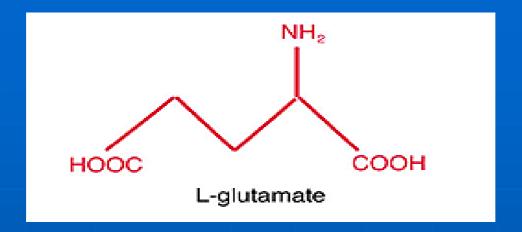
# GLUTAMATE RECEPTORS

SUBMITTED BY

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- Glutamate is the major excitatory neurotransmitter in the mammalian CNS, acting through both ligand gated ion channels (ionotropic receptors) and Gprotein coupled (metabotropic) receptors
- Glutamate receptors play a vital role in the mediation of excitatory synaptic transmission

	receptors	NMDA-receptor family {	NMDAR1 NMDAR2A,
		AMPA-receptor family	GluR1 GluR2 GluR3
Glutamate	Ionotropic	Kainate-receptor family Expression 1985	GluR6
acts on:	ptors	Group I (Phospholipase C activation)	mGlu1A mGlu5A
	Metabotropic receptors	Group II (Adenylyl cyclase inhibition)	mGlu2 mGlu3
	Metabo	Group III (Adenylyl cyclase inhibition)	mGlu4, mGlu8 mGlu6, mGlu6

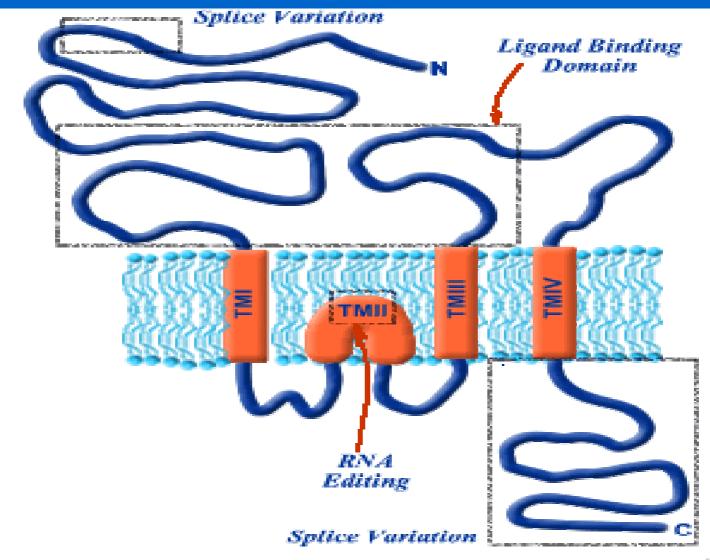
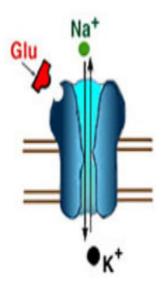


Figure 2. General structure of an ionotropic glutamate receptor subunit. Ionotropic glutamate receptor subunits follow the same basic structural pattern with an extracellular N-terminus and large loop that together form the ligand binding domain. The C-terminus is intracellular and is often the site of splice variation and interaction sites with intracellular proteins.

#### non NMDA receptor



### **NMDA** receptor

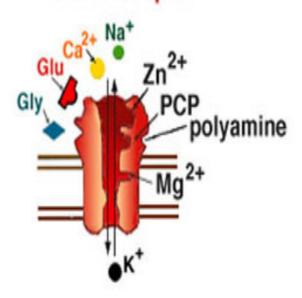


Fig. 6a. Non-NMDA receptors are selectively agonized by kainate, AMPA and quisqualate. The associated ion channels are more perm eable to Na+ and K+ ions tha Ca+2 (from Kandel et al, 1991).

Fig. 6b. NMDA receptors are structurally complex, with separate binding sites for glutamate, glycine MG+2, Zn+2 and polyamines. NMDA-gated channels are more permeable to Ca+2 than Na+ ions (from Kandel et al., 1991).

# NON-NMDA RECEPTORS

- Give immediate response upon binding of glutamate
- More permeable to Na+ & K+ than Ca+ +
- Influx of Na+ causes depolarization of the membrane which is the major factor for activating NMDA type receptors

### NMDA RECEPTORS

Unlike non NMDA receptors, they require a co-agonist
 GLYCINE

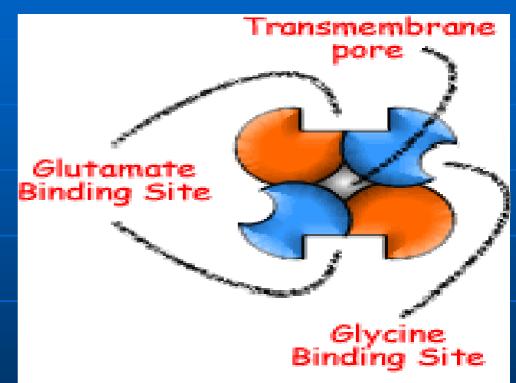
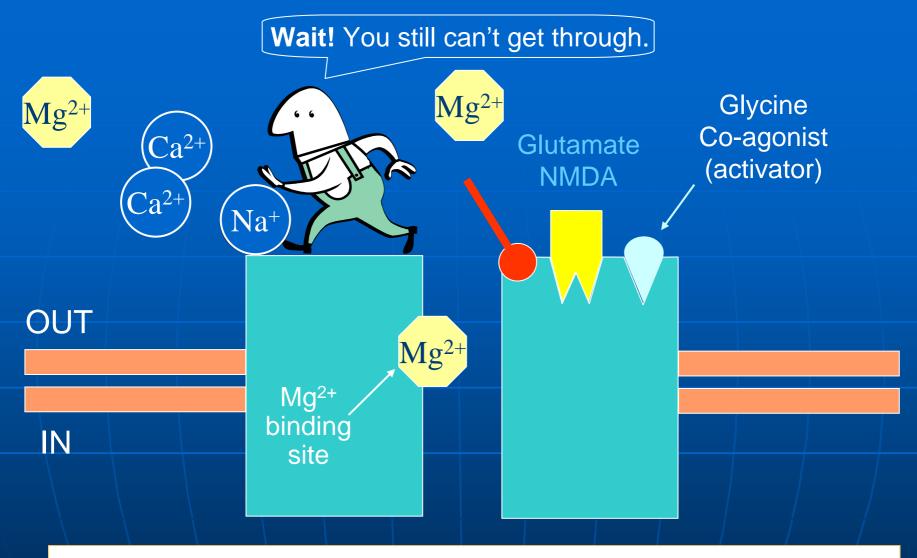


Figure 3. Blnding sites on the NMDA receptor. Note - the stoichiometry of the reeptor complex is schematic only. It is not known as yet whether the complex is a tetramer or a pentamer



Because, as long as the membrane remains polarised, the pore of the channel is blocked by physiological, extracellular concentration of Mg<sup>2+</sup>.

## Metabotropic Glutamate (mGlu) Receptors

 Metabotropic glutamate (mGlu) receptors are G-protein coupled receptors (GPCRs)

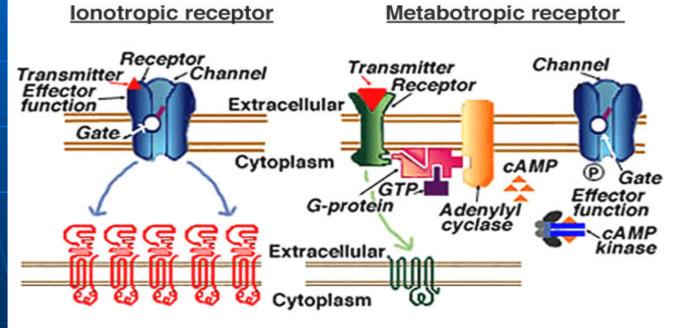


Fig. 5a. Ionotropic receptors and their associated ion channels form one complex (top). Each iGluR is formed from the co-assembly of multiple (4-5) subunits (From Kandel et al., 1991).

Fig. 5b. Metabotropic receptors are coupled to their associated ion channels by a second messenger cascade (top). Each mGluR is composed of one polypeptide, which is coupled to a G-protein (from Kandel et al., 1991).

 They are not co-assembled from multiple subunits, but are one polypeptide with common 7 transmembrane domain

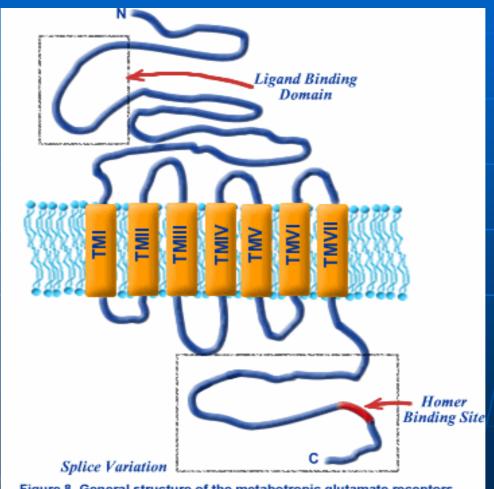


Figure 8. General structure of the metabotropic glutamate receptors. The mGlu receptors are coupled to G-proteins and possess the common 7-transmembrane domain motifly found in these receptor types. However, the N-terminus is much larger than in the adrenergic family of GPCRs and contains the ligand binding domain. The C-terminus generally undergoes extensive splice variation.

# They have been subdivided into three groups, based on

- sequence similarity,
- pharmacology
- intracellular signalling mechanisms

### Metabotropic Glutamate Receptors

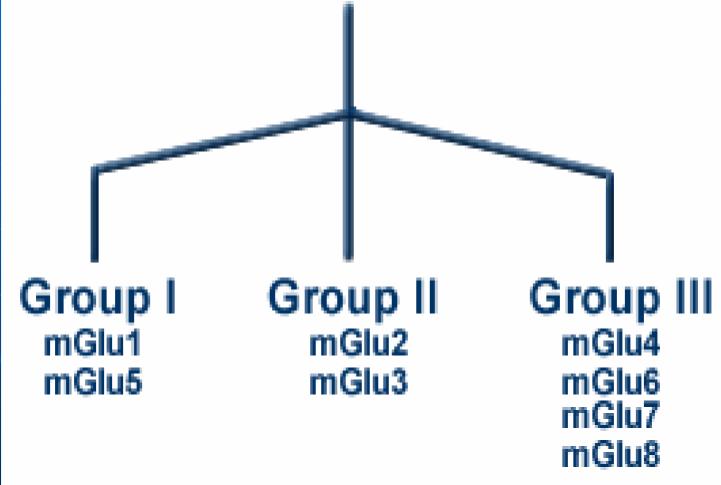


Figure 7. Classification of the metabotropic glutamate receptors.

### Metabotropic glutamate receptor groups (from Pin and Duvoisin, 1995).

group	mGluR	agonist(s)	intracellular pathway
1	mGluR1, mGluR5	quisqualate, ACPD	increase phospholipase C activity, increase cAMP levels, increase protein kinase A activity
II	mGluR2, mGluR3	L-CCG-1, ACPD	decrease cAMP levels
III	mGluR4, mGluR6. mGluR7, mGluR8	L-AP4 (APB)	decrease cAMP or cGMP levels

### REFERENCES

http://www.webvision.med.utah.edu/GLU.ht ml

http://www.bris.ac.uk/Depts/Synaptic/info/glutamate.html

# THANK YOU FOR YOUR ATTENTION