

Supplementary Material

1 Supplementary Data

Table 1 Cannabinoid effects on cys-loop receptors:

Anandamide (AEA) Arachidonylethanolamide					
Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GlyR $\alpha 1$	PAM	EC50 319 nM $\pm$ 31 nM		X. laevis oocytes	(Hejazi et al. 2006)
GlyR $\alpha 1\beta 1$	PAM	EC50 318 nM $\pm$ 24 nM		X. laevis oocytes	(Hejazi et al. 2006)
GlyRa1	PAM	EC50 38 nM $\pm$ 11 nM		HEK293 cells	(Yang et al. 2008)
GlyRa1 β	PAM	EC50 75 nM $\pm$ 20 nM	128 $\pm$ 33%	HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 2$		-		HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 3$		-		HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 1$	PAM		$\sim$ 160 %*	HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 2$	PAM		$\sim$ 90 %*	HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 3$	PAM		$\sim$ 110 %*	HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR	PAM		I/CTRL(*100%) 20 $\pm$ 7% at 1 μ M	hippocampal CA1 and CA3 pyramidal neurons and Purkinje neurons	(Lozovaya et al. 2005)
GlyR $\alpha 1$	PAM				
GlyR $\alpha 1$	PAM	EC50 Glycine + AEA: 5.5 $\pm$ 2.0 μ M	86 $\pm$ 24 % at 1 μ M AEA 800 $\pm$ 71% at 30 μ M AEA	spinal neurons	(Xiong et al. 2012)
GlyR $\alpha 1$ / GlyR $\alpha 1\beta$	PAM	EC50 Glycine + AEA: 4.2 $\pm$ 1.95 μ M	97 $\pm$ 16% / 85 $\pm$ 12% at 1 μ M AEA	HEK293 cells	(Xiong et al. 2012)
GABAAR $\alpha 2\beta 3\gamma 2$		-		X. laevis oocytes	(Hejazi et al. 2006)
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	minor PAM effects at 1 μ M and 3 μ M		X. laevis oocytes	(Sigel et al. 2011)

5HT3A	NAM	inhibition	X. laevis oocytes, HEK293 cells	(Xiong et al. 2008)
5HT3A	NAM	IC50 129.6 nM	HEK293 cells	(Barann et al. 2002)
5HT3A		1 μ M 46% 10 μ M 41% 100 μ M 29%	binding assay	(Kimura et al. 1998)
nAChR α 4 β 2	NAM	IC50 (at 20 min) of approximately 300 nM	SH-EP1 cells	(Spivak et al. 2007)
nAChR α 4 β 2	NAM	IC50 value of 0.9 ± 2 μ M	thalamic synaptosomes	(Butt et al. 2008)
nAChR α 7	NAM	10 nM to 30 μ M IC50 229.7nM $\pm$ 20.4 nM	X. laevis oocytes	(Oz et al. 2003)

2-Arachidonoylglycerol (2-AG)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR α 1 β 2 γ 2	PAM	EC50 2.1 ± 0.5 μ M subunit selectivity β 2 > β 3 (1/3) > β 1 (no effect) superadditivity between THDOC and 2-AG.	$138 \pm 21\%$	X. laevis oocytes	(Sigel et al. 2011)
GABAAR α 1 β 2 δ	PAM	EC50 2.9 ± 1.8 μ M		X. laevis oocytes	(Sigel et al. 2011)
GABAAR α 1 β 2 γ 2	PAM	binding site at M3-M4 interface of β 2 subunit		X. laevis oocytes Homology modeling and docking site-directed mutagenesis	(Baur et al. 2013)
GABAAR α 1 β 2 γ 2L	PAM	EC50 GABA: 181.3 μ M (127.0 – 258.7) EC50 GABA + 10 μ M 2-AG: 2.4 μ M (0.8 – 6.7)	88.9 % (70.6 – 99.4)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR α 2 β 2 γ 2L	PAM	EC50 GABA: 214.5 μ M (157.7 – 255.3) EC50 GABA + 10 μ M 2-AG: 15.7 μ M (8.2 – 30.0)	290.6 % (223.5 – 357.7)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR α 2 β 2(V436T) γ 2L	PAM	EC50 GABA: 219.2 μ M (185.5 – 259.0) EC50 GABA + 10 μ M 2-AG: 6.0 μ M (1.4 – 11.9)	99.1 % (74.1 – 124)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR α 3 β 2 γ 2L	PAM	EC50 GABA: 155.1 μ M (96.1 – 250.6) EC50 GABA + 10 μ M 2-AG: 14.7 μ M (6.8 – 31.3)	127.0 % (93.0 – 161.0)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR α 4 β 2 γ 2L	PAM	EC50 GABA: 84.6 μ M (62.7 – 114.1) EC50 GABA + 10 μ M 2-AG: 3.9 μ M (2.1 – 7.3)	114.7 % (95.3 – 134.1)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR α 5 β 2 γ 2L	PAM	EC50 GABA: 24.2 μ M (21.4 – 27.3) EC50 GABA + 10 μ M 2-AG: 1.5 μ M (0.3 – 8.2)	98.3 % (64.1 – 132.5)	X. laevis oocytes	(Bakas et al. 2017)

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GABAAR $\alpha 6\beta 2\gamma 2L$	PAM	EC50 GABA: 13.4 μM (7.5 – 24.3) EC50 GABA + 10 μM 2-AG: 6.4 μM (1.2 – 35.5)	118.6 % (62.8 – 174.4)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 1\gamma 2L$	PAM	EC50 GABA + 10 μM 2-AG: 13.3 μM (8.2 – 21.3)	142.1 % (113.2 – 169.4)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 3\gamma 2L$	PAM	EC50 GABA + 10 μM 2-AG: 9.8 μM (6.5 – 14.6)	239.7 % (204.4 – 275.1))	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 4\beta 2\delta$	PAM	EC50 GABA + 10 μM 2-AG: 4.8 μM (3.4 – 6.8)	479.6 % (436.3 – 527.4)	X. laevis oocytes	(Bakas et al. 2017)
GlyR	NAM		I/ICTRL(*100%) 40 $\pm$ 7% 1 μM	hippocampal CA1 and CA3 pyramidal neurons and Purkinje neurons	(Lozovaya et al. 2005)
GlyR $\alpha 1$	NAM			CHO cells	(Lozovaya et al. 2011)
nAChR $\alpha 7$	NAM	IC50 168 nM		X. laevis oocytes	(Oz et al. 2004)

Noladineether (NE) 2-AGE

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	~65 % compared to 2-AG		X. laevis oocytes	(Sigel et al. 2011)

N-Arachidonoyl Serine (NASer)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	~200 % compared to 2-AG		X. laevis oocytes	(Baur et al. 2013)
GlyR $\alpha 1$	PAM			HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 2$	NAM			HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 3$	NAM			HEK293 cells	(Yévenes and Zeilhofer 2011)

N-arachidonyl-glycine (NAGly)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM		~300 % compared to 2-AG	X. laevis oocytes	(Baur et al. 2013)
GABAAR $\alpha 1\beta 2\gamma 2$	PAM		~450 % at 3 μM NA-Gly	X. laevis oocytes	(Baur et al. 2013)
GABAAR $\alpha 1\beta 1\gamma 2$	PAM		~40 % at 3 μM NA-Gly	X. laevis oocytes	(Baur et al. 2013)

GlyR $\alpha 1$	NAM	IC50 2,4 μM $\pm$ 0,7 μM at EC80 IC50 24 μM $\pm$ 6,2 μM at EC10 "bell-shaped"	HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 2$	NAM	IC50 3,03 μM $\pm$ 0,09 μM	HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 3$	NAM	IC50 1.32 μM 0.10 μM	HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 1$	PAM	101 $\pm$ 11% at 10 μM NA-Gly	HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 2$	NAM	-56 $\pm$ 5% at 10 μM NA-G	HEK293 cells	(Yévenes and Zeilhofer 2011)
GlyR $\alpha 3$	NAM	-32 $\pm$ 3% at 10 μM NA-G	HEK293 cells	(Yévenes and Zeilhofer 2011)

Tetrahydrocannabinol (THC)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	~30 % compared to 2-AG at 3 μM THC		X. laevis oocytes	(Sigel et al. 2011)
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	THC at 3 μM significantly enhanced GABA-activated currents		HEK293 cells	(Yao et al. 2020)
GABAAR $\alpha 2\beta 3\gamma 2$	-	no effects at 300 nM THC		X. laevis oocytes	(Hejazi et al. 2006)
GABAAR $\alpha 2\beta 3\gamma 2$	PAM	weak effects at 10 μM THC***		X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 4\beta 1\delta$	NAM	GABA EC50 0,02 μM (0,01 - 0,02) GABA + 10 μM THC EC50 1,2 (0,20 - 6,66)***		X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 4\beta 3\delta$	PAM	GABA EC50 3,0 μM (2,42 - 3,72) GABA + 10 μM THC EC50 0,4 μM (0,19 - 0,99)***		X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 6\beta 3$	PAM		~500% at 1 μM THC***	X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 4\beta 3$	PAM		~300% at 1 μM THC***	X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 2\beta 3$	PAM	GABA EC50 6,0 μM GABA + 10 μM THC EC50 1,39 μM ***		X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 2\beta 3$	PAM	GABA EC50 10,6 μM GABA + 10 μM THC EC50 3,7 μM ***		X. laevis oocytes	(Schmiedhofer 2017)
GlyR $\alpha 1\beta$	PAM		44 $\pm$ 13% at 30 nM THC 82 $\pm$ 4% at 100 nM THC 136 $\pm$ 11% at 300 nM THC	cultured spinal neurons	(Xiong et al. 2011)
GlyR $\alpha 2$	PAM		232 $\pm$ 35% at 1 μM THC	HEK293 cells	(Xiong et al. 2011)
GlyR $\alpha 3$	PAM		97 $\pm$ 7% at 100 nM THC 1127 $\pm$ 142% at 1 μM THC	HEK293 cells	(Xiong et al. 2011)
GlyR $\alpha 1$	PAM		1156 $\pm$ 472% 1 μM THC	HEK293 cells	(Xiong et al. 2011)

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GlyR $\alpha 1$	PAM		~1000 % at 1 μ M THC****	HEK293 cells	(Yao et al. 2020)
GlyR $\alpha 3$	PAM		~1000 % at 1 μ M THC****	HEK293 cells	(Yao et al. 2020)
GlyR $\alpha 1\beta$	PAM		~400 % at 1 μ M THC****	HEK293 cell	(Yao et al. 2020)
GlyR $\alpha 1$	PAM	162 % $\pm$ 12 % at 1 μ M THC EC50 1.3 μ M $\pm$ 0.6 μ M	260% $\pm$ 30 % at max.	X. laevis oocytes	(Wells et al. 2015)
5HT3A	NAM	THC at 1 μ M significantly reduced the 5-HT-activated current		HEK293 cells	(Yao et al. 2020)
5HT3A	NAM	IC50 38,4 nM		HEK293 cells	(Barann et al. 2002)
5HT3A	NAM	IC50 119 nM $\pm$ 13 nM	97 % $\pm$ 5% at 1 μ M THC	HEK293 cells	(Xiong et al. 2011)
5HT3A	NAM	IC50 285 nm $\pm$ 23 nM 1 ng of cRNA IC50 1.2 μ M $\pm$ 0.3 μ M 3 ng of cRNA.	68,5% at 3 μ M THC with 3 ng cRNA	X. laevis oocytes	(Yang et al. 2010)
nACh $\alpha 7$	-	10 μ M THC show no effect		X. laevis oocytes	(Mahgoub et al. 2013)
nAChR $\alpha 4\beta 2$	-	THC has no impact, even coapplied wit AEA		SH-EP1 cells	(Spivak et al. 2007)
nAChR $\alpha 7$	-	THC has no impact		X. laevis oocytes	(Oz et al. 2004)

Cannabidiol (CBD)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GlyR $\alpha 3$	PAM		491 $\pm$ 101% at 1 μ M CBD	HEK293 cells	(Xiong et al. 2012)
GlyR $\alpha 1\beta$	PAM	EC50 Glycine + CBD: 12.3 $\pm$ 3.8 μ mol/l EC50 Glycine: 132.4 $\pm$ 12.3 μ mol/l		HEK293 cells	(Ahrens et al. 2009)
GlyR $\alpha 1$	PAM	EC50 Glycine + CBD: 18.1 $\pm$ 6.2 μ mol/l EC50 Glycine: 144.3 $\pm$ 22.7 μ mol/l		HEK293 cells	(Ahrens et al. 2009)
GlyR $\alpha 1$ S267I	-	S267I in TM2 diminishes CBD modulatory effects of GlyR		HEK293 cells	(Foadi et al. 2010)
GABAAR $\alpha 1\beta 2$	PAM	EC50 GABA: 5.6 μ M (3.6 – 6.9) EC50 GABA + 10 μ M CBD: 3.7 μ M (1.7 – 8.0)	217.3 % (174.7 – 259.9)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 1\beta 2\gamma 2L$	PAM	EC50 GABA: 181.3 μ M (127.0 – 258.7) EC50 GABA + 10 μ M CBD: 6.5 μ M (3.5 – 12.3)	153.9 % (132.6 – 175.1)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 2$	PAM	EC50 GABA: 22.9 μ M (15.4 – 34.1) EC50 GABA + 10 μ M CBD: 2.0 μ M (1.0 – 4.1)	262.6 % (219.2 – 305.9)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 2\gamma 2L$	PAM	EC50 GABA: 214.5 μ M (157.7 – 255.3) EC50 GABA + 10 μ M CBD: 3.7 μ M (1.7 – 8.0)	331.5 % (277.2 – 391.1)	X. laevis oocytes	(Bakas et al. 2017)

GABAAR $\alpha 2\beta 2(V436T)\gamma 2L$	PAM	EC50 GABA: 219.2 μM (185.5 – 259.0) EC50 GABA + 10 μM CBD: 13.3 μM (4.6 – 35.8)	123.6 % (79.9 – 167.2)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 3\beta 2\gamma 2L$	PAM	EC50 GABA: 155.1 μM (96.1 – 250.6) EC50 GABA + 10 μM CBD: 10.0 μM (6.0 – 16.7)	129.4 % (109.9 – 148.8)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 4\beta 2\gamma 2L$	PAM	EC50 GABA: 84.6 μM (62.7 – 114.1) EC50 GABA + 10 μM CBD: 0.9 μM (0.4 – 1.8)	90.4 % (74.8 – 105.9)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 5\beta 2\gamma 2L$	PAM	EC50 GABA: 24.2 μM (21.4 – 27.3) EC50 GABA + 10 μM CBD: 1.4 μM (0.6 – 3.2)	73.6 % (57.0 – 90.3)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 6\beta 2\gamma 2L$	PAM	EC50 GABA: 13.4 μM (7.5 – 24.3) EC50 GABA + 10 μM CBD: 8.2 μM (2.4 – 28.1)	71.9 % (42.6 – 101.2)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 1\gamma 2L$	PAM	EC50 GABA: 142.2 μM (105.4 – 191.7) EC50 GABA + 10 μM CBD: 17.4 μM (9.4 – 23.0)	151.4 % (123.5 – 189.9)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 2\beta 3\gamma 2L$	PAM	EC50 GABA: 50.8 μM (32.7 – 78.9) EC50 GABA + 10 μM CBD: 4.4 μM (3.1 – 6.3)	268.4 % (249.6 – 287.6)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 4\beta 2\delta$	PAM	EC50 GABA + 10 μM CBD: 23.1 μM (15.8 – 33.7)	752.4 % (649.2 – 881.5)	X. laevis oocytes	(Bakas et al. 2017)
GABAAR $\alpha 4\beta 3\delta$	PAM	EC50 GABA: 3.0 μM (2.42 - 3.72) EC50 GABA + CBD: 0.41 μM (0.10 - 1.69)	1170 % $\pm$ 127.2 at 10 μM CBD at EC 3-5 GABA 259.7 % $\pm$ 17.42 at 10 μM CBD at EC 90-100 GABA	X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 4\beta 1\delta$	NAM	EC50 GABA + 10 μM CBD 0.034 μM^{***} EC50 GABA 0.015 μM^{***}		X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 6\beta 3\delta$	PAM	EC50 GABA + 10 μM CBD EC50 1.2 μM (0.16 - 8.65) EC50 GABA 3.16 μM (1.91 - 5.23)		X. laevis oocytes	(Schmiedhofer 2017)
5HT3A	NAM	IC50 329 nM $\pm$ 19 nM	82 % $\pm$ 13% at 1 μM CBD	HEK293 cells	(Xiong et al. 2011)
5HT3A	NAM	IC50 0.6 $\mu M \pm 0.1 \mu M$	81% $\pm$ 5% at 1 μM CBD injected with 1 ng of 5-HT3A receptor cRNA	X. laevis oocytes	(Yang et al. 2010)
nAChR $\alpha 7$	NAM	IC50 11.3 $\mu M \pm 1.8 \mu M$		Whole-cell patch clamp recordings in rat hippocampal slices	(Mahgoub et al. 2013)

DH-Cannabidiol (DH-CBD)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GlyR $\alpha 1$	PAM	DH-CBD significantly reduced glycine EC50 values in seven of nine hyperekplexia mutant receptors		HEK293 cells	(Xiong et al. 2014)

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GlyR $\alpha 1$ R271Q	PAM	EC50 Glycine: 21 μ M $\pm$ 3.2 μ M EC50 Glycine + DH-CBD: 1.2 μ M $\pm$ 0.1 μ M		HEK293 cells	(Xiong et al. 2014)
GlyR $\alpha 3$	PAM	EC50 Glycine: 377 μ M $\pm$ 45 μ M EC50 Glycine + DH-CBD: 195 μ M $\pm$ 51 μ M at 1 μ M DH-CBD EC50 Glycine + DH-CBD: 58 μ M $\pm$ 13 μ M at 10 μ M DH-CBD	989 $\pm$ 171% at 1 μ M DH-CBD	HEK293 cells	(Xiong et al. 2012)
GlyR $\alpha 1$	PAM	DH-CBD at 1 mM remarkably enhanced IGly		HEK293 cells	(Lu et al. 2018)
GABAAR $\alpha 1\beta 2\gamma 2$	-	no effect on GABAAR alone, DH-CBD-induced disruption of the interaction between GlyR $\alpha 1$ R271Q and GABAAR		HEK293 cells	(Zou et al. 2020)

Cannabigerol (CBG)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 3\gamma 2$	PAM	EC50 GABA: 21.3 μ M (18,67 - 24,4) *** EC50 GABA + 10 μ M CBG: 12.49 μ M (3,83 - 40,79) ***	~200 % ***	X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 6\beta 3\delta$	PAM		~300 % ***	X. laevis oocytes	(Schmiedhofer 2017)
GABAAR $\alpha 6\beta 3\gamma 2$	PAM	EC50 GABA: 3.638 μ M (2,44 - 5,42) *** EC50 GABA + 10 μ M CBG: 3.084 μ M (0,97 - 9,77 μ M) ***	~150 % ***	X. laevis oocytes	(Schmiedhofer 2017)

Ajulemic Acid

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GlyR $\alpha 1$ S267I	-			HEK293 cells	(Foadi et al. 2010)

beta-Caryophyllen (BCP)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 4\beta 3\delta$	NAM		80 % $\pm$ 33 at 600 μ M**	HEK293 cells	(Janzen et al. 2021)
GABAAR $\alpha 6\beta 3\delta$	NAM		76 % $\pm$ 26 at 600 μ M**	HEK293 cells	(Janzen et al. 2021)
GABAAR $\alpha 1\beta 2\gamma 2$	NAM		90 % $\pm$ 37 at 600 μ M**	HEK293 cells	(Janzen et al. 2021)
GABAAR $\alpha 1\beta 2$	NAM		98 % $\pm$ 13 at 600 μ M**	HEK293 cells	(Janzen et al. 2021)

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Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GlyR $\alpha 1$	PAM	EC50 co-activation: $5.1 \mu\text{M} \pm 2.6$ EC50 direct activation: $188.7 \mu\text{M} \pm 46.2$		HEK293 cells	(Demir et al. 2009)
GlyR $\alpha 1$ S267I	-			HEK293 cells	(Foadi et al. 2010)
GlyR $\alpha 1$	PAM	EC50: $0.27 \mu\text{M} \pm 0.05$		HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 1\beta$	PAM		$78 \% \pm 5$ at $30 \mu\text{M}$	HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 2$	NAM	IC50: $0.090 \mu\text{M} \pm 0.021$		HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 3$	NAM	IC50: $0.050 \mu\text{M} \pm 0.006$		HEK293 cells	(Yang et al. 2008)

Rimonabant (SR141716A)

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	EC50: $7.3 \mu\text{M} \pm 0.5 \mu\text{M}$	$3381 \% \pm 165$	X. laevis oocytes	(Baur et al. 2012)

AM251

Receptor	Effects	Results	max. potentiation / max. inhibition	System	Source
GABAAR $\alpha 1\beta 2\gamma 2$	PAM	EC50 $0.40 \mu\text{M} \pm 0.13$ EC50 GABA: $15.4 \mu\text{M} \pm 0.8 \mu\text{M}$ EC50 GABA + $1 \mu\text{M}$ AM251: $5.5 \mu\text{M} \pm 0.4 \mu\text{M}$	$881 \% \pm 167$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 1\beta 2$	PAM		$\sim 550 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 1\beta 3\gamma 2$	PAM		$\sim 1250 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 2\beta 2\gamma 2$	PAM		$\sim 850 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 3\beta 2\gamma 2$	PAM		$\sim 400 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 5\beta 2\gamma 2$	PAM		$\sim 500 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 6\beta 2\gamma 2$	PAM		$\sim 200 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)
GABAAR $\alpha 1\beta 1\gamma 2$	PAM		$\sim 20 \%$ at $3 \mu\text{M}$ AM251 + $0.5 \mu\text{M}$ GABA*	X. laevis oocytes	(Baur et al. 2012)

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GABAAR $\alpha 4\beta 2\delta$	PAM	~350 % at 3 μ M AM251 + 0,5 μ M GABA*	X. laevis oocytes	(Baur et al. 2012)
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Receptor	Effects	Results	max. potentiation / max. inhibition	System	
GlyR $\alpha 1$	-			HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 1\beta$	-			HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 2$	NAM	IC50: 0.22 μ M $\pm$ 0.05		HEK293 cells	(Yang et al. 2008)
GlyR $\alpha 3$	NAM	IC50: 0.050 μ M $\pm$ 0.006		HEK293 cells	(Yang et al. 2008)
5HT3A	NAM	IC50: 103.5 nM		HEK293 cells	(Barann et al. 2002)
nAChR $\alpha 7$	-	no effects		X. laevis oocytes	(Oz et al. 2004)

*read from graph

**high concentration of substance

*** low sample size (n < 4)

**** $\Delta 9$ -THC effects are cholesterol dependent

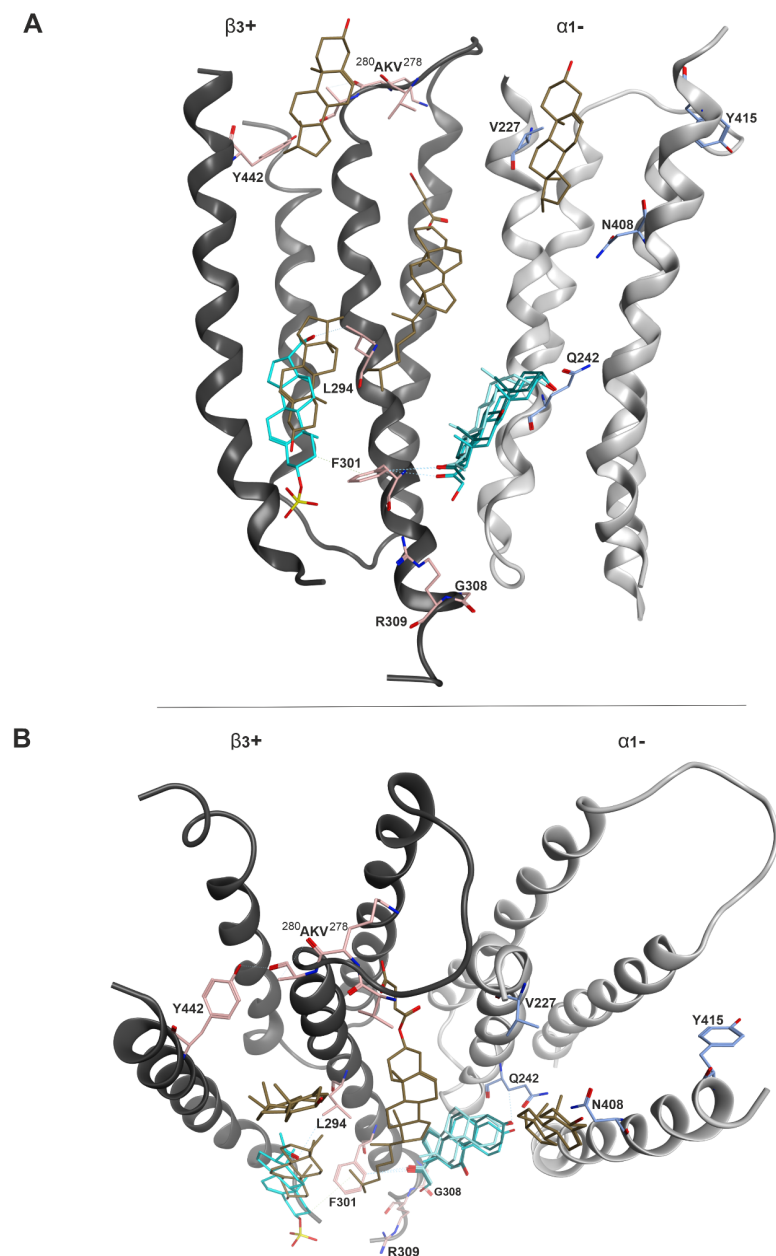
Supplementary Table 2. Cys-loop receptors genetic variants in epilepsies. Cys-loop receptors genetic variants in epilepsies. This table provides an overview of variants in Cys-loop receptor subunit encoding genes identified in patients.

Epilepsy syndrom	genetic basis
DS (Dravet syndrome)	GABRA1 (S76R, R112Q, L146M, R214C, R214H, L215P, G251S, V287I, K306T) GABRB2 (A159S, Y181F, F331I, F331S) GABRB3 (T157M, R232Q, T281I) GABRG2 (Q40X, T90R, P302L, Q390X) (Fu et al., 2022)
LGS (Lennox gastaut syndrome)	GABRB3 (D120N) (Qu et al., 2020) GABRB3(D120N, E180G, Y302C) (Janve et al., 2016) GABRA1 (T292I) GABRB2 (I246T, P252L, I288S) GABRB3 (D120N, E180G, Y302C, A305T, N328D) GABRG2 (P83S) (Fu et al., 2022)
WS (West syndrome) and IS (infantile spasm)	GABRA1 (R112Q, P260S, P260L, M263T, M263I, T292I, L296S, W315L) GABRB2 (T184I, R240T, F245S, P252L, I299S) GABRB3 (L52V, I69T, E77K, M80L, N110D, L256Q, L278F, Y302C) (Fu et al., 2022) GABRB3(N110D, E180G) GABRB1(F246S) (Janve et al., 2016)
EIEE (early infantile epileptic encephalopathy)	GABRA1 (P260L, T289P, T289A) GABRB2 (K303N) GABRB3 (T287I) (Fu et al., 2022)
EMA (early myoclonic encephalopathy)	GABRB2 (I246T, T284K, T287P) (Fu et al., 2022)
EOEE (early onset epileptic encephalopathy)	GABRA2 (T292K) (Butler et al., 2018) GABRA1 (R112Q, N115D, V287L, A332V) GABRB2 (P252L, K298G, K303R) GABRB3 (N110D, K127R, L170R, T185I, S254F, L256Q, T288N, L293H, A305V) (Fu et al., 2022)
EIMFS (epilepsy of infancy with migrating focal seizures)	GABRA1 (P280T) GABRB3 (L124F, Y245H, S254F, T281A, L284M)
CAE (Childhood absence epilepsy) JAE (Juvenile absence epilepsy)	GABRB3 (G32R) (Gurba et al., 2012) GABRA1 (R214C, L267I, S326fs328X) GABRB2 (V316I) GABRB3 (P11S, S15F, G32R, V37G, E357K) GABRG2 (R82Q, T90M, R177fs) (Fu et al., 2022)
SHE (sleep-related hypermotor epilepsy) ADSHE formerly autosomal dominant nocturnal frontal lobe epilepsy (ADNFLE)	GABRB2 V337G CHRNA4 R336H CHRNA2 V287L CHRNA2 I279A CHRNA4 S284L

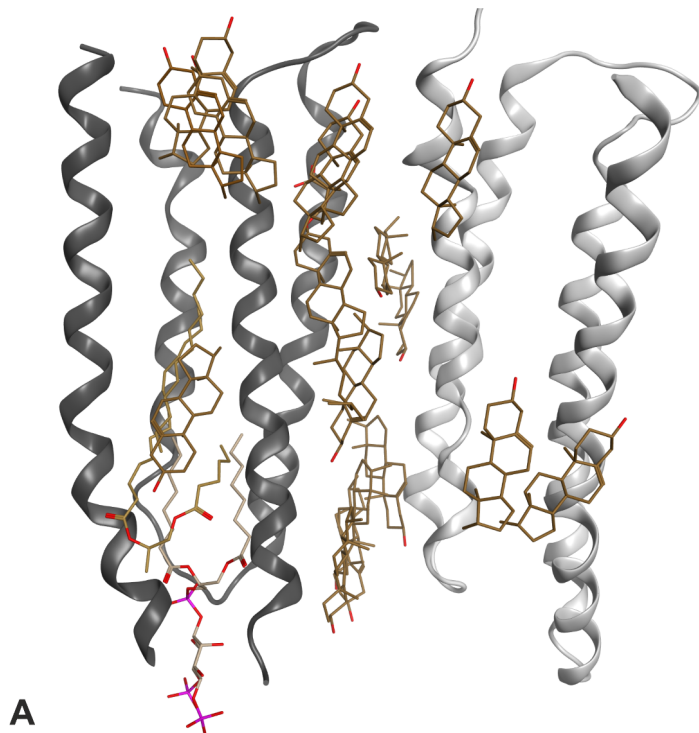
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	CHRNA4 S280P CHRNA4 insL (Becchetti et al., 2020)
MAE (epilepsy with myoclonic-atonic seizures) JME (juvenile myoclonic epilepsy) MSE (myoclonic status epilepsy)	GABRA1 (F104C, R214C, K306T, A322D) GABRB2 (V262F) GABRB3 (S76C, R111X, D120N, R142L, Y184H) GABRG2 (R323Q) (Fu et al., 2022)
IGE (idiopathic generalized epilepsies)	GABRA1 (A322D) GABRA4 (T320A, L26M) GABRA5 (I48L) GABRA6 (R46W, P385S) GABRP (V10M) GABRE (G66S, Y38S) GABRG2 (R43Q, K289M, Q351X) GABRD (E177A, R220H) (Dibbens et al., 2009)
Unspecified EE	GABRA1 (G251D) GABRB2 (M79T, D125N, Y244H, P252A, L277S, T287P, I288S) GABRB3 (Y182F, R232X, R232Q, Q249K, P253L, P301L, Y302C) GABRG2 (A106T, I107T, P282T, P282S, R323W, F343L) (Fu et al., 2022)
DEE (developmental and epileptic encephalopathy) EDD (epileptogenic developmental disorders) GDD (global development delay) NDD (neurodevelopmental disorder) NDDE (neurodevelopmental disorder with epilepsy)	GABRA1 (M79T, A112E, D125N, Y181F, Y183H, T184I, F224C, R240T, Y244H, F245S, P252A, P252T, V262F, L277S, V282A, T284K, R293P, K298G, Y301C, K303N, A304V) GABRB1 (V78L, M80L, Q249H, P253S, T288I, L321P) (Fu et al., 2022)
Generalized epilepsy with febrile seizures	GABRD (G177A) variant (Dibbens et al., 2004) GABRA1 (V74I9) GABRB2 (D108Y, V133M, M161L, N350_del) GABRB3 (P54L, T157M, R429Q) GABRG2 (Q40X, N79S, P83S, T90M, R136X, R323Q, K328M, Q390X, W429X) (Fu et al., 2022)
FS (febrile seizure)	GABRB2 (R354C) GABRB3 (T157M) GABRG2 (R82Q, R136X, R177G, R177fs, K328M) (Fu et al., 2022)
RE (rolandic epilepsy)	GABRG2 (G257R, R323Q, I389V) (Fu et al., 2022)
Complex epilepsy	GABRD (E177A, R220H) (Feng et al., 2006)

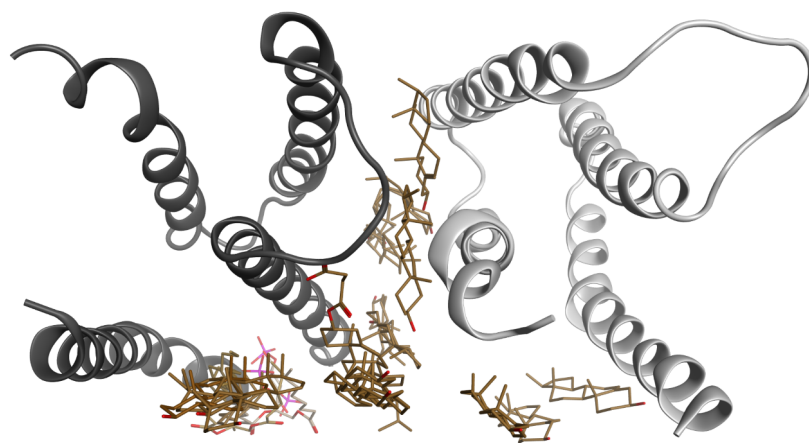
Development of epilepsy	GABRA1 (A294D) mutation (Fisher, 2004)
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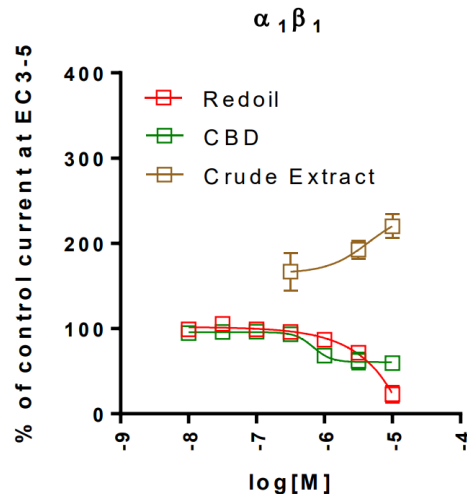
Supplementary Figure 1. Steroid binding sites on a $\beta 3+$ (dark grey) $\alpha 1-$ (light grey) GABA_A receptor dimer (PDB ID 6HUO - Masiulis et al., 2019). Amino acids which have been proposed based on photoaffinity labeling by Chen et al., 2019 are rendered in stick representation. Color coding: Steroid binding site at the lower interface site: THDOC (5OSB in cyan - Lavery et al., 2017), alphaxalone (6CDU in dark cyan - Chen et al., 2018), pregnanolone (5OJM in light cyan - Miller et al., 2017); lipid associated lower TMD site: pregnenolone sulfate (5OSC in cyan - Lavery et al., 2017) and cholesterol (6D6T in brown - Zhu et al., 2018); lipid associated upper TMD site at the $\beta 3+$ subunit: cholesterol (6D6T in brown); lipid associated upper TMD site at the $\alpha 1-$ subunit: cholesterol (6D6T in brown); lipid associated TMD interface site: cholesterol hemisuccinate (5OSC in brown - Lavery et al., 2017)



B



Supplementary Figure 2. Cholesterol, cholesterol derivatives and other endogenous lipid molecules: PDB files with models of cholesterol (and derivatives), and PIP2 (6I53 - Lavery et al., 2019, 6D6T, 6D6U and 5OSC) have been superposed to render the localizations of the lipid molecules in direct comparison. A: Dimer seen from the outside. B: dimer seen from the perspective of the ECD. Dark grey ribbon: principal subunit, light grey ribbon: complementary subunit. Ribbon rendering of 6D6U. All cholesterol are from 6D6U and 6D6T. The cholesterol hemisuccinate at the interface site is from 5OSC.

Pilot experiment: plant matrix vs. extract fractions

Supplementary Figure 3. Pilot Experiment. This Figure shows distinct effects of cannabis extract fractions on $\alpha_1\beta_1$ GABAAR. **Crude extract** produced with CO₂ extraction contains most substances from cannabis, including the cannabinoid part (25% CBD, 4% Δ^9 -THC). **Redoil** a fraction of the crude extract consists of a fraction containing mostly cannabinoids (~90% CBD, ~10% Δ^9 -THC, and other plant compounds), **CBD** was used in purified form (99,87 %, BSPG, UK) The Figure was adopted from ([Schmiedhofer, 2017](#)).

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