced frequency of the mEPSPs is due to the gradual reduction in the mEPSP amplitude, such that they are not discernable from noise in the baseline and thus are not detected to monitor their frequency.

Similar results were obtained for domoic acid as for kainate. The mEPSP amplitudes and the frequency are reduced (Figure 2-4A, B). Also the resting membrane potential is not significantly altered. In examining subtle effects on single quantal events the shape of the quantal events were examined and significant effects in rise time or decay were measured (data not shown). The amplitude and area (i.e., area under the curve) of the mEPSPs both decreased in a dose-dependent manner for 1mM and 500μM (Figure 2-4C). With exposure to kainate, the smaller mEPSPs appeared to fade into the background noise while the previously larger mEPSPs events were now observed as smaller mEPSPs.

Since the postsynaptic glutamate receptors on the muscle appear to bind kainate, I expected the known agonist of kainate receptors, ATPA, to act similarly, despite knowing that the receptors are not a kainate subtype. ATAP at 1mM did not reduce the evoked EPSPs (Figure 2-5A, B) nor did ATPA have any effect on the resting membrane potential (Figure 2-5C). In fact, some preparations increased in the EPSP amplitude while others decreased, thus producing no net significant effect.

If there are subtle effects of kainate in depressing presynaptic vesicular docking or altering [Ca^{2+}]i dynamics, one would expect that facilitation or the short-term depression (i.e., negative facilitation) index might detect such an effect. A short four pulse train of 20Hz produces the characteristic response known for this preparation, as shown in Figure 2-6A. The initial EPSP is normally used as a relative reference to the 4th EPSP in the train. It has been previously demonstrated that 1mM [Ca^{2+}]o will produce a marked depression in the response train where as 500μM [Ca^{2+}]o produces less depression and sometimes a slight facilitation. Thus, both 1mM and 500μM [Ca^{2+}]o were examined with 1mM and 500μM kainate to unmask any ceiling effect in the depression by 1mM [Ca^{2+}]o exposure. The 1st and 4th EPSPs rapidly attenuate in amplitude upon exposure to 1mM kainate and 0.5mM [Ca^{2+}]o (Figure 2-6B). The EPSPs also
show a large reduction to kainate when bathed with 1mM \([\text{Ca}^{2+}]_o\) (Figure 2-6C). However, no significant differences, by parametric analysis (Student’s \(t\)-test), in the facilitation index are observed among the four conditions for exposure to kainate as determined by the difference in the median FI values. The difference is measured from saline to exposure of kainate within each preparation (n=5; 500μM kainate-1mM \([\text{Ca}^{2+}]_o\); 1mM kainate-1mM \([\text{Ca}^{2+}]_o\); 500μM kainate-500μM \([\text{Ca}^{2+}]_o\); 1mM kainate-500μM \([\text{Ca}^{2+}]_o\); Figure 2-6D). If the effect to exposure of kainate (1mM) with 1mM \([\text{Ca}^{2+}]_o\) is analyzed by non-parametric statistics, then five out five preparations showed an increase in the mean FI \((p<0.05, \text{Wilcoxon Rank Sum})\). However, two preparations showed only small increases of 10 and 12%. Thus, I do not feel confident in stating that there is a significant increase in the mean FI with kainate by non-parametric analysis. On the other hand, box-whisker plots of the FI values before and after exposure to kainate, for each preparation, in the various paradigms revealed a trend again related to the same data set of kainate (1mM) with 1mM \([\text{Ca}^{2+}]_o\) (Figure 2-7). The box-whisker plots represent the median, 25th, 75th, and range. One can observe that the variability in the distributions is larger for the kainate (1mM) with 1mM \([\text{Ca}^{2+}]_o\) as compared to the same preparations before the exposure. In the other paradigms the variability is about the same for saline and kainate exposure. To further examine this variability in FI, the standard deviation for each preparation before and after exposure to kainate was examined. Only the kainate (1mM) with 1mM \([\text{Ca}^{2+}]_o\) showed a consistent trend (five out of five; \(p<0.05, \text{Wilcoxon Rank Sum Test}\); also by Student’s \(t\)-test \(p<0.05\)) with an increase in the variability while exposed to kainate as compared to the saline baseline. The changes in the individual preparations for this paradigm are shown in Figure 2-8.

In order to further investigate possible presynaptic, metabotropic glutamate receptor action on the motor nerve terminals, the common metabotropic glutamate receptor agonist trans-1-aminocyclopentane-1, 3-dicarboxylic acid (t-ACPD) was used. Potentially, an increase or a decrease in the evoked EPSP would be possible at this insect NMJ since at the crayfish NMJ (also an arthropod species) an effect was reported (Schramm and Dudel, 1997). However, no significant alterations on evoked EPSPs amplitudes (Figure 2-9A, B) or on the resting membrane potential of the muscle occurred during exposure to t-ACPD (500μM) at the Drosophila NMJ. In comparison to the sham saline control (Figure 2-2D), the distributions in EPSP amplitudes overlap; thus, the slight reduction in EPSP amplitudes with t-ACPD are not significant.

**Discussion**

In this study I showed, in a concentration-dependant manner, that kainate rapidly reduced the amplitude of evoked EPSPs as well as the spontaneous quantal events. The lack in any response of t-ACPD and ATPA as well as a lack in these compounds mimicking kainate’s action suggest that there is not a presynaptic mechanism of a kainate-like or metabotropic autoreceptors on these motor nerve terminals. However, the kainate receptor agonist, domoic acid, did follow the actions of kainate in reducing the evoked EPSPs. I suggest this action
is due to domoic acid and kainate acting as an antagonist on the postsynaptic quisqualate receptors present on muscle and that no presynaptic contribution can be proposed for the rapid reduced amplitude in the evoked EPSP and quantal responses. The reduced frequency in the spontaneous quantal responses in the presence of kainate or domoic acid appears to be due to the responses decreasing in size to a point that they are not detectable from the noise or that the postsynaptic receptors are fully blocked.

The responses observed at the crayfish NMJ by Schramm and Dudel (1997) for analogous types of NMJs as I used in the Drosophila larval NMJ (Atwood and Cooper, 1995, 1996) must be different in regards to presynaptic autoreceptors. In their study, t-ACPD and glutamate itself produced varied responses in mean quantal content. Some preparations they reported showed an enhancement while others a decrease in evoked postsynaptic events. They concluded that there may be differential regulation in the type and density of the metabotropic presynaptic autoreceptors to account for variation among preparations and effects to t-ACPD and glutamate. There are methodological differences in stimulating the nerve terminals. In my studies, I elicited action potentials within the nerve to result in Ca\textsuperscript{2+} influx and vesicular fusion. However, Schramm and Dudel used depolarization pulses delivered over the terminal, through the focal electrode, to directly open voltage-gated channels. Seal stability with the focal electrode over the NMJ, longer than normal action potential depolarization pulses and tweaking the current pulses to a fine degree in order to balance a percentage of activating voltage dependent ion channels, would appear to induce potentially more experimental variability among preparations than inducing a standard action potential within the nerve for evoking synaptic transmission. The large field effect of a current pulse, to induce presynaptic depolarization, through a macropatch electrode could have an effect on the postsynaptic glutamate receptors such that the receptor interaction with pharmacological agents of even glutamate itself might be altered. To resolve these issues in the crayfish would require selectively stimulating the motor nerve and repeating the study.

As for domoic acid, the results suggest kainate is an antagonist to the postsynaptic glutamate receptors in a dose-dependent manner. It appears kainate can behave as an antagonist but not ATPA for these postsynaptic muscle receptors. In addition, since ATPA did not produce any significant alteration in the evoked EPSP amplitude, there does not appear to be any pre- or postsynaptic action of ATPA. Quisqualate has been suggested to activate ionotropic as well as metabotropic glutamate receptors just as glutamate does itself, which suggests quisqualate is a non-selective glutamate receptor agonist (Schoepp et al., 1990). However, AMPA, which is significantly similar to quisqualate in structure and is a weak kainate receptor agonist, cannot activate nor even bind to homomeric GluR6 kainate receptor (Egebjerg et al., 1991). Such differences may be induced by steric hindrance near the ligand binding domain. In fact, AMPA did not change the EPSP amplitude in my studies, but kainate resulted in reduced amplitude by acting as an antagonist. Therefore, there seems to be other factors besides steric hindrance. It would seem likely that a ligand binding site of kainate
at the Drosophila NMJ would be highly conserved to the vertebrate kainate receptor, since kainate is blocking evoked responses; however, the similarity of extracellular domains are not very high. Recent studies suggested that the properties of ionotropic glutamate receptor subunits can be changed by mutating the amino acid at the second transmembrane segment which forms the lining of the pore region (Bochet et al., 1991; Curutchet et al., 1992; Verdoorn et al., 1991). For the functional muscle-expressed glutamate receptor GluR-IIC, GluR-IID, GluR-IIE and either GluR-IIA or GluR-IIB is essential for function. Therefore, only based on the ligand binding domain tetrameric structure was suggested. The correct subunit stoichiometry is not known. The receptor can be pentameric or higher such as (IIA)(IIB)(IIC)(IID)(IIE) or (IIA)(IIA)(IIC)(IID)(IIE) to be functional. Also, glutamate receptor complexes which contain either GluR-IIA or GluR-IIB have opposite physiological properties by showing slow and fast desensitization, respectively (DiAntonio et al., 1999). Thus, as constituted receptor subunits are put together it is likely that the receptor is functionally changed. There could also be mixed subunit combinations along the nerve terminal among the many synaptic sites, since the terminal is developmental different in age along its length (Atwood and Cooper, 1995; He et al., 2009; Li et al., 2002) and function (Bogdanik et al., 2004; Guerrero et al., 2005; Marrus et al., 2004; Liebl and Featherstone, 2005;). Developmental alteration in kainate receptors are known in vertebrates (Lee et al., 2001). Thus, variability in whole muscle responses to pharmacological agents might be masked, but if older and younger regions of the terminals were monitored with focal macropatch electrodes larger differences in the pharmacological profiles and sensitivities might be observed.

It was proposed that the five subunits consisting of postsynaptic receptors in Drosophila NMJ are somewhat similar in homology to kainate receptors in vertebrates and thus, they were termed kainate-like (Betz et al., 1993; Guerrero et al., 2005; Marrus et al., 2004; Qin et al., 2005; Völkner et al., 2000). In comparing the postsynaptic receptor molecular sequence of Drosophila NMJ with a kainate receptor in Homo sapiens, which contains the highest homology in the vertebrates, each subunit shows a moderate similarity between two species. Extracellular domains of the receptors are not significantly higher in similarity compared to total sequence (Figure 2-10). In a phylogeny tree with 36 glutamate receptor protein sequences in Drosophila melanogaster and human’s NMDA, kainate and AMPA receptors, the five subunits of postsynaptic receptors in Drosophila NMJ appear to be in a different subgroup to even the closest of the vertebrate kainate/AMPA type receptors (Figure 2-11).

There are multiple examples of pre- and/or post-synaptic action of kainate in enhancing or suppressing synaptic transmission in vertebrate neuronal circuits (Grabauskas et al., 2007; Hegarty et al., 2007; Kamiya and Ozawa, 2000; Pinheiro and Mulle, 2008; Rozas et al., 2003; Schoepf, 2002); and it has been reported earlier that kainate blocked synaptic transmission at the Drosophila NMJ (Chang and Kidokoro, 1996; McLarnon and Quastel, 1988). However, the earlier studies with Drosophila did use Schneider’s Drosophila medium which is known to have a depressing action on synaptic transmission itself (Ball et al., 2003; Stewart et al., 1994). Mechanistically I do not know how kainate is blocking
transmission, but it appears likely that kainate acts as an antagonist in this
preparation. Possible kainate could act on the receptor to desensitize it to
 glutamate or in addition act as a non-competitive antagonist. Quisqualate
receptors are known to be prone to desensitization at the crayfish NMJ (Tour et
al., 2000) as well as other glutamate receptors (Sun et al., 2002).

The results with kainate at different concentrations and the two
concentrations of bathing [Ca\textsuperscript{2+}]\textsubscript{o} is interesting since only the one condition of
1mM kainate and 1mM [Ca\textsuperscript{2+}]\textsubscript{o} produced a significant trend in both an increase in
the median FI value as well variability in the responses as compared to baseline
saline exposure (Figures 2-7 and 2-8). As stated earlier the increase in median
values is small in some preparations so I am cautious in bearing too much emphasis
on this result; however, the increase in variability when exposed to
kainate is of interest. If kainate is acting as a competitive antagonist then it is
feasible to speculate with each pulse in the stimulus train the glutamate is
resulting in a greater displacement of the kainate and possibly resulting in more
competitive-competition. As the glutamate is being taken back up in the nerve
terminal by the glutamate transporters, kainate’s action can dominate producing
very variable responses on the amplitudes of the EPSPs throughout the four
pulse train. Considering when saline is exchanged for saline containing kainate,
the 1\textsuperscript{st} EPSP within the train is dampened significantly by the antagonistic action
and during the evoked release of glutamate then the kainate is displaced would
possible explain the slight, but consistent, increase in the median FI values as
well for this experimental paradigm. This also follows for the concept that when
the [Ca\textsuperscript{2+}]\textsubscript{o} is reduced to 0.5mM, the evoked amount of glutamate is not as
pronounced so less displacement of the kainate would occur, which might
explain why there is not as change on FI or on the median as well as the
variability in the FI (Figure 2-7). I am not aware of reports addressing
pharmacological examination of competitive antagonists during pulse trains to
account for fluctuation in facilitation measures to compare my findings. More
studies are warranted to examine this tentative working model in competitive
antagonism during synaptic facilitation/depression to explain the mechanistic
details behind these findings.

Since I have not been able to detect any presynaptic autoreceptor function
at these NMJs, it would be interesting to know if these motor neurons produce
mRNA for any glutamate receptor subtype. There has yet to be investigation of
possible presynaptic actions at other NMJs in the larvae or adult of Drosophila.
The subset of junctions that I examined may not be indicative of all the
Drosophila NMJs, so further investigations are warranted.
REFERENCES

Chapter 1


Schramm, M., Dudel, J., 1997. Metabotropic glutamate autoreceptors on nerve


Chapter 2


nociceptors rapidly switch from high to low Ca2+ permeability. J. Neurosci. 21(13), 4572-4581.


Figure 2-1: The dissected 3rd instar Drosophila larva preparation. (A) The preparation is pinned at the four corners to keep the preparation taut. The ventral abdominal muscle m6 is used in this study. The segmental nerves are stimulated by placing the nerve into a suction electrode and recruiting the Is and Ib motor neurons. (B) The terminals of Ib and Is on m6 and m7 are readily observed after treatment with fluorescently tagged anti-HRP antibody (Scale: 50μm). (C) Elicited excitatory postsynaptic potentials (EPSPs) in m6 are measured with an intracellular recording. Representative individual responses from the Ib and Is motor axons as well as the composite Ib and Is response are shown for a early 3rd instar.
Figure 2-2: The effect of kainate on the evoked EPSP amplitudes. (A) Application of 1mM kainate reduces the EPSP rapidly and the effect can be partially washed out. (B) The dose-response effect in damping the EPSP amplitude. For all concentrations of kainate the EPSP amplitudes are reduced. (C) The dose-response effects on the resting membrane potential for m6. The resting membrane potentials do not appear to be changed. With the maintaining resting membrane potential, reduced EPSP amplitudes are induced by the antagonistic action of kainate. (D) Sham control for exchanging the saline bath and time revealed a 13.5% reduction in the EPSP amplitude. When compared to sham control, both 500μM and 1mM still show significantly reduced EPSPs (around 46% decrease).
Figure 2-3: Effects of domoic acid (agonist for AMPA and kainate receptors) at the NMJ. Domoic acid is a potent kainate receptor agonist. (A) Spontaneous or mEPSPs (miniature EPSPs) are reduced in amplitude upon exposure to 1mM of domoic acid. With washing out the domoic acid, the spontaneous responses gradually come up again. (B) The frequency of mEPSPs is reduced by domoic acid in a dose-dependent manner. Glutamate mimics the effect. (C) Domoic acid did not produce a consistent effect on the resting membrane potential; however glutamate produced a drastic depolarization fitting for agonistic action. Domoic acid attenuates the effect of glutamate on membrane depolarization. So, domoic acid appears to be blocking the glutamate action on the muscle.
Figure 2-4: Effects of kainate (1 and 0.5 mM) on mEPSPs properties. (A) Spontaneous or mEPSPs (miniature EPSPs) are reduced in amplitude upon exposure. (B) The frequency of mEPSPs is reduced by domoic acid in a dose-dependent manner, as well as the (C) amplitude and area of the mEPSPs. Kainate seems to be acting presynaptically as well as postsynaptically. However, the problem of this measuring is that the mEPSPs can be buried within the noise as the amplitude of mEPSPs are reduced.
Figure 2-5: Effects of ATPA (1 mM; a selective agonist of vertebrate GluR5, kainate receptors) on the amplitude of evoked EPSPs (A), (B) as well as the resting membrane potential of the muscle (C). ATPA does not appear to alter the EPSP amplitude or the resting membrane potential.
Figure 2-6: Effects of kainate on short-term depression (i.e., negative facilitation) with varied \([\text{Ca}^{2+}]_o\). (A) A four pulse stimulus train delivered at 20Hz results in an initial large EPSP followed by depressed responses. The procedure for measuring EPSP amplitudes is indicated. Facilitation index (FI) is then calculated from these measures. (B) A typical response for the effect of kainate (KA; 1 mM) on the pulse train in the presence of 0.5 mM \([\text{Ca}^{2+}]_o\). (C) A typical response for the effect of KA (1 mM) on the pulse train in the presence of 1 mM \([\text{Ca}^{2+}]_o\). (D) The difference in facilitation index (FI) is shown for both 1mM and 0.5mM KA with either 1mM or 0.5mM \([\text{Ca}^{2+}]_o\). The differences in the median values are obtained from the difference in FI saline to exposure of KA within each preparation. (n=5 for each paradigm; * p<0.05 Wilcoxon Rank Sum Test for the effect of KA as compared to saline).
Figure 2-7: Effects of kainate (KA) and calcium on FI. Box and whisker plots for FI in various conditions are shown as before and after exposure to kainate, for each of the five preparations in the four experimental conditions. The plots show the median, 25\textsuperscript{th}, 75\textsuperscript{th} percentiles and range. In almost all preparations, median values of FI index seem to be the same before and after exposure to kainate. So, kainate does not appear to have a presynaptic action. With 1mM Ca\textsuperscript{2+} and 1mM kainate, both median and variation of FI index are increased in five out of five preparations after exposure to kainate. However, after changing the [kainate] into 0.5mM, neither median nor variation of facilitation index shows a consistent increase in values. So, kainate seems to be acting as a competitive antagonist.
Figure 2-8: The standard deviation (SD) for each preparation before and during exposure to kainate (1mM) with 1mM [Ca\(^{2+}\)]\textsubscript{o} produced a consistent trend in increasing in the variability among the data sets while exposed to kainate (KA) as compared the saline (S) baseline (p<0.05, Wilcoxon Rank Sum Test; also p<0.5 Student’s t-test).
Figure 2-9: Action of t-ACPD on evoked EPSPs and resting membrane potential. (A) Evoked EPSPs in saline and during exposures to t-ACPD (50 μM). (B) The mean percent change in the amplitude of the evoked EPSPs with exposure to t-ACPD. (C) The mean percent change in the resting membrane potential with exposure to t-ACPD. t-ACPD does not appear to have an action on the postsynaptic muscle or the glutamate receptors.
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Figure 2-10: Sequence comparison of five postsynaptic receptor subunits identified from *Drosophila* muscle mRNA to the five known human kainate receptor subunits. Each two sequences were compared by using a blastP alignment. The top bar is a comparison for the total protein sequence. The bottom bar is a comparison of the putative extracellular domains among the proteins. The gray area is the percentage of identical sequence and the white area is the percent in similarity based on the blastP (NCBI) algorithm. The five human receptor subunits used are: (A) GluR5; (B) GluR6; (C) GluR7; (D) KA1; and (E) KA2.

Figure 2-11: Phylogeny tree comparing all known putative 36 ionotropic glutamate receptor subunits in the *Drosophila* genome with human glutamate receptors (kainate receptor subunits GluR5, GluR6 and KA2; AMPA receptor subunit GluR1; NMDA receptor subunits NR-1 and NR-2). The comparison was performed by ClustalW2 (EMBL-EBI).
Vita

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