

# 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

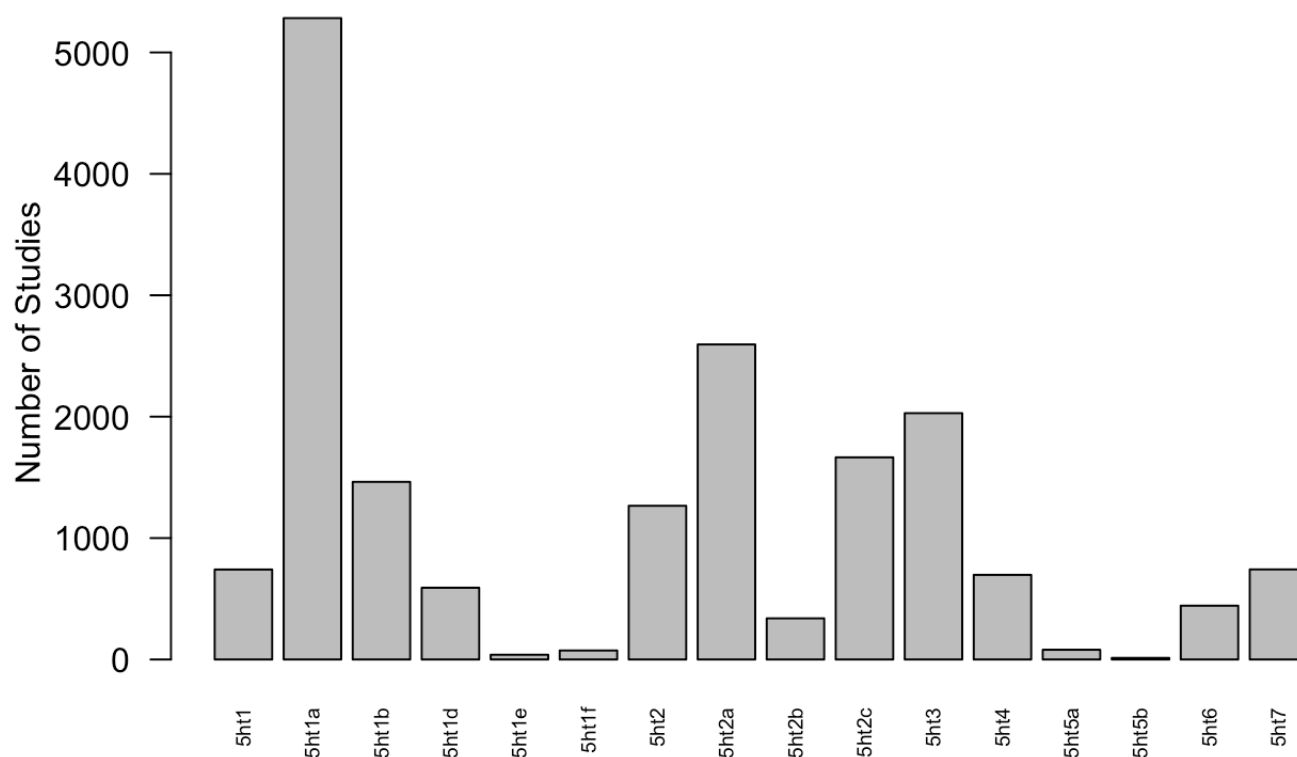
```

```

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

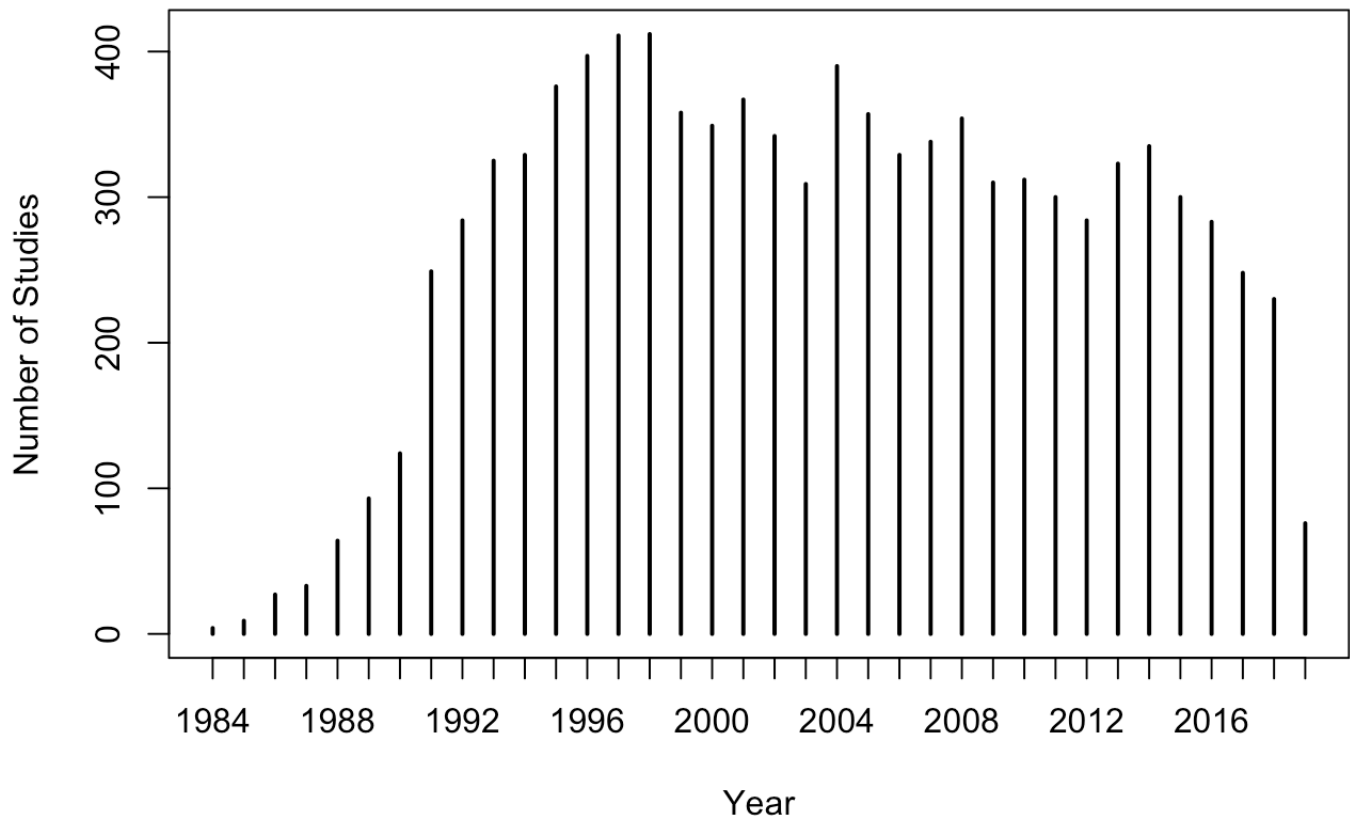
```

## 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)

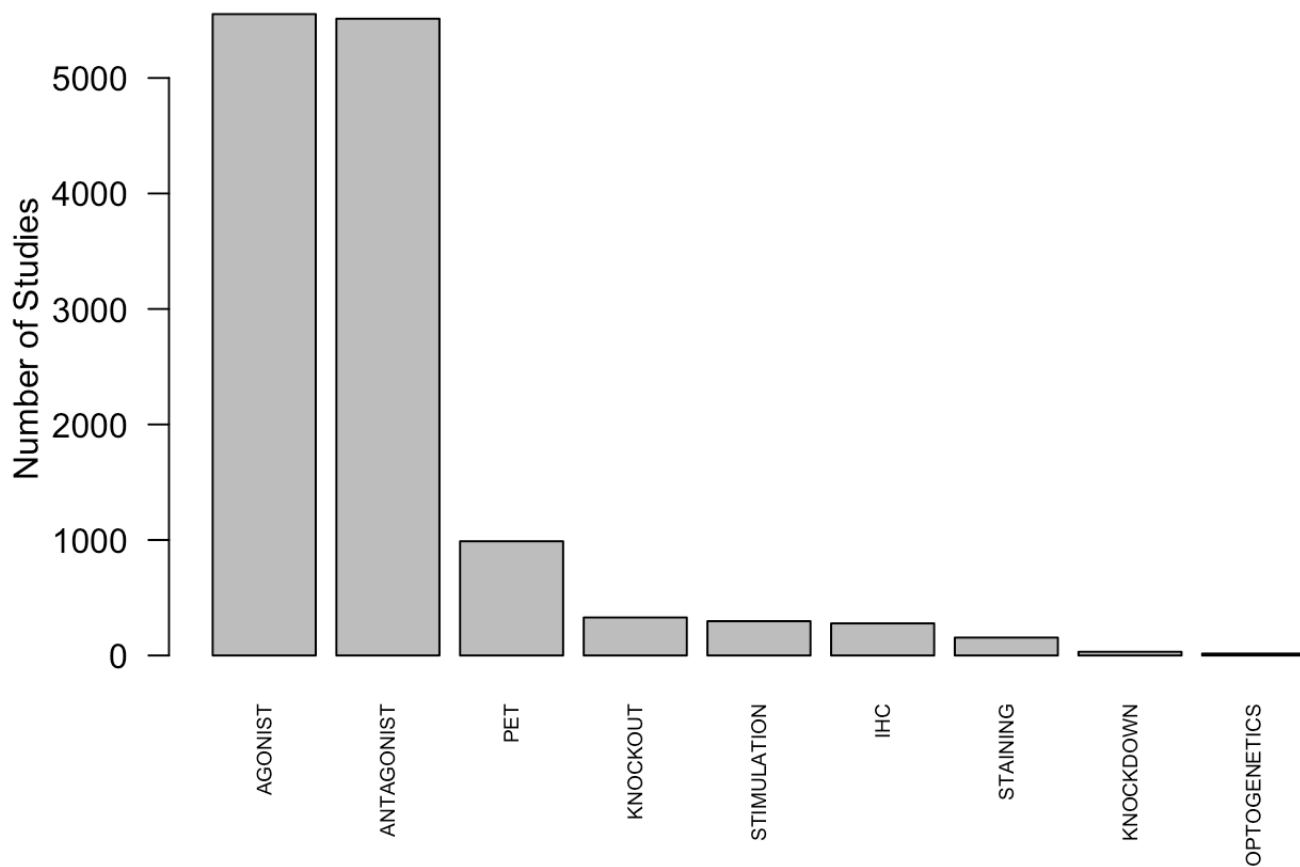
Methods <- gsub("Method.", "", Methods)
Methods <- gsub("IMMUNOHISTOCHEMISTRY", "IHC", Methods)

(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)

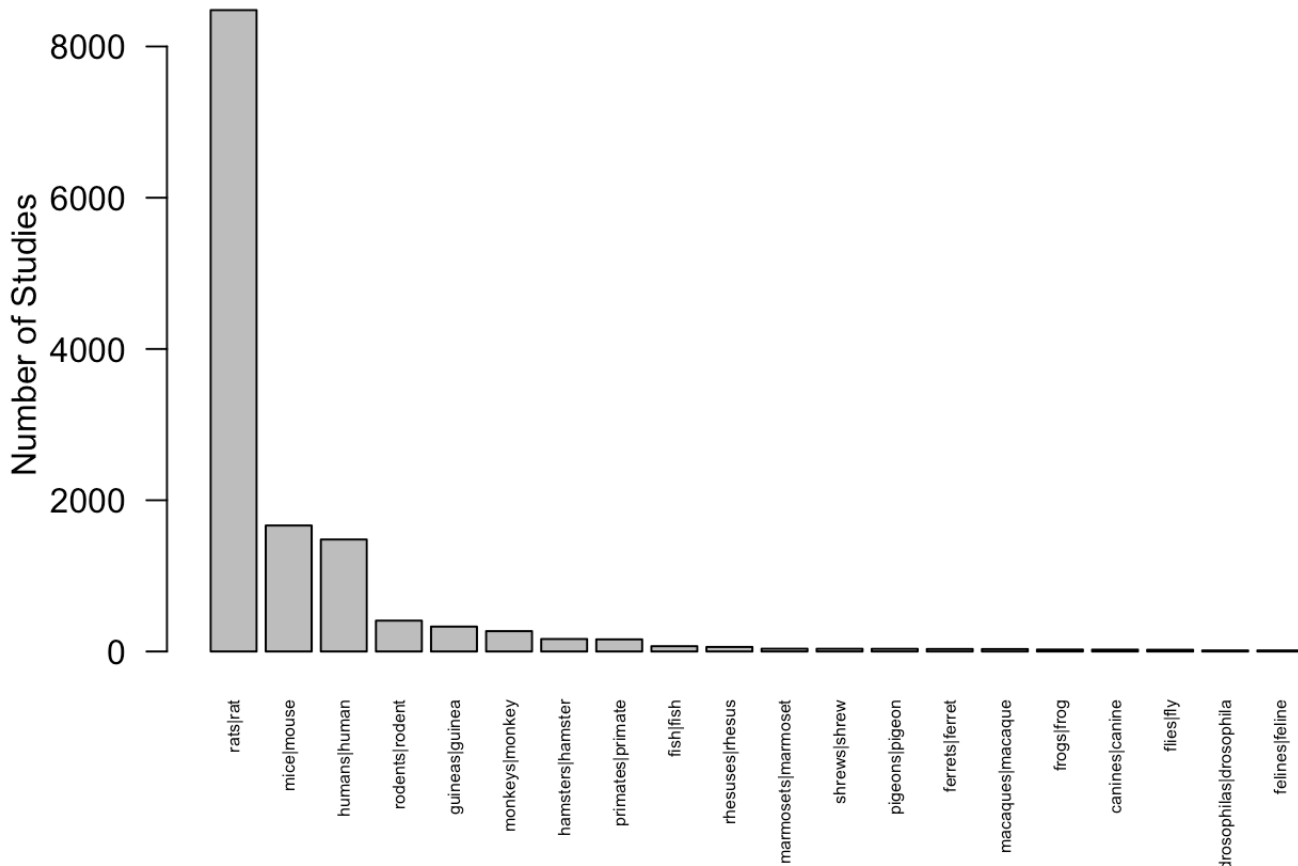
Species <- gsub(".*b", "", Species)
Species <- gsub('.{2}$', '', Species)
Species <- gsub(".*\\(", "", Species)
Species <- gsub('.{1}$', '', Species)

(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

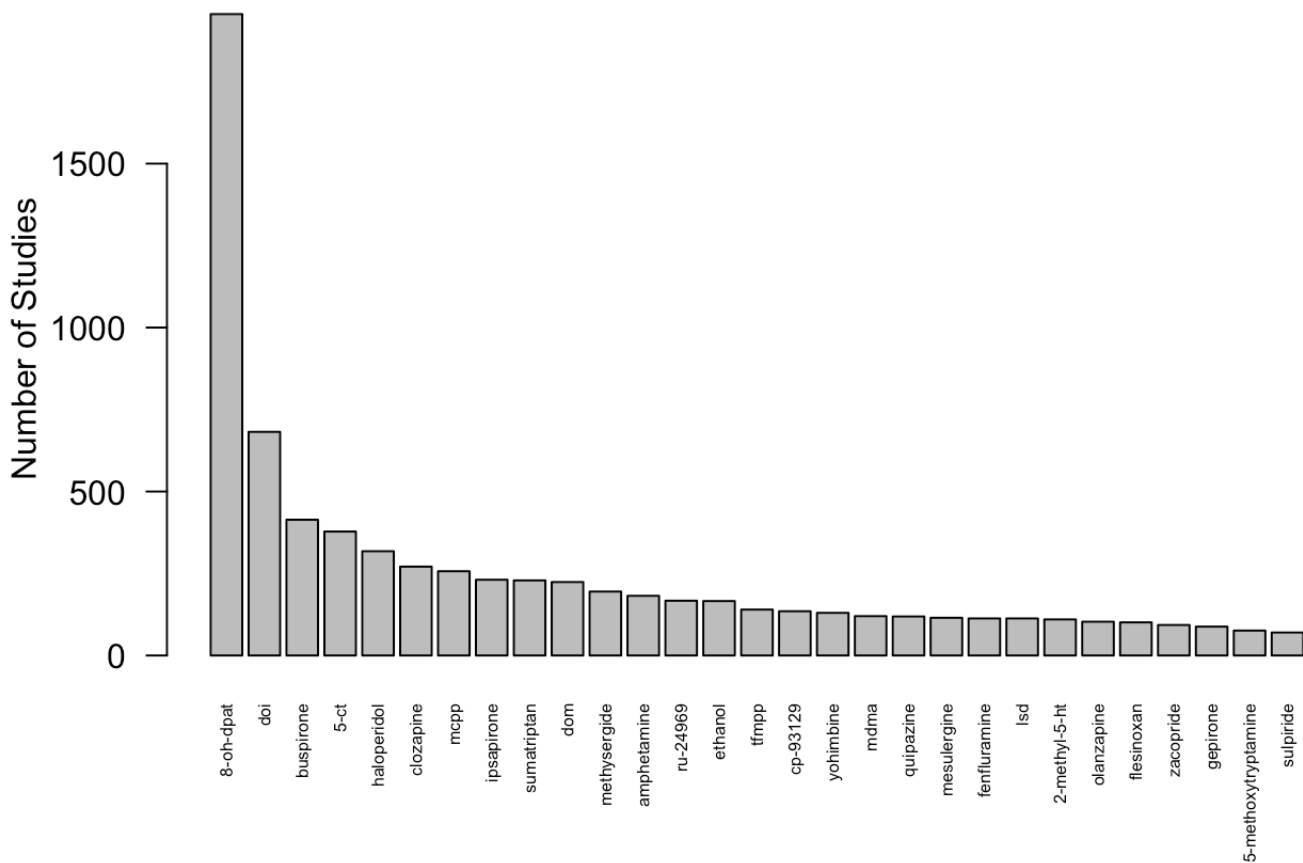
```

```

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))

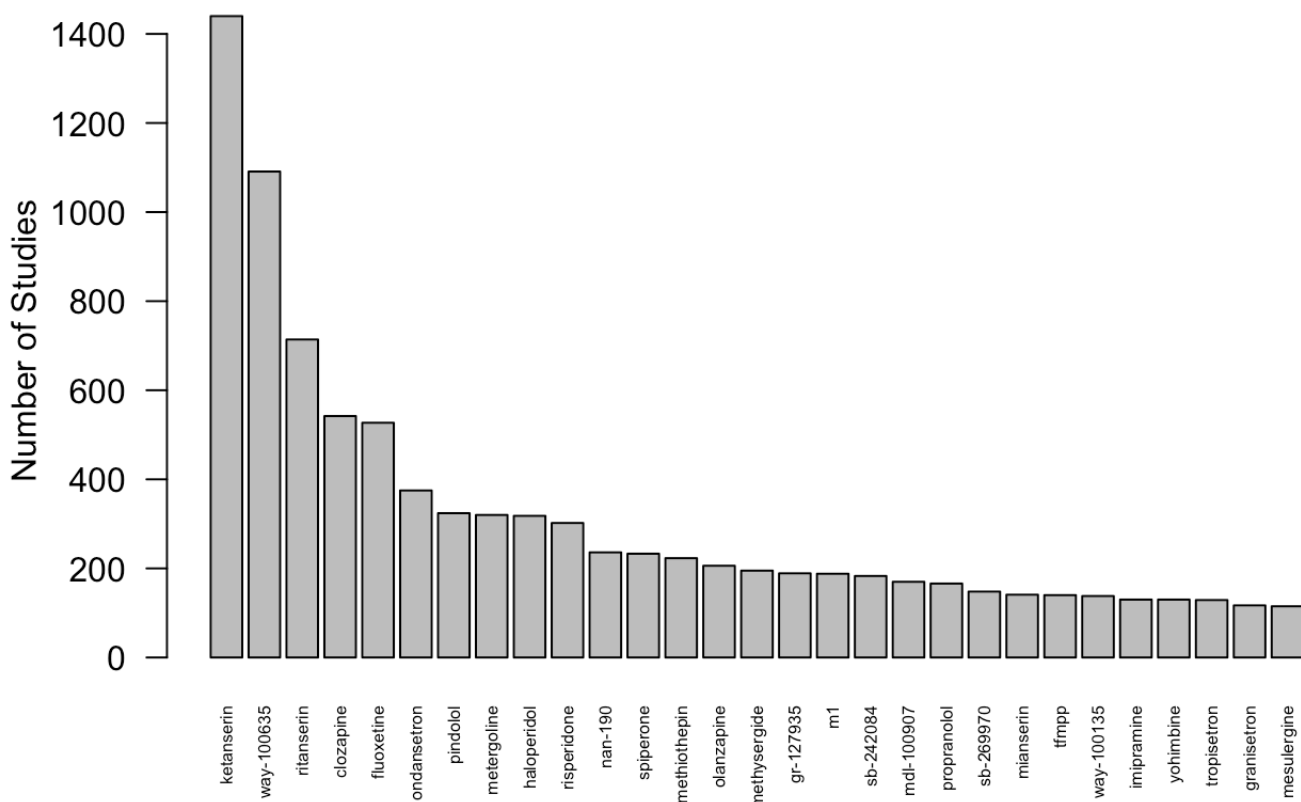
(antagonists <- sort(table(Antagonists), decreasing = TRUE)[c(1:29)])
```



```
## Antagonists
## ketanserin way-100635 ritanserin clozapine fluoxetine
## 1440 1091 714 542 527
## ondansetron pindolol metergoline haloperidol risperidone
## 375 324 320 318 302
## nan-190 spiperone methiothepin olanzapine methysergide
## 236 233 223 206 195
## gr-127935 m1 sb-242084 mdl-100907 propranolol
## 189 188 183 170 166
## sb-269970 mianserin tfmpp way-100135 imipramine
## 148 141 140 138 130
## yohimbine tropisetron granisetron mesulergine
## 130 129 117 115
```

```
barplot(antagonists, 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(2:31)])
```

```
## Regions
##          raphes|raphe|nuclei|nucleus
##                                973
##          striatums|striatum
##                                676
##    pres|pre|frontals|frontal|cortexes|cortex
##                                664
##    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
## ventrals|ventral|tegmentals|tegmental|areas|area
##                                181
##    cingulates|cingulate|cortexes|cortex
```

```

##                                     132
##                                putamens|putamen
##                                     132
## paraventriculars|paraventricular|nuclei|nucleus
##                                     129
##            anteriors|anterior|cingulates|cingulate
##                                     119
##                                posteriors|posterior
##                                     103
##                basals|basal|ganglias|ganglia
##                                     102
##            entorhinals|entorhinal|cortexes|cortex
##                                     100
##                loci|locus|coeruleuses|coeruleus
##                                     98
##    periaqueductals|periaqueductal|grays|gray
##                                     98
## suprachiasmatics|suprachiasmatic|nuclei|nucleus
##                                     90
##            olfactoryes|olfactory|bulbs|bulb
##                                     81
##            solitaires|solitary|tracts|tract
##                                     68
##            laterals|lateral|amygdalas|amygdala
##                                     67
##    orbitofrontals|orbitofrontal|cortexes|cortex
##                                     65
##                areas|area|postremas|postrema
##                                     58
##    basolaterals|basolateral|amygdalas|amygdala
##                                     57

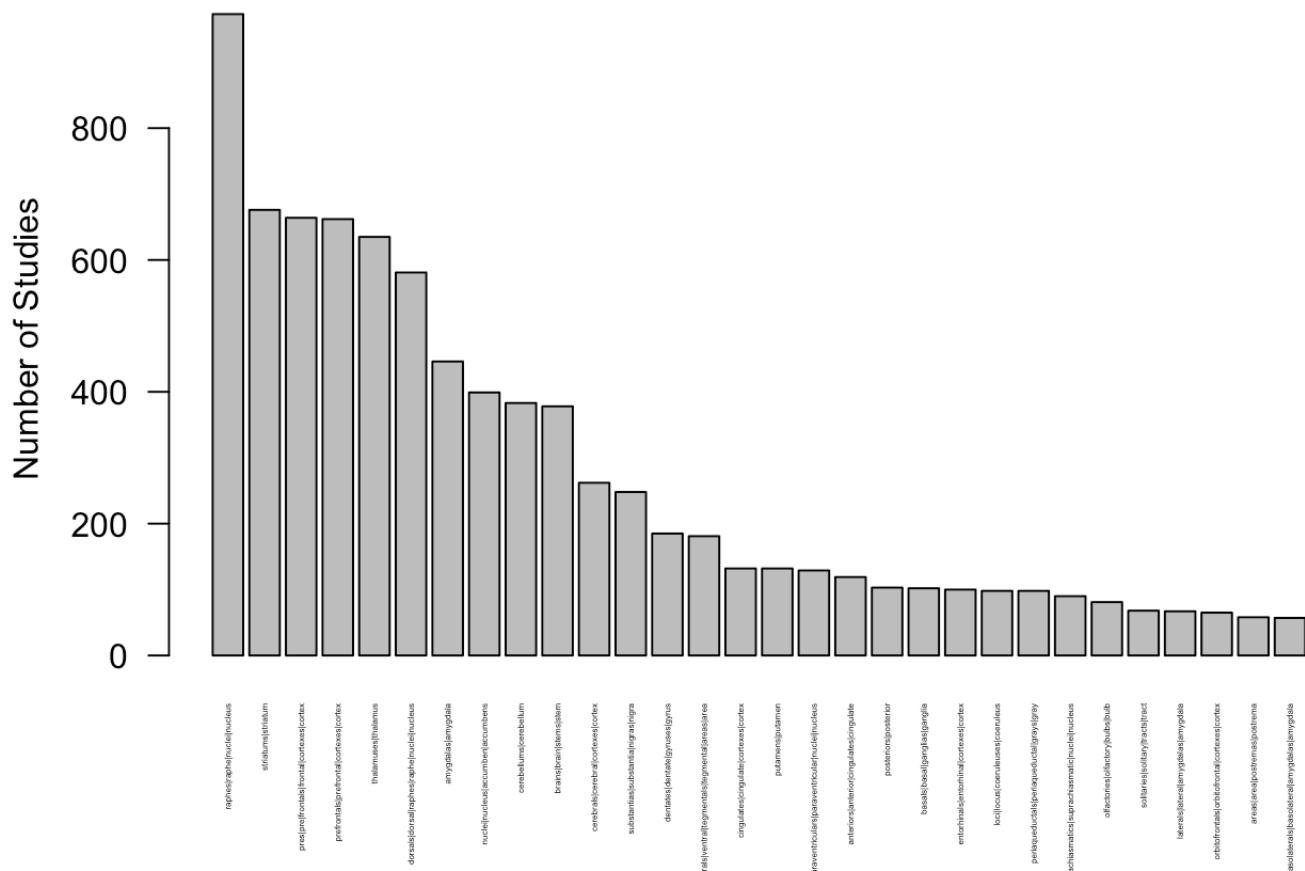
```

```

barplot(regions, las = 2, cex.names=.25, 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))

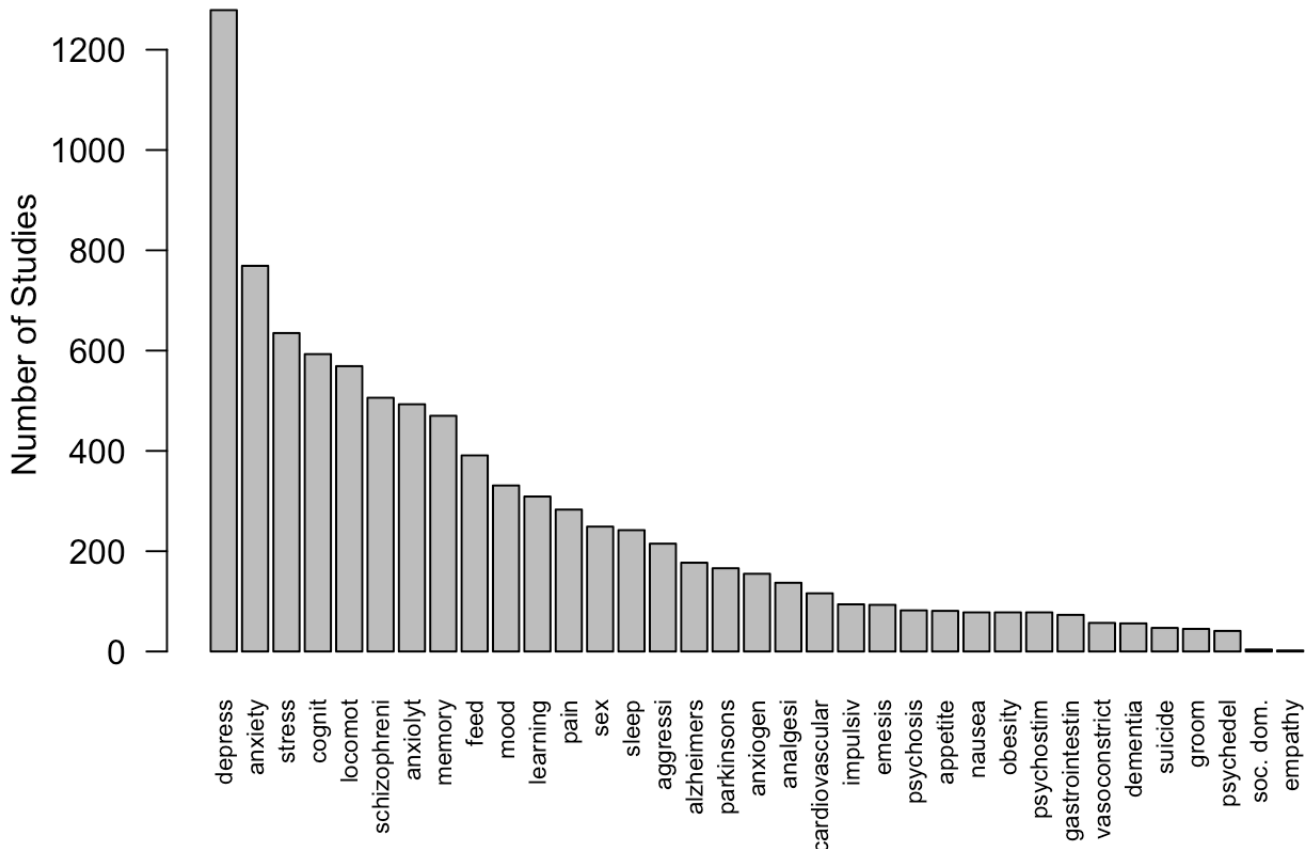
```

## ## 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-HT1A", ylab = "Number of Studies")
```

## Topics Studied with 5-HT1A

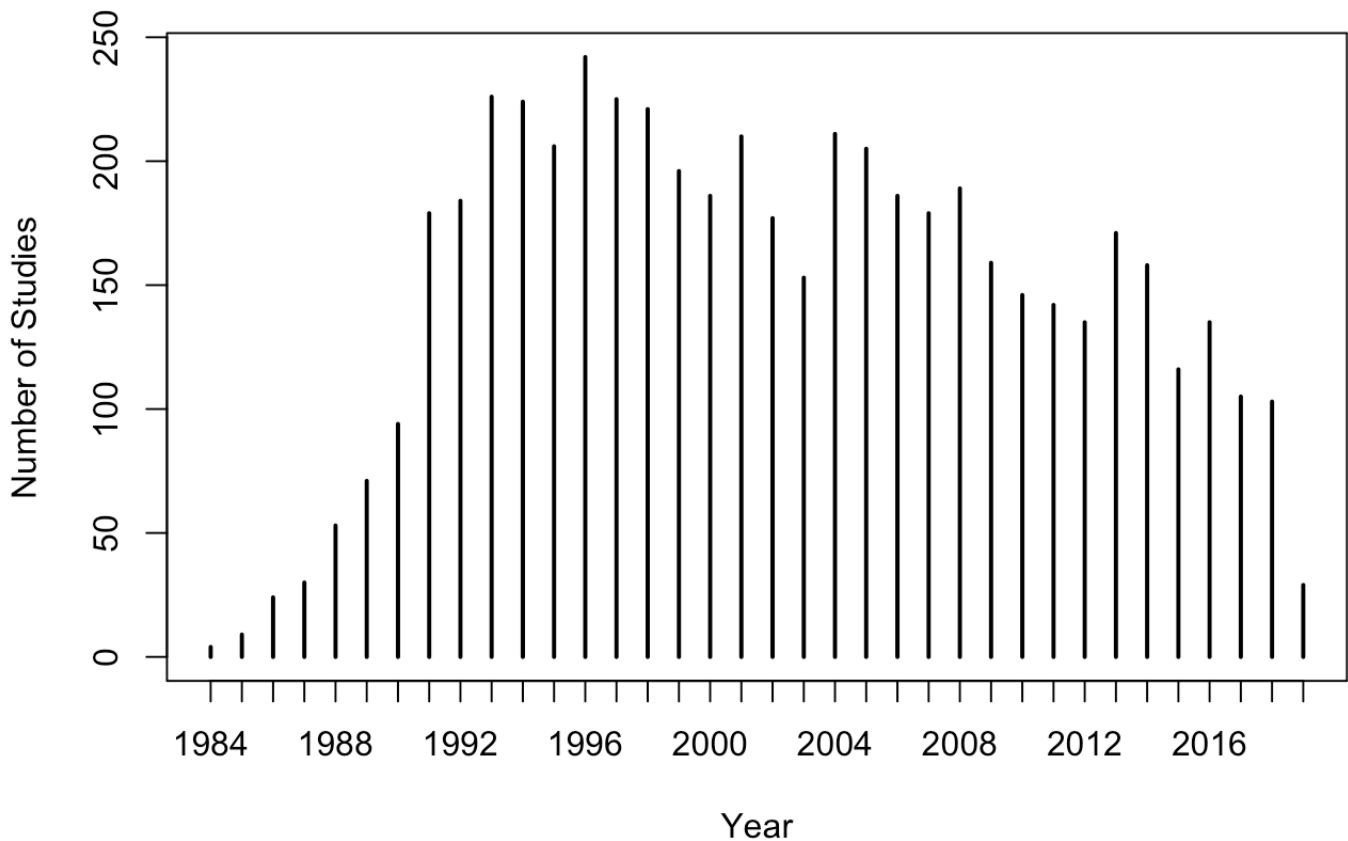


# 5-HT1A

```
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(xla$Methods, ";")

Methods1a <- unlist(Methods1a)

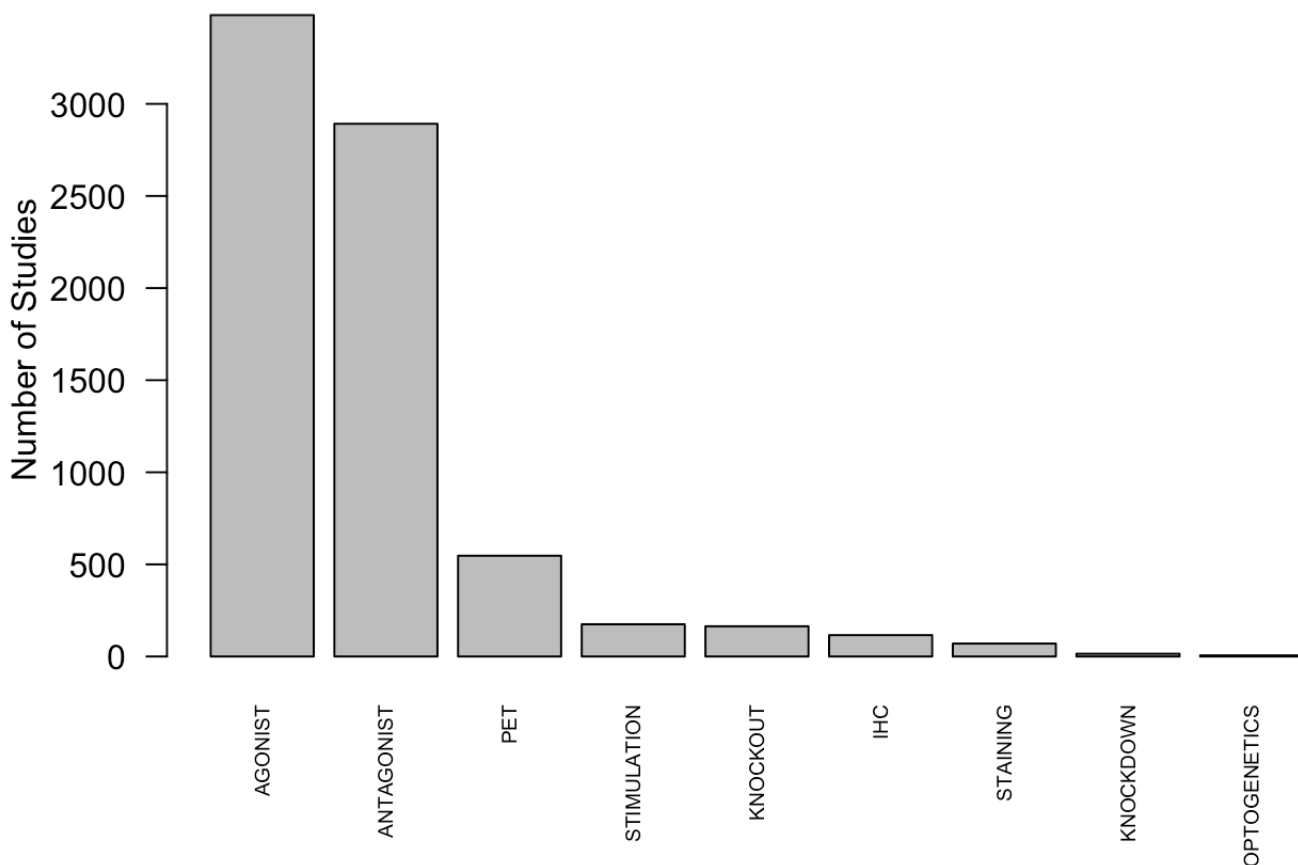
Methods1a <- gsub("Method.", "", Methods1a)
Methods1a <- gsub("IMMUNOHISTOCHEMISTRY", "IHC", Methods1a)

(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)

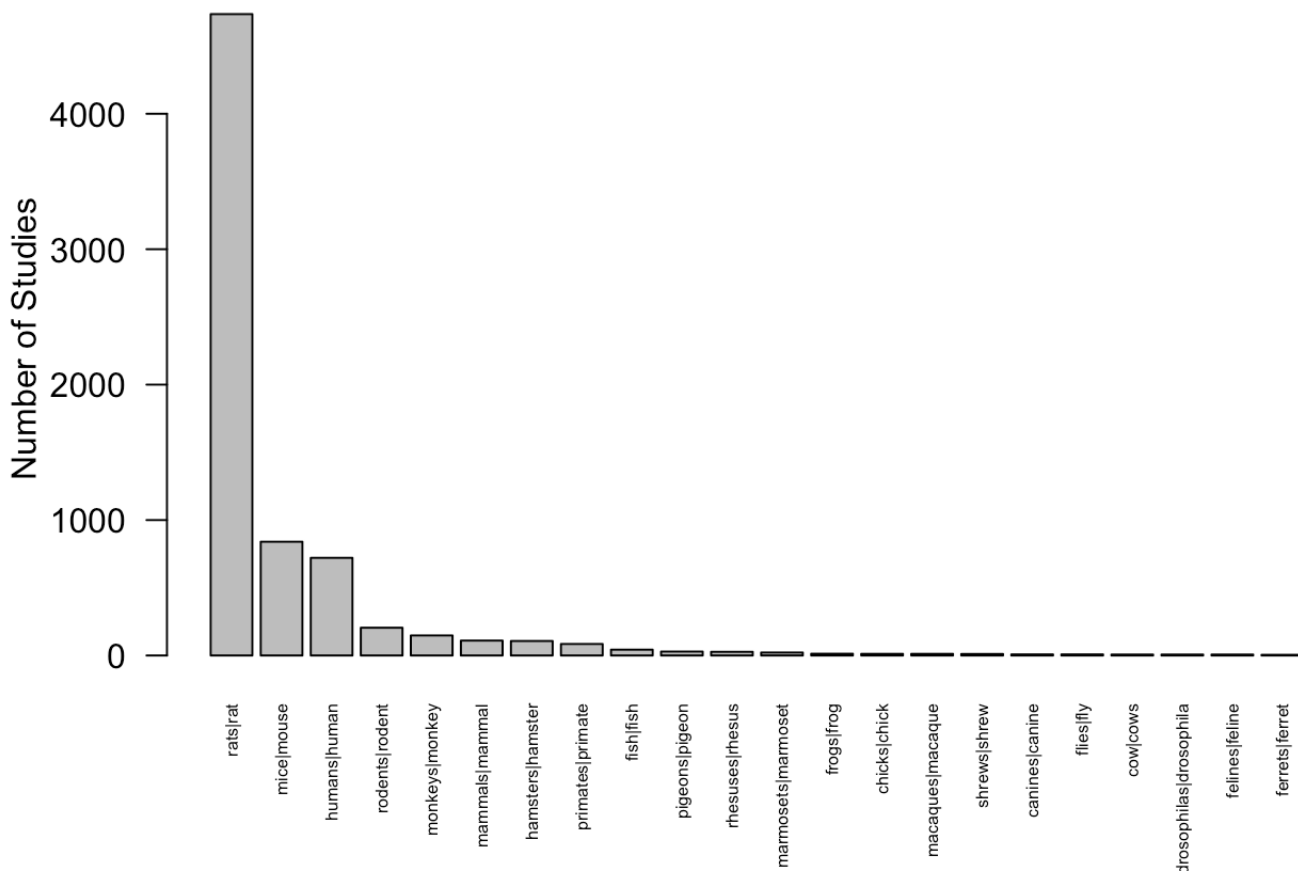
(speciesla <- sort(table(Speciesla), decreasing = TRUE)[c(1, 24, 31, 54, 62, 71, 73,
83, 108, 120, 124, 133, 154, 163, 166, 168, 181, 185, 190, 191, 192, 207)])
```

```
## 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(x1a$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)

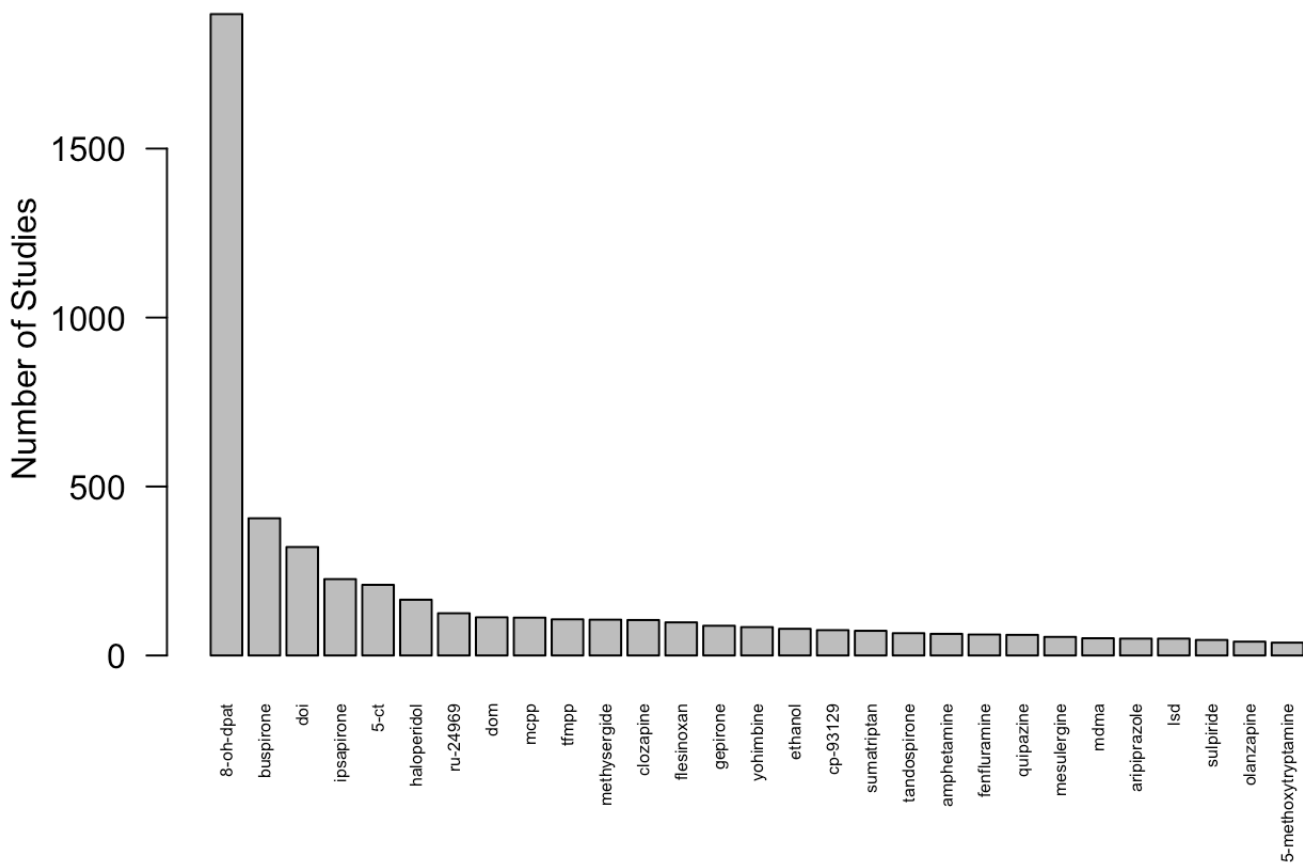
#sort(table(Agonists1a), decreasing = TRUE)
(agonists1a <- sort(table(Agonists1a), decreasing = TRUE)[c(1, 3:30)])

```

```
## 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)

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

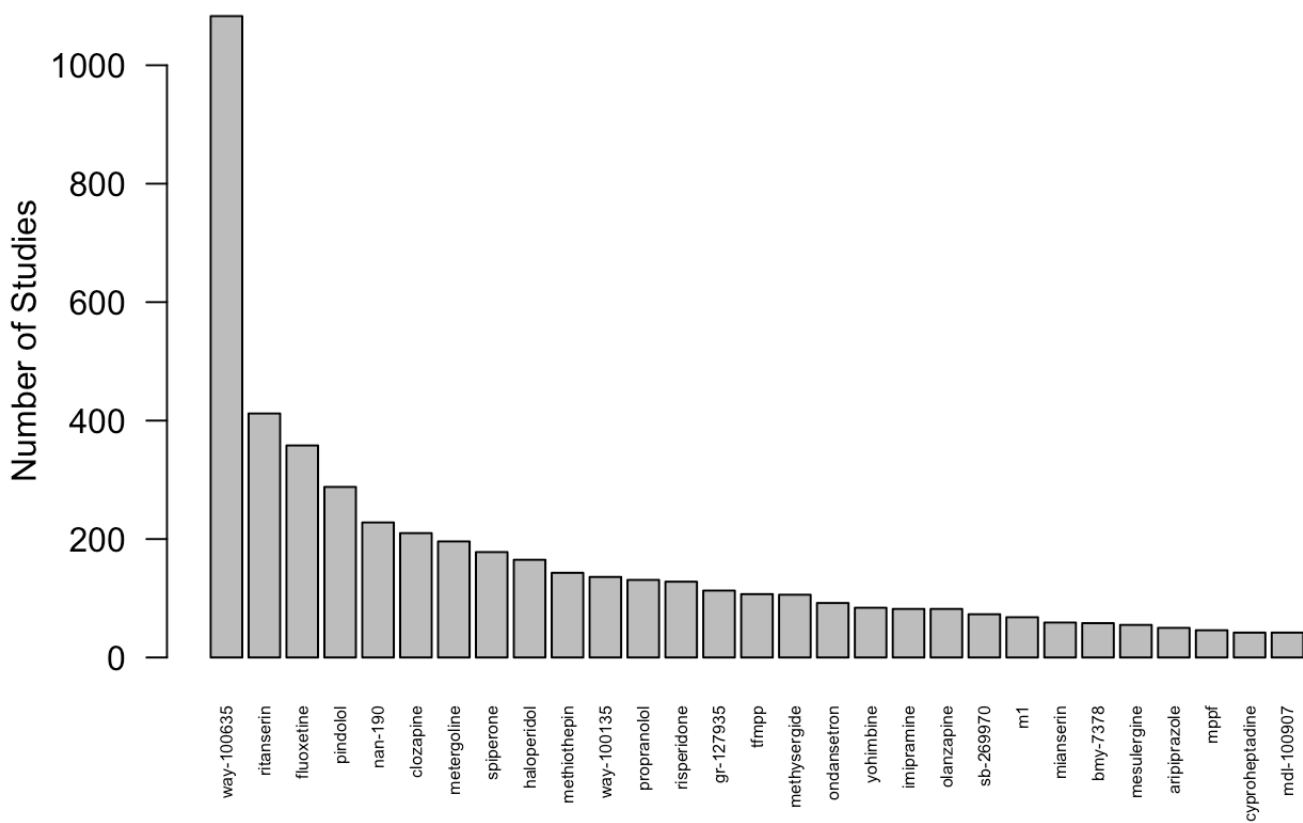
(antagonists1a <- sort(table(Antagonists1a), decreasing = TRUE)[c(1, 3:30)])

```

```
## Antagonists1a
##      way-100635      ritanserin      fluoxetine      pindolol      nan-190
##      1083          412          358          288          228
##      clozapine      metergoline      spiperone      haloperidol      methiothepin
##      210          196          178          165          143
##      way-100135      propranolol      risperidone      gr-127935      tfmpp
##      136          131          128          113          107
##      methysergide      ondansetron      yohimbine      imipramine      olanzapine
##      106          92          84          82          82
##      sb-269970          m1          mianserin      bmy-7378      mesulergine
##      73          68          59          58          55
##      aripiprazole      mppf      cyproheptadine      mdl-100907
##      50          46          42          42
```

```
barplot(antagonists1a, las = 2, cex.names=.5, main = "Antagonists Used to Study 5-HT1A", ylab = "Number of Studies")
```

## Antagonists Used to Study 5-HT1A



```

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(2:31)])

```

```

## Regionsla
##          raphes|raphe|nuclei|nucleus
##                                807
##      dorsals|dorsal|raphes|raphe|nuclei|nucleus
##                                476
##          thalamuses|thalamus
##                                387
##      prefrontals|prefrontal|cortexes|cortex
##                                358
##      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
##      cerebrals|cerebral|cortexes|cortex
##                                124
##      paraventriculars|paraventricular|nuclei|nucleus

```

```

##                                     81
##          cingulates|cingulate|cortexes|cortex
##                                     78
##          loci|locus|coeruleuses|coeruleus
##                                     74
##          entorhinals|entorhinal|cortexes|cortex
##                                     69
## ventrals|ventral|tegmentals|tegmental|areas|area
##                                     68
##          periaqueductals|periaqueductal|grays|gray
##                                     61
##          anteriors|anterior|cingulates|cingulate
##                                     60
## suprachiasmatics|suprachiasmatic|nuclei|nucleus
##                                     54
##          posteriors|posterior
##                                     53
##          putamens|putamen
##                                     50
##          basals|basal|ganglias|ganglia
##                                     47
##          temporals|temporal|lobes|lobe
##                                     42
##          laterals|lateral|amygdalas|amygdala
##                                     33
##          olfactoryies|olfactory|bulbs|bulb
##                                     33
## orbitofrontals|orbitofrontal|cortexes|cortex
##                                     30
##          preoptics|preoptic|areas|area
##                                     30
##          basolaterals|basolateral|amygdalas|amygdala
##                                     26

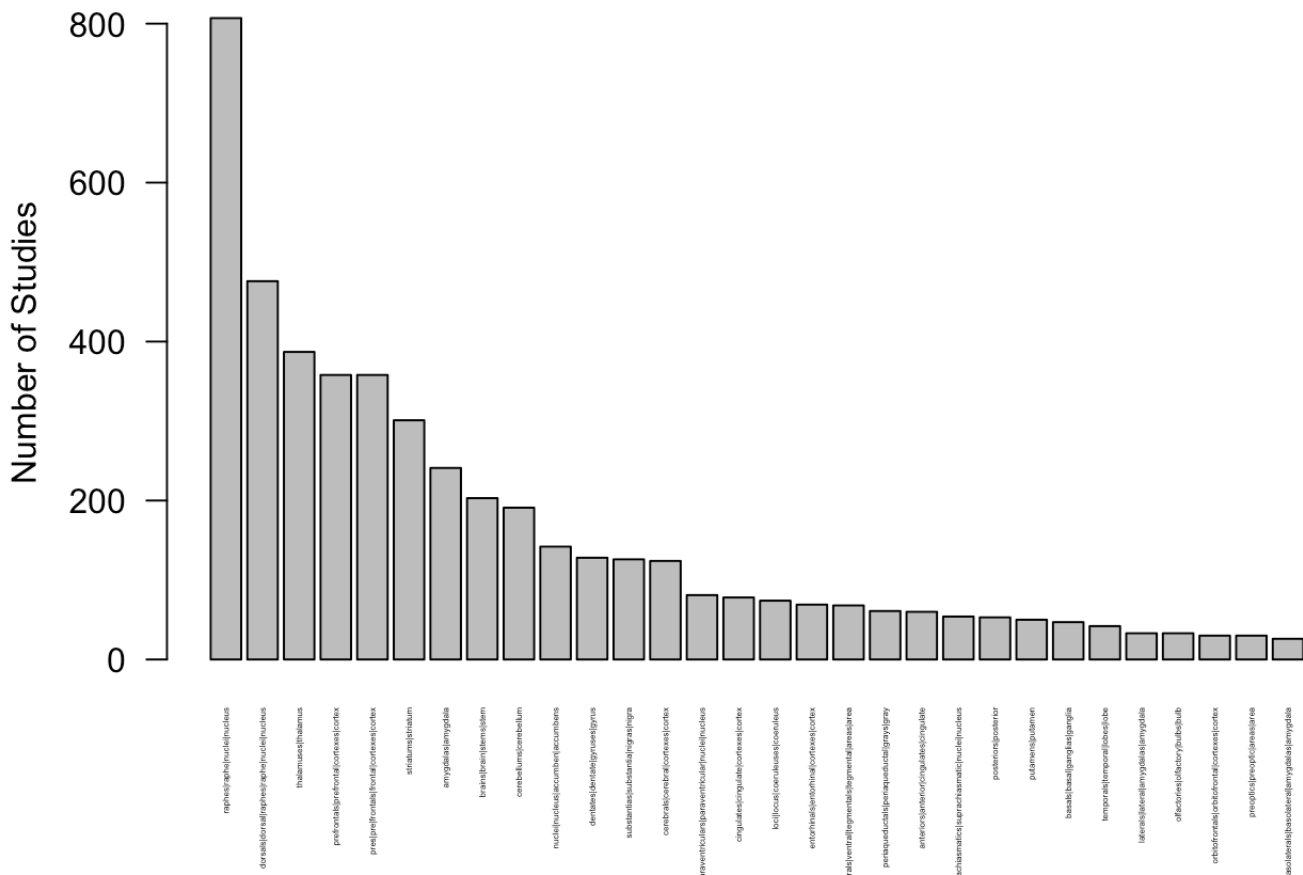
```

```

barplot(regions1a, las = 2, cex.names=.25, main = "Brain Regions Studied in Conjuncti
on 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

