

TRANSMITTER-GATED CHANNELS

Acetylcholine (nicotinic)

Overview: Nicotinic acetylcholine receptors are members of the Cys-loop family of transmitter-gated ion channels that includes the GABA_A, strychnine-sensitive glycine and 5-HT₃ receptors. All nicotinic receptors are formed as pentamers of subunits. Genes (ENSF0000000049) encoding a total of 17 subunits ($\alpha 1$ –10, $\beta 1$ –4, δ , ϵ and γ) have been identified. All subunits are of mammalian origin with the exception of $\alpha 8$ (avian). Each subunit possesses four TM domains. All α subunits possess two tandem cysteine residues near to the site involved in acetylcholine binding, and subunits not named α lack those tandem cysteines. The acetylcholine-binding site is formed by at least three peptide loops on the α subunit (principal component), and three on the adjacent subunit (complementary component). The determination of a high-resolution (2.7 Å) crystal structure of the acetylcholine-binding protein from *Lymanaea stagnalis*, a structural homologue of the extracellular binding domain of a nicotinic receptor pentamer, has revealed the binding site in detail (reviewed by Karlin, 2002, Smit *et al.*, 2003). Nicotinic receptors at the somatic neuromuscular junction of adult animals have the stoichiometry ($\alpha 1$)₂ $\beta 1\epsilon\delta$, whereas an extrajunctional ($\alpha 1$)₂ $\beta 1\gamma\delta$ receptor predominates in embryonic and denervated skeletal muscle. Other nicotinic receptors are assembled as combinations of $\alpha(2$ –6) and $\beta(2$ –4) subunits. For $\alpha 2$, $\alpha 3$, $\alpha 4$ and $\beta 2$ and $\beta 4$ subunits, pairwise combinations of α and β (e.g. $\alpha 3\beta 4$, $\alpha 2\beta 4$) are sufficient to form a functional receptor *in vitro*, but more complex isoforms may exist *in vivo*. $\alpha 5$ and $\beta 3$ subunits lack function when expressed alone or pairwise, but participate in the formation of functional hetero-oligomeric receptors (e.g. $\alpha 4\alpha 5\alpha 6\beta 2$, $\alpha 6\beta 2\beta 3$) when coexpressed with at least two other subunits. The $\alpha 6$ subunit can form a functional receptor when coexpressed with $\beta 4$ *in vitro*, but more efficient expression ensues from incorporation of a third partner, such as $\beta 3$. The $\alpha 7$, $\alpha 8$ and $\alpha 9$ subunits form functional homo-oligomers, and can also combine with a second α subunit to constitute a hetero-oligomeric assembly (e.g. avian $\alpha 7\alpha 8$). For functional expression of the $\alpha 10$ subunit, coassembly with $\alpha 9$ is necessary. The latter, along with the $\alpha 10$ subunit, appears to be largely confined to cochlear and vestibular hair cells.

The nicotinic receptor subcommittee of NC-IUPHAR has recommended a nomenclature and classification scheme for nicotinic acetylcholine (nACh) receptors based on the subunit composition of known, naturally and/or heterologously expressed nACh receptor subtypes (Lukas *et al.*, 1999). Headings for this table reflect abbreviations designating nACh receptor subtypes based on the predominant α subunit contained in that receptor subtype. An asterisk following the indicated α subunit denotes that other subunits are known to, or may, assemble with the indicated α subunit to form the designated nACh receptor subtype(s). Where subunit stoichiometries within a specific nACh receptor subtype are known, numbers of a particular subunit larger than 1 are indicated by a subscript following the subunit (enclosed in parentheses).

Nomenclature	$\alpha 1^*$	$\alpha 2^*$	$\alpha 3^*$
Previous names	Muscle type, muscle	—	Autonomic, ganglionic
Potency order of commonly used agonists	($\alpha 1$) ₂ $\beta 1\gamma\delta$ (embryonic): sub > epi > DMPP > ACh > carb ~ sux > nico ~ cyt > cho ($\alpha 1$) ₂ $\beta 1\epsilon\delta$ (adult): sux > cyt = DMPP > nic	$\alpha 2\beta 2$: epi > ana-a > DMPP > nic = cyt > ACh $\alpha 2\beta 4$: epi > DMPP = nic = cyt [†] > ACh	$\alpha 3\beta 2$: epi > DMPP = cyt > nic > ACh $\alpha 3\beta 4$: epi > ana-a > DMPP > cyt [†] = nic > ACh
Selective antagonists	α -Bungarotoxin, α -conotoxin GI, α -conotoxin MI, pancuronium	—	$\alpha 3\beta 2$: α -conotoxin MII (also blocks $\alpha 6\beta 2^*$), α -conotoxin-GIC $\alpha 3\beta 4$: α -conotoxin AuIB
Commonly used antagonists	($\alpha 1$) ₂ $\beta 1\gamma\delta$: Bgt > pan > (+)-Tc (high-affinity $\alpha 1/\delta$ -binding site, low-affinity α/γ site) $\alpha 1$ ($\beta 1\epsilon\delta$): Bgt > pan > (+)-Tc	$\alpha 2\beta 2$: DH β E (K_B = 0.9 μ M), (+)-Tc (K_B = 1.4 μ M) $\alpha 2\beta 4$: DH β E (K_B = 3.6 μ M), (+)-Tc (K_B = 4.2 μ M)	$\alpha 3\beta 2$: DH β E (K_B = 1.6 μ M), (+)-Tc (K_B = 2.4 μ M) $\alpha 3\beta 4$: DH β E (K_B = 19 μ M), (+)-Tc (K_B = 2.2 μ M)
Channel blockers	Gallamine	—	Mecamylamine, hexamethonium
Probes	[³ H]/[¹²⁵ I]- α -bungarotoxin	[³ H]/[¹²⁵ I]-epibatidine (hz2 β 4, 42 pM; rz2 β 2, 10 pM; rz2 β 4, 87 pM), [³ H]-cytisine	[³ H]/[¹²⁵ I]-epibatidine (hz3 β 2, 7 pM; hz3 β 4, 230 pM; rz3 β 2, 14 pM; rz3 β 4, 300 pM), [³ H]-cytisine
Functional characteristics	$\alpha 1$ ($\beta 1\gamma\delta$): P_{Ca}/P_{Na} = 0.16–0.2, P_f = 2.1%; $\alpha 1$ ($\beta 1\epsilon\delta$): P_{Ca}/P_{Na} = 0.65–1.38, P_f = 4.1–4.2%	$\alpha 2\beta 2$: P_{Ca}/P_{Na} ~ 1.5	$\alpha 3\beta 2$: P_{Ca}/P_{Na} = 1.5; $\alpha 3\beta 4$: P_{Ca}/P_{Na} = 0.78–1.1, P_f = 2.7–4.6%

Nomenclature	$\alpha 4^*$	$\alpha 6^*$	$\alpha 7^*$
Previous names	Neuronal, α -bungarotoxin-insensitive	—	Neuronal, α -bungarotoxin-sensitive
Selective agonists	$\alpha 4\beta 2$: TC-2559 (Chen <i>et al.</i> , 2003), TC-2403 (RJR-2403, Papke <i>et al.</i> , 2000),	—	AR-R1779 (Mullen <i>et al.</i> , 2000), PSAB-OFB (Broad <i>et al.</i> , 2002), PNU-282987 (Bodnar <i>et al.</i> , 2005)
Potency order of commonly used agonists	$\alpha 4\beta 2$: epi > nic > cyt > ACh > DMPP = sub > carb > cho > sux $\alpha 4\beta 4$: epi > cyt > nic > DMPP > ACh	rx6h β 4: ACh > cyt > nic > DMPP cx6h β 4: epi > cyt > nic > ACh [†] $\alpha 6/\alpha 3\beta 2\beta 3$ chimera: α -conotoxin PIA	($\alpha 7$) _s : ana-a > epi > DMAC > OH-GTS-21 = DMPP [†] > cyt [†] > nic [†] = GTS-21 > ACh > cho
Selective antagonists	—	$\alpha 6\beta 2^*$: α -conotoxin MII (also blocks $\alpha 3\beta 2$) cx6h β 4: mec, (+)-Tc, hex rx6h β 4: (+)-Tc	($\alpha 7$) _s : α -bungarotoxin, methyllycaconitine, α -conotoxin ImI
Commonly used antagonists	$\alpha 4\beta 2$: DH β E (K_B = 0.1 μ M), (+)-Tc (K_B = 3.2 μ M) $\alpha 4\beta 4$: DH β E (K_B = 0.01 μ M), (+)-Tc (K_B = 0.2 μ M)	—	($\alpha 7$) _s : Bgt > MLA > (+)-Tc [†] > atr > DH β E
Channel blockers	—	Mecamylamine, hexamethonium	—
Probes	[³ H]/[¹²⁵ I]-epibatidine (hz4 β 2, 10–33 pM; hz4 β 4, 187 pM; rz4 β 2, 30 pM, rz4 β 4, 85 pM), [³ H]-cytisine, [³ H]-nicotine	[³ H]-epibatidine (native chick cx6 β 4*, 35 pM)	[³ H]/[¹²⁵ I]- α -bungarotoxin ((hz7) _s , 700–800 pM), [³ H]-methyllycaconitine (native rz7*, 1.9 nM)
Functional characteristics	$\alpha 4\beta 2$: P_{Ca}/P_{Na} = 1.65, P_f = 2.6–2.9%; $\alpha 4\beta 4$: P_f = 1.5–3.0%	—	P_{Ca}/P_{Na} = 6.6–20, P_f = 8.8–11.4%

PSAB-OFB also activates 5-HT₃ receptors

Nomenclature	α8* (avian)	α9*	α10*
Previous names	Neuronal, α-bungarotoxin-sensitive	—	—
Potency order of commonly used agonists	(α8) ₅ : cyt ~ nic ≥ ACh > DMPP	(α9) ₅ : cho > ACh > sub > car	ACh
Selective antagonists	—	(α9) ₅ : α-bungarotoxin, strychnine, nicotine, muscarine	α10α9: α-bungarotoxin, strychnine, nicotine, muscarine
Commonly used antagonists	(α8) ₅ : Bgt > atr ≥ (+)-Tc ≥ str	(α9) ₅ : Bgt > MLA > str ~ tropisetron > (+)-Tc > bic ≥ atr ~ epi > mec > DHβE > cyt > nic > mus	α10α9: Bgt > tropisetron = str > (+)-Tc > bic = atr > nic > mus
Channel blockers	—	—	—
Probes	[³ H]/[¹²⁵ I]-α-bungarotoxin	[³ H]/[¹²⁵ I]-α-bungarotoxin	—
Functional characteristics	—	α9: P _{Ca} /P _{Na} = 9; α9α10: P _{Ca} /P _{Na} = 9	—

A firm consensus has yet to emerge concerning the pharmacological profiles at different nACh receptor subtypes. There are differences in profiles for a given receptor subtype across species. Moreover, measures of agonist potencies and efficacies, or antagonist affinities, are confounded by differences in experimental design across studies (oocyte or mammalian cell heterologous expression systems or natural expression; test agonist concentrations; competitive/noncompetitive modes of antagonism; electrophysiological, ion flux or calcium ion mobilization measurements; etc.). Therefore, provisional and incomplete information about pharmacological rank order potency profiles (no efficacy data) is provided in the table based largely on data from studies of heterologously expressed, human nACh receptors. The dagger (†) as superscript designates ligands whose rank order placement differs across species and/or experimental design.

Abbreviations: **ABT-594**, (R)-5-(2-azetidylmethoxy)-2-chloropyridine; **ACh**, acetylcholine; **ana-a**, anatoxin-a; **AR-R17779**, (–)-spiro[1-azabicyclo[2.2.2]octane-3,5'-oxazolidin-2'-one; **atr**, atropine; **Bgt**, α-bungarotoxin; **bic**, bicuculline; **car**, carbamylcholine; **cho**, choline; **cyt**, cytosine; **DHβE**, dihydro-β-erythroidine; **DMAC**, 3-(4)-dimethylaminocinnamylidene anabaseine; **DMPP**, 1,1-dimethyl-4-phenylpiperazinium; **epi**, epibatidine; **GTS-21**, 3-(2,4)-dimethoxybenzylidene anabaseine (DMXB); **hex**, hexamethonium; **mec**, mecamlamine; **MLA**, methyllycaconitine; **mus**, muscarine; **nic**, nicotine; **OH-GTS-21**, 3-(4-hydroxy, 2-methoxy)benzylidene anabaseine; **pan**, pancuronium; **PNU-282987**, N-[(3R)-1-azabicyclo[2.2.2]oct-3-yl]-4-chlorobenzamide hydrochloride; **PSAB-OPF**, (R)-(–)-5'-phenylspiro[1-azabicyclo[2.2.2]octane-3,2'-(3'H)furo[2,3-b]pyridine; **str**, strychnine; **sub**, suberyldicholine; **sux**, succinylcholine; **TC-2403**, (E)-N-methyl-4-(3-pyridinyl)-3-butene-1-amine; **TC-2559**, also known as (RJR-2403), (E)-N-methyl-4-[3-(5-ethoxypyridinyl)]-3-buten-1-amine; **(+)-Tc**, (+)-tubocurarine

Further Reading:

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GABA_A (γ-aminobutyric acid)

Overview: The GABA_A receptor is a transmitter-gated ion channel of the Cys-loop family that includes the nicotinic acetylcholine, 5-HT₃ and strychnine-sensitive glycine receptors. The receptor exists as a pentamer of 4Tm subunits that form an intrinsic anion channel. Sequences of six α, three β, three γ one δ, three ρ, one ε, one π and one θ GABA_A receptor subunits (ENSF0000000053) have been reported in mammals (Barnard, 2000; Korpi *et al.*, 2002). The π-subunit is restricted to reproductive tissue. Alternatively, spliced versions of α4- and α6- (both not functional), α5-, β2-, β3- and γ2-subunits exist (see Barnard, 2000). In addition, three ρ-subunits (ρ1–3) function as either homo- or hetero-oligomeric assemblies (Bormann & Feigenspan, 2000; Zhang *et al.*, 2001). Although receptors formed from ρ-subunits have sometimes been termed GABA_C receptors (Zhang, 2001), they represent a subpopulation of GABA_A receptor, classed as the GABA_{A0r} subtype, under NC-IUPHAR proposals (Barnard *et al.*, 1998). Many GABA_A receptor subtypes contain α-, β- and γ-subunits with the likely stoichiometry 2α.2β.1γ (Korpi *et al.*, 2002; Fritschy & Brünig, 2003). It is thought that the majority of GABA_A receptors harbour a single type of α- and β-subunit variant. The α1β2γ2 hetero-oligomer constitutes the largest population of GABA_A receptors in the CNS, followed by the α2β3γ2 and α3β3γ2 isoforms. Receptors that incorporate the α4- α5- or α6-subunit, or the β1-, γ1-, γ3-, δ-, ε- and θ-subunits, are less numerous, but they may nonetheless serve important functions. For example, extrasynaptically located receptors that contain α6- and δ-subunits in cerebellar granule cells, or an α4- and δ-subunit in dentate gyrus granule cells and thalamic neurones, mediate a non-desensitising tonic current that is important for neuronal excitability in response to ambient concentrations of GABA (see Mody & Pearce, 2004; Semyanov *et al.*, 2004; Farrant & Nusser, 2005). The α- and β-subunits contribute to the GABA binding site and both the α- and γ-subunits are required for the benzodiazepine (BZ) site. The particular α- and γ-subunit isoforms exhibit marked effects on recognition and/or efficacy at the BZ site. Thus, receptors incorporating either α4- or α6-subunits are not recognised by 'classical' benzodiazepines, such as flunitrazepam. It is beyond the scope of this supplement to discuss the pharmacology of individual GABA receptor isoforms in detail; such information can be gleaned in the reviews by Barnard *et al.* (1998), Frolund *et al.* (2002), Korpi *et al.* (2002), Krogsgaard-Larsen *et al.* (2002) and Johnston (2005). Agents that discriminate between α-subunit isoforms are noted in the table and additional agents that demonstrate selectivity between receptor isoforms are indicated in the text below.

The classification of GABA_A receptors has been addressed by NC-IUPHAR (Barnard *et al.*, 1998). The proposed scheme utilises subunit structure and receptor function as the basis for classification. In view of the fact that a benzodiazepine (BZ) binding site is not unique to the GABA_A receptor, and that certain receptor isoforms (i.e. those incorporating α4- or α6-subunits) are insensitive to classical benzodiazepines, it is recommended that the term 'GABA_A/benzodiazepine receptor complex' should no longer be used and be replaced by 'GABA_A receptor'. The term benzodiazepine receptor itself is contentious because receptors should generally be named to reflect their endogenous ligand and many discriminatory ligands acting at this site are generally not benzodiazepines (e.g. zolpidem, an imidazopyridine). Here, the term 'BZ site of the GABA_A receptor' is adopted as one of the two alternatives proposed by NC-IUPHAR.

Nomenclature	GABA_A
Ensembl Gene family ID	ENSF0000000053
Selective agonists (GABA site)	Muscimol, isoguvacine, THIP, piperidine-4-sulphonic acid (low efficacy at α4 and α6 subunits), isonipecotic acid (α4 and α6 subunit selective <i>via</i> relatively high efficacy)
Selective antagonists (GABA site)	Bicuculline, gabazine
Selective agonists (BZ site)	Diazepam (not α4- or α6-subunits), flunitrazepam (not α4- or α6-subunits), zolpidem, zaleplon and indiplon (α1 subunit selective <i>via</i> high affinity), L838417 (α2, α3 and α5 subunit selective as a partial agonist <i>versus</i> antagonist at α1-subunit-containing receptors), Ro154513 (selective for α4- and α6-subunit-containing receptors as an agonist <i>versus</i> inverse agonist at α1-, α2-, α3- and α5-subunit-containing receptors), TP003 (selective for α3-subunit-containing receptors as a high efficacy partial agonist <i>versus</i> essentially antagonist activity at α1- α2- and α5-subunit-containing receptors), TPA023 (selective for α2- and α3-subunit-containing receptors as a low efficacy partial agonist <i>versus</i> essentially antagonist activity at α1- and α5-subunit-containing receptors)
Selective antagonists (BZ site)	Flumazenil (low affinity for α4- or α6-subunits), ZK93426, L838417 (α1 subunit selective <i>via</i> antagonist activity <i>versus</i> partial agonist at α2-, α3- and α5-subunit subunit containing receptors)
Inverse agonists (BZ site)	DMCM, Ro194603, L655708 (α5 selective <i>via</i> high affinity), RY024 (α5 selective <i>via</i> high affinity)
Endogenous allosteric modulators	5α-pregnan-3α-ol-20-one (potentiation), tetrahydrodeoxycorticosterone (potentiation), Zn ²⁺ (potent inhibition of receptors formed from binary combinations of α and β subunit, incorporation of a γ subunit reduces inhibitory potency, Krishek <i>et al.</i> , 1998), extracellular protons (subunit dependent activity, Krishek <i>et al.</i> , 1996)
Channel blockers	Picrotoxin, TBPS
Probes	
GABA site	[³ H]-muscimol, [³ H]-gabazine
BDZ site	[³ H]-Flunitrazepam (not α4- or α6-subunit), [³ H]-zolpidem (α1-subunit selective), [³ H]-L655708 (α5-subunit selective), [³ H]-RY80 (α5-subunit selective), [³ H]-Ro154513 [selectively labels α4- and α6-subunit-containing receptors in the presence of a saturating concentration of a 'classical' benzodiazepine (e.g. diazepam)], [³ H]-CGS8216, [¹¹ C]-flumazenil (low affinity for α4- or α6-subunits), [¹⁸ F]-fluoroethylflumazenil
Anion channel	[³⁵ S]-TBPS

The potency and efficacy of many GABA agonists varies between receptor GABA_A receptor isoforms (Frolund *et al.*, 2002; Krogsgaard-Larsen *et al.*, 2002). For example, THIP is a partial agonist at receptors with the subunit composition α4β3γ2, but elicits currents in excess of those evoked by GABA at the α4β3δ receptor where GABA itself is a low efficacy agonist (Brown *et al.*, 2002; Bianchi & MacDonald, 2003). Recent data suggest that the presence of the γ subunit within the heterotrimeric complex impairs responsiveness to agonists (Störustovu & Ebert, 2005). The GABA_A receptor contains distinct allosteric sites that bind barbiturates and endogenous (e.g. 5α-pregnan-3α-ol-20-one) and synthetic (e.g. alphaxalone) neuroactive steroids in a diastereo- or enantio-selective manner (see Belelli & Lambert 2005). Picrotoxinin and TBPS act at an allosteric site within the chloride channel pore to negatively regulate channel activity; negative allosteric regulation by γ-butyrolactone derivatives also involves the picrotoxinin site, whereas positive allosteric regulation by such compounds is proposed to occur at a distinct locus. Many intravenous (e.g. etomidate, propofol) and volatile (e.g. halothane, isoflurane) anaesthetics and alcohols also exert a regulatory influence upon GABA_A receptor activity. Specific amino-acid residues within GABA_A receptor α- and β-subunits that influence allosteric regulation by anaesthetic and nonanaesthetic compounds have been identified (see Belelli *et al.*, 1999; Krazowski *et al.*, 2000; Thompson & Wafford, 2001; Hemmings *et al.*, 2005). An array of natural products including flavonoid and terpenoid compounds exert varied actions at GABA_A receptors (reviewed in detail by Johnston, 2005).

In addition to the agents listed in the table, modulators of GABA_A receptor activity that exhibit subunit dependent activity include: salicylidene salicylhydrazide (negative allosteric modulator selective for β1- *versus* β2-, or β3-subunit-containing receptors (Thompson *et al.*, 2004)); loreclezole, etomidate, tracazolate and mefenamic acid (positive allosteric modulators with selectivity for β2/β3- over β1-subunit-containing receptors, see Korpi *et al.*, 2002); tracazolate (intrinsic efficacy, i.e. potentiation, or inhibition, is dependent upon the identity of the γ1-3-, δ-, or ε-subunit co-assembled with α1- and β1-subunits (Thompson *et al.*, 2002)); amiloride (selective blockade of receptors containing an α6-subunit (Fisher, 2002)); frusemide (selective blockade of receptors containing an α6-subunit coassembled with β2/β3-, but not β1-subunit (see Korpi *et al.*, 2002); La³⁺ (potentiates responses mediated by α1β3γ2L receptors, weakly inhibits α6β3γ2L receptors, and strongly blocks α6β3δ and α4β3δ receptors (Saxena *et al.*, 1997; Brown *et al.*, 2002)); ethanol (selectively potentiates responses mediated by α4β3δ and α6β3δ receptors *versus* receptors in which β2 replaces δ (e.g. γ replaces δ (Wallner *et al.*, 2003, but see also Borghese *et al.*, 2005)). It should be noted that the apparent selectivity of some positive allosteric modulators (e.g. neurosteroids such as 5α-pregnan-3α-ol-20-one for δ-subunit-containing receptors (e.g. α1β3δ) may be a consequence of the unusually low efficacy of GABA at this receptor isoform (Bianchi *et al.*, 2003).

A subpopulation of retinal GABA receptors (activated by *trans*-4-aminocrotonic acid) assembled from ρ subunits is bicuculline-insensitive and gates Cl⁻ channels that are insensitive to barbiturates and benzodiazepines and selectively blocked by TPMPA. Isoguvacine and piperidine-4-sulphonic acid do not activate GABA_A

receptors assembled from ρ subunits, and THIP is a moderately potent antagonist. Receptors formed from ρ subunits have often been found to be insensitive to neuroactive steroids, but relatively high concentrations of such compounds can modulate the activity of the $\rho 1$ subunit in a stereoselective manner, 5α -pregnanes potentiating, and 5β -pregnanes inhibiting, responses elicited by low concentrations of GABA (Morris & Amin, 2004). Although these receptors have sometimes been termed GABA_C receptors (see Zhang *et al.*, 2001), this appellation is not endorsed by NC-IUPHAR and they are currently viewed as a sub-class of GABA_A receptor. This position is strengthened by the observation that single amino-acid mutations can impart some typical features of GABA_A receptor pharmacology upon the GABA_{A α 6} subtype (Belelli *et al.*, 1999; Walters *et al.*, 2000).

Abbreviations: **CGS8216**, 2-phenylpyrazolo[4,3-c]quinolin-3(5)-one; **DMCM**, methyl-6,7-dimethoxy-4-ethyl- β -carboline-3-carboxylate; **Gabazie**, also known as **SR95531**; **L655708**, ethyl(s)-(11,12,13,13a-tetrahydro-7-methoxy-9-oxo)-imidazo[1,5-a]pyrrolo[2,1-c][1,4]benzodiazepine-1-carboxylate; **L838417**, 7-*tert*-butyl-3-(2,5-difluoro-phenyl)-6-(2-methyl-2*H*-[1,2,4]triazol-3-ylmethoxy)-[1,2,4]triazolo[4,3-b]pyridazine; **Ro154513**, ethyl-8-azido-5,6-dihydro-5-methyl-6-oxo-4*H*-imidazo[1,5-a][1,4]benzodiazepine-3-carboxylate; **Ro194603**, imidazo[1,5-a][1,4]thienodiazepinone; **RY024**, *tert*-butyl-8-ethynyl-5,6-dihydro-5-methyl-6-oxo-4*H*-imidazo[1,5-*z*][1,4]benzodiazepine-3-carboxylate; **RY80**, ethyl-8-acetylene-5,6-dihydro-5-methyl-6-oxo-4*H*-imidazo[1,5-a][1,4]benzodiazepine-3-carboxylate; **SR95531**, 2-(3'-carboxy-2'-propyl)-3-amino-6-*p*-methoxyphenylpyridazinium bromide; **TBPS**, *tert*-butylbicyclophosphorothionate; **THIP**, also known as gaboxadol; **TP003**, 4,2'-Difluoro-5'-[8-fluoro-7-(1-hydroxy-1-methylethyl)imidazo[1,2-*a*]pyridine-3-yl]biphenyl-2-carbonitrile; **TPA023**, 7-(1,1-dimethylethyl)-6-(2-ethyl-2*H*-1,2,4-triazol-3-ylmethoxy)-3-(2-fluorophenyl)-1,2,4-triazolo[4,3-*b*]pyridazine; **TPMPA**, (1,2,5,6-tetrahydropyridine-4-yl)methylphosphinic acid; **ZK93423**, 6-benzyloxy-4-methoxy-methyl- β -carboline-3-carboxylate ethyl ester; **ZK93426**, 5-isopropyl-4-methyl- β -carboline-3-carboxylate ethyl ester

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Glutamate (ionotropic)

Overview: The ionotropic glutamate receptors comprise members of the NMDA (*N*-methyl-D-aspartate), AMPA (α -amino-3-hydroxy-5-methyl-4-isoxazole propionic acid) and kainate receptor classes, named originally according to their preferred, synthetic, agonist (see Dingledine *et al.* (1999) for a comprehensive review). Receptor heterogeneity within each class arises from the homo-oligomeric, or hetero-oligomeric, assembly of distinct subunits into cation-selective tetramers. All glutamate receptor subunits have the membrane topology of an extracellular N-terminus, three transmembrane domains (TM1, TM3 and TM4), a channel lining re-entrant 'p-loop' (MD2) located between TM1 and TM3 that enters and exits the membrane at its cytoplasmic surface, and an intracellular C-terminus (see Madden, 2002). It is beyond the scope of this supplement to discuss the pharmacology of individual ionotropic glutamate receptor isoforms in detail; such information can be gleaned in the reviews by Dingledine *et al.* (1999), Yamakura & Shimoji (1999), Jane *et al.* (2000), Huettner (2003) and Cull-Candy & Leszkiewicz (2004). Agents that discriminate between subunit isoforms are, where appropriate, noted in the tables and additional compounds that distinguish between receptor isoforms are indicated in the text below.

The classification of glutamate receptors has been addressed by NC-IUPHAR (Lodge & Dingledine, 2000). The proposed scheme, which recommends a revised nomenclature for ionotropic glutamate receptor subunits, is adopted here. Commonly used alternative appellations are indicated in parenthesis.

NMDA receptors: NMDA receptors assemble as heteromers that may be drawn from GLU_{N1} (NMDA-R1, NR1, GluR ξ 1), GLU_{N2A} (NMDA-R2A, NR2A, GluR ϵ 1), GLU_{N2B} (NMDA-R2B, NR2B, GluR ϵ 2), GLU_{N2C} (NMDA-R2C, NR2C, GluR ϵ 3), GLU_{N2D} (NMDA-R2D, NR2D, GluR ϵ 4), GLU_{N3A} (NMDA-R3A) and GLU_{N3B} (NMDA-R3B) subunits. Alternative splicing can generate eight isoforms of GLU_{N1} with differing pharmacological properties. Various splice variants of GLU_{N2B,2C,2D} and GLU_{N3A} have also been reported. Activation of NMDA receptors requires the binding of two agonists, glutamate to the S1 and S2 regions of the GLU_{N2} subunit and glycine to S1 and S2 regions of the GLU_{N1} subunit (Erreger *et al.*, 2004). The minimal requirement for efficient functional expression of NMDA receptors *in vitro* is a di-heteromeric assembly of GLU_{N1} and at least one GLU_{N2} subunit variant, as a dimer of dimers arrangement (Madden, 2002; Furukawa *et al.*, 2005). However, more complex tri-heteromeric assemblies, incorporating multiple subtypes of GLU_{N2} subunit, or GLU_{N3} subunits, can be generated *in vitro* and occur *in vivo*. The NMDA receptor channel commonly has a high relative permeability to Ca²⁺ and is blocked, in a voltage-dependent manner, by Mg²⁺ at resting potential.

Nomenclature	NMDA
Ensembl Gene family ID	ENSF00000000436
Selective agonists (glutamate site)	NMDA, aspartate, D,L(tetrazol-5-yl)glycine, homoquinolinic acid (partial agonist)
Selective antagonists (glutamate site)	D-AP5, CGS19755, CGP37849, LY233053, D-CCPene (GLU _{N2A} = GLU _{N2B} > GLU _{N2C} = GLU _{N2D}), PPDA (GLU _{N2C} = GLU _{N2D} > GLU _{N2B} = GLU _{N2A} , Feng <i>et al.</i> , 2004), PEAQX (GLU _{N2A} > GLU _{N2C} > GLU _{N2D} > GLU _{N2B} , Auberson <i>et al.</i> , 2002; Feng <i>et al.</i> , 2004), conantokin-G (GLU _{N2B} > GLU _{N2D} = GLU _{N2C} = GLU _{N2A})
Selective agonists (glycine site)	Glycine, D-serine, (+)-HA966 (partial agonist)
Selective antagonists (glycine site)	5,7-Dichlorokynurenate, L689560, L701324, GV196771A
Channel blockers	Mg ²⁺ , dizocilpine, ketamine, phencyclidine, memantine, amantidine
Probes	
Glutamate site	[³ H]-CPP, [³ H]-CGS19755, [³ H]-CGP39653
Glycine site	[³ H]-Glycine, [³ H]-L689560, [³ H]-MDL105519
Cation channel	[³ H]-Dizocilpine

In addition to the glutamate and glycine binding sites documented in the table, physiologically important inhibitory modulatory sites exist for Mg²⁺, Zn²⁺, and protons (see Dingledine *et al.*, 1999; Yamakura & Shimoji, 1999; Cull-Candy & Leszkiewicz, 2004). The receptor is also allosterically modulated, in both positive and negative directions, by endogenous neuroactive steroids in a subunit-dependent manner. For example, pregnenolone sulfate potentiates di-heteromeric assemblies of GLU_{N1}/GLU_{N2A} and GLU_{N1}/GLU_{N2B} subunits, but inhibits receptors assembled as GLU_{N1}/GLU_{N2C}, or GLU_{N1}/GLU_{N2D}, heteromers (Malayev *et al.*, 2002). Tonic proton blockade of NMDA receptor function is alleviated by polyamines and the inclusion of exon 5 within GLU_{N1} subunit splice variants, whereas the non-competitive antagonist ifenprodil increases the fraction of receptors blocked by protons at ambient concentration. Inclusion of exon 5 also abolishes potentiation by polyamines and inhibition by Zn²⁺. Receptors assembled from GLU_{N1} and GLU_{N2C} subunits are unusually insensitive to proton blockade. Ifenprodil, its analogue CP101606, haloperidol, felbamate and Ro84304 discriminate between recombinant NMDA receptors assembled from GLU_{N1} and either GLU_{N2A}, or GLU_{N2B}, subunits by acting as selective, non-competitive, antagonists of hetero-oligomers incorporating GLU_{N2B}. LY233536 is a competitive antagonist that also displays selectivity for GLU_{N2B} over GLU_{N2A} subunit-containing receptors. Similarly, CGP61594 is a photoaffinity label that interacts selectively with receptors incorporating GLU_{N2B} *versus* GLU_{N2A}, GLU_{N2D} and, to a lesser extent, GLU_{N2C} subunits. Conversely, the voltage-independent component of NMDA receptor inhibition by Zn²⁺ is most pronounced for receptors that contain the GLU_{N2A} *versus* GLU_{N2B} subunit. In addition to influencing the pharmacological profile of the NMDA receptor, the identity of the GLU_{N2} subunit co-assembled with GLU_{N1} is an important determinant of biophysical properties that include sensitivity to block by Mg²⁺, single-channel conductance and channel deactivation time (Cull-Candy & Leszkiewicz, 2004). Incorporation of the GLU_{N3A} subunit into tri-heteromers containing GLU_{N1} and GLU_{N2} subunits is associated with decreased single channel conductance, reduced permeability to Ca²⁺ and decreased susceptibility to block by Mg²⁺. Reduced permeability to Ca²⁺ has also been observed following the inclusion of GLU_{N3B} in tri-heteromers. The expression of GLU_{N3A}, or GLU_{N3B}, with GLU_{N1} alone is reported to form a cation channel with unique properties that include activation by glycine (but not NMDA), lack of permeation by Ca²⁺ and resistance to blockade by Mg²⁺ and NMDA receptor antagonists (Chatterton *et al.*, 2002).

AMPA and Kainate receptors: AMPA receptors assemble as homomers, or heteromers, that may be drawn from GLU_{A1} (GluR1, GluRA, GluR-A, GluR-K1), GLU_{A2} (GluR2, GluRB, GluR-B, GluR-K2), GLU_{A3} (GluR3, GluRC, GluR-C, GluR-K3), or GLU_{A4} (GluR4, GluRD, GluR-D) subunits. Homotetramers formed from GLU_{A2} subunits express relatively poorly due to their retention within the endoplasmic reticulum (see Bredt & Nicoll, 2003). Functional kainate receptors can be expressed as homomers of GLU_{K5} (GluR5, GluR-5, EAA3), GLU_{K6} (GluR6, GluR-6, EAA4), or GLU_{K7} (GluR7, GluR-7, EAA5) subunits. GLU_{K5-7} subunits are also capable of assembling into heterotetramers (see Lerma, 2003). Two additional kainate receptor subunits, GLU_{K1} (KA1, KA-1, EAA1) and GLU_{K2} (KA2, KA-2, EAA2), when expressed individually, form high affinity binding sites for kainate, but lack function (see Huettner, 2003). GLU_{K1} and GLU_{K2} can form heteromers when co-expressed with GLU_{K5-7} subunits (Lerma, 2003). RNA encoding the GLU_{A2} subunit undergoes extensive RNA editing in which the codon encoding a p-loop glutamine residue (Q) is converted to one encoding arginine (R). This Q/R site strongly influences the biophysical properties of the receptor. Recombinant AMPA receptors lacking RNA edited GLU_{A2} subunits are: (1) permeable to Ca²⁺; (2) blocked by intracellular polyamines at depolarized potentials causing inward rectification; (3) blocked by extracellular argitoxin and Joro spider toxins and (4) demonstrate higher channel conductances than receptors containing the edited form of GLU_{A2} (Seeburg & Hartner, 2003). GLU_{K5} and GLU_{K6}, but not other kainate receptor subunits, are similarly edited and broadly similar functional characteristics apply to kainate receptors lacking either an RNA edited GLU_{K5}, or GLU_{K6}, subunit (Lerma, 2003). Native AMPA and kainate receptors displaying differential channel conductances, Ca²⁺ permeabilities and sensitivity to block by intracellular polyamines have been identified. GLU_{A1-4} can exist as two variants generated by alternative splicing (termed 'flip' and 'flop') that differ in their desensitization kinetics and their desensitization in the presence of cyclothiazide. Splice variants of GLU_{K5-7} also exist, but their functional significance is unknown (Lerma, 2003).

Nomenclature	AMPA	Kainate
Ensembl Gene family ID	ENSF00000000118	ENSF00000000118
Selective agonists	AMPA, (S)-5-fluorowillardiine	ATPA, (S)-5-iodowillardiine, (2S,4R)-4-methyl glutamate (SYM2081), LY339434, domoic acid (except homomeric GLU _{K7}), kainite
Selective antagonists	NBQX, ATPO, LY293558, GYKI53655/LY300168 (active isomer GYKI53784/LY303070) (non-competitive)	UBP296 (More <i>et al.</i> , 2004), LY294486, LY382884, NS3763 (non-competitive, Christensen <i>et al.</i> , 2004)
Channel blockers	Intracellular polyamines, extracellular argitoxin, extracellular Joro toxin, (all subtype selective)	Intracellular polyamines (subtype selective)
Probes	[³ H]-AMPA, [³ H]-CNQX	[³ H]-Kainate, [³ H]-(2S,4R)-4-methyl glutamate

All AMPA receptors are additionally activated by kainate (and domoate) with relatively low potency ($EC_{50} \sim 100 \mu M$). AMPA receptor activity is potentiated by several classes of agent that are not tabulated above including: pyrrolidones (piracetam, aniracetam); benzothiazides (cyclothiazide); benzylpiperidines (CX-516 (BDP-12), CX-546) and biarylpropylsulfonamides (LY392098, LY404187 and LY503430) (O'Neill *et al.*, 2004). Activation of kainate receptors by AMPA shows subunit dependency (e.g. heteromers containing GLU_{K6} and GLU_{K2} subunits are sensitive; homomers assembled from the GLU_{K6} subunit, or GLU_{K7} subunit, are insensitive). Quinoxalinediones such as CNQX and NBQX show limited selectivity between AMPA and kainate receptors. LY293558 also has kainate (GLU_{K5}) receptor activity. ATPO is a potent competitive antagonist of AMPA receptors, has a weaker antagonist action at kainate receptors comprising GLU_{K5} subunits, but is devoid of activity at kainate receptors formed from GLU_{K6} or GLU_{K6}/GLU_{K2} subunits. The pharmacological activity of ATPO resides with the (S)-enantiomer. ATPA, UBP296, LY294486, LY339434, LY382884 and (S)-5-iodowillardiine interact selectively with kainate receptors containing a GLU_{K5} subunit. (2S,4R)-4-methyl glutamate (SYM2081) is equipotent in activating (and desensitizing) GLU_{K5} and GLU_{K6} receptor isoforms and, *via* the induction of desensitization at low concentrations, has been used as a functional antagonist of kainate receptors. Both (2S,4R)-4-methyl glutamate and LY339434 have agonist activity at NMDA receptors. (2S,4R)-4-methyl glutamate is also an inhibitor of the glutamate transporters EAAT1 and EAAT2.

Abbreviations: AMPA, (RS)- α -amino-3-hydroxy-5-methyl-4-isoxazole propionic acid; APTA, (RS)-2-amino-3-(3-hydroxy-5-tert-butylisoxazol-4-yl)propionic acid; ATPO, (RS)-2-amino-3-(3-[5-tert-butyl-3-(phosphonomethoxy)-4-isoxazolyl]propionic acid; CGP37849, (RS)-(E)-2-amino-4-methylphosphono-3-pentanoic acid; CGP39653, (RS)-(E)-2-amino-4-propylphosphono-3-pentanoic acid; CGSI9755, 4-phosphonomethyl-2-piperidinecarboxylic acid; CNQX, 6-cyano-7-nitroquinoxaline-2,3-dione; CP101606, (1S,2S)-1-(4-hydroxyphenyl)-2-(4-hydroxy-4-phenylpiperidino)-1-propanol; CPP, (±)-2-carboxypiperazine-4-yl)propyl-1-phosphonic acid; CX-516, 1-(quinoxalin-6-ylcarbonyl)piperidine; CX-546, 1-(1,4-benzodioxan-6-ylcarbonyl)piperidine; D-AP5, D(2)-2-amino-5-phosphonopentanoate; D-CCPene, 3-(2-carboxypiperazine-4-yl)-propenyl-1-phosphonic acid; GV196771A, E-4,6-dichloro-3-(2-oxo-1-phenyl-pyrrolidin-3-ylidenemethyl)-1H-indole-2-carboxylic acid; GYKI53655, 1-(4-aminophenyl)-4-methyl-7,8-methylenedioxy-5H-(3N-methylcarbamate)-2,3-benzodiazepine, also known as LY300168; GYKI53784, (–)-1-(4-aminophenyl)-4-methyl-7,8-methylenedioxy-4,5-dihydro-3-methylcarbamoyl-2,3-benzodiazepine, also known as LY303070; HA966, 3-amino-1-hydroxypyrrolid-2-one; L689560, *trans*-2-carboxy-5,7-dichloro-4-phenylaminocarbonylamino-1,2,3,4-tetrahydroquinoline; L701324, 7-chloro-4-hydroxy-3-(3-phenoxy)phenyl-2(H)quinolone; LY233053, *cis*(1)-4-[(2H-tetrazol-5-yl)methyl]piperidine-2-carboxylic acid; LY233536, (RS)-6-(1H-tetrazol-5-ylmethyl)decahydroisoquinoline-3-carboxylic acid; LY293558, 3S,4zR,6R,8zR-6-[2-(1(2H)-tetrazol-5-yl)ethyl]-decahydroisoquinoline-3-carboxylate; LY294486, (3SR,4zRS,6SR,8RS)-6-[(1H-tetrazol-5-yl)methyl]oxy)methyl]-1,2,3,4z,5,6,7,8,8z-decahydroisoquinoline-3-carboxylic acid; LY339434, (2S,4R,6E)-2-amino-4-carboxy-7-(2-naphthyl)hept-6-enoic acid; LY382884, (3S, 4aR, 6S, 8aR)-6-[(4-carboxyphenyl)methyl-1,2,3,4,4a,5,6,7,8,8a-decahydro isoquinoline-3-carboxylic acid; LY392098, propane-2-sulfonic acid [2-(4-thiophen-3-yl-phenyl)-propyl]-amide; LY404187, Propane-2-sulfonic acid [2-(4'-cyano-biphenyl-4-yl)-propyl]-amide; LY503430, (R)-4'-[1-fluoro-1-methyl-2-(propane-2-sulfonylamino)-ethyl]-biphenyl-4-carboxylic acid methylamide; MDL105519, (E)-3-(2-phenyl-2-carboxyethyl)-4,6-dichloro-1H-indole-2-carboxylic acid; NBQX, 6-nitro-7-sulfamoyl-benz(f)quinoxaline-2,3-dione; NS3763, 5-carboxyl-2,4-di-benzamido-benzoic acid; PEAQX, 5-phosphonomethyl-1,4-dihydroquinoxaline-2,3-dione, also known as NVP-AAM077; PPDA, (2S*,3R*)-1-(phenanthrene-2-carbonyl)piperazine-2,3-dicarboxylic acid; Ro8-4304, 4-3-[4-(4-fluoro-phenyl)-3,6-dihydro-2H-pyridin-1-yl]-2-hydroxy-propoxy-benzamide; UBP296, (RS)-3-2-carboxybenzyl)willardiine

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Glycine receptors

Overview: The inhibitory glycine receptor (provisional nomenclature adopted here classifies glycine receptor isoforms by their α subunit) is a member of the Cys-loop superfamily of transmitter-gated ion channels that includes the GABA_A, nicotinic acetylcholine and 5-HT₃ receptors. Structurally and functionally, the glycine receptor is most closely related to the GABA_A receptor. The receptor is expressed either as a homopentamer of α subunits, or a complex now thought to harbour 2 α and 3 β subunits (Grudzinska *et al.*, 2005), that contain an intrinsic Cl[−] channel. Four differentially expressed isoforms of the α subunit ($\alpha 1$ – $\alpha 4$) and one variant of the β subunit ($\beta 1$) have been identified by genomic and cDNA cloning. Further diversity originates from alternative splicing of the primary gene transcripts for $\alpha 1$ ($\alpha 1^{\text{INS}}$ and $\alpha 1^{\text{del}}$), $\alpha 2$ ($\alpha 2A$ and $\alpha 2B$) and $\alpha 3$ ($\alpha 3S$ and $\alpha 3L$) subunits and by RNA editing of the $\alpha 3$ subunit (Meier *et al.*, 2005). In particular, the $\alpha 2B$ subunit has a 2–4-fold higher sensitivity to glycine, β -alanine and taurine. Predominantly, the mature form of the receptor contains $\alpha 1$ (or $\alpha 3$) and β subunits while the immature form is mostly composed of only $\alpha 2$ subunits. RNA transcripts encoding the $\alpha 4$ subunit have not been detected in adult humans. The $\alpha 4$ subunit may be a pseudogene in man and is not tabulated here. The N-terminal domain of the α subunit contains both the agonist and strychnine-binding sites that consist of several discontinuous regions of amino acids. Inclusion of the β subunit in the pentameric glycine receptor reduces single channel conductance and alters pharmacology. It also anchors the receptor, *via* an amphipathic sequence within the intracellular loop region, to gephyrin, a cytoskeletal attachment protein, that binds to tubulin and thus clusters and anchors hetero-oligomeric receptors to the synapse (see Kneussel and Betz, 2000; Moss and Smart, 2001). G-protein $\beta\gamma$ subunits enhance the open state probability of native and recombinant glycine receptors most probably *via* a direct association (Yevenes *et al.*, 2003).

Nomenclature	$\alpha 1$	$\alpha 2$	$\alpha 3$
Ensembl ID	ENSG00000145888	ENSG00000101958	ENSG00000145451
Selective agonists (potency order)	Glycine > β -alanine > taurine	Glycine > β -alanine > taurine	Glycine > β -alanine > taurine
Selective antagonists and modulators with subunit selectivity	Strychnine, PMBA, picrotoxin (+ β weakens block), ginkgolide B (IC ₅₀ = 0.6 μ M + β = 0.18 μ M), pregnenolone sulphate (K _i = 1.9 μ M; + β = 2.7 μ M), tropisetron (K _i = 84 μ M; + β = 44 μ M), colchicine (IC ₅₀ = 324 μ M), Zn ²⁺ (IC ₅₀ = 15 μ M)	Strychnine, PMBA, picrotoxin (+ β weakens block), ginkgolide B (IC ₅₀ = 3.7 μ M + β = 0.14 μ M), pregnenolone sulphate (K _i = 5.5 μ M; + β = 10.1 μ M), tropisetron (K _i = 13 μ M; + β = 5.4 μ M), colchicine (IC ₅₀ = 64 μ M), DCKA (IC ₅₀ = 188 μ M), Zn ²⁺ (IC ₅₀ = 360 μ M)	Strychnine, picrotoxin (+ β weakens block), ginkgolide B (IC ₅₀ = 1.8 μ M + β = 0.55 μ M), α EMBTl (+ β converts block to potentiation), Zn ²⁺ (IC ₅₀ = 150 μ M)
Selective potentiators	α EMBTl	—	(α EMBTl reduces $\alpha 3$ -mediated responses)
Channel blockers (IC ₅₀)	Cyanotriphenylborate (1.3 μ M + β = 2.8 μ M)	Cyanotriphenylborate (\gg 20 μ M; + β = 7.5 μ M)	—
Probes	[³ H]-strychnine	[³ H]-strychnine	[³ H]-strychnine
Functional characteristics	γ = 86 pS (main state) (+ β = 44 pS)	γ = 111 pS (main state) (+ β = 54 pS)	γ = 105 pS (main state) (+ β = 48)

Data in the table refer to homo-oligomeric assemblies of the α subunit, significant changes introduced by coexpression of the $\beta 1$ subunit (ENSG00000109738) are indicated in parenthesis. Not all glycine receptor ligands are listed within the table, but those that may be useful in distinguishing between glycine receptor isoforms are indicated. Pregnenolone sulphate, tropisetron and colchicine, for example, although not selective antagonists of glycine receptors, are included for this purpose. Strychnine is a potent and selective competitive glycine receptor antagonist with affinities in the range of 5–15 nM. RU5135 demonstrates comparable potency, but additionally blocks GABA_A receptors. Both the endocannabinoids, anandamide and 2-arachidonylglycerol, block neuronal glycine receptors at physiological concentrations (Lozavaya *et al.*, 2005). Several analogues of muscimol and piperidine act as agonists and antagonists of both glycine and GABA_A receptors. Picrotoxin acts as an allosteric inhibitor with strong selectivity towards homomeric receptors composed of α subunits and its components, picrotoxinin and picrotin, have similar inhibitory potencies (reviewed by Lynch, 2004). In addition to the compounds listed in the table, numerous agents act as allosteric regulators of glycine receptors (comprehensively reviewed by Laube *et al.*, 2002; Lynch, 2004). Zn²⁺ acts through distinct binding sites of high- and low affinity to allosterically enhance channel function at low (< 10 μ M) concentrations and inhibits responses at higher concentrations in a subunit selective manner (Miller *et al.*, 2005). The effect of Zn²⁺ is somewhat mimicked by Ni²⁺. Elevation of intracellular Ca²⁺ produces fast potentiation of glycine receptor-mediated responses. Dideoxyforskolin (4 μ M) and tamoxifen (0.2–5 μ M) both potentiate responses to low glycine concentrations (15 μ M), but act as inhibitors at higher glycine concentrations (100 μ M). Additional modulatory agents that enhance glycine receptor function include inhalational, and several intravenous general anaesthetics (e.g. minaxolone, propofol and pentobarbitone) and certain neurosteroids. Ethanol and higher order *n*-alcohols also act allosterically to enhance glycine receptor function. Solvents inhaled as drugs of abuse (e.g. toluene, 1-1-1-trichloroethane) may act at sites that overlap with those recognising alcohols and volatile anaesthetics to produce potentiation of glycine receptor function. The function of glycine receptors formed as homomeric complexes of $\alpha 1$ or $\alpha 2$ subunits, or hetero-oligomers of $\alpha 1/\beta$ or $\alpha 2/\beta$ subunits, is differentially affected by the 5-HT₃ receptor antagonist tropisetron (ICS 205–930), which may evoke potentiation or inhibition depending upon the subunit composition of the receptor and the concentrations of the modulator and glycine employed. Additional tropines, including atropine, modulate glycine receptor activity.

Abbreviations: α EMBTl, α -ethyl, α -methyl- γ -thiobutylolactone; DCKA, dichlorokynurenine acid; PMBA, 3-[2'-phosphonomethyl[1,1'-biphenyl]-3-yl]alanine; RU5135, 3 α -hydroxy-16-imino-5 β -17-azaandrostan-11-one

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5-Hydroxytryptamine₃

Overview: The 5-HT₃ receptor (nomenclature as agreed by NC-IUPHAR Subcommittee on 5-hydroxytryptamine (serotonin) receptors (Hoyer *et al.*, 1994)) is a transmitter-gated ion channel of the Cys-loop family that includes the nicotinic acetylcholine, GABA_A and strychnine-sensitive glycine receptors. The receptor exists as a pentamer of 4TM subunits that form an intrinsic cation-selective channel. Three 5-HT₃ receptor subunits (5-HT_{3A}, 5-HT_{3B} and 5-HT_{3C}) have been cloned, but only homo-oligomeric assemblies of 5-HT_{3A} and hetero-oligomeric assemblies of 5-HT_{3A} and 5-HT_{3B} subunits have been characterised in detail. Putative *HTR3D* and *HTR3E* genes have also been described (Niesler *et al.*, 2003) but there is presently no evidence that they encode *bone fide* 5-HT₃ receptor subunits that are functional. The hetero-oligomeric receptor has recently been reported to contain two copies of the 5-HT_{3A} subunit and three copies of the 5-HT_{3B} subunit in the order B-B-A-B-A (Barrera *et al.*, 2005). The 5-HT_{3B} subunit imparts distinctive biophysical properties upon hetero-oligomeric (5-HT_{3A}/5-HT_{3B}) *versus* homo-oligomeric (5-HT_{3A}) recombinant receptors (Davies *et al.*, 1999; Dubin *et al.*, 1999; Hanna *et al.*, 2000; Kelley *et al.*, 2003; Stewart *et al.*, 2003; Peters *et al.*, 2005), but generally has little effect upon the apparent affinity of agonists, or the affinity of antagonists (Brady *et al.*, 2001; but see Dubin *et al.*, 1999). However, homo- and hetero-oligomeric 5-HT₃ receptors differ in their allosteric regulation by some general anaesthetic agents (Solt *et al.*, 2005). The diversity of 5-HT₃ receptors is increased by alternative splicing of the 5-HT_{3A} subunit. To date, inclusion of the 5-HT_{3A} subunit appears imperative for 5-HT₃ receptor function.

Nomenclature	5-HT ₃
Former names	M
Ensembl ID	5-HT _{3A} ENSG00000166736, 5-HT _{3B} ENSG00000149305
Selective agonists (pEC ₅₀)	2-Methyl-5-HT (5.3–5.5), 3-chlorophenyl-biguanide (5.4–5.7)
Selective antagonists (pIC ₅₀)	Granisetron (9.5), ondansetron (9.5), tropisetron (9.2)
Channel blockers	Diltiazem, TMB-8, picrotoxin [+ 5-HT _{3B} potency reduced, Das & Dillon, 2003]
Probes	[³ H]-ramosetron (0.15 nM), [³ H]-granisetron (1.2 nM), [³ H]-(S)-zacopride (2.0 nM), [³ H]-GR65630 (2.6 nM), [³ H]-LY278584 (3 nM)
Functional characteristics	γ = 0.4–0.8 ps [+ 5-HT _{3B} , γ = 16 ps]; inwardly rectifying current [+ 5-HT _{3B} , rectification reduced]; relative permeability to divalent cations reduced by coexpression of the 5-HT _{3B} subunit

Quantitative data in the table refer to homo-oligomeric assemblies of the human 5-HT_{3A} subunit, or the receptor native to human tissues. Significant changes introduced by coexpression of the 5-HT_{3B} subunit are indicated in parenthesis. Human (Belelli *et al.*, 1995; Miyaki *et al.*, 1995), rat (Isenberg *et al.*, 1993), mouse (Maricq *et al.*, 1991), guinea-pig (Lankiewicz *et al.*, 1998) and ferret (Mochizuki *et al.*, 2000) orthologues of the 5-HT_{3A} receptor subunit have been cloned that exhibit intraspecies variations in receptor pharmacology. Notably, most ligands display significantly reduced affinities at the guinea-pig 5-HT₃ receptor in comparison with other species. In addition to the agents listed in the table, native and recombinant 5-HT₃ receptors are subject to allosteric modulation by extracellular divalent cations, alcohols, several general anaesthetics and 5-hydroxy and halide-substituted indoles (see reviews by Parker *et al.*, 1996; Peters *et al.*, 1997 and Lovinger, 1999).

Abbreviations: GR65630, 3-(5-methyl-1*H*-imidazol-4-yl)-1-(1-methyl-1*H*-indol-3-yl)-1-propanone; LY278584, 1-methyl-*N*-(8-methyl-8-azabicyclo[3.2.1]oct-3-yl)-1*H*-indazole-3-carboxamide; TMB-8, 8-(diethylamino)octyl-3,4,5-trimethoxybenzoate

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P2X

Overview: P2X receptors (nomenclature as agreed by NC-IUPHAR Subcommittee on P2X Receptors, Khakh *et al.*, 2001) are putative trimeric (Jiang *et al.*, 2003, Nicke *et al.*, 1998) transmitter-gated channels, conducting Na⁺, K⁺ and Ca²⁺, with two putative TM domains, where the endogenous ligand is ATP. The relationship of many of the cloned receptors to endogenously expressed receptors is not yet established. The Nomenclature Subcommittee has recommended that for P2X receptors, structural criteria should be the initial criteria for nomenclature where possible. Functional P2X receptors exist as polymeric transmitter-gated channels; the native receptors may occur as homopolymers (e.g. P2X₁ in smooth muscle) or heteropolymers (e.g. P2X₂:P2X₃ in the nodose ganglion). P2X₇ receptors have been shown to form functional homopolymers which, in turn, activate pores permeable to low molecular weight solutes (Donnelly-Roberts *et al.*, 2004).

Nomenclature	P2X ₁	P2X ₂	P2X ₃	P2X ₄
Ensembl ID	ENSG00000108405	ENSG00000177026	ENSG00000109991	ENSG00000135124
Selective agonists	L-βγ-meATP, αβ-meATP	—	αβ-meATP	—
Selective antagonists	TNP-ATP (pIC ₅₀ 8.9, Virginio <i>et al.</i> , 1998), Ip ₅ I (pIC ₅₀ 8.5), NF023 (pIC ₅₀ 6.7); NF449 (pIC ₅₀ 6.3, Kassack <i>et al.</i> , 2004)	—	TNP-ATP (pIC ₅₀ 8.9, Virginio <i>et al.</i> , 1998), A317491 (7.5, Jarvis <i>et al.</i> , 2002)	—

Nomenclature	P2X ₅	P2X ₆	P2X ₇
Other names	—	—	P _{2Z}
Ensembl ID	ENSG00000083454	ENSG00000099957	ENSG00000089041
Selective antagonists	—	—	Brilliant Blue G (pIC ₅₀ 8.0, Jiang <i>et al.</i> , 2000)

Agonists listed show selectivity within recombinant P2X receptors of *ca.* one order of magnitude. Several P2X receptors (particularly P2X₁ and P2X₃) may be inhibited by desensitisation using stable agonists (e.g. αβ-meATP); suramin and PPADS are non-selective antagonists at rP2X_{1–3,5} and hP2X₄, but not rP2X_{4,6,7} (Buell *et al.*, 1996), and can also inhibit ATPase activity (Crack *et al.*, 1994). Ip₅I is inactive at rP2X₂, an antagonist at rP2X₃ (pIC₅₀ 5.6) and enhances agonist responses at rP2X₄ (King *et al.*, 1999). Antagonist potency of NF023 at recombinant P2X₂, P2X₃ and P2X₅ is two orders of magnitude lower than that at P2X₁ receptors (Soto *et al.*, 1999). The P2X₇ receptor may be inhibited in a non-competitive manner by the protein kinase inhibitors KN-62 and chelerythrine (Shemon *et al.*, 2004), while P2X₇ receptor-induced pore formation may be blocked by the p38 MAP kinase inhibitor SB202190 (Donnelly-Roberts *et al.*, 2004). Some recombinant P2X receptors expressed to high density bind [³⁵S]-ATPγS and [³H]-αβ-meATP, although the latter can also bind to 5'-nucleotidase (Michel *et al.*, 1995).

Abbreviations: **A317491**, 5-([3-phenoxybenzyl][(1*S*)-1,2,3,4-tetrahydro-1-naphthalenyl]amino)carbonyl)-1,2,4-benzenetricarboxylic acid; **ATPγS**, adenosine 5'-(3-thio)triphosphate; **Ip₅I**, diinosine-5',5''-pentaphosphate; **αβ-meATP**, αβ-methylene-adenosine 5'-triphosphate; **βγ-meATP**, βγ-methylene-adenosine 5'-triphosphate; **KN-62**, 1-(*N*,*O*-bis[5-isoquinolinesulphonyl]-*N*-methyl-1-tyrosyl)-4-phenylpiperazine; **NF023**, 8,8'-(carbonylbis[imino-3,1-phenylene carbonylimino])bis-1,3,5-naphthalenetrisulfonic acid; **NF449**, 4,4',4'',4'''-(carbonylbis[imino-5,1,3-benzenetriyl-bis(carbonylimino)])tetrakisbenzene-1,3-disulfonic acid octasodium salt; **PPADS**, pyridoxalphosphate-6-azophenyl-2',4'-disulphonate; **SB202190**, 4-[4-(4-fluorophenyl)-5-pyridin-4-yl-1*H*-imidazol-2-yl]phenol; **TNP-ATP**, 2',3'-*O*-(2,4,6-trinitrophenyl)-ATP

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ZAC (zinc-activated channel)

Overview: The zinc-activated channel (ZAC, provisional nomenclature and alternatively termed L2) is a recently identified member of the Cys-loop family that includes the nicotinic acetylcholine, 5-HT₃, GABA_A and strychnine-sensitive glycine receptors (Davies *et al.*, 2003; Houtani *et al.*, 2005). The channel is likely to exist as a homopentamer of 4TM subunits that form an intrinsic cation-selective channel displaying constitutive activity that is blocked by (+)-tubocurarine (Davies *et al.*, 2003). ZAC is present in the human, chimpanzee, dog, cow and opossum genomes, but is functionally absent from mouse, or rat, genomes (Davies *et al.*, 2003; Houtani *et al.*, 2005).

Nomenclature	ZAC
Ensembl ID	ENSG00000186919
Selective agonists (pEC ₅₀)	Zn ²⁺ (3.3)
Selective antagonists (pIC ₅₀)	(+)-Tubocurarine (5.2)
Functional characteristics	Outwardly rectifying current (both constitutive and evoked by Zn ²⁺)

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