

Serotonin_DataAnalysis_Apr9

```
# Read in data
setwd("~/Desktop/Serotonin_Paper")
Sys.setlocale("LC_ALL", "C")
```

```
## [1] "C/C/C/C/C/en_US.UTF-8"
```

```
x <- read.csv("5-HT_Extract_Results_Apr8.csv", as.is = TRUE)

library(dplyr)
```

```
##
## Attaching package: 'dplyr'
```

```
## The following objects are masked from 'package:stats':
##
##      filter, lag
```

```
## The following objects are masked from 'package:base':
##
##      intersect, setdiff, setequal, union
```

Overall

```
nrow(x)
```

```
## [1] 9631
```

```

Receptors <- strsplit(x$Receptor, ";")

Receptors <- unlist(Receptors)

# 1c is really 2c
Receptors <- gsub("5ht1c","5ht2c", Receptors)

Receptors <- gsub("5ht3a", "5ht3", Receptors)
Receptors <- gsub("5ht3b", "5ht3", Receptors)
Receptors <- gsub("5ht3c", "5ht3", Receptors)
Receptors <- gsub("5ht3d", "5ht3", Receptors)
Receptors <- gsub("5ht3e", "5ht3", Receptors)
Receptors <- gsub("5ht3f", "5ht3", Receptors)

Receptors <- gsub("5ht4a", "5ht4", Receptors)
Receptors <- gsub("5ht4b", "5ht4", Receptors)
Receptors <- gsub("5ht4c", "5ht4", Receptors)
Receptors <- gsub("5ht4d", "5ht4", Receptors)
Receptors <- gsub("5ht4e", "5ht4", Receptors)
Receptors <- gsub("5ht4f", "5ht4", Receptors)

Receptors <- gsub("5ht6a", "5ht6", Receptors)
Receptors <- gsub("5ht6b", "5ht6", Receptors)
Receptors <- gsub("5ht6c", "5ht6", Receptors)
Receptors <- gsub("5ht6d", "5ht6", Receptors)
Receptors <- gsub("5ht6e", "5ht6", Receptors)
Receptors <- gsub("5ht6f", "5ht6", Receptors)

Receptors <- gsub("5ht7a", "5ht7", Receptors)
Receptors <- gsub("5ht7b", "5ht7", Receptors)
Receptors <- gsub("5ht7c", "5ht7", Receptors)
Receptors <- gsub("5ht7d", "5ht7", Receptors)
Receptors <- gsub("5ht7e", "5ht7", Receptors)
Receptors <- gsub("5ht7f", "5ht7", Receptors)

(receptors <- table(Receptors)[c(1:10,14,15,17,18,22,23)])

```

```

## Receptors
## 5ht1 5ht1a 5ht1b 5ht1d 5ht1e 5ht1f 5ht2 5ht2a 5ht2b 5ht2c 5ht3 5ht4
## 741 5283 1463 591 39 74 1266 2595 339 1665 2029 697
## 5ht5a 5ht5b 5ht6 5ht7
## 80 13 443 742

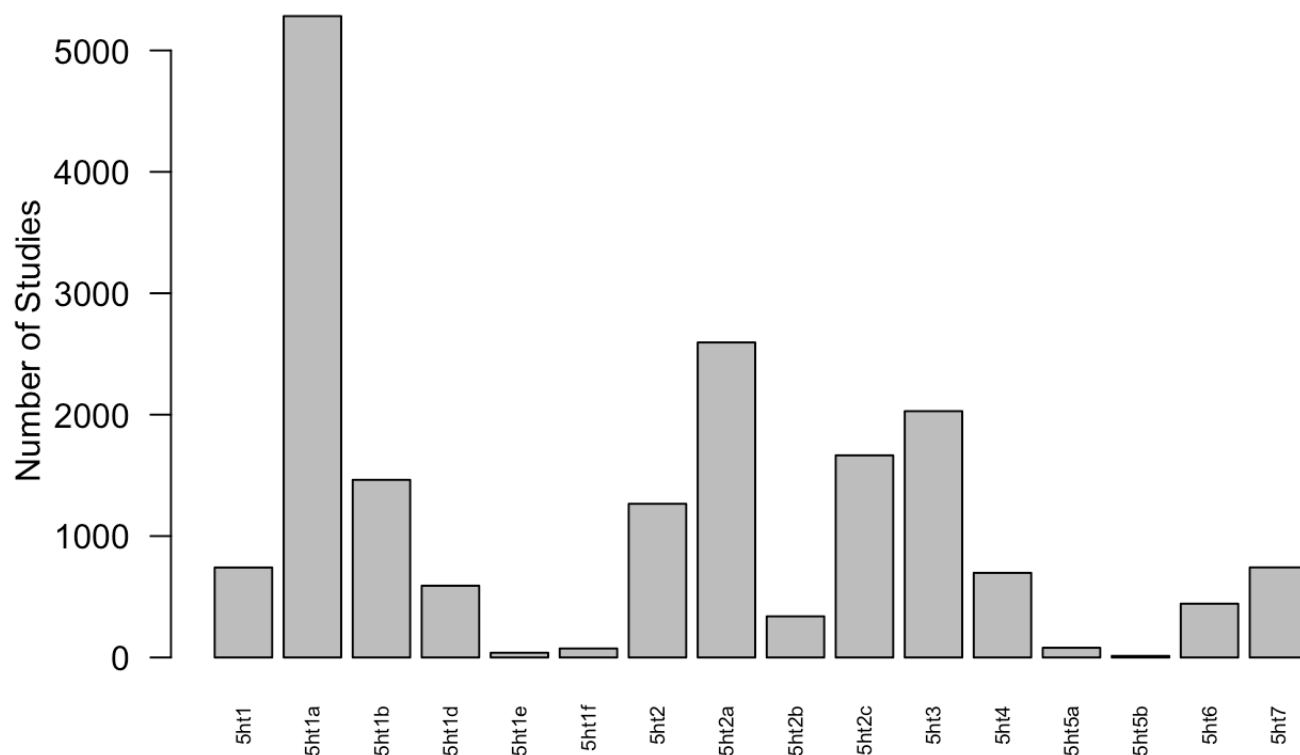
```

```

barplot(receptors, las = 2, cex.names=.6, main = "Number of Studies by 5-HT Receptor"
, ylab = "Number of Studies")

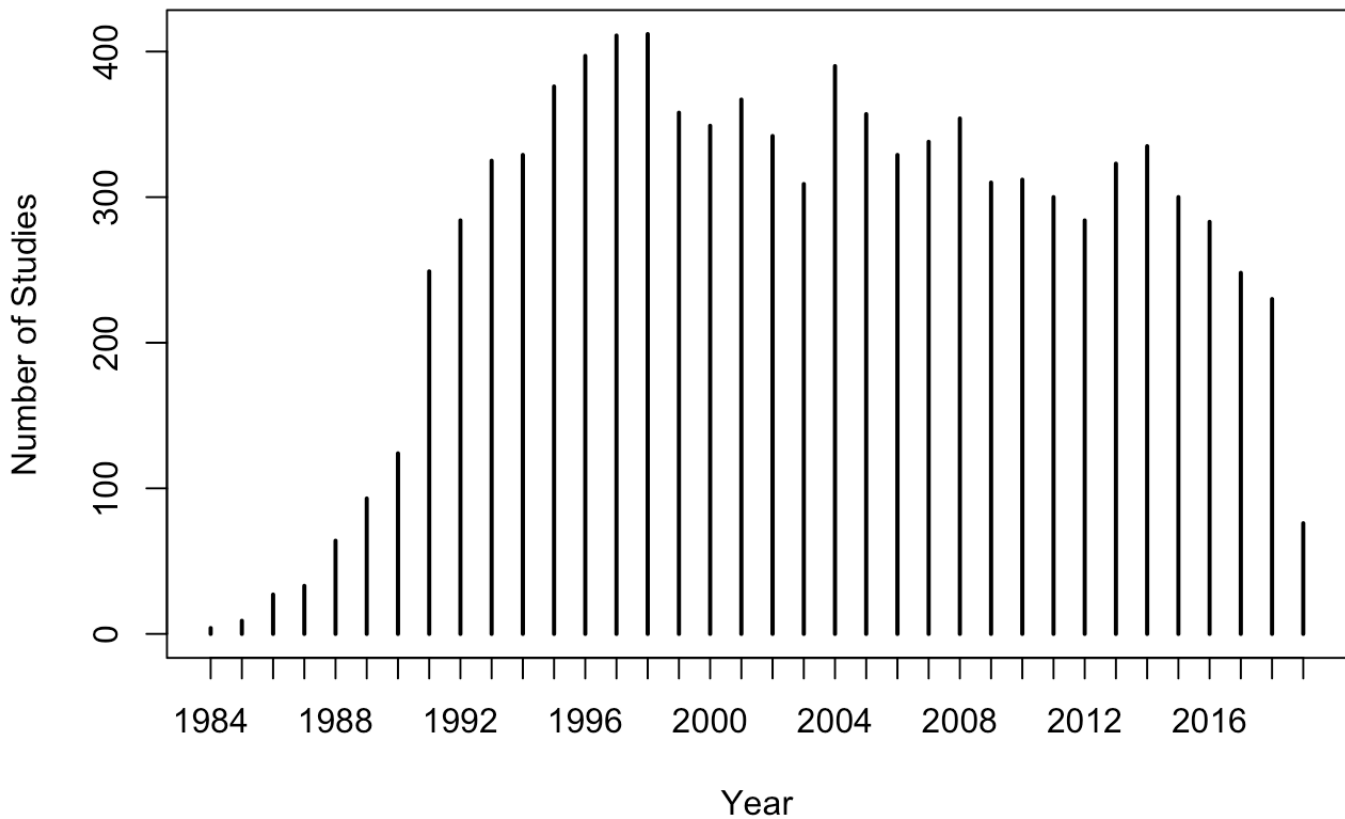
```

Number of Studies by 5-HT Receptor



```
plot(table(x$Year), main = "Number of 5-HT Receptor Subtype Studies by Year", xlab =  
"Year", ylab = "Number of Studies")
```

Number of 5-HT Receptor Subtype Studies by Year



```
Methods <- strsplit(x$Methods, ";")

Methods <- unlist(Methods)

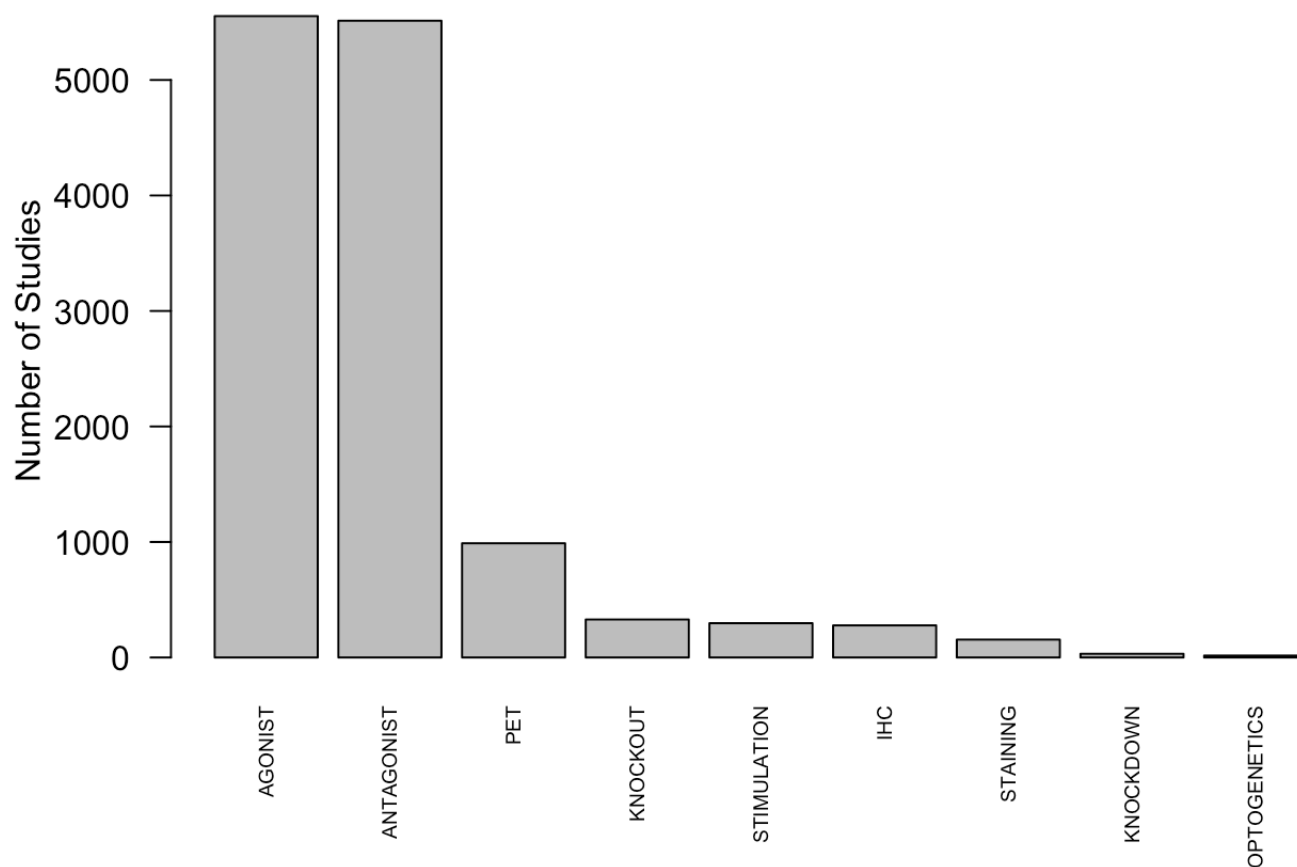
Methods <- gsub("Method.", "", Methods)
Methods <- gsub("IMMUNOHISTOCHEMISTRY", "IHC", Methods)

(methods <- sort(table(Methods), decreasing = TRUE))
```

```
## Methods
##      AGONIST      ANTAGONIST      PET      KNOCKOUT      STIMULATION
##      5553         5513         989         329         297
##      IHC      STAINING      KNOCKDOWN      OPTOGENETICS
##      278         155         32         17
```

```
barplot(methods, las = 2, cex.names=.6, main = "Methods Used to Study 5-HT Receptors"
, ylab = "Number of Studies")
```

Methods Used to Study 5-HT Receptors



```
Species <- strsplit(x$Species, ";")

Species <- unlist(Species)

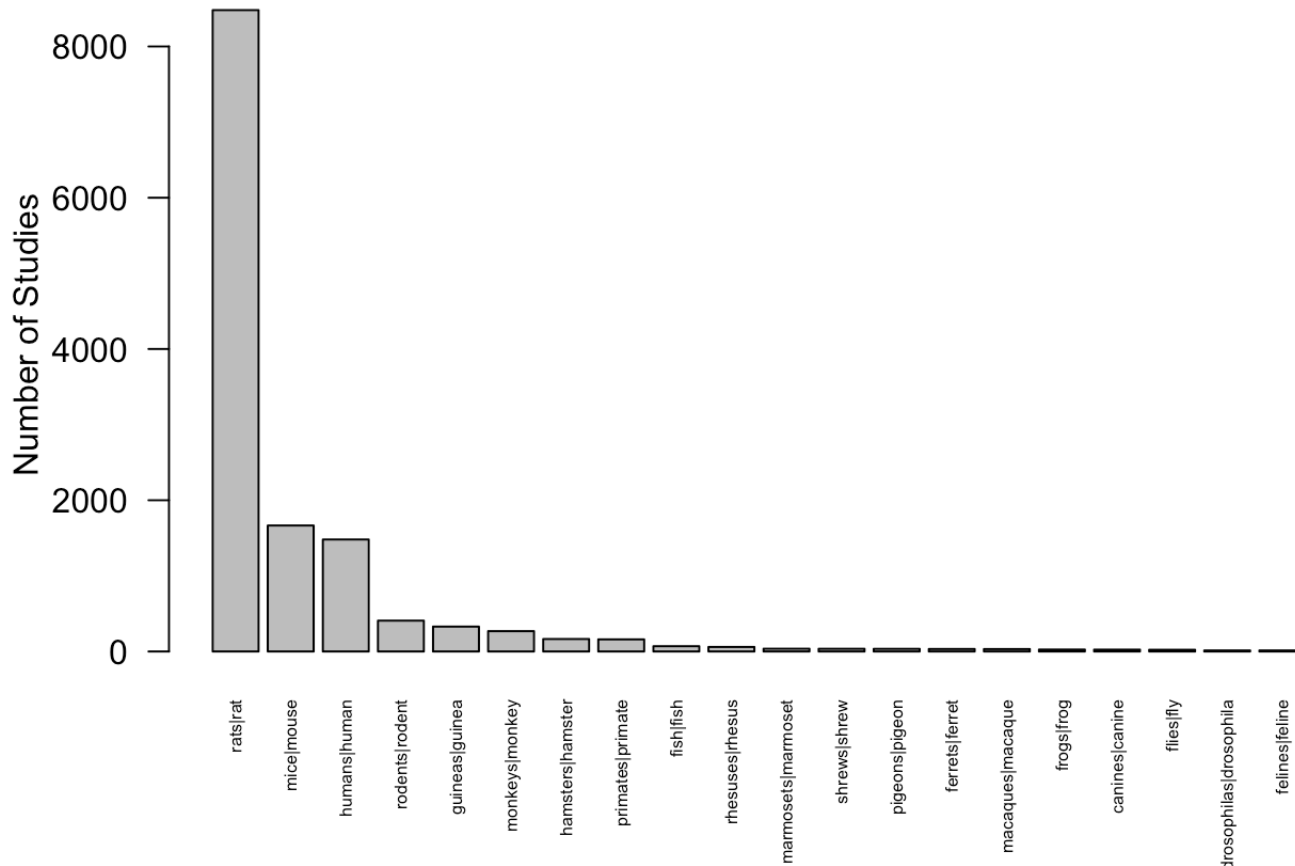
Species <- gsub(".*b", "", Species)
Species <- gsub('.{2}$', '', Species)
Species <- gsub(".*\\(", "", Species)
Species <- gsub('.{1}$', '', Species)

(species <- sort(table(Species), decreasing = TRUE)[c(2,21,27,53,59,64,80,84,112,118,
138,140,142,145,146,154,157,164,191,192)])
```

```
## Species
##          rats|rat          mice|mouse          humans|human
##          8481          1666          1481
##          rodents|rodent          guineas|guinea          monkeys|monkey
##          408          329          269
##          hamsters|hamster          primates|primate          fish|fish
##          165          160          70
##          rhesuses|rhesus          marmosets|marmoset          shrews|shrew
##          60          37          36
##          pigeons|pigeon          ferrets|ferret          macaques|macaque
##          35          33          32
##          frogs|frog          canines|canine          flies|fly
##          26          25          23
## drosophilas|drosophila          felines|feline
##          14          14
```

```
barplot(species, las = 2, cex.names=.5, main = "Species Used to Study 5-HT Receptors"
, ylab = "Number of Studies")
```

Species Used to Study 5-HT Receptors



```

Agonists <- strsplit(x$Agonist, ";")

Agonists <- unlist(Agonists)
Agonists <- gsub(".*\\\\\\\\", "", Agonists)
Agonists <- substring(Agonists, 2)
Agonists <- gsub('.{2}$', '', Agonists)
Agonists <- gsub("\\\\?", "-", Agonists)
Agonists <- gsub(" ", "", Agonists)

Agonists <- gsub("5-carboxamidotryptamine", "5-ct", Agonists)

(agonists <- sort(table(Agonists), decreasing = TRUE)[c(1, 3:30)])

```

```

## Agonists
##          8-oh-dpat          doi          buspirone
##          1956          682          414
##          5-ct          haloperidol          clozapine
##          378          318          271
##          mcpp          ipsapirone          sumatriptan
##          257          231          229
##          dom          methysergide          amphetamine
##          224          195          182
##          ru-24969          ethanol          tfmpp
##          167          166          140
##          cp-93129          yohimbine          mdma
##          135          130          120
##          quipazine          mesulergine          fenfluramine
##          119          115          113
##          lsd          2-methyl-5-HT          olanzapine
##          113          110          103
##          flesinoxan          zacopride          gepirone
##          101          93          88
## 5-methoxytryptamine          sulpiride
##          76          70

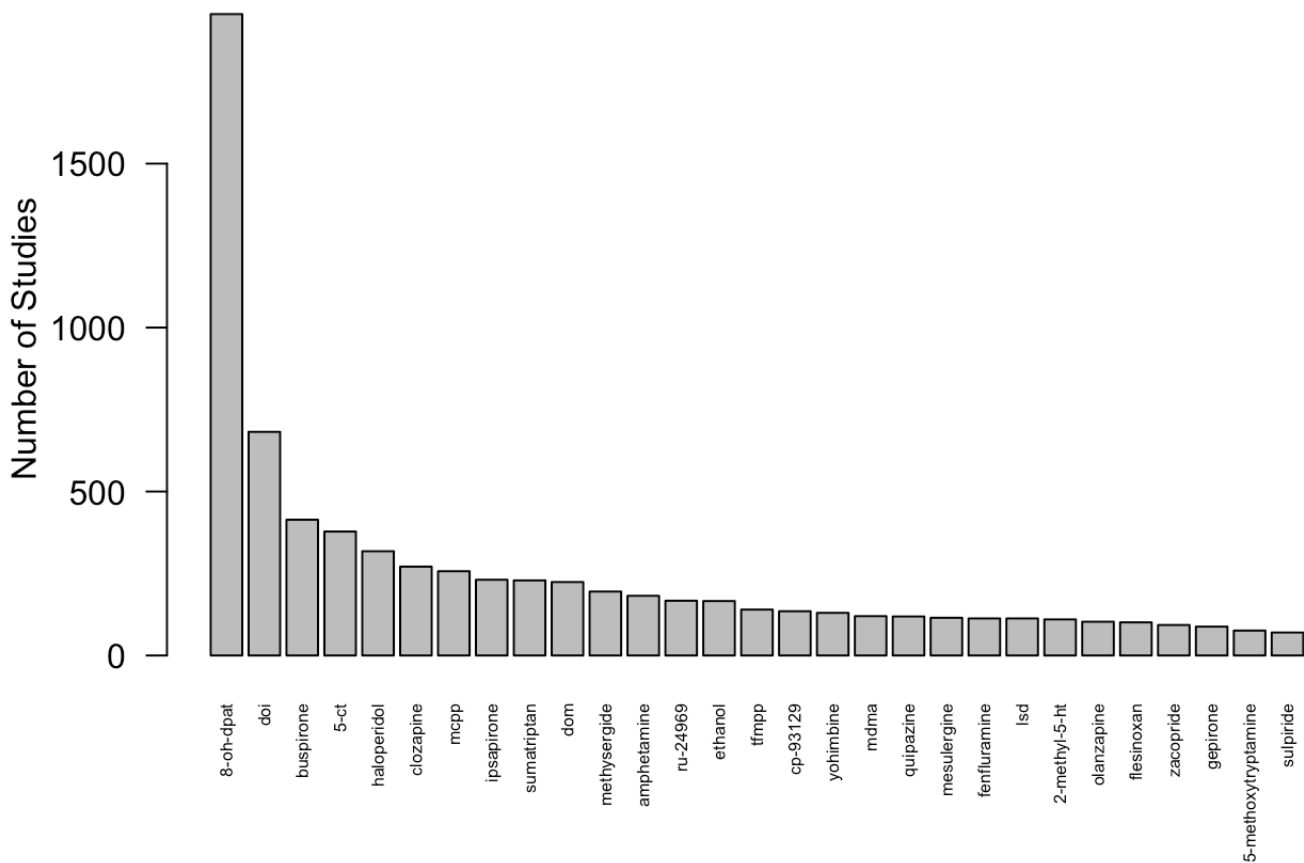
```

```

barplot(agonists, las = 2, cex.names=.5, main = "Agonists Used to Study 5-HT Receptor
s", ylab = "Number of Studies")

```

Agonists Used to Study 5-HT Receptors



```
Antagonists <- strsplit(x$Antagonist, ";")

Antagonists <- unlist(Antagonists)

Antagonists <- gsub(".*\\\\\\\\", "", Antagonists)
Antagonists <- substring(Antagonists, 2)
Antagonists <- gsub('.{2}$', '', Antagonists)
Antagonists <- gsub("\\\\?", "-", Antagonists)
Antagonists <- gsub(" ", "", Antagonists)

#a <- read.csv("Antagonists_List.csv", as.is = TRUE)
#sort(table(a$Antagonist))

(antagonists <- sort(table(Antagonists), decreasing = TRUE))
```

```
## Antagonists
##          ketanserin          way-100635          ritanserin
##              1440              1091              714
##          clozapine          fluoxetine          ondansetron
```


##	542	527	375
##	pindolol	metergoline	haloperidol
##	324	320	318
##	risperidone	nan-190	spiperone
##	302	236	233
##	methiothepin	olanzapine	methysergide
##	223	206	195
##	gr-127935	m1	sb-242084
##	189	188	183
##	mdl-100907	propranolol	sb-269970
##	170	166	148
##	mianserin	tfmpp	way-100135
##	141	140	138
##	imipramine	yohimbine	tropisetron
##	130	130	129
##	granisetron	mesulergine	zacopride
##	117	115	93
##	cyproheptadine	sb-224289	sb-206553
##	90	75	72
##	quetiapine	aripiprazole	bmy-7378
##	70	58	58
##	cyanopindolol	agomelatine	metoclopramide
##	57	51	51
##	rs-102221	mppf	ly-53857
##	49	46	45
##	gr-113808	spiroxatrine	ziprasidone
##	44	42	40
##	chlorpromazine	amitriptyline	mirtazapine
##	39	38	38
##	alprenolol	clomipramine	eltoprazine
##	37	37	34
##	renzapride	sb-271046	sertindole
##	34	34	34
##	gr-125487	sb-399885	isamoltane
##	31	30	28
##	sb-258585	trazodone	sb-216641
##	28	28	27
##	nad-299	cinanserin	brl-15572
##	25	22	21
##	mefway	sb-243213	vortioxetine
##	20	20	20
##	nefazodone	alosetron	lisuride
##	19	18	18
##	sb-200646	sb-204070	lurasidone
##	18	18	17
##	rs-39604	pindobind	ro04-6790
##	17	16	15

##	rs-127445	sdz-216525	fluphenazine
##	15	15	14
##	gr-46611	ergotamine	maprotiline
##	14	13	13
##	memantine	pimavanserin	thioridazine
##	13	13	13
##	tramadol	bromocriptine	sb-258719
##	13	12	12
##	pimozide	sdzser-082	tegaserod
##	11	11	11
##	ar-a000002	idalopirdine	amisulpride
##	10	10	9
##	cariprazine	quinine	sb-204741
##	9	9	9
##	sb-699551	asenapine	luae58054
##	9	8	8
##	norfluoxetine	pizotifen	deramciclane
##	8	8	7
##	mosapride	nortriptyline	sb-215505
##	7	7	7
##	sb-656104-a	sb-742457	uh-301
##	7	7	7
##	xylamidine	zotepine	doxepin
##	7	7	6
##	iodocyanopindolol	loxapine	ro-4368554
##	6	6	6
##	robalzotan	amoxapine	promethazine
##	6	5	5
##	sb-357134	volinanserin	acp-103
##	5	5	4
##	eplivanserin	icil69369	sarpogrelate
##	4	4	4
##	sb-221284	lecozotan	ly-310762
##	4	3	3
##	piboserod	sb-649915	trifluoperazine
##	3	3	3
##	ac-90179	asp-5736	chloroquine
##	2	2	2
##	fr-260010	ly-215840	ly-367265
##	2	2	2
##	ms-245	paliperidone	sb-203186
##	2	2	2
##	sb-236057	apd-125	as-2030680
##	2	1	1
##	as-2674723	cerlapirdine	etoperidone
##	1	1	1
##	fg5983	fluperlapine	hydroxyzine

##	1	1	1
##	intepirdine	lamotrigine	ly-272015
##	1	1	1
##	ly-367642	ly-456219	ly-456220
##	1	1	1
##	o-desmethyiltramadol	oxprenolol	rosarugosa
##	1	1	1
##	rvt-101	s-32212	sb-258741
##	1	1	1

```

antagonists["metergoline"] <- antagonists["metergoline"]/4
antagonists["methiothepin"] <- antagonists["methiothepin"]/3
antagonists["clozapine"] <- antagonists["clozapine"]/3
antagonists["ziprasidone"] <- antagonists["ziprasidone"]/2
antagonists["vortioxetine"] <- antagonists["vortioxetine"]/2
antagonists["spiperone"] <- antagonists["spiperone"]/2
antagonists["sdzser-082"] <- antagonists["sdzser-082"]/2
antagonists["sb-242,084"] <- antagonists["sb-242,084"]/2
antagonists["sb-288,357"] <- antagonists["sb-288,357"]/2
antagonists["sb-206,553"] <- antagonists["sb-206,553"]/2
antagonists["sb-200,646"] <- antagonists["sb-200,646"]/2
antagonists["ritanserine"] <- antagonists["ritanserine"]/2
antagonists["risperidone"] <- antagonists["risperidone"]/2
antagonists["olanzapine"] <- antagonists["olanzapine"]/2
antagonists["mirtazapine"] <- antagonists["mirtazapine"]/2
antagonists["mianserine"] <- antagonists["mianserine"]/2
antagonists["methysergide"] <- antagonists["methysergide"]/2
antagonists["loxapine"] <- antagonists["loxapine"]/2
antagonists["latmepirdine"] <- antagonists["latmepirdine"]/2
antagonists["ketanserine"] <- antagonists["ketanserine"]/2
antagonists["isamoltane"] <- antagonists["isamoltane"]/2
antagonists["imipramine"] <- antagonists["imipramine"]/2
antagonists["fluphenazine"] <- antagonists["fluphenazine"]/2
antagonists["egis-12233"] <- antagonists["egis-12233"]/2
antagonists["chlorpromazine"] <- antagonists["chlorpromazine"]/2
antagonists["aripiprazole"] <- antagonists["aripiprazole"]/2
antagonists["amoxapine "] <- antagonists["amoxapine "]/2
antagonists["amisulpride "] <- antagonists["amisulpride "]/2
antagonists["ziprasidone"] <- antagonists["ziprasidone"]/2

(antagonists <- sort(antagonists, decreasing = TRUE)[1:30])

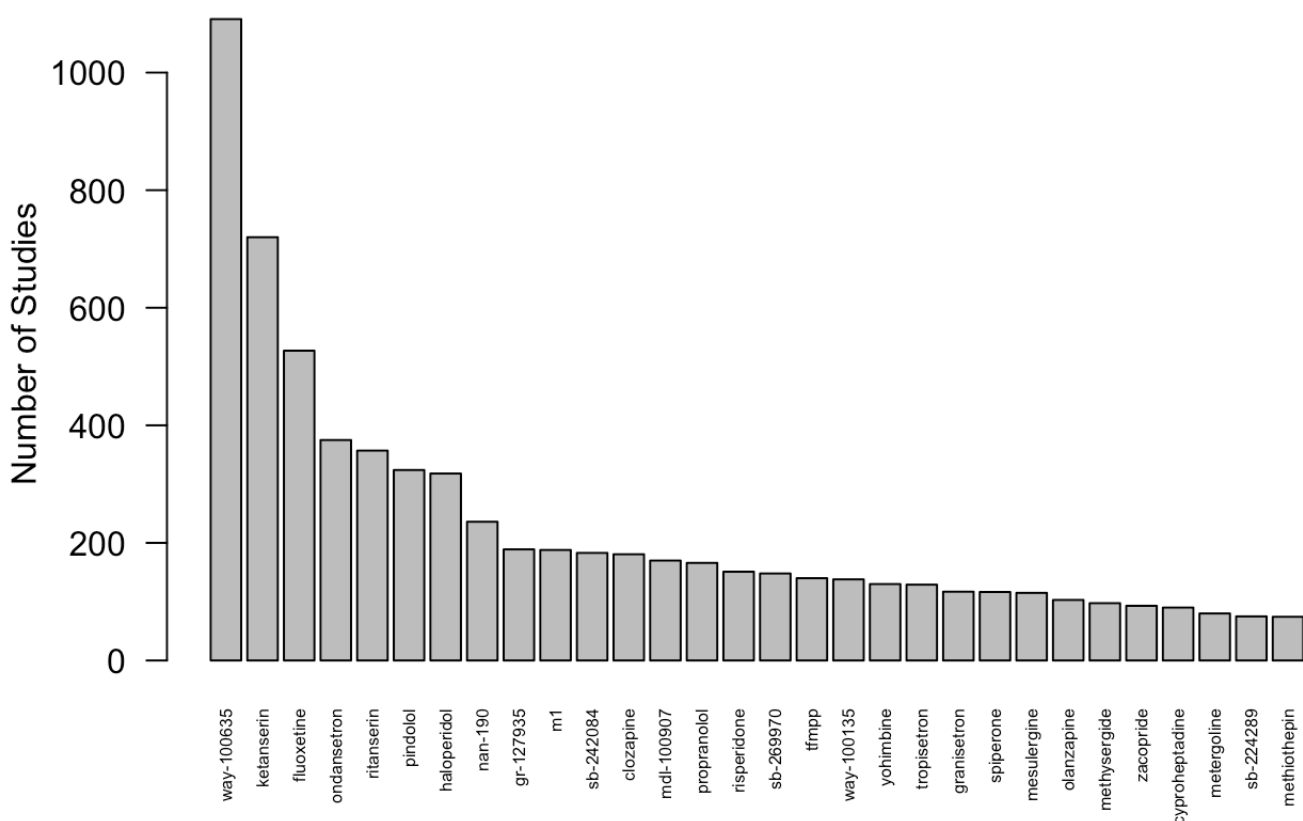
```

##	way-100635	ketanserin	fluoxetine	ondansetron	ritanserin
##	1091.00000	720.00000	527.00000	375.00000	357.00000
##	pindolol	haloperidol	nan-190	gr-127935	m1
##	324.00000	318.00000	236.00000	189.00000	188.00000
##	sb-242084	clozapine	mdl-100907	propranolol	risperidone
##	183.00000	180.66667	170.00000	166.00000	151.00000
##	sb-269970	tfmpp	way-100135	yohimbine	tropisetron
##	148.00000	140.00000	138.00000	130.00000	129.00000
##	granisetron	spiperone	mesulergine	olanzapine	methysergide
##	117.00000	116.50000	115.00000	103.00000	97.50000
##	zacopride	cyproheptadine	metergoline	sb-224289	methiothepin
##	93.00000	90.00000	80.00000	75.00000	74.33333

```
antagonists <- as.matrix(antagonists)
```

```
barplot(antagonists[,1], las = 2, cex.names=.5, main = "Antagonists Used to Study 5-HT Receptors", ylab = "Number of Studies")
```

Antagonists Used to Study 5-HT Receptors



```
Regions <- strsplit(x$Brain_Regions, ";")

Regions <- unlist(Regions)

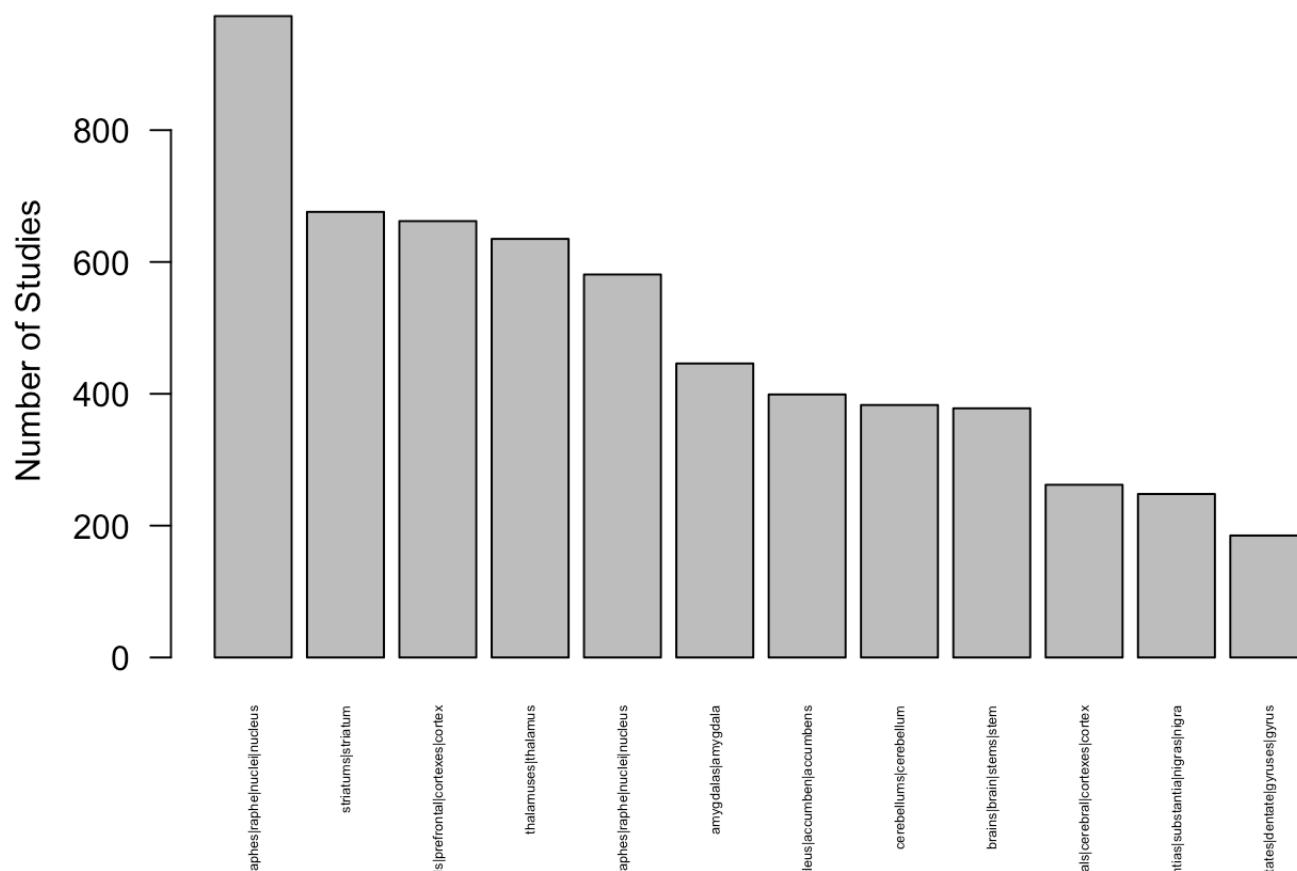
Regions <- gsub(".*i\\)\\(", "", Regions)
Regions <- gsub("\\)\\(-", "", Regions)
Regions <- gsub('.{3}$', '', Regions)
Regions <- gsub("\\\\", "", Regions)
Regions <- gsub("\\\\+", "", Regions)
Regions <- gsub("s\\)", "", Regions)
Regions <- gsub("?\\(", "", Regions)
Regions <- gsub("\\?", "", Regions)

(regions <- sort(table(Regions), decreasing = TRUE)[c(2,3,5:14)])
```

```
## Regions
##          raphes|raphe|nuclei|nucleus
##                                973
##          striatums|striatum
##                                676
##    prefrontals|prefrontal|cortexes|cortex
##                                662
##          thalamuses|thalamus
##                                635
## dorsals|dorsal|raphes|raphe|nuclei|nucleus
##                                581
##          amygdalas|amygdala
##                                446
##    nuclei|nucleus|accumben|accumbens
##                                399
##          cerebellums|cerebellum
##                                383
##          brains|brain|stems|stem
##                                378
##    cerebrals|cerebral|cortexes|cortex
##                                262
##    substantias|substantia|nigras|nigra
##                                248
##          dentates|dentate|gyruses|gyrus
##                                185
```

```
barplot(regions, las = 2, cex.names=.4, main = "Brain Regions Studied in Conjunction
with 5-HT Receptors", ylab = "Number of Studies")
```

Brain Regions Studied in Conjunction with 5-HT Receptors



```
Topics <- strsplit(x$Topic_Spec, ";")

Topics <- unlist(Topics)

Topics <- gsub(".*\\\\\\\\", "", Topics)
Topics <- substring(Topics, 2)
Topics <- gsub('.{2}$', '', Topics)
Topics <- gsub("social dominance", "soc. dom.", Topics)

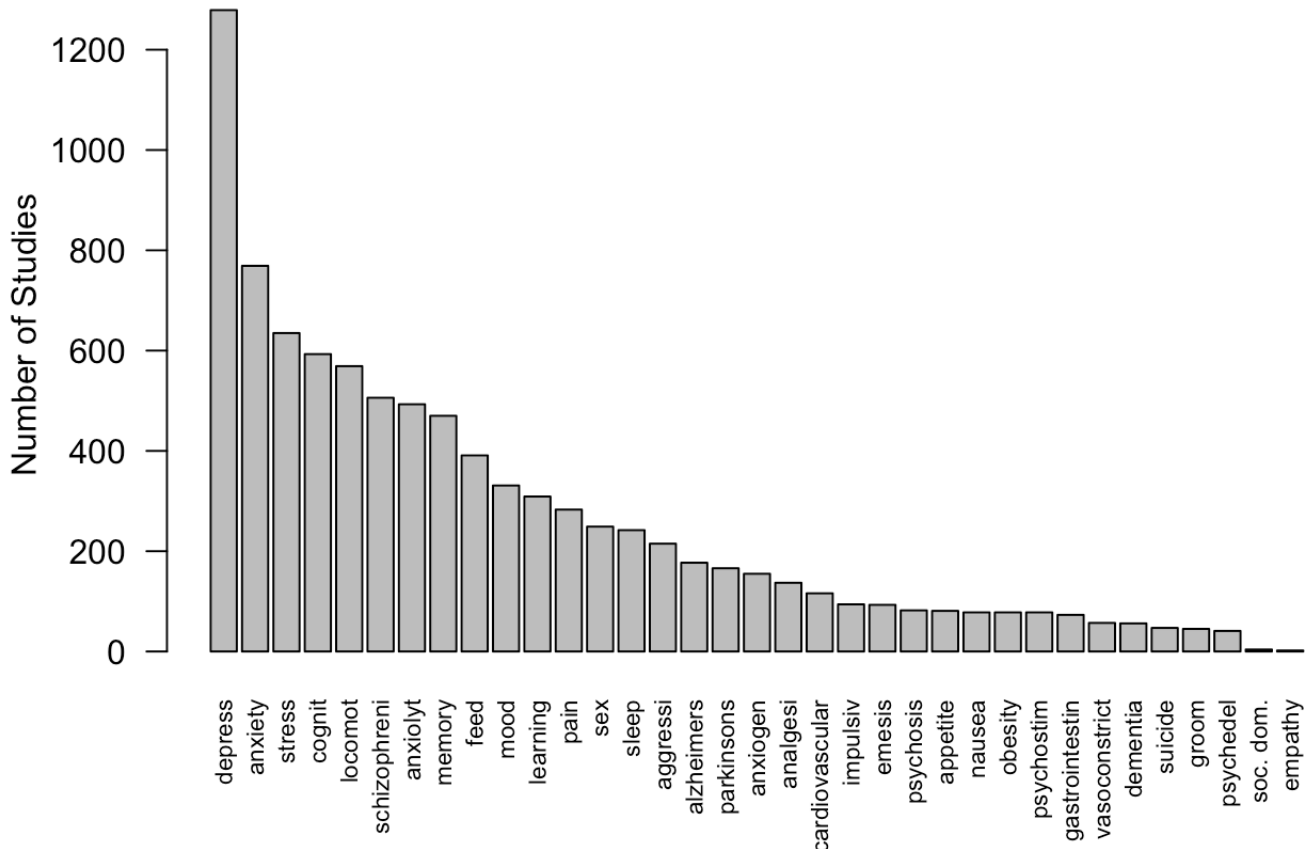
(topics <- sort(table(Topics), decreasing = TRUE))
```

Topics

##	depress	anxiety	stress	cognit	locomot
##	1279	769	635	593	569
##	schizophreni	anxiolyt	memory	feed	mood
##	506	493	470	391	331
##	learning	pain	sex	sleep	aggressi
##	309	283	249	242	215
##	alzheimers	parkinsons	anxiogen	analgesi	cardiovascular
##	177	166	155	137	116
##	impulsiv	emesis	psychosis	appetite	nausea
##	94	93	82	81	78
##	obesity	psychostim	gastrointestin	vasoconstrict	dementia
##	78	78	73	57	56
##	suicide	groom	psychedel	soc. dom.	empathy
##	47	45	41	4	2

```
barplot(topics, las = 2, cex.names=.7, main = "Topics Studied with 5-HT Receptors", y
lab = "Number of Studies")
```

Topics Studied with 5-HT Receptors

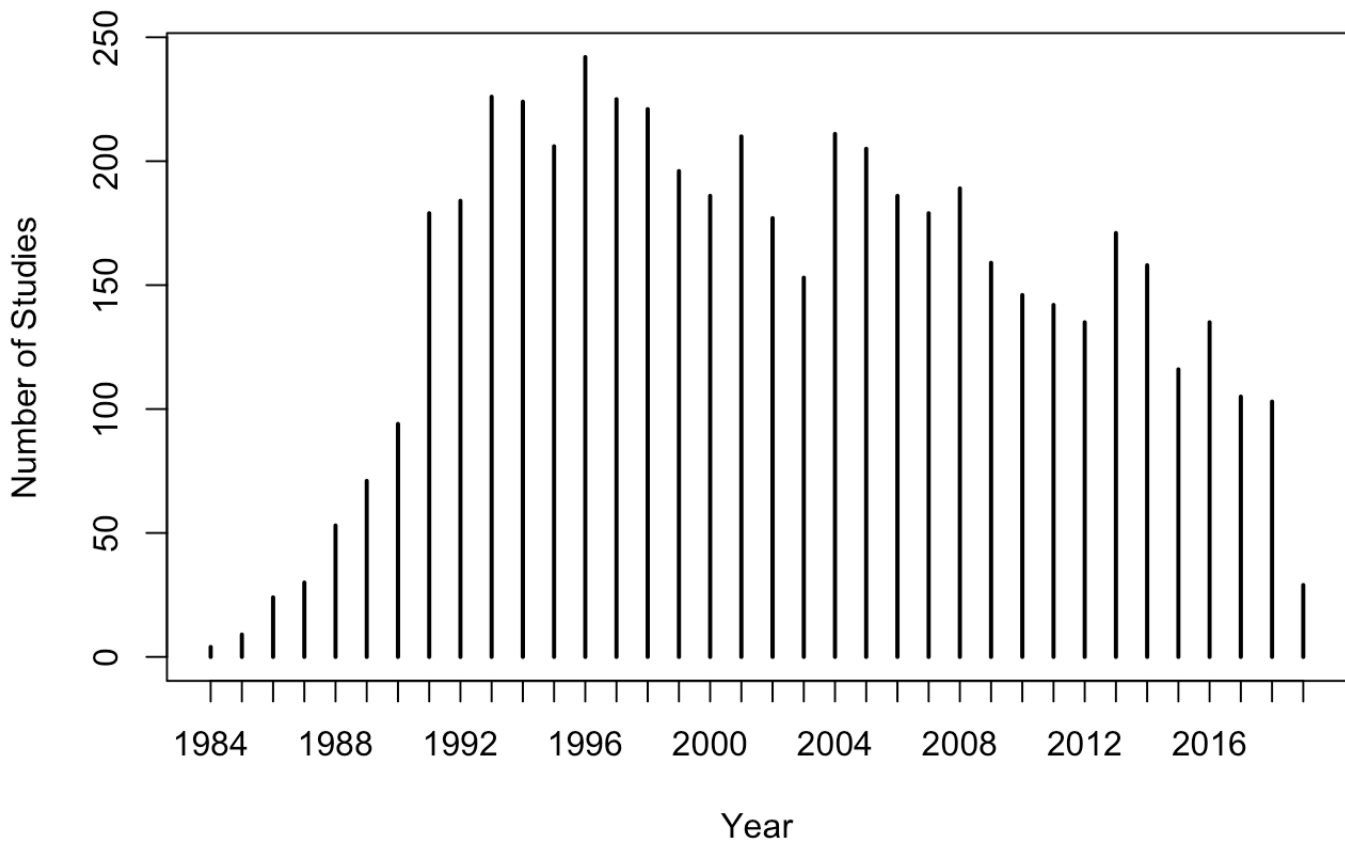


5-HT1A

```
x1a <- subset(x, grepl("5ht1a", x$Receptor))
```

```
plot(table(x1a$Year), main = "Number of 5-HT1A Studies by Year", xlab = "Year", ylab = "Number of Studies")
```

Number of 5-HT1A Studies by Year



```
sort(table(x1a$Year), decreasing = TRUE)
```

```
##
## 1996 1993 1997 1994 1998 2004 2001 1995 2005 1999 2008 2000 2006 1992 1991
## 242 226 225 224 221 211 210 206 205 196 189 186 186 184 179
## 2007 2002 2013 2009 2014 2003 2010 2011 2012 2016 2015 2017 2018 1990 1989
## 179 177 171 159 158 153 146 142 135 135 116 105 103 94 71
## 1988 1987 2019 1986 1985 1984
## 53 30 29 24 9 4
```



```
Methods1a <- strsplit(x1a$Methods, ";")

Methods1a <- unlist(Methods1a)

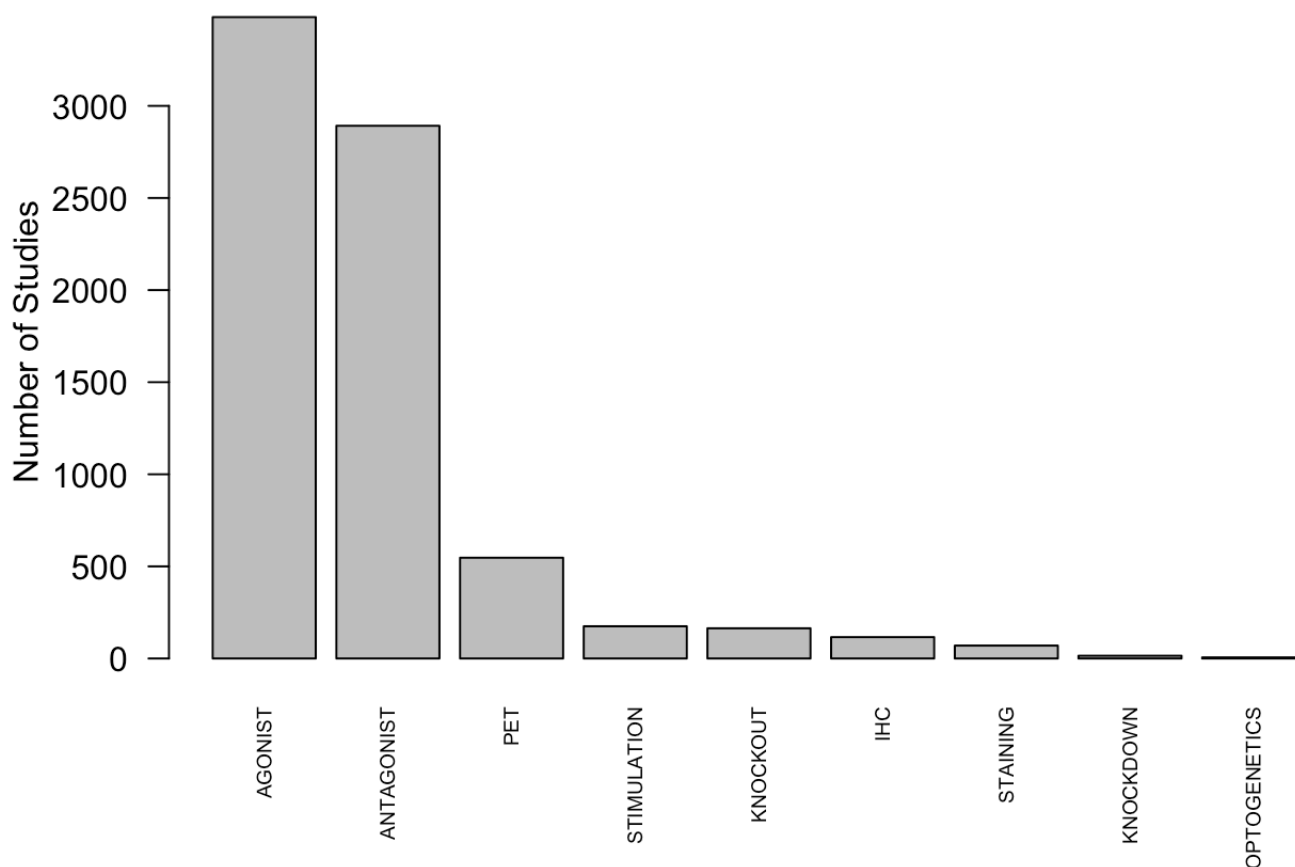
Methods1a <- gsub("Method.", "", Methods1a)
Methods1a <- gsub("IMMUNOHISTOCHEMISTRY", "IHC", Methods1a)

(methods1a <- sort(table(Methods1a), decreasing = TRUE))
```

```
## Methods1a
##      AGONIST      ANTAGONIST      PET      STIMULATION      KNOCKOUT
##      3482      2892      547      175      164
##      IHC      STAINING      KNOCKDOWN      OPTOGENETICS
##      116      70      15      6
```

```
barplot(methods1a, las = 2, cex.names=.6, main = "Methods Used to Study 5-HT1A", ylab = "Number of Studies")
```

Methods Used to Study 5-HT1A



```
Speciesla <- strsplit(xla$Species, ";")

Speciesla <- unlist(Speciesla)

Speciesla <- gsub(".*b","", Speciesla)
Speciesla <- gsub('.{2}$', '', Speciesla)
Speciesla <- gsub(".*\\(", "", Speciesla)
Speciesla <- gsub('.{1}$', '', Speciesla)

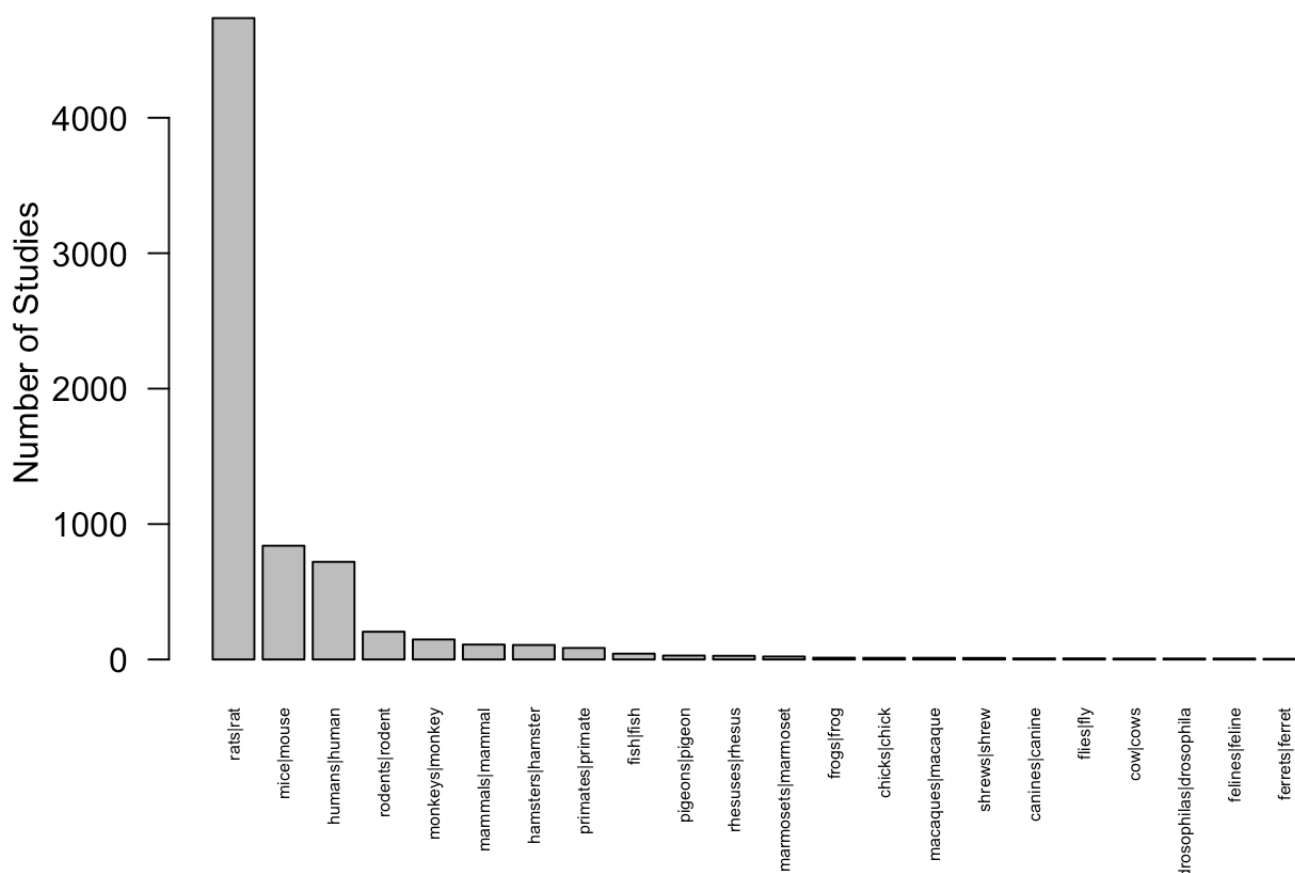
#sort(table(Speciesla), decreasing = TRUE)

(speciesla <- sort(table(Speciesla), decreasing = TRUE)[c(1, 24, 31, 54, 62, 71, 73,
83, 108, 120, 124, 133, 154, 163, 166, 168, 181, 185, 190, 191, 192, 207)])
```

```
## Speciesla
##          rats|rat          mice|mouse          humans|human
##          4736          840          721
##      rodents|rodent      monkeys|monkey      mammals|mammal
##          205          148          110
##      hamsters|hamster      primates|primate          fish|fish
##          107          85          43
##      pigeons|pigeon      rhesuses|rhesus      marmosets|marmoset
##          29          27          22
##          frogs|frog          chicks|chick      macaques|macaque
##          13          12          12
##      shrews|shrew      canines|canine          flies|fly
##          11          8          8
##          cow|cows      drosophilas|drosophila      felines|feline
##          7          7          7
##      ferrets|ferret
##          5
```

```
barplot(speciesla, las = 2, cex.names=.5, main = "Species Used to Study 5-HT1A", ylab
= "Number of Studies")
```

Species Used to Study 5-HT1A



```

Agonists1a <- strsplit(xla$Agonist, ";")

Agonists1a <- unlist(Agonists1a)
Agonists1a <- gsub(".*\\\\\\", "", Agonists1a)
Agonists1a <- substring(Agonists1a, 2)
Agonists1a <- gsub('.{2}$', '', Agonists1a)
Agonists1a <- gsub("\\\\?", "-", Agonists1a)
Agonists1a <- gsub(" ", "", Agonists1a)

Agonists1a <- gsub("5-carboxamidotryptamine", "5-ct", Agonists1a)

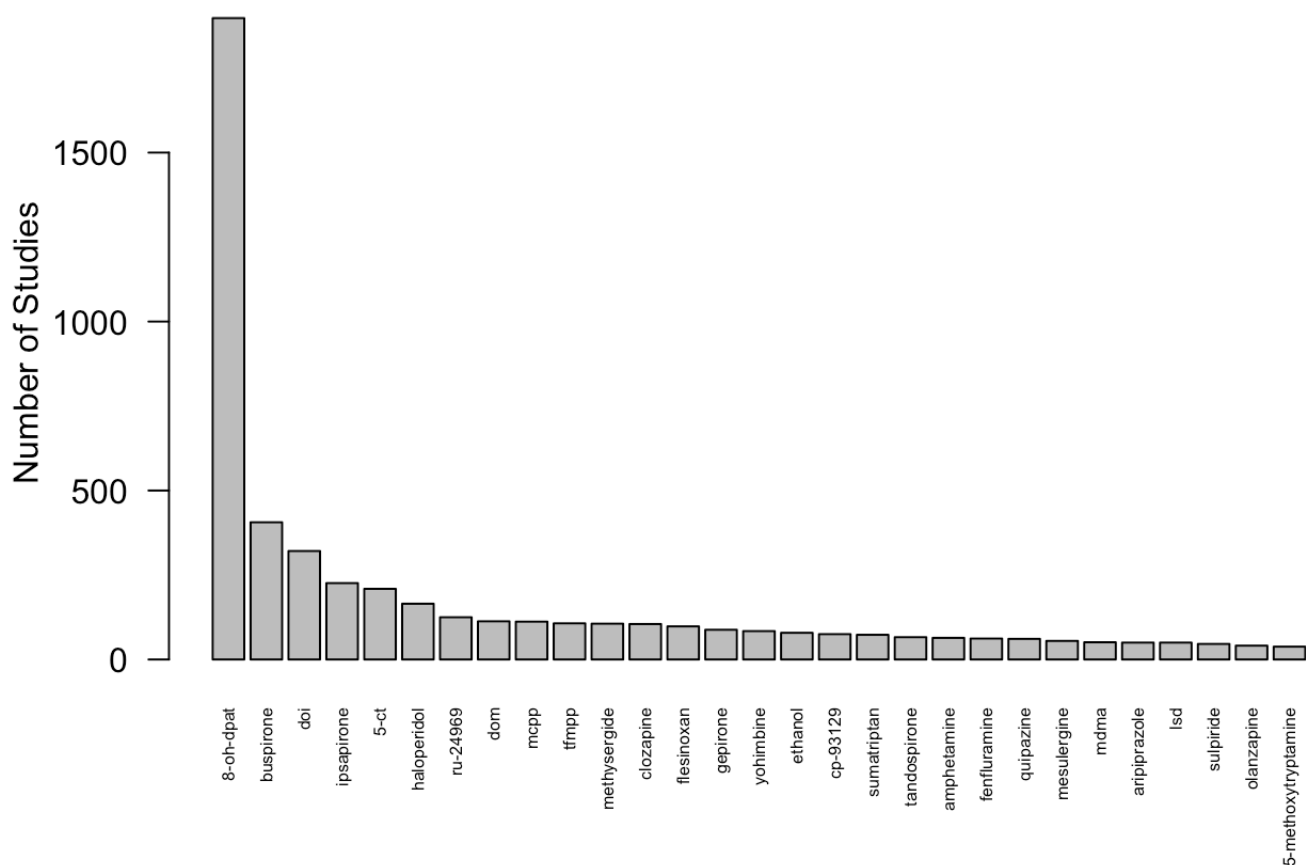
#sort(table(Agonists1a), decreasing = TRUE)
(agonists1a <- sort(table(Agonists1a), decreasing = TRUE)[c(1, 3:30)])

```

```
## Agonistsla
##      8-oh-dpat      buspirone      doi
##      1898      406      321
##      ipsapirone      5-ct      haloperidol
##      226      209      165
##      ru-24969      dom      mcpp
##      125      113      112
##      tfmpp      methysergide      clozapine
##      107      106      105
##      flesinoxan      gepirone      yohimbine
##      98      88      84
##      ethanol      cp-93129      sumatriptan
##      79      75      73
##      tandospirone      amphetamine      fenfluramine
##      66      64      62
##      quipazine      mesulergine      mdma
##      61      55      51
##      aripiprazole      lsd      sulpiride
##      50      50      46
##      olanzapine 5-methoxytryptamine
##      41      38
```

```
barplot(agonistsla, las = 2, cex.names=.5, main = "Agonists Used to Study 5-HT1A", ylab = "Number of Studies")
```

Agonists Used to Study 5-HT1A



```
Antagonists1a <- strsplit(xla$Antagonist, ";")
```

```
Antagonists1a <- unlist(Antagonists1a)
```

```
Antagonists1a <- gsub(".*\\\\\\\\", "", Antagonists1a)
```

```
Antagonists1a <- substring(Antagonists1a, 2)
```

```
Antagonists1a <- gsub('.{2}$', '', Antagonists1a)
```

```
Antagonists1a <- gsub("\\?", "-", Antagonists1a)
```

```
Antagonists1a <- gsub(" ", "", Antagonists1a)
```

```
#sort(table(Antagonists1a), decreasing = TRUE)
```

```
(antagonists1a <- sort(table(Antagonists1a), decreasing = TRUE))
```

```
## Antagonists1a
```

##	way-100635	ketanserin	ritanserin	fluoxetine
##	1083	686	412	358
##	pindolol	nan-190	clozapine	metergoline
##	288	228	210	196

##	spiperone	haloperidol	methiothepin	way-100135
##	178	165	143	136
##	propranolol	risperidone	gr-127935	tfmpp
##	131	128	113	107
##	methysergide	ondansetron	yohimbine	imipramine
##	106	92	84	82
##	olanzapine	sb-269970	m1	mianserin
##	82	73	68	59
##	bmy-7378	mesulergine	aripiprazole	mppf
##	58	55	50	46
##	cyproheptadine	mdl-100907	spiroxatrine	quetiapine
##	42	42	41	38
##	sb-242084	sb-224289	cyanopindolol	alprenolol
##	38	37	34	33
##	tropisetron	eltoprazine	ziprasidone	zacopride
##	33	32	31	28
##	granisetron	ly-53857	clomipramine	nad-299
##	27	27	25	25
##	mefway	amitriptyline	vortioxetine	chlorpromazine
##	20	19	18	16
##	pindobind	sb-216641	lurasidone	mirtazapine
##	16	16	15	15
##	sdz-216525	trazodone	brl-15572	isamoltane
##	15	15	14	14
##	sb-206553	cinanserin	renzapride	metoclopramide
##	14	11	11	10
##	bromocriptine	cariprazine	gr-113808	gr-125487
##	9	9	9	8
##	lisuride	sertindole	gr-46611	maprotiline
##	8	8	7	7
##	rs-102221	sb-258719	thioridazine	tramadol
##	7	7	7	7
##	uh-301	pimozide	quinine	robalzotan
##	7	6	6	6
##	rs-39604	sb-399885	sb-699551	asenapine
##	6	6	6	5
##	nefazodone	ro04-6790	sb-258585	agomelatine
##	5	5	5	4
##	fluphenazine	norfluoxetine	rs-127445	sb-656104-a
##	4	4	4	4
##	amoxapine	ergotamine	lecozotan	nortriptyline
##	3	3	3	3
##	pizotifen	sb-200646	sb-204070	sb-204741
##	3	3	3	3
##	sb-649915	sdzser-082	xylamidine	zotepine
##	3	3	3	3
##	amisulpride	deramciclane	doxepin	icil69369

##	2	2	2	2
##	iodocyanopindolol	loxapine	paliperidone	sb-215505
##	2	2	2	2
##	sb-221284	sb-243213	sb-271046	alosetron
##	2	2	2	1
##	ar-a000002	etoperidone	fg5983	fluperlapine
##	1	1	1	1
##	fr-260010	hydroxyzine	lamotrigine	ly-215840
##	1	1	1	1
##	memantine	oxprenolol	pimavanserin	promethazine
##	1	1	1	1
##	ro-4368554	sarpogrelate	sb-203186	sb-258741
##	1	1	1	1
##	sb-357134	tegaserod	volinanserin	
##	1	1	1	

```

antagonistsla["metergoline"] <- antagonistsla["metergoline"]/4
antagonistsla["methiothepin"] <- antagonistsla["methiothepin"]/3
antagonistsla["clozapine"] <- antagonistsla["clozapine"]/3
antagonistsla["ziprasidone"] <- antagonistsla["ziprasidone"]/2
antagonistsla["vortioxetine"] <- antagonistsla["vortioxetine"]/2
antagonistsla["spiperone"] <- antagonistsla["spiperone"]/2
antagonistsla["sdzser-082"] <- antagonistsla["sdzser-082"]/2
antagonistsla["sb-242,084"] <- antagonistsla["sb-242,084"]/2
antagonistsla["sb-288,357"] <- antagonistsla["sb-288,357"]/2
antagonistsla["sb-206,553"] <- antagonistsla["sb-206,553"]/2
antagonistsla["sb-200,646"] <- antagonistsla["sb-200,646"]/2
antagonistsla["ritanserine"] <- antagonistsla["ritanserine"]/2
antagonistsla["risperidone"] <- antagonistsla["risperidone"]/2
antagonistsla["olanzapine"] <- antagonistsla["olanzapine"]/2
antagonistsla["mirtazapine"] <- antagonistsla["mirtazapine"]/2
antagonistsla["mianserine"] <- antagonistsla["mianserine"]/2
antagonistsla["methysergide"] <- antagonistsla["methysergide"]/2
antagonistsla["loxapine"] <- antagonistsla["loxapine"]/2
antagonistsla["latmepirdine"] <- antagonistsla["latmepirdine"]/2
antagonistsla["ketanserine"] <- antagonistsla["ketanserine"]/2
antagonistsla["isamoltane"] <- antagonistsla["isamoltane"]/2
antagonistsla["imipramine"] <- antagonistsla["imipramine"]/2
antagonistsla["fluphenazine"] <- antagonistsla["fluphenazine"]/2
antagonistsla["egis-12233"] <- antagonistsla["egis-12233"]/2
antagonistsla["chlorpromazine"] <- antagonistsla["chlorpromazine"]/2
antagonistsla["aripiprazole"] <- antagonistsla["aripiprazole"]/2
antagonistsla["amoxapine "] <- antagonistsla["amoxapine "]/2
antagonistsla["amisulpride "] <- antagonistsla["amisulpride "]/2
antagonistsla["ziprasidone"] <- antagonistsla["ziprasidone"]/2

(antagonistsla <- sort(antagonistsla, decreasing = TRUE)[1:30])

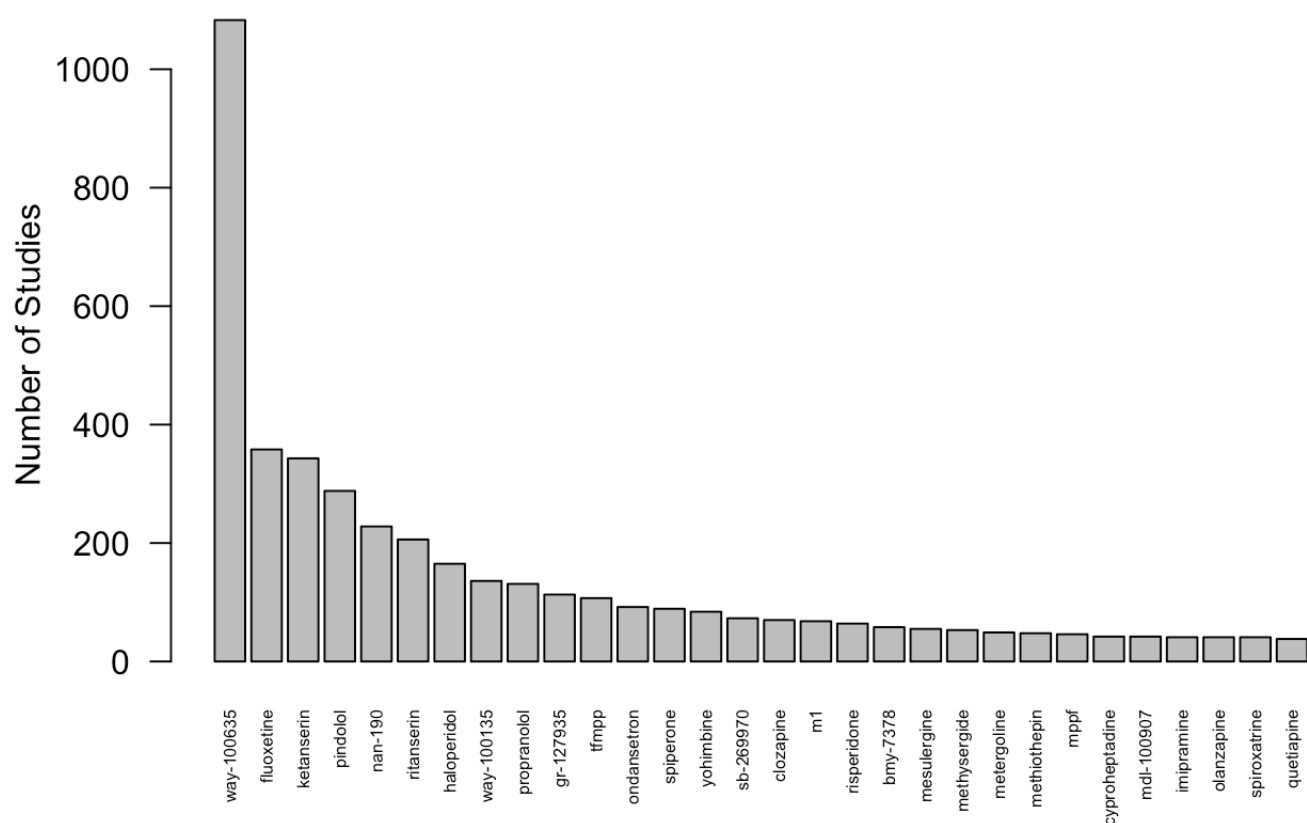
```

##	way-100635	fluoxetine	ketanserine	pindolol	nan-190
##	1083.00000	358.00000	343.00000	288.00000	228.00000
##	ritanserine	haloperidol	way-100135	propranolol	gr-127935
##	206.00000	165.00000	136.00000	131.00000	113.00000
##	tfmpp	ondansetron	spiperone	yohimbine	sb-269970
##	107.00000	92.00000	89.00000	84.00000	73.00000
##	clozapine	m1	risperidone	bmy-7378	mesulergine
##	70.00000	68.00000	64.00000	58.00000	55.00000
##	methysergide	metergoline	methiothepin	mpf	cypheptadine
##	53.00000	49.00000	47.66667	46.00000	42.00000
##	mdl-100907	imipramine	olanzapine	spiroxatrine	quetiapine
##	42.00000	41.00000	41.00000	41.00000	38.00000


```
antagonistsla <- as.matrix(antagonistsla)
```

```
barplot(antagonistsla[,1], las = 2, cex.names=.5, main = "Antagonists Used to Study 5-HT1A Receptors", ylab = "Number of Studies")
```

Antagonists Used to Study 5-HT1A Receptors



```
Regionsla <- strsplit(xla$Brain_Regions, ";")

Regionsla <- unlist(Rregionsla)

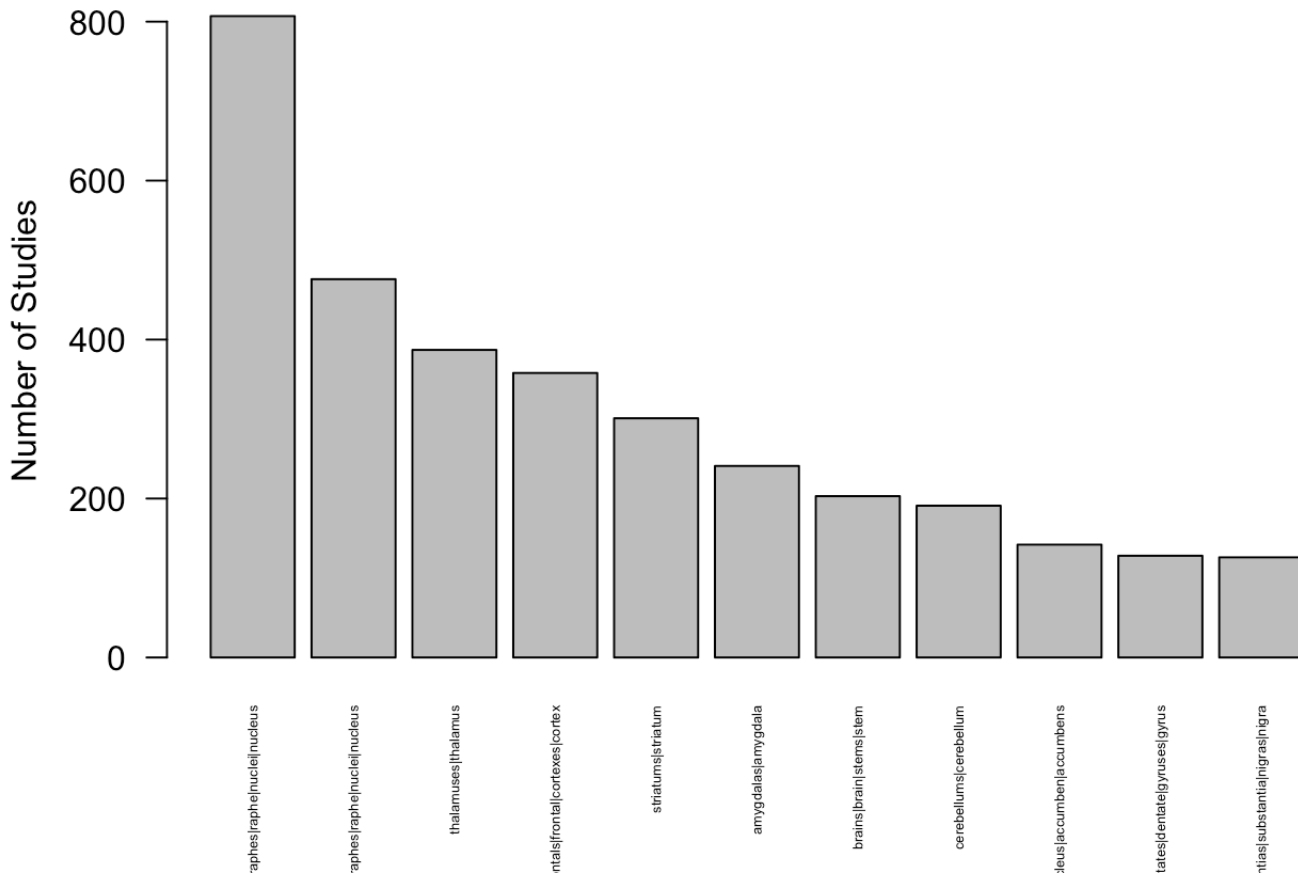
Regionsla <- gsub(".*i\\)\\(", "", Regionsla)
Regionsla <- gsub("\\)\\(-", "", Regionsla)
Regionsla <- gsub('.{3}$', '', Regionsla)
Regionsla <- gsub("\\\\", "", Regionsla)
Regionsla <- gsub("\\+", "", Regionsla)
Regionsla <- gsub("s\\)", "", Regionsla)
Regionsla <- gsub("?\\(", "", Regionsla)
Regionsla <- gsub("\\?", "", Regionsla)

#sort(table(Rregionsla), decreasing = TRUE)
(regionsla <- sort(table(Rregionsla), decreasing = TRUE)[c(2:4,6:13)])
```

```
## Regionsla
##          raphes|raphe|nuclei|nucleus
##                                807
## dorsals|dorsal|raphes|raphe|nuclei|nucleus
##                                476
##          thalamuses|thalamus
##                                387
## pres|pre|frontals|frontal|cortexes|cortex
##                                358
##          striatums|striatum
##                                301
##          amygdalas|amygdala
##                                241
##          brains|brain|stems|stem
##                                203
##          cerebellums|cerebellum
##                                191
##          nuclei|nucleus|accumben|accumbens
##                                142
##          dentates|dentate|gyruses|gyrus
##                                128
##          substantias|substantia|nigras|nigra
##                                126
```

```
barplot(regionsla, las = 2, cex.names=.4, main = "Brain Regions Studied in Conjunction with 5-HT1A", ylab = "Number of Studies")
```

Brain Regions Studied in Conjunction with 5-HT1A



```
Topics1a <- strsplit(x1a$Topic_Spec, ";")

Topics1a <- unlist(Topics1a)

Topics1a <- gsub(".*\\\\\\\\", "", Topics1a)
Topics1a <- substring(Topics1a, 2)
Topics1a <- gsub('.{2}$', '', Topics1a)
Topics1a <- gsub("social dominance", "soc. dom.", Topics1a)

(topics1a <- sort(table(Topics1a), decreasing = TRUE))
```

```
## Topics1a
##      depress      anxiety      stress      anxiolyt      locomot
##      821          486          401          365          299
##      cognit      feed      schizophreni      memory      mood
##      256          229          218          215          192
##      sex      learning      sleep      aggressi      pain
##      178          140          140          128          106
##      anxiogen      parkinsons      cardiovascular      analgesi      alzheimers
##      94          94          71          66          53
##      impulsiv      dementia      suicide      psychosis      psychostim
##      35          34          33          28          27
##      groom      appetite      vasoconstrict      nausea      gastrointestin
##      24          22          19          15          13
##      obesity      psychedel      emesis      soc. dom.
##      11          7          6          1
```

```
barplot(topics1a, las = 2, cex.names=.7, main = "Topics Studied with 5-HT1A", ylab =
"Number of Studies")
```

Topics Studied with 5-HT1A

