

Mindset Manipulation Experiment 2 Bayes Supplement

RW Hass

8/1/2024

Notes on the Supplement

This was designed to illustrate our compliance with recommendations of [van de Schoot, et al \(2020\)](#). Importantly, that tutorial utilizes the advanced tools made possible by `rjags` and `rstan`. Some of the information is particular to the MCMC methods used by `rstan`, and we tried our best to produce information that was relevant given the `BayesFactor` package's limited capabilities. The data and the underlying RMarkdown for this and the second experiment are available [here](#).

Packages used: `psych`, `car`, `dplyr`, `BayesFactor`

session info:

```
## - Session info -----
## setting value
## version R version 4.3.2 (2023-10-31)
## os      macOS Sonoma 14.6
## system  aarch64, darwin20
## ui      X11
## language (EN)
## collate en_US.UTF-8
## ctype   en_US.UTF-8
## tz      America/New_York
## date    2024-08-01
## pandoc  3.2.1 @ /opt/homebrew/bin/ (via rmarkdown)
##
## - Packages -----
## package      * version      date (UTC) lib source
## abind         1.4-5        2016-07-21 [1] CRAN (R 4.3.0)
## BayesFactor  * 0.9.12-4.7  2024-01-24 [1] CRAN (R 4.3.1)
## cachem       1.1.0        2024-05-16 [1] CRAN (R 4.3.3)
## car           * 3.1-2        2023-03-30 [1] CRAN (R 4.3.0)
## carData      * 3.0-5        2022-01-06 [1] CRAN (R 4.3.0)
## cli          3.6.3        2024-06-21 [1] CRAN (R 4.3.3)
## coda         * 0.19-4.1     2024-01-31 [1] CRAN (R 4.3.1)
## colorspace   2.1-0        2023-01-23 [1] CRAN (R 4.3.0)
## devtools     2.4.5        2022-10-11 [1] CRAN (R 4.3.0)
## digest       0.6.36       2024-06-23 [1] CRAN (R 4.3.3)
## dplyr        * 1.1.4        2023-11-17 [1] CRAN (R 4.3.1)
## ellipsis     0.3.2        2021-04-29 [1] CRAN (R 4.3.0)
## evaluate     0.24.0       2024-06-10 [1] CRAN (R 4.3.3)
## fansi        1.0.6        2023-12-08 [1] CRAN (R 4.3.1)
## fastmap      1.2.0        2024-05-15 [1] CRAN (R 4.3.3)
## forcats     * 1.0.0        2023-01-29 [1] CRAN (R 4.3.0)
## fs           1.6.4        2024-04-25 [1] CRAN (R 4.3.1)
```

##	generics	0.1.3	2022-07-05	[1]	CRAN	(R 4.3.0)
##	ggplot2	* 3.5.1	2024-04-23	[1]	CRAN	(R 4.3.1)
##	glue	1.7.0	2024-01-09	[1]	CRAN	(R 4.3.1)
##	gtable	0.3.5	2024-04-22	[1]	CRAN	(R 4.3.1)
##	hms	1.1.3	2023-03-21	[1]	CRAN	(R 4.3.0)
##	htmltools	0.5.8.1	2024-04-04	[1]	CRAN	(R 4.3.1)
##	htmlwidgets	1.6.4	2023-12-06	[1]	CRAN	(R 4.3.1)
##	httpuv	1.6.15	2024-03-26	[1]	CRAN	(R 4.3.1)
##	knitr	1.48	2024-07-07	[1]	CRAN	(R 4.3.3)
##	later	1.3.2	2023-12-06	[1]	CRAN	(R 4.3.1)
##	lattice	0.22-6	2024-03-20	[1]	CRAN	(R 4.3.1)
##	lifecycle	1.0.4	2023-11-07	[1]	CRAN	(R 4.3.1)
##	lubridate	* 1.9.3	2023-09-27	[1]	CRAN	(R 4.3.1)
##	magrittr	2.0.3	2022-03-30	[1]	CRAN	(R 4.3.0)
##	Matrix	* 1.6-5	2024-01-11	[1]	CRAN	(R 4.3.1)
##	MatrixModels	0.5-3	2023-11-06	[1]	CRAN	(R 4.3.1)
##	memoise	2.0.1	2021-11-26	[1]	CRAN	(R 4.3.0)
##	mime	0.12	2021-09-28	[1]	CRAN	(R 4.3.0)
##	miniUI	0.1.1.1	2018-05-18	[1]	CRAN	(R 4.3.0)
##	mnormt	2.1.1	2022-09-26	[1]	CRAN	(R 4.3.0)
##	munsell	0.5.1	2024-04-01	[1]	CRAN	(R 4.3.1)
##	mvtnorm	1.2-5	2024-05-21	[1]	CRAN	(R 4.3.3)
##	nlme	3.1-165	2024-06-06	[1]	CRAN	(R 4.3.3)
##	pbapply	1.7-2	2023-06-27	[1]	CRAN	(R 4.3.0)
##	pillar	1.9.0	2023-03-22	[1]	CRAN	(R 4.3.0)
##	pkgbuild	1.4.4	2024-03-17	[1]	CRAN	(R 4.3.1)
##	pkgconfig	2.0.3	2019-09-22	[1]	CRAN	(R 4.3.0)
##	pkgload	1.4.0	2024-06-28	[1]	CRAN	(R 4.3.3)
##	profvis	0.3.8	2023-05-02	[1]	CRAN	(R 4.3.0)
##	promises	1.3.0	2024-04-05	[1]	CRAN	(R 4.3.1)
##	psych	* 2.4.6.26	2024-06-27	[1]	CRAN	(R 4.3.3)
##	purrr	* 1.0.2	2023-08-10	[1]	CRAN	(R 4.3.0)
##	R6	2.5.1	2021-08-19	[1]	CRAN	(R 4.3.0)
##	Rcpp	1.0.12	2024-01-09	[1]	CRAN	(R 4.3.1)
##	readr	* 2.1.5	2024-01-10	[1]	CRAN	(R 4.3.1)
##	remotes	2.5.0	2024-03-17	[1]	CRAN	(R 4.3.1)
##	rlang	1.1.4	2024-06-04	[1]	CRAN	(R 4.3.3)
##	rmarkdown	2.27	2024-05-17	[1]	CRAN	(R 4.3.3)
##	rstudioapi	0.16.0	2024-03-24	[1]	CRAN	(R 4.3.1)
##	scales	1.3.0	2023-11-28	[1]	CRAN	(R 4.3.1)
##	sessioninfo	1.2.2	2021-12-06	[1]	CRAN	(R 4.3.0)
##	shiny	1.8.1.1	2024-04-02	[1]	CRAN	(R 4.3.1)
##	stringi	1.8.4	2024-05-06	[1]	CRAN	(R 4.3.1)
##	stringr	* 1.5.1	2023-11-14	[1]	CRAN	(R 4.3.1)
##	tibble	* 3.2.1	2023-03-20	[1]	CRAN	(R 4.3.0)
##	tidyr	* 1.3.1	2024-01-24	[1]	CRAN	(R 4.3.1)
##	tidyselect	1.2.1	2024-03-11	[1]	CRAN	(R 4.3.1)
##	tidyverse	* 2.0.0	2023-02-22	[1]	CRAN	(R 4.3.0)
##	timechange	0.3.0	2024-01-18	[1]	CRAN	(R 4.3.1)
##	tzdb	0.4.0	2023-05-12	[1]	CRAN	(R 4.3.0)
##	urlchecker	1.0.1	2021-11-30	[1]	CRAN	(R 4.3.0)
##	usethis	2.2.3	2024-02-19	[1]	CRAN	(R 4.3.1)
##	utf8	1.2.4	2023-10-22	[1]	CRAN	(R 4.3.1)
##	vctrs	0.6.5	2023-12-01	[1]	CRAN	(R 4.3.1)

```
## withr          3.0.0      2024-01-16 [1] CRAN (R 4.3.1)
## xfun           0.45       2024-06-16 [1] CRAN (R 4.3.3)
## xtable         1.8-4      2019-04-21 [1] CRAN (R 4.3.0)
## yaml           2.3.9      2024-07-05 [1] CRAN (R 4.3.3)
##
## [1] /Library/Frameworks/R.framework/Versions/4.3-arm64/Resources/library
##
## -----

Descriptive statistics and correlation (correlations not used in manuscript)

# r.out <- corr.test(d[,c("JoanScore", "DTScore", "DTFlu", "Fixed", "Growth", "CSE")])
# desc.out <- describe(d[,c("JoanScore", "DTScore", "DTFlu", "Fixed", "Growth", "CSE")])

# per group
describeBy(d[,c("JoanScore", "DTScore", "DTFlu", "Fixed", "Growth", "CSE")], group = d$Condition)

##
## Descriptive statistics by group
## group: Fixed
##      vars  n mean   sd median trimmed  mad min max range  skew kurtosis
## JoanScore   1 55 2.85 0.73   3.00   2.82 1.48 2.0   4   2.0  0.22   -1.15
## DTScore     2 53 2.82 0.72   2.86   2.82 0.64 1.0   5   4.0  0.04    0.63
## DTFlu       3 55 6.60 3.41   6.00   6.40 2.97 0.0  17  17.0  0.63    0.45
## Fixed       4 55 3.13 1.05   3.25   3.14 1.48 1.0   5   4.0 -0.06   -1.15
## Growth      5 55 3.52 0.87   3.50   3.55 0.74 1.5   5   3.5 -0.32   -0.66
## CSE         6 55 3.37 0.95   3.33   3.39 0.99 1.0   5   4.0 -0.37   -0.19
##
##      se
## JoanScore 0.10
## DTScore   0.10
## DTFlu     0.46
## Fixed     0.14
## Growth    0.12
## CSE       0.13
## -----
## group: Growth
##      vars  n mean   sd median trimmed  mad min  max range  skew kurtosis
## JoanScore   1 56 2.68 0.90   3.00   2.65 1.48 1.0  5.0   4.0  0.36   -0.55
## DTScore     2 56 2.79 0.59   2.75   2.79 0.46 1.5  4.5   3.0  0.26    0.47
## DTFlu       3 56 7.98 3.76   7.50   7.63 3.71 2.0 17.0  15.0  0.82   -0.02
## Fixed       4 56 2.60 0.92   2.42   2.58 0.86 1.0  5.0   4.0  0.38   -0.59
## Growth      5 56 3.98 0.57   4.00   3.98 0.37 2.5  5.0   2.5 -0.15   -0.44
## CSE         6 56 3.18 1.03   3.33   3.20 0.99 1.0  5.0   4.0 -0.21   -0.90
##
##      se
## JoanScore 0.12
## DTScore   0.08
## DTFlu     0.50
## Fixed     0.12
## Growth    0.08
## CSE       0.14
```

Bayesian Estimation for Hypotheses 1 and 2

Note, the following sections follow the same layout of Supplement 1, and more info on the checks are available there.

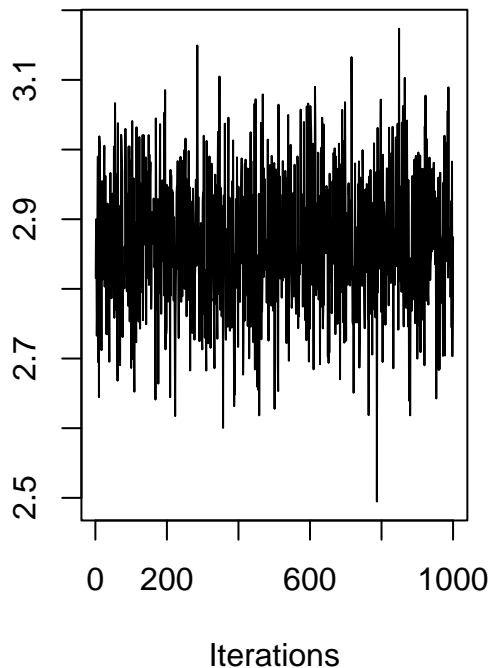
Fixed Mindset

```
# default is rscale = medium or sqrt(2)/2
ttestBF(formula = Fixed ~ Condition, data = d)

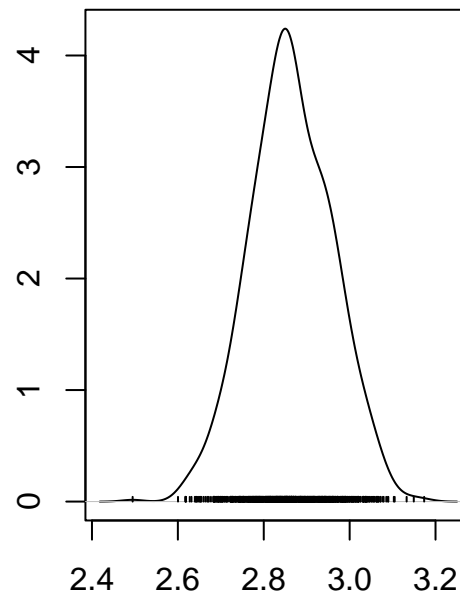
## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 6.569093 ±0%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

fixedpost <- ttestBF(formula = Fixed ~ Condition, data = d, posterior = TRUE, iterations = 1000)
plot(fixedpost[, "mu"])
```

Trace of var1



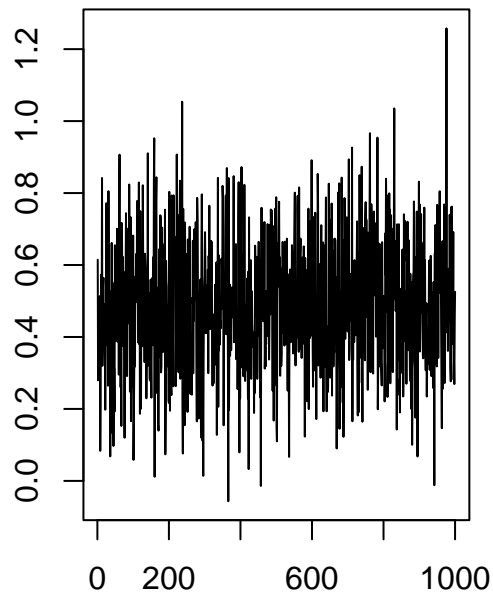
Density of var1



N = 1000 Bandwidth = 0.02553

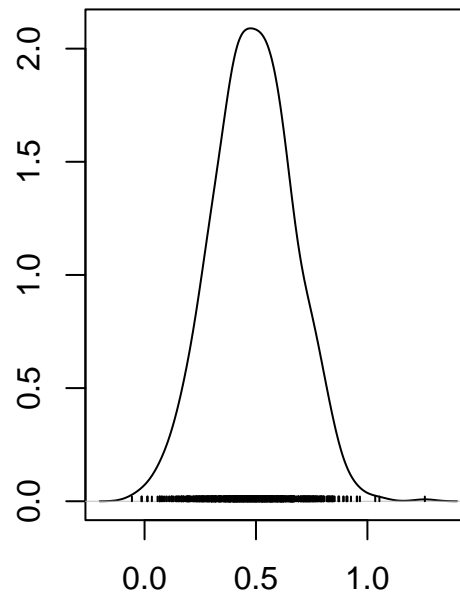
```
plot(fixedpost[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04819

```
quantile(fixedpost[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## 0.1394619 0.4874589 0.8344100
```

```
### with wide rscale = wide or 1
```

```
ttestBF(formula = Fixed ~ Condition, data = d, rscale = "wide")
```

```
## Bayes factor analysis
```

```
## -----
```

```
## [1] Alt., r=1 : 5.574359 ±0%
```

```
##
```

```
## Against denominator:
```

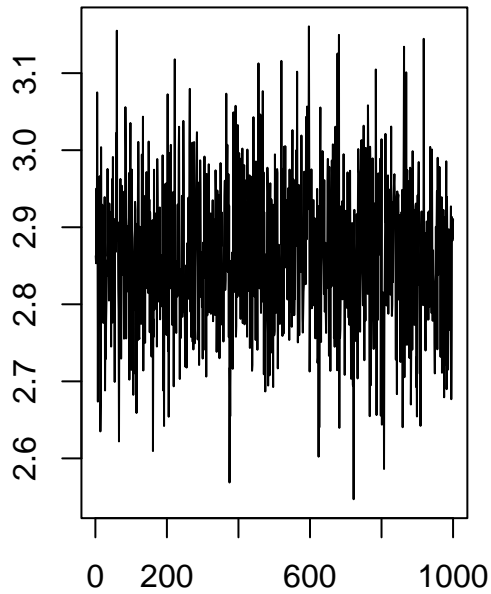
```
## Null, mu1-mu2 = 0
```

```
## ---
```

```
## Bayes factor type: BFindepSample, JZS
```

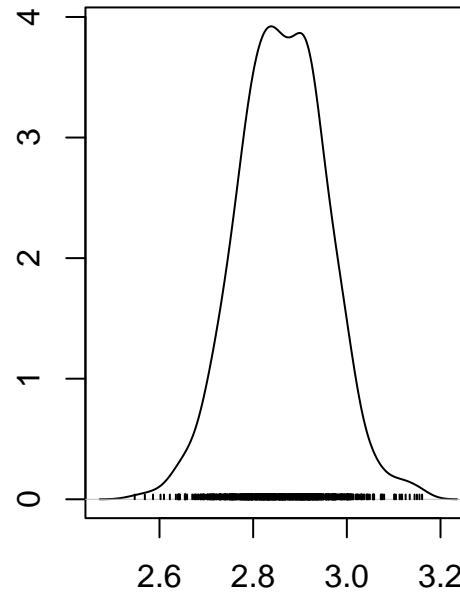
```
fixedpost.w <- ttestBF(formula = Fixed ~ Condition, data = d, rscale = "wide", posterior = TRUE, iterat.
plot(fixedpost.w[, "mu"])
```

Trace of var1



Iterations

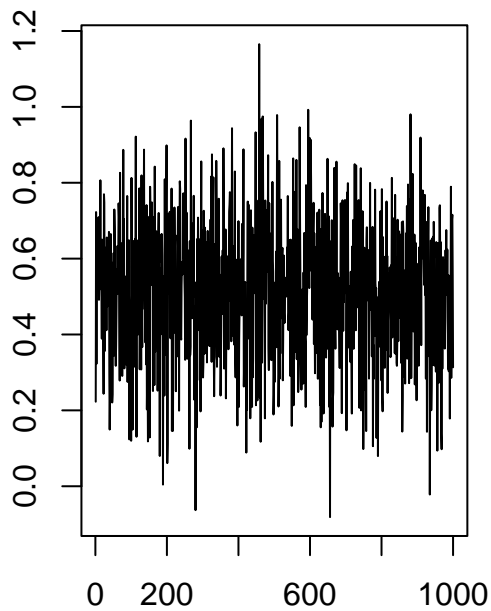
Density of var1



N = 1000 Bandwidth = 0.02488

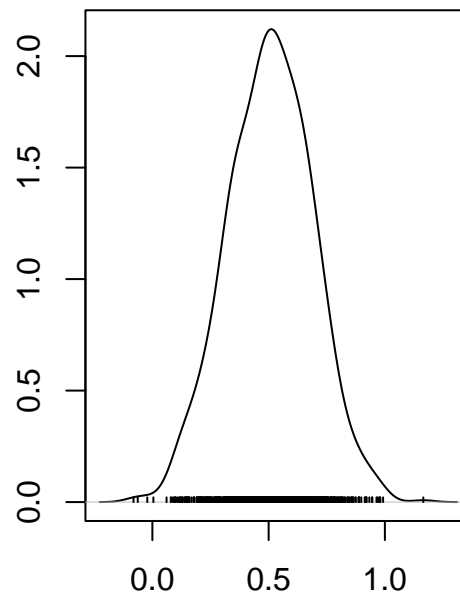
```
plot(fixedpost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04838

```
quantile(fixedpost.w[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## 0.1440049 0.5086717 0.8577710
```

```

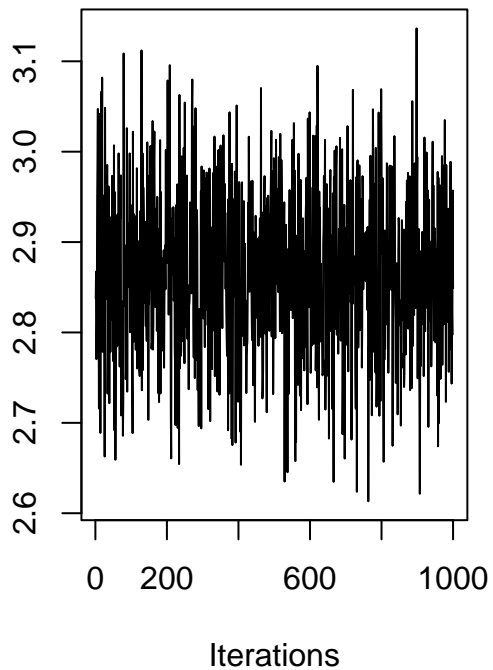
### with ultrawide rscale = ultrawide or sqrt(2)
ttestBF(formula = Fixed ~ Condition, data = d, rscale = "ultrawide")

## Bayes factor analysis
## -----
## [1] Alt., r=1.414 : 4.42181 ±0.01%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

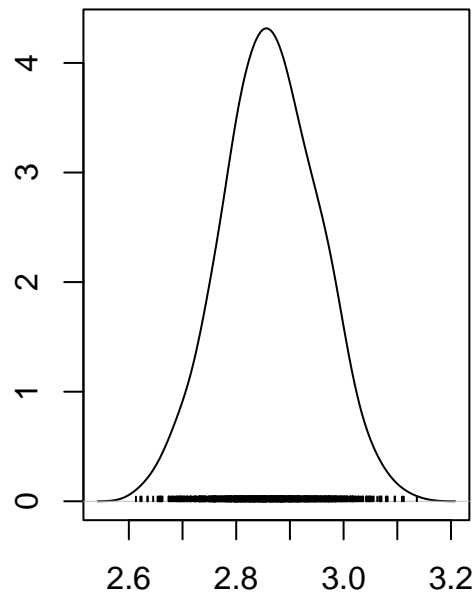
fixedpost.uw <- ttestBF(formula = Fixed ~ Condition, data = d, rscale = "ultrawide", posterior = TRUE,
plot(fixedpost.uw[, "mu"])

```

Trace of var1



Density of var1

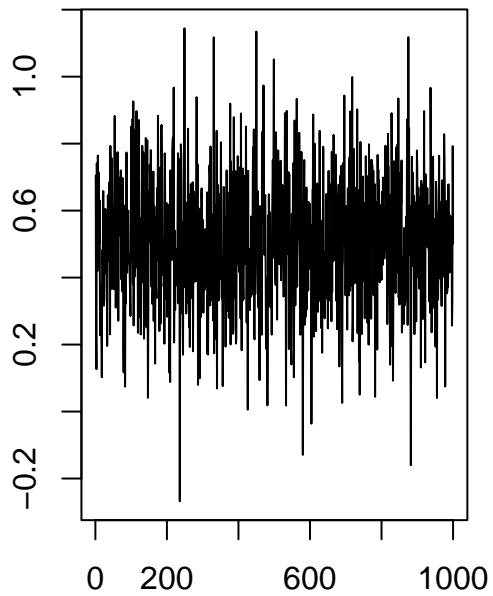


```

plot(fixedpost.uw[, "beta (Fixed - Growth)"])

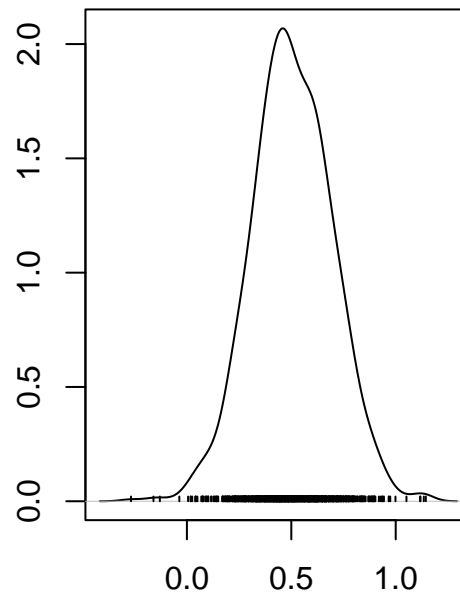
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04989

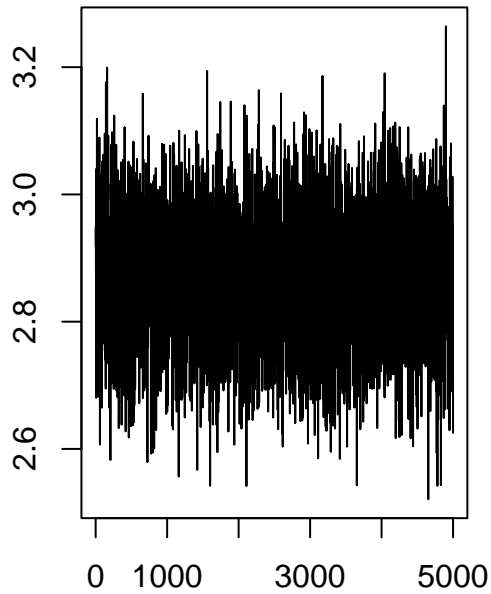
```
quantile(fixedpost.uw[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## 0.1263479 0.4986083 0.8826288
```

```
### More iterations
```

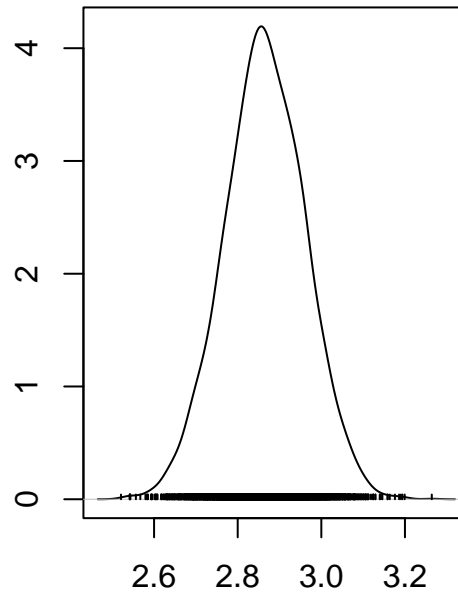
```
fixedpost.check <- ttestBF(formula = Fixed ~ Condition, data = d, posterior = TRUE, iterations = 5000)  
plot(fixedpost.check[, "mu"])
```


Trace of var1



Iterations

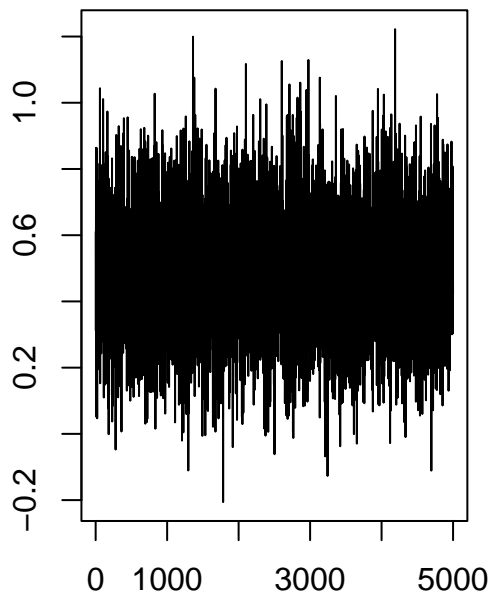
Density of var1



N = 5000 Bandwidth = 0.01857

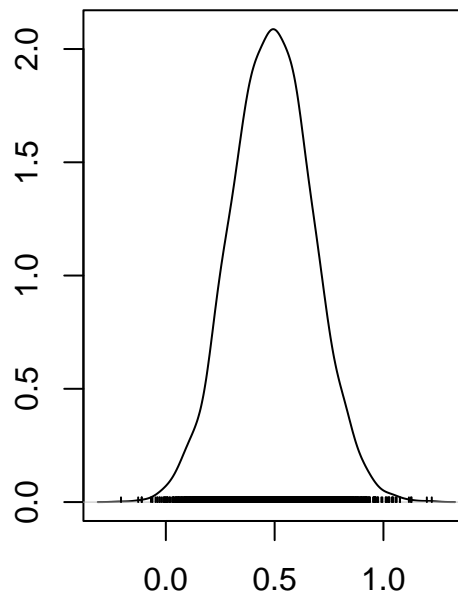
```
plot(fixedpost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.03566

```
quantile(fixedpost.check[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## 0.1269584 0.4884059 0.8476794
```

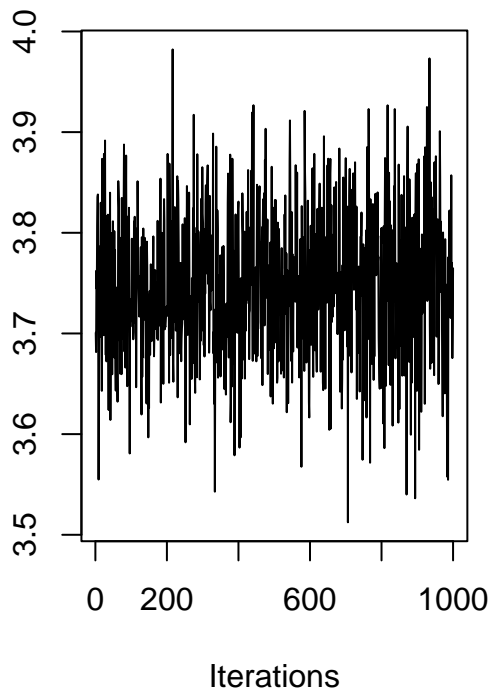
Growth Mindset as outcome

```
## default / medium
ttestBF(formula = Growth ~ Condition, data = d)

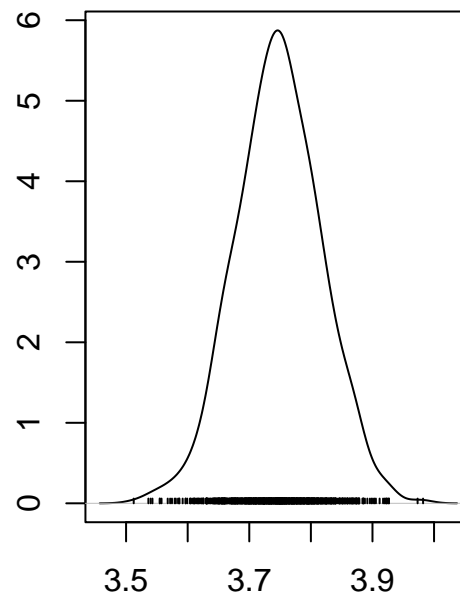
## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 21.44464 ±0%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

gpost <- ttestBF(formula = Growth ~ Condition, data = d, posterior = TRUE, iterations = 1000)
plot(gpost[, "mu"])
```

Trace of var1



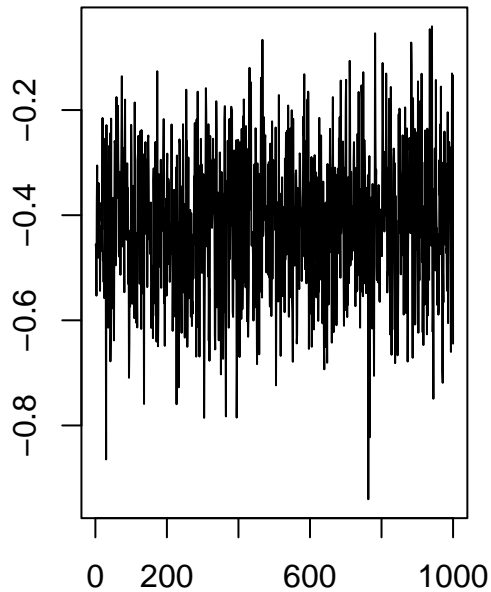
Density of var1



N = 1000 Bandwidth = 0.01839

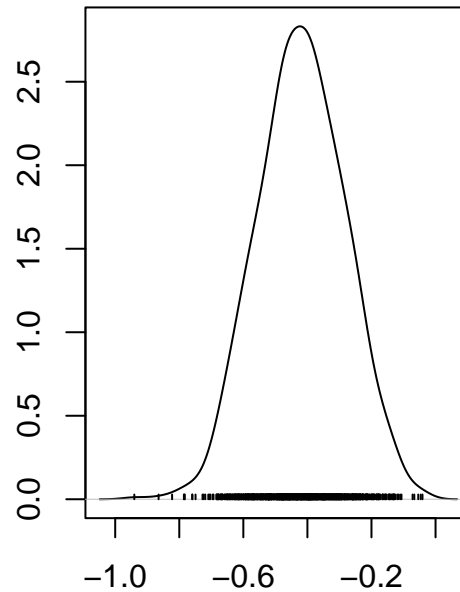
```
plot(gpost[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.03599

```
quantile(gpost[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.6733249 -0.4188110 -0.1560531
```

```
# Wide
```

```
ttestBF(formula = Growth ~ Condition, data = d, rscale = "wide")
```

```
## Bayes factor analysis
```

```
## -----
```

```
## [1] Alt., r=1 : 18.94251 ±0%
```

```
##
```

```
## Against denominator:
```

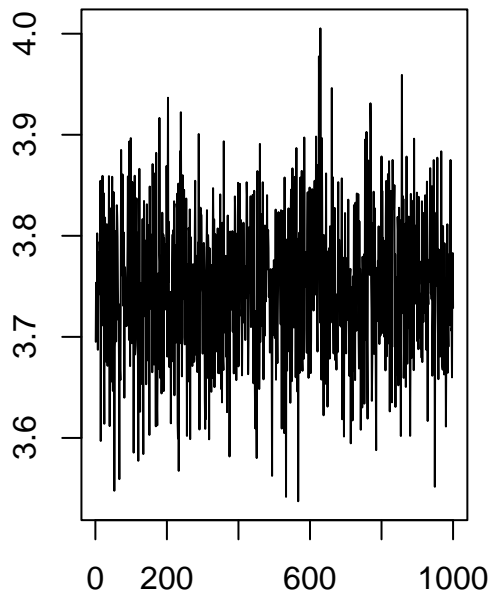
```
## Null, mu1-mu2 = 0
```

```
## ---
```

```
## Bayes factor type: BFindepSample, JZS
```

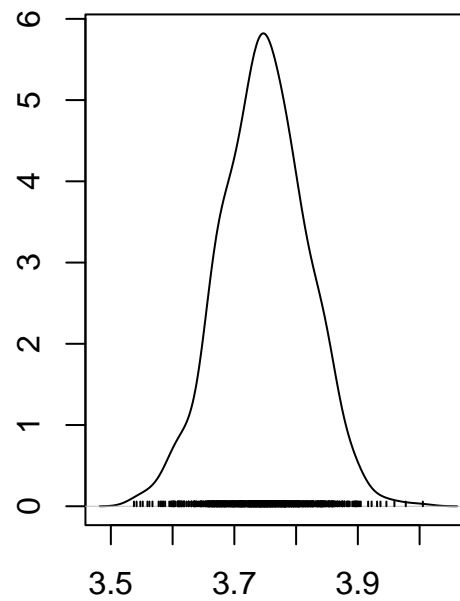
```
gpost.w <- ttestBF(formula = Growth ~ Condition, data = d, rscale = "wide", posterior = TRUE, iteration = 1000)
plot(gpost.w[, "mu"])
```

Trace of var1



Iterations

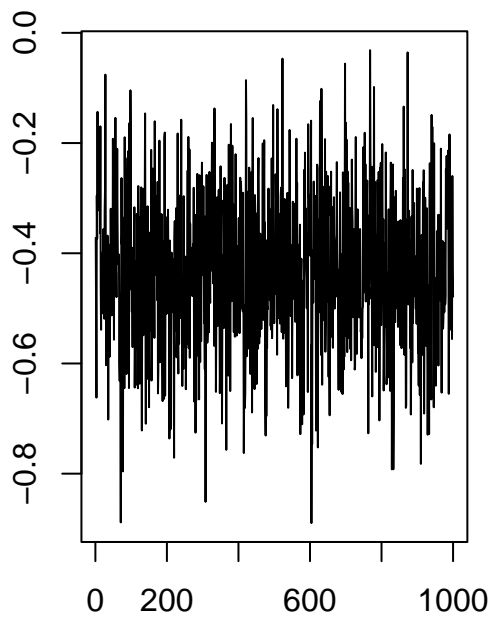
Density of var1



N = 1000 Bandwidth = 0.01846

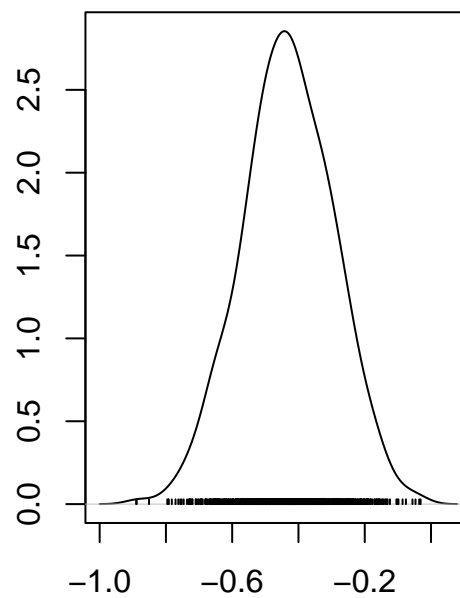
```
plot(gpost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.03671

```
quantile(gpost.w[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## -0.7034592 -0.4325637 -0.1653726
```

```
# UltraWide
```

```
ttestBF(formula = Growth ~ Condition, data = d, rscale = "ultrawide")
```

```
## Bayes factor analysis
```

```
## -----
```

```
## [1] Alt., r=1.414 : 15.46237 ±0%
```

```
##
```

```
## Against denominator:
```

```
## Null, mu1-mu2 = 0
```

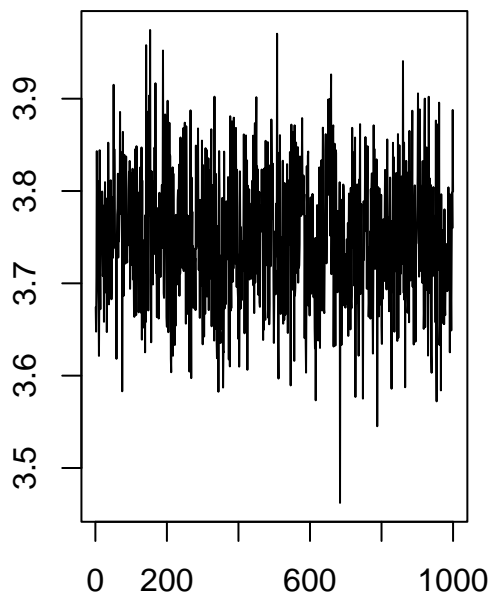
```
## ---
```

```
## Bayes factor type: BFindepSample, JZS
```

```
gpost.uw <- ttestBF(formula = Growth ~ Condition, data = d, rscale = "ultrawide", posterior = TRUE, ite
```

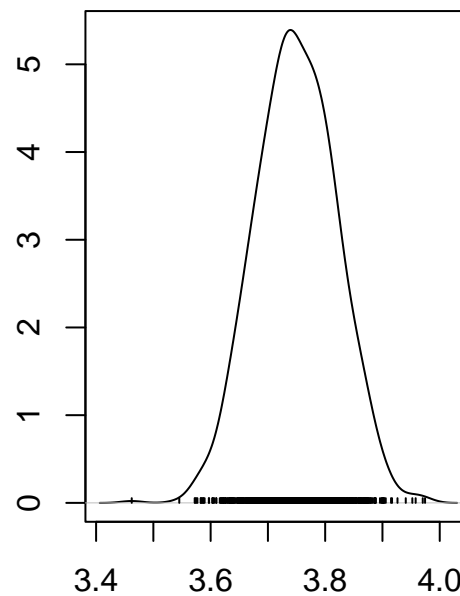
```
plot(gpost.uw[, "mu"])
```

Trace of var1



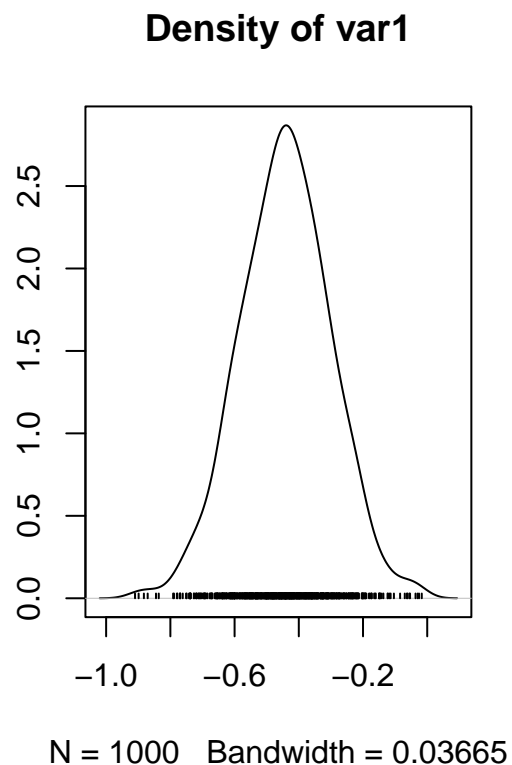
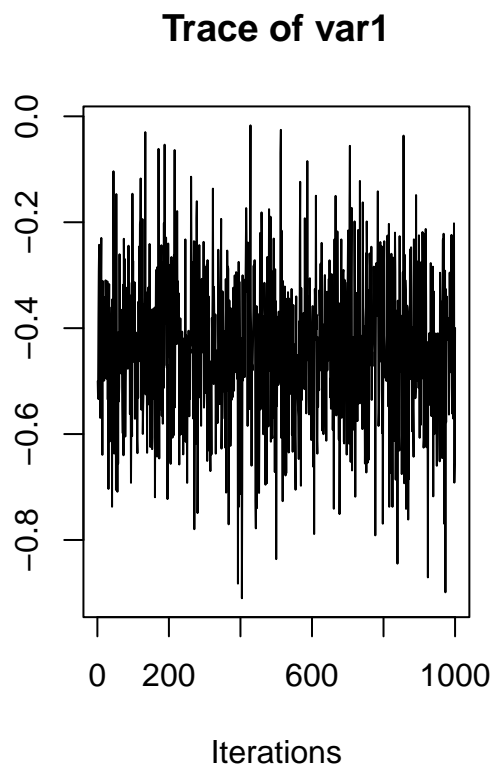
Iterations

Density of var1



N = 1000 Bandwidth = 0.01862

```
plot(gpost.uw[, "beta (Fixed - Growth)"])
```



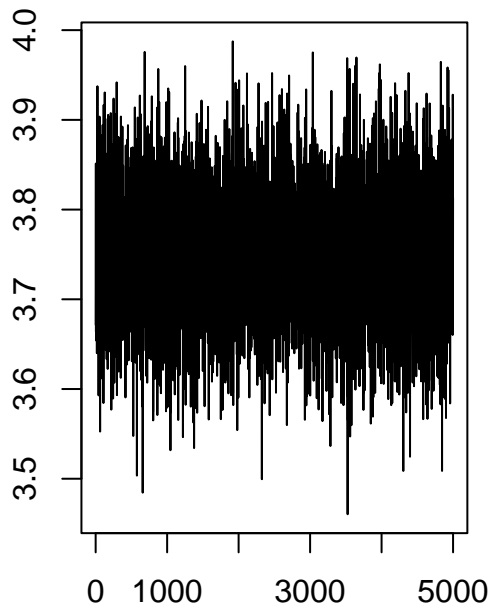
```
quantile(gpost.uw[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.7228277 -0.4413185 -0.1727467
```

```
### More iterations
```

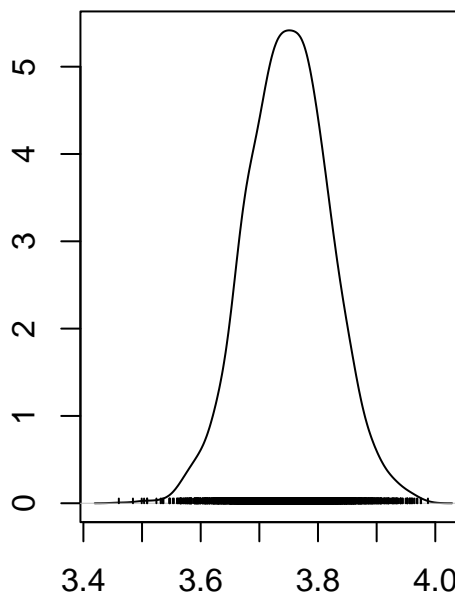
```
gpost.check <- ttestBF(formula = Growth ~ Condition, data = d, posterior = TRUE, iterations = 5000)
plot(gpost.check[, "mu"])
```

Trace of var1



Iterations

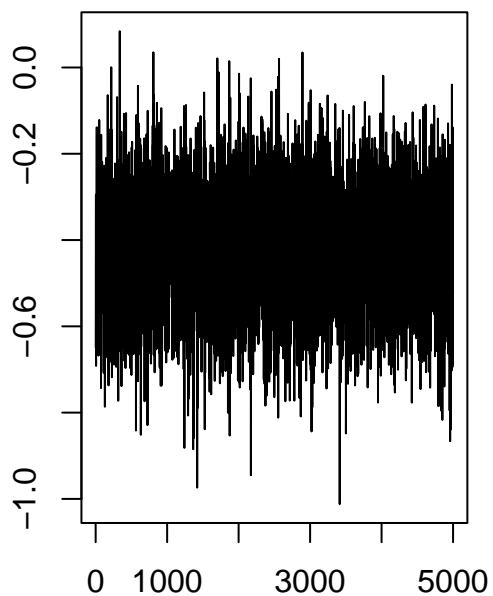
Density of var1



N = 5000 Bandwidth = 0.0137

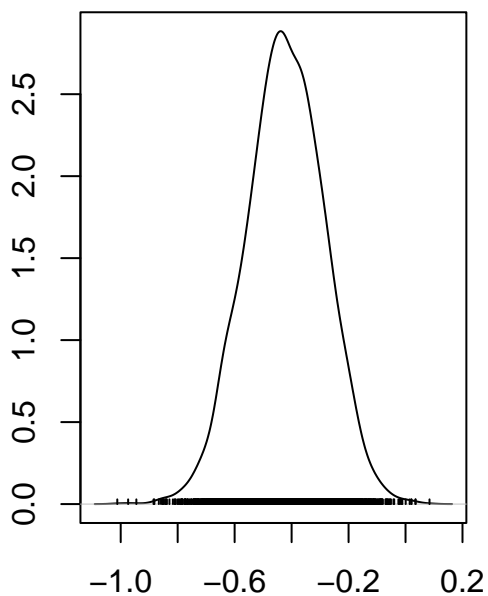
```
plot(gpost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.02641

```
quantile(gpost.check[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.6918381 -0.4225028 -0.1643170
```

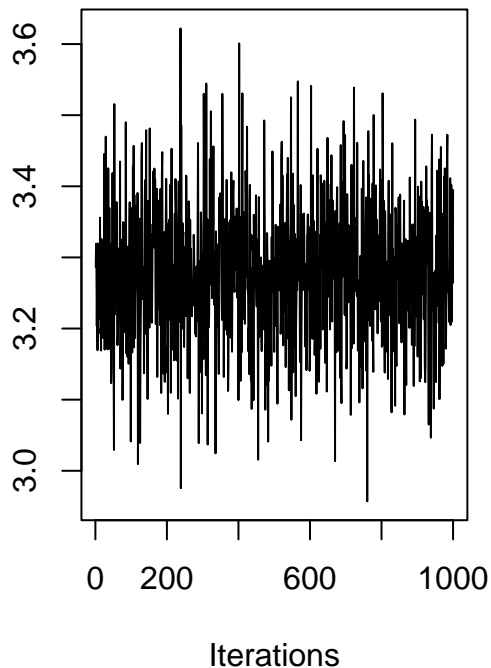
Creative Self-efficacy as outcome

```
## Default / medium
ttestBF(formula = CSE ~ Condition, data = d)

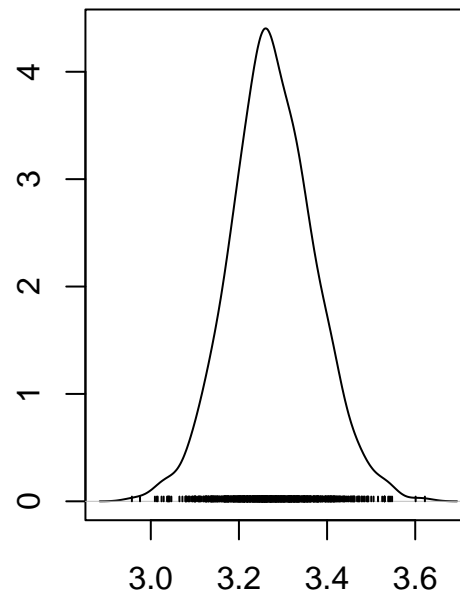
## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 0.3184985 ±0.02%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

csepost <- ttestBF(formula = CSE ~ Condition, data = d, posterior = TRUE, iterations = 1000)
plot(csepost[, "mu"])
```

Trace of var1

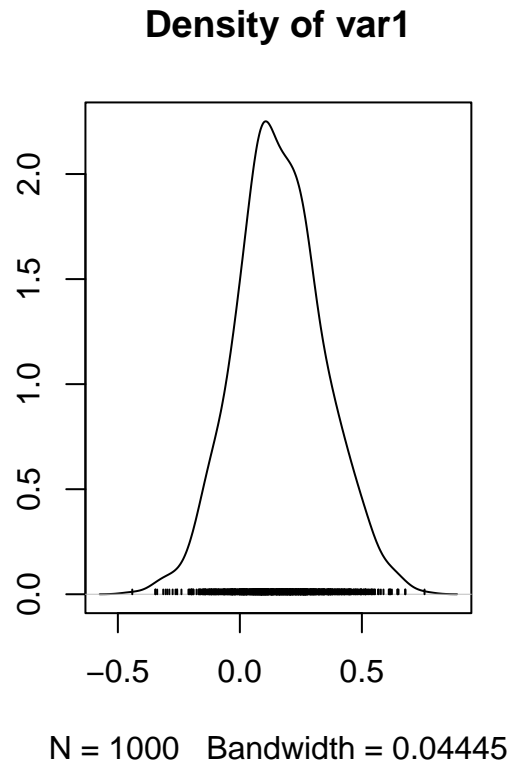
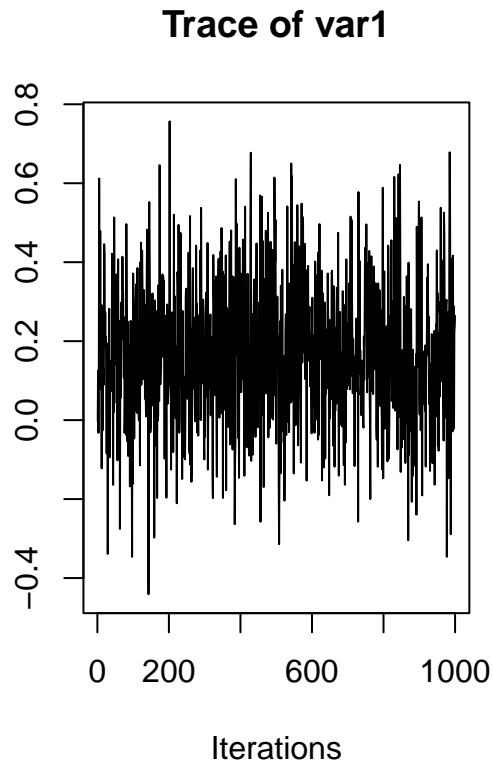


Density of var1



N = 1000 Bandwidth = 0.02432

```
plot(csepost[, "beta (Fixed - Growth)"])
```

```
quantile(csepost[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.1679336  0.1584152  0.5259200
```

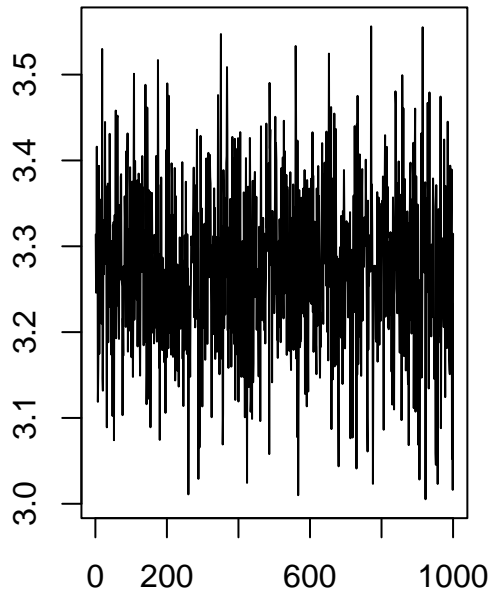
```
# Wide
```

```
ttestBF(formula = CSE ~ Condition, data = d, rscale = "wide")
```

```
## Bayes factor analysis
## -----
## [1] Alt., r=1 : 0.2374207 ±0.06%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS
```

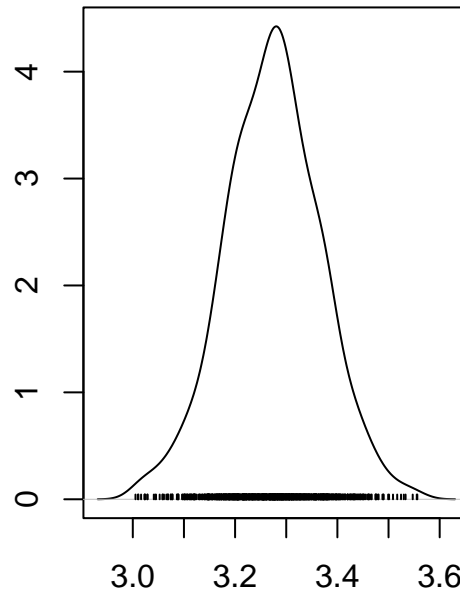
```
csepost.w <- ttestBF(formula = CSE ~ Condition, data = d, rscale = "wide", posterior = TRUE, iterations
plot(csepost.w[, "mu"])
```

Trace of var1



Iterations

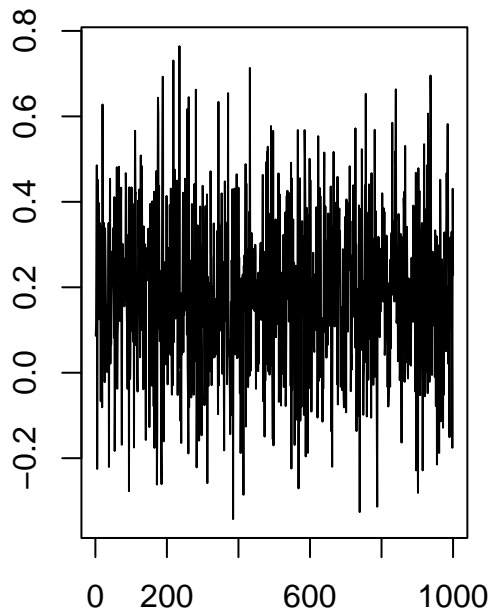
Density of var1



N = 1000 Bandwidth = 0.02461

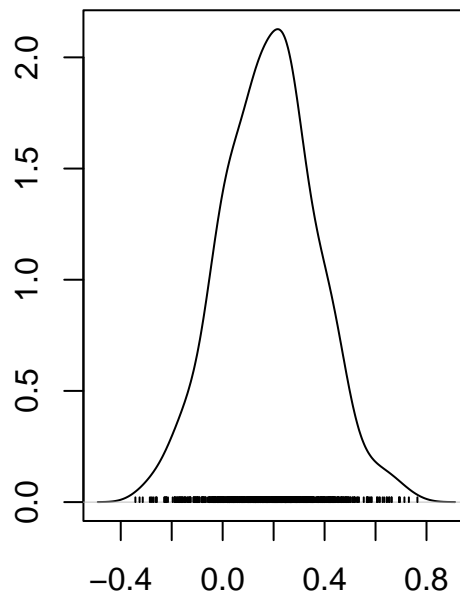
```
plot(csepost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04926

```
quantile(csepost.w[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

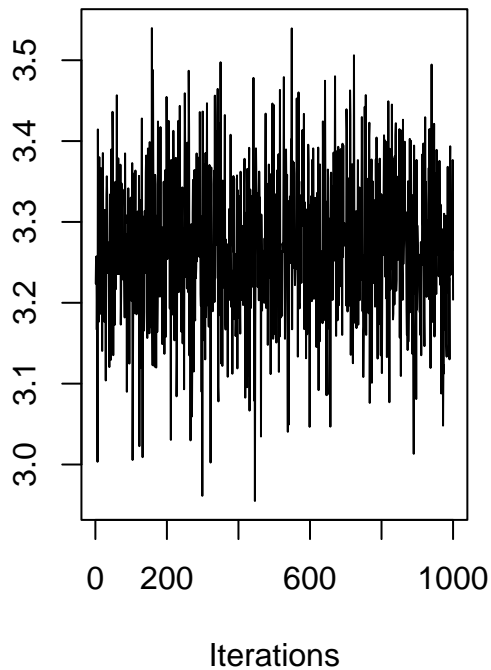
```
##      2.5%      50%      97.5%  
## -0.1760123  0.1823150  0.5662757
```

```
# ultra Wide
ttestBF(formula = CSE ~ Condition, data = d, rscale = "ultrawide")

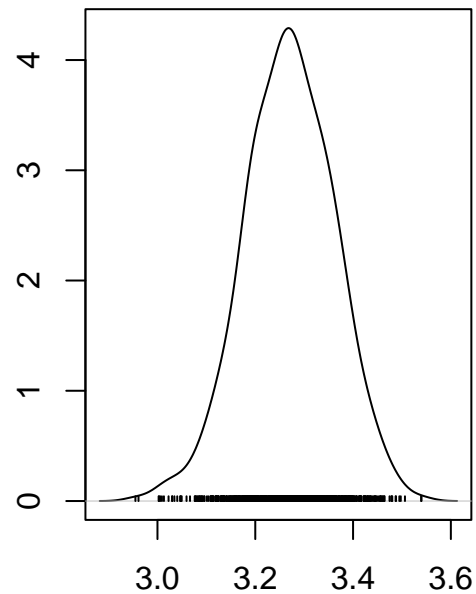
## Bayes factor analysis
## -----
## [1] Alt., r=1.414 : 0.173071 ±0.1%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

csepost.uw <- ttestBF(formula = CSE ~ Condition, data = d, rscale = "ultrawide", posterior = TRUE, iter
plot(csepost.uw[, "mu"])
```

Trace of var1



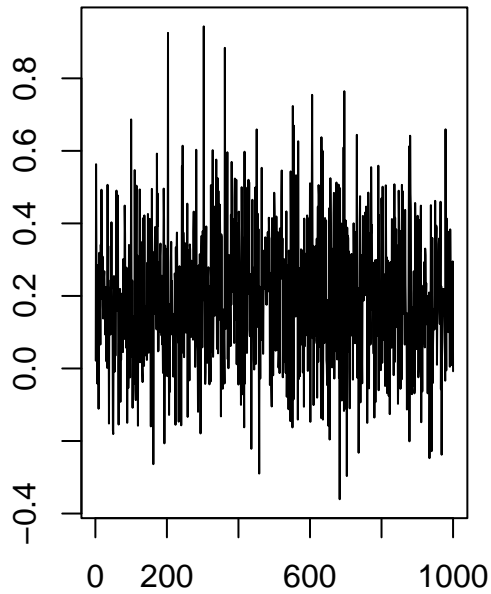
Density of var1



N = 1000 Bandwidth = 0.02432

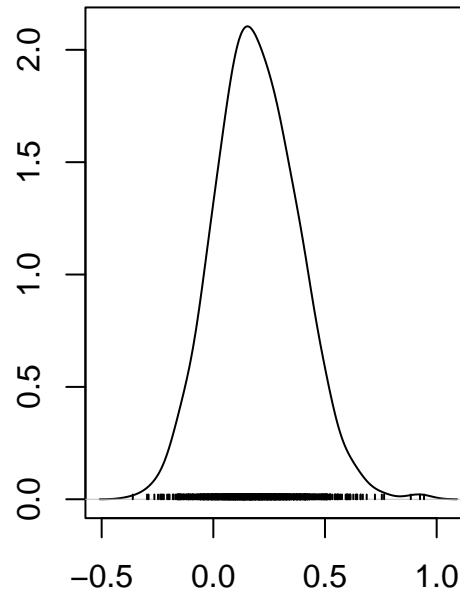
```
plot(csepost.uw[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04933

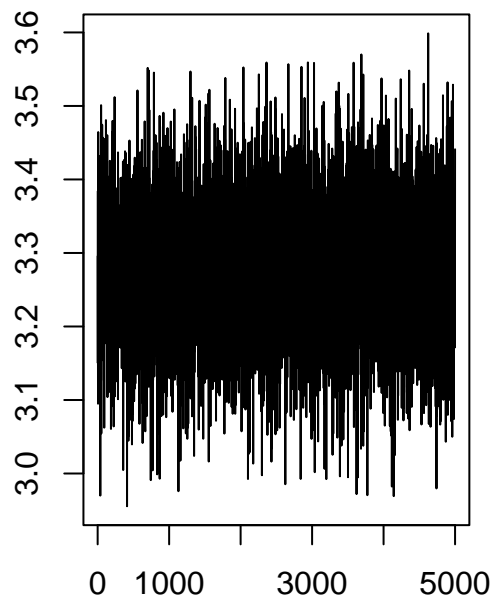
```
quantile(csepost.uw[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.1499714  0.1867698  0.5634292
```

```
### more iterations
```

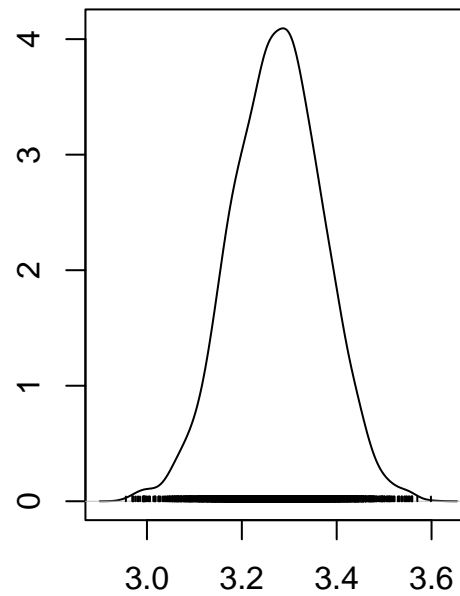
```
csepost.check <- ttestBF(formula = CSE ~ Condition, data = d, posterior = TRUE, iterations = 5000)
plot(csepost.check[, "mu"])
```

Trace of var1



Iterations

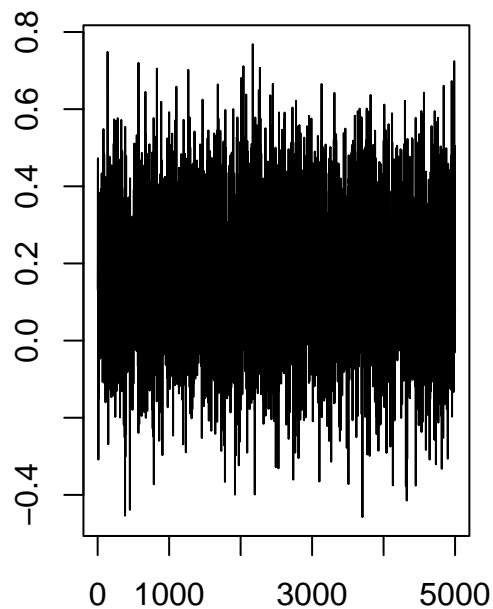
Density of var1



N = 5000 Bandwidth = 0.01836

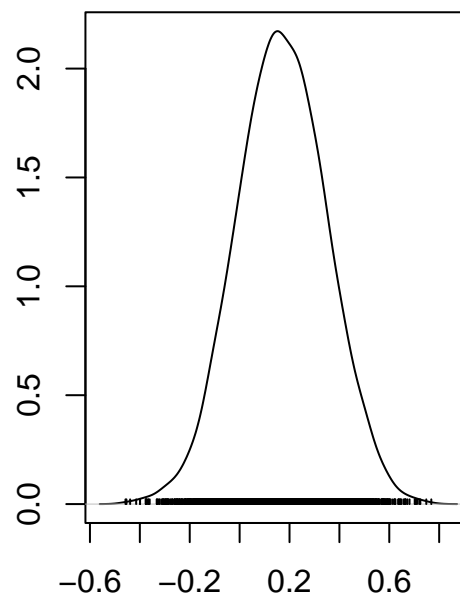
```
plot(csepost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.03443

```
quantile(csepost.check[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.1781354  0.1673873  0.5169461
```

Joan's problem ratings

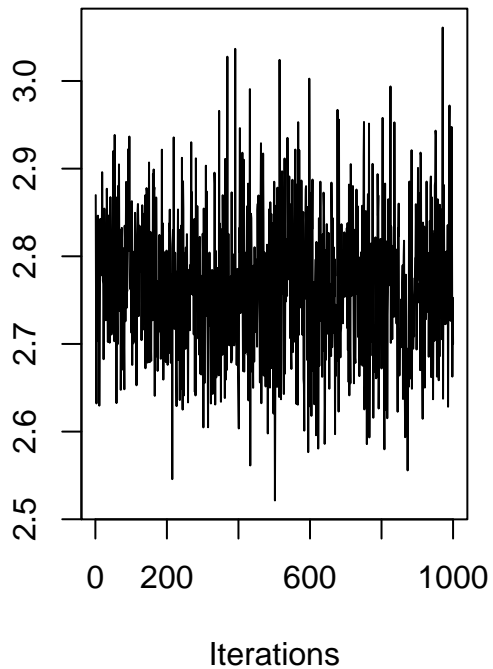
```
# default / medium
ttestBF(formula = JoanScore ~ Condition, data = d)

## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 0.3574592 ±0.02%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

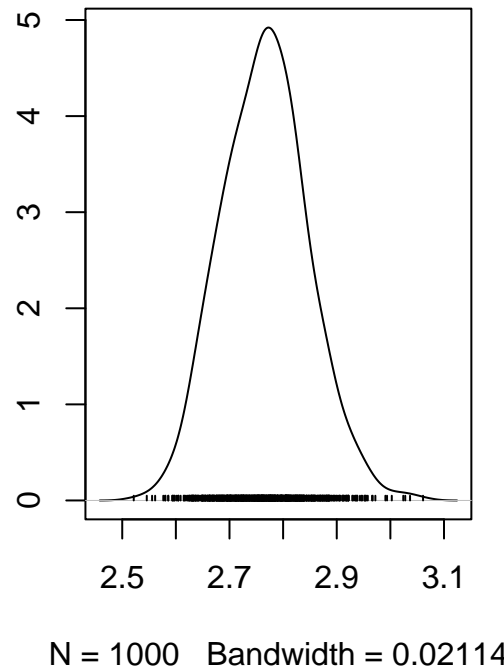
joanpost <- ttestBF(formula = JoanScore ~ Condition, data = d, posterior = TRUE, iterations = 1000)

plot(joanpost[, "mu"])
```

Trace of var1

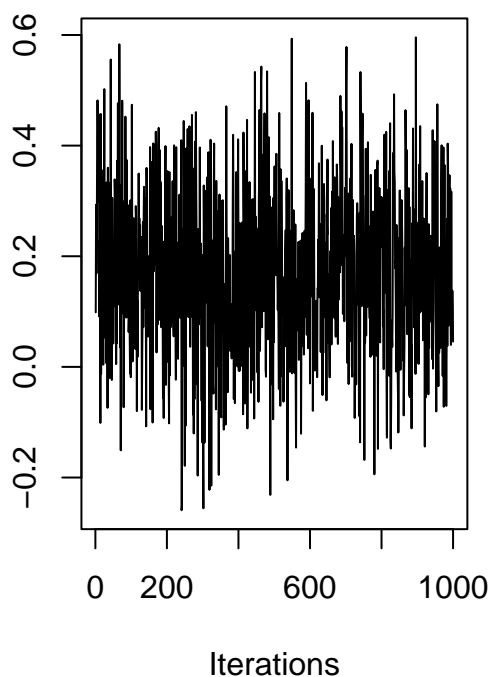


Density of var1

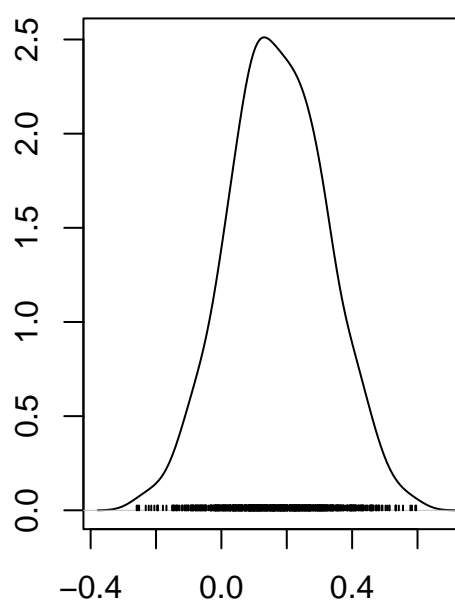


```
plot(joanpost[, "beta (Fixed - Growth)"])
```

Trace of var1



Density of var1



N = 1000 Bandwidth = 0.03973

```
quantile(joanpost[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -0.1186595  0.1673654  0.4605831
```

```
# wide
```

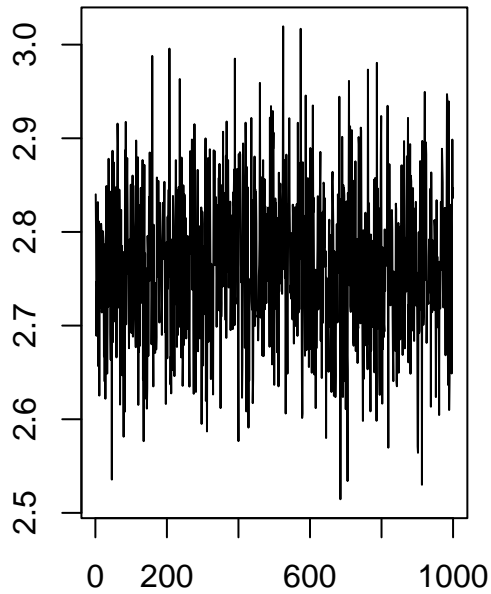
```
ttestBF(formula = JoanScore ~ Condition, data = d, rscale = "wide")
```

```
## Bayes factor analysis
## -----
## [1] Alt., r=1 : 0.2680115 ±0.05%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS
```

```
joanpost.w <- ttestBF(formula = JoanScore ~ Condition, data = d, rscale = "wide", posterior = TRUE, ite
```

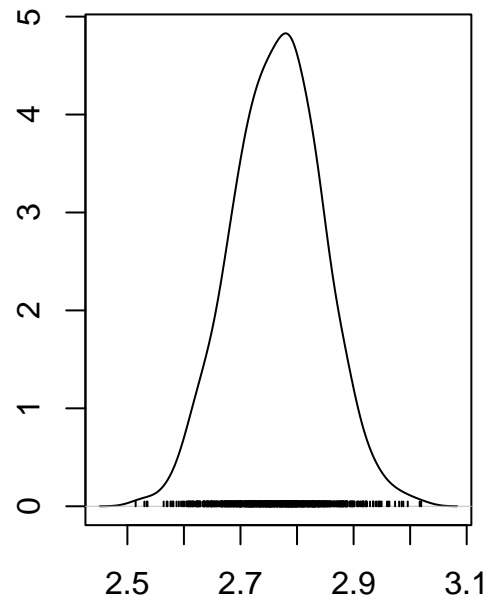
```
plot(joanpost.w[, "mu"])
```

Trace of var1



Iterations

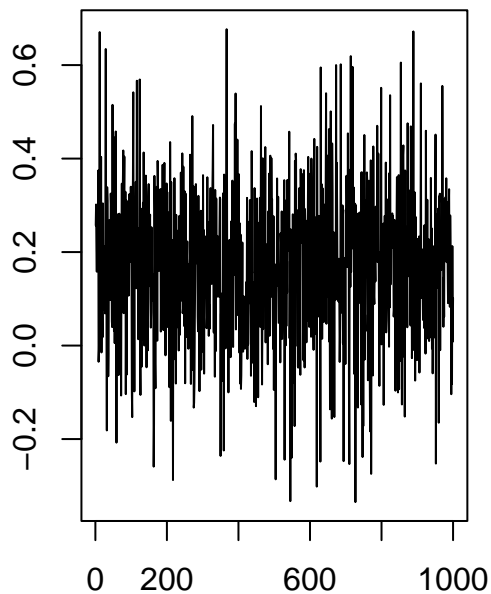
Density of var1



N = 1000 Bandwidth = 0.02115

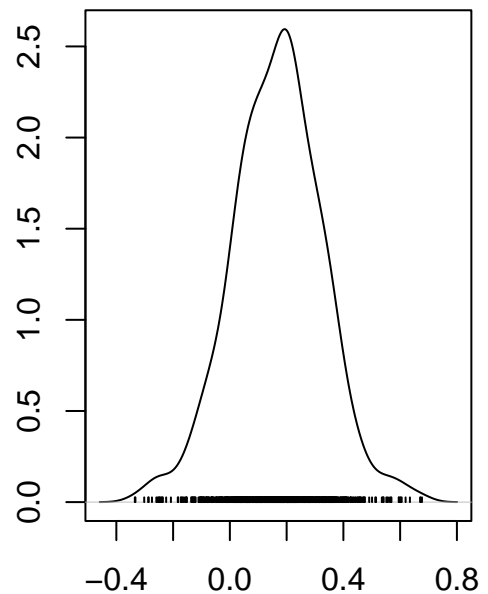
```
plot(joanpost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.04125

```
quantile(joanpost.w[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## -0.1531376  0.1739365  0.4720155
```



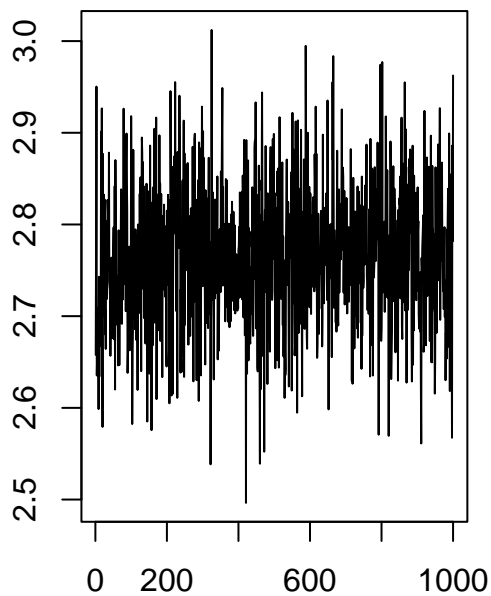
```
# ultrawide
ttestBF(formula = JoanScore ~ Condition, data = d, rscale = "ultrawide")

## Bayes factor analysis
## -----
## [1] Alt., r=1.414 : 0.1960675 ±0.09%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

joanpost.uw <- ttestBF(formula = JoanScore ~ Condition, data = d, rscale = "ultrawide", posterior = TRUE)

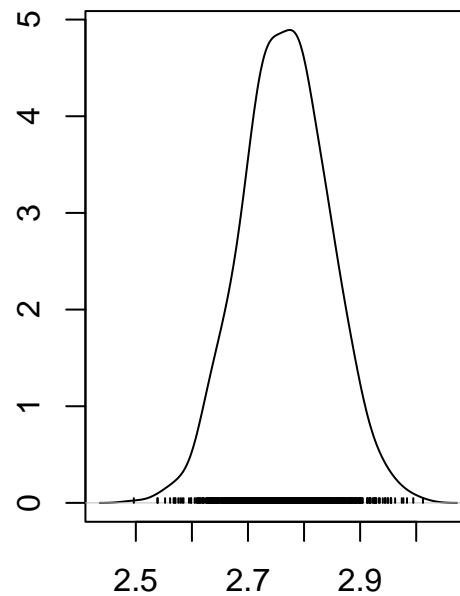
plot(joanpost.uw[, "mu"])
```

Trace of var1



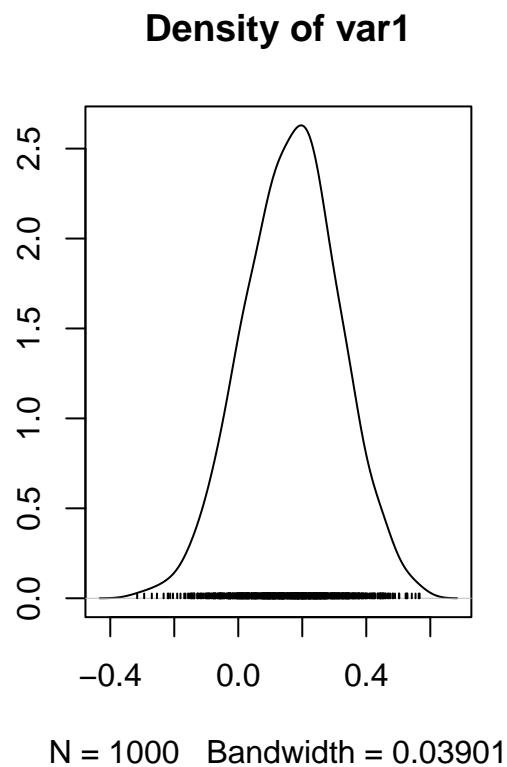
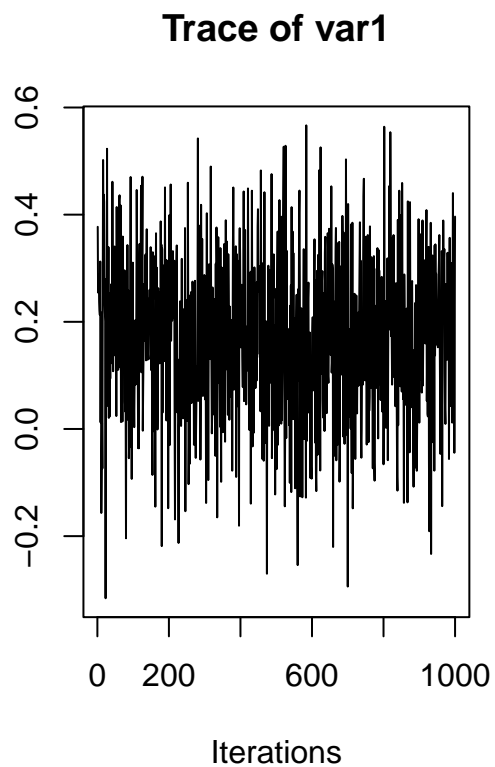
Iterations

Density of var1



N = 1000 Bandwidth = 0.02026

```
plot(joanpost.uw[, "beta (Fixed - Growth)"])
```



```
quantile(joanpost.uw[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

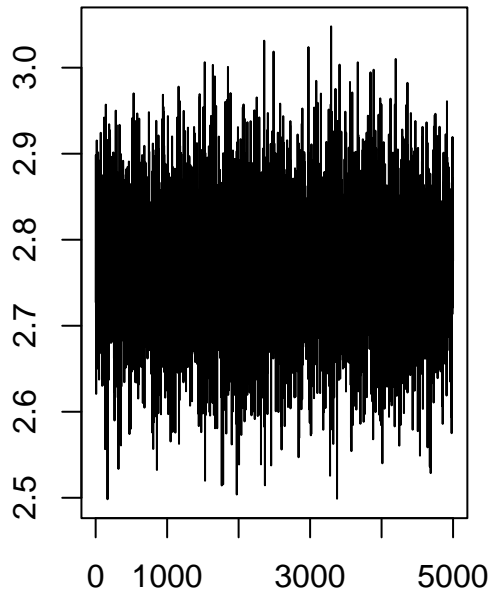
```
##      2.5%      50%      97.5%
## -0.1274836  0.1734112  0.4507211
```

```
### more iterations
```

```
joanpost.check <- ttestBF(formula = JoanScore ~ Condition, data = d, posterior = TRUE, iterations = 500)
```

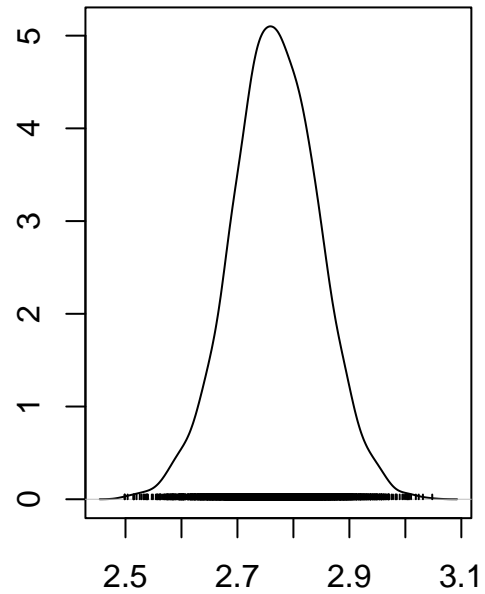
```
plot(joanpost.check[, "mu"])
```

Trace of var1



Iterations

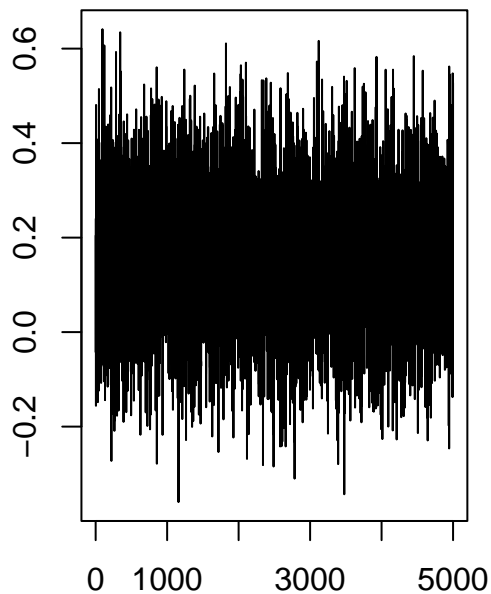
Density of var1



N = 5000 Bandwidth = 0.01478

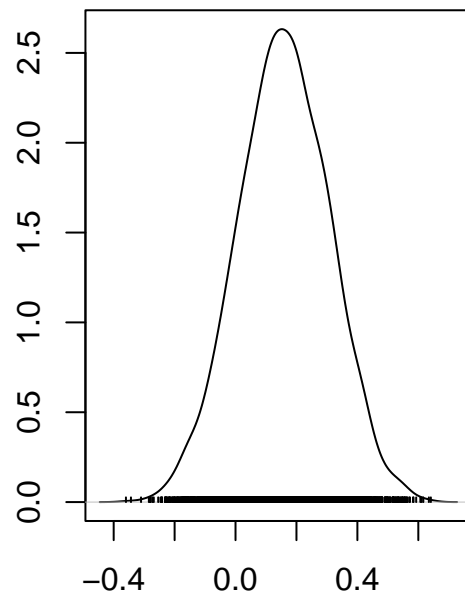
```
plot(joanpost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.02868

```
quantile(joanpost.check[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## -0.1395923  0.1578671  0.4416287
```

Divergent thinking

note, two participants failed to respond to the DT prompts (row 17 and 25) and were not included in this analysis

Divergent Thinking Originality

```
## first make the subset data
dt.subset <- na.omit(d[,c("DTScore", "DTFlu", "Condition")])

## DT originality ratings

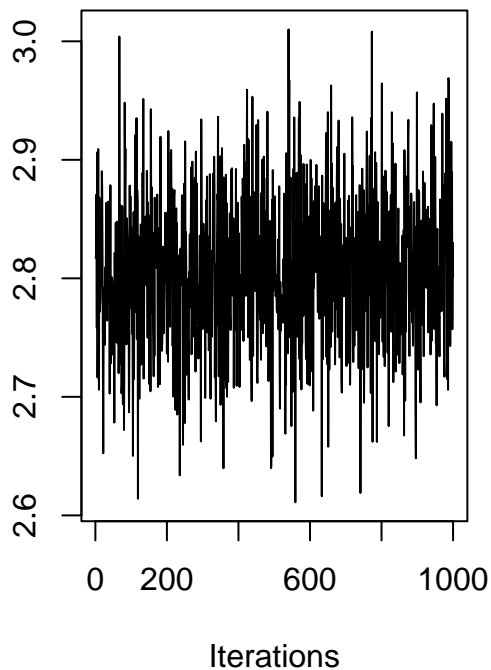
# Default / medium
ttestBF(formula = DTScore ~ Condition, data = dt.subset)

## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 0.2056849 ±0.03%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

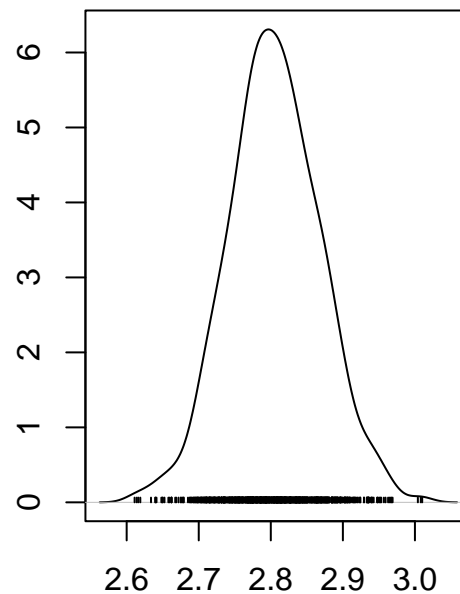
dtpost <- ttestBF(formula = DTScore ~ Condition, data = dt.subset, posterior = TRUE, iterations = 1000)

plot(dtpost[, "mu"])
```

Trace of var1



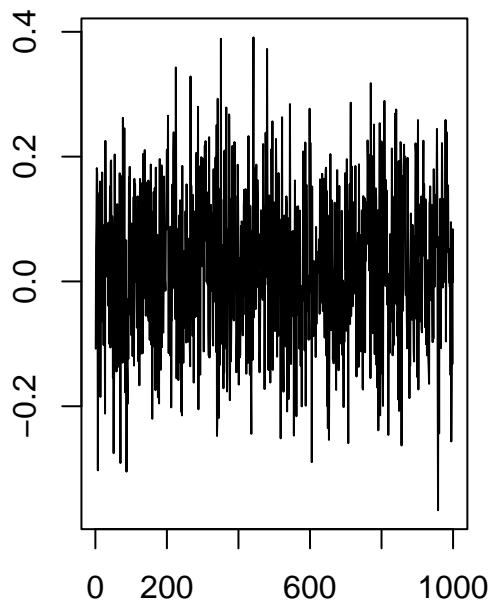
Density of var1



N = 1000 Bandwidth = 0.01614

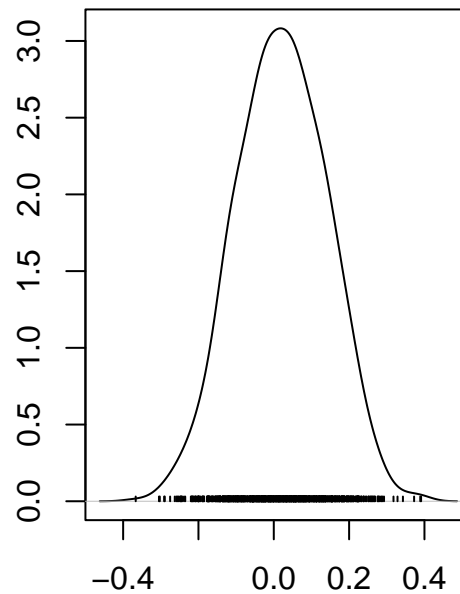
```
plot(dtpost[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.03183

```
quantile(dtpost[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##          2.5%          50%          97.5%
## -0.20890209  0.02049619  0.24566789
```

```
# wide
```

```
ttestBF(formula = DTScore ~ Condition, data = dt.subset, rscale = "wide")
```

```
## Bayes factor analysis
```

```
## -----
```

```
## [1] Alt., r=1 : 0.1498916 ±0.08%
```

```
##
```

```
## Against denominator:
```

```
## Null, mu1-mu2 = 0
```

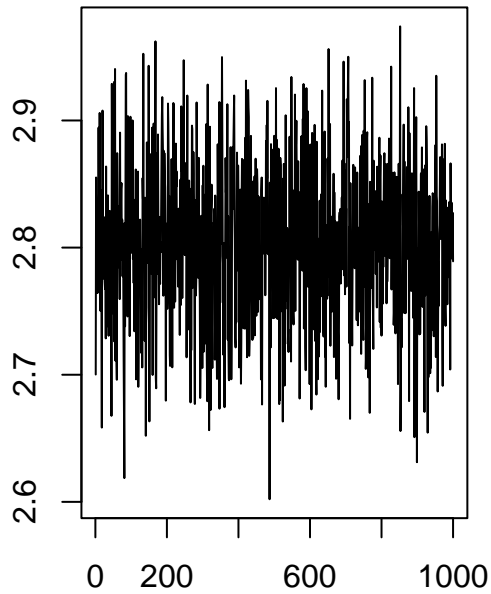
```
## ---
```

```
## Bayes factor type: BFindepSample, JZS
```

```
dtpost.w <- ttestBF(formula = DTScore ~ Condition, data = dt.subset, rscale = "wide", posterior = TRUE,
```

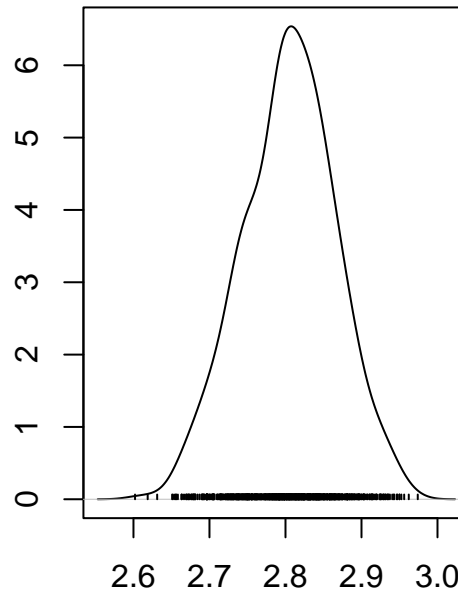
```
plot(dtpost.w[, "mu"])
```

Trace of var1



Iterations

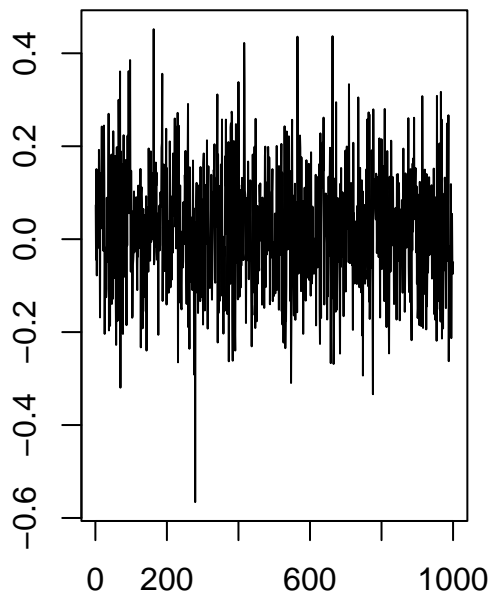
Density of var1



N = 1000 Bandwidth = 0.01636

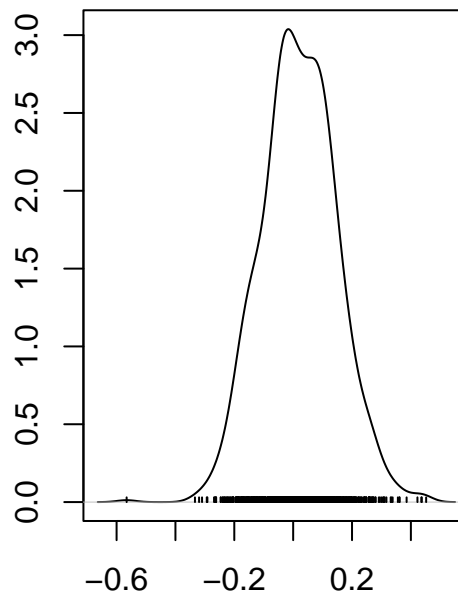
```
plot(dtpost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.03275

```
quantile(dtpost.w[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

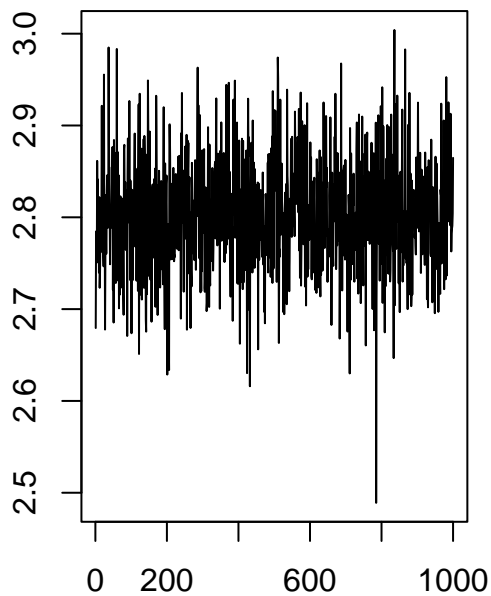
```
##      2.5%      50%      97.5%  
## -0.2174833  0.0188377  0.2685011
```

```
# ultra wide
ttestBF(formula = DTScore ~ Condition, data = dt.subset, rscale = "ultrawide")

## Bayes factor analysis
## -----
## [1] Alt., r=1.414 : 0.1077728 ±0.15%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS

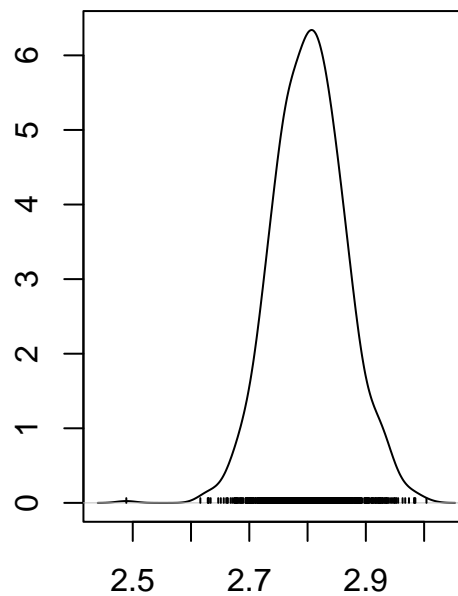
dtpost.uw <- ttestBF(formula = DTScore ~ Condition, data = dt.subset, rscale = "ultrawide", posterior =
plot(dtpost.uw[, "mu"])
```

Trace of var1



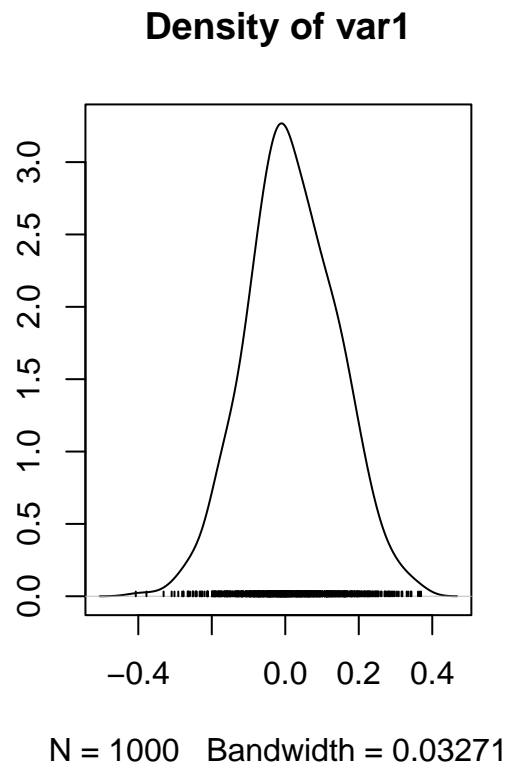
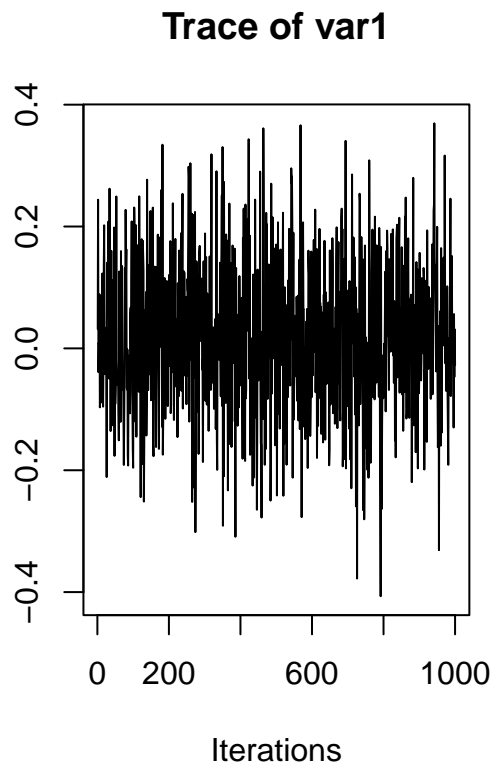
Iterations

Density of var1



N = 1000 Bandwidth = 0.01635

```
plot(dtpost.uw[, "beta (Fixed - Growth)"])
```



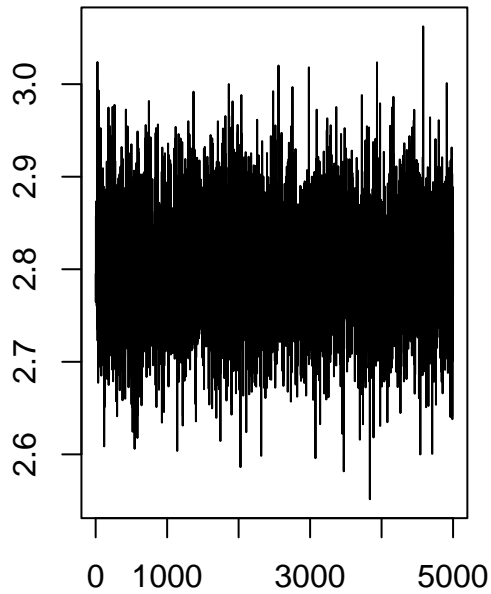
```
quantile(dtpost.uw[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##          2.5%          50%          97.5%
## -0.21249611  0.01019647  0.25403206
```

```
# More iterations
```

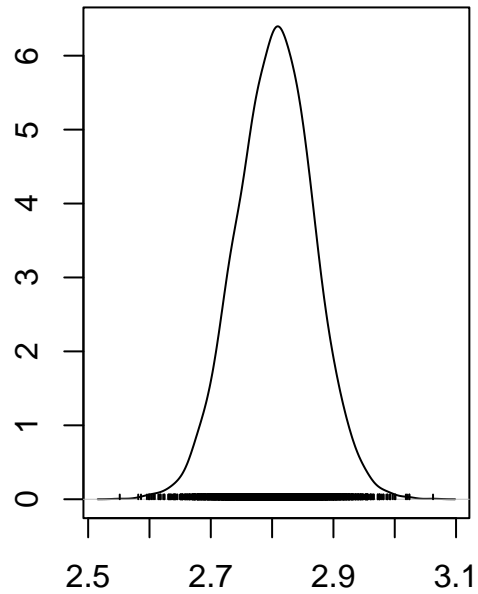
```
dtpost.check <- ttestBF(formula = DTScore ~ Condition, data = dt.subset, posterior = TRUE, iterations = 
plot(dtpost.check[, "mu"])
```


Trace of var1



Iterations

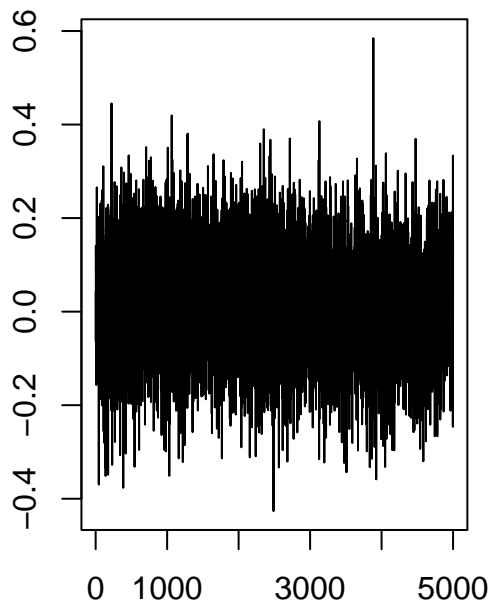
Density of var1



N = 5000 Bandwidth = 0.01198

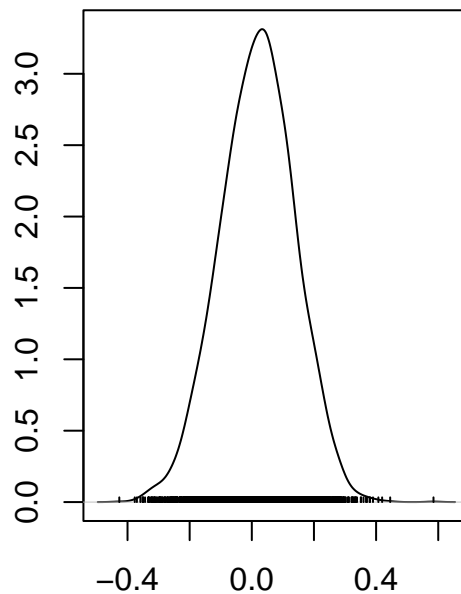
```
plot(dtpost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.02316

```
quantile(dtpost.check[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

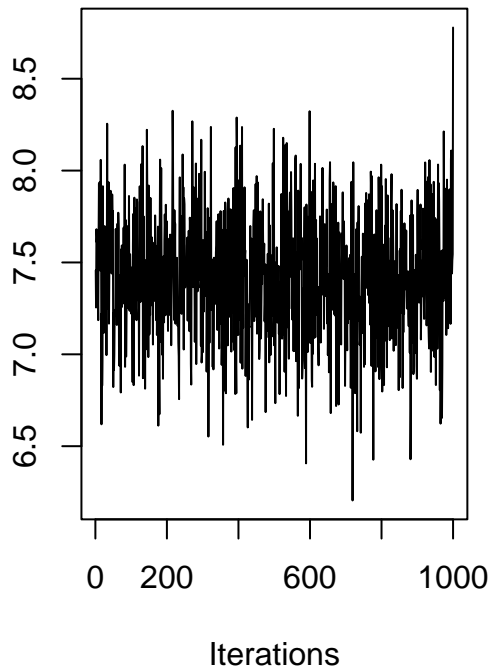
```
##          2.5%          50%          97.5%  
## -0.21780014  0.02103769  0.24612178
```

DT fluency

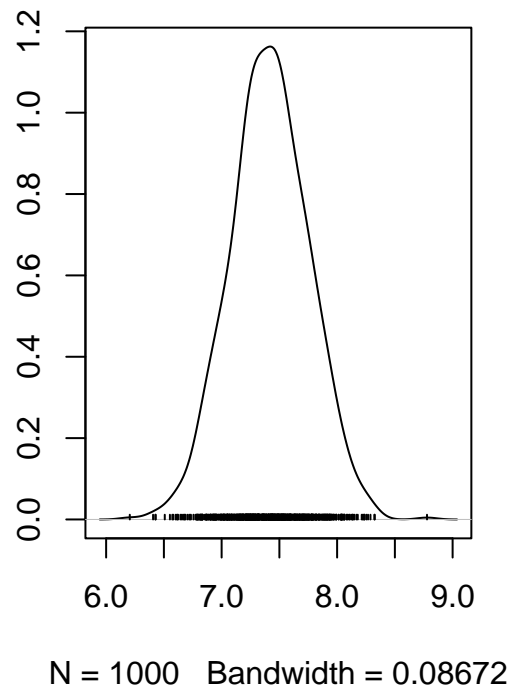
```
# default / medium
ttestBF(formula = DTFlu ~ Condition, data = dt.subset)

## Bayes factor analysis
## -----
## [1] Alt., r=0.707 : 0.7210424 ±0.02%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS
flupost <- ttestBF(formula = DTFlu ~ Condition, data = dt.subset, posterior = TRUE, iterations = 1000)
plot(flupost[, "mu"])
```

Trace of var1

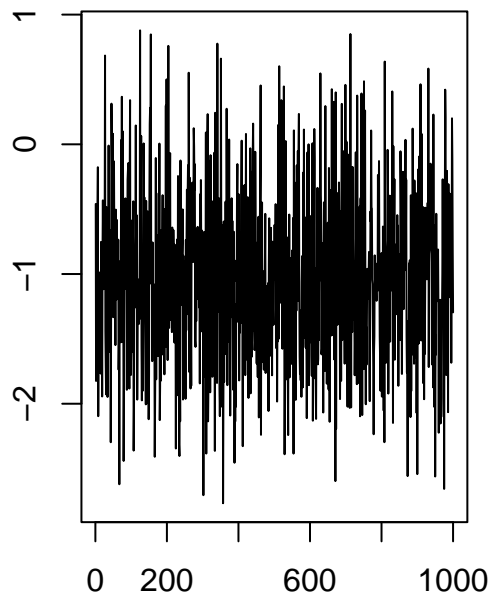


Density of var1



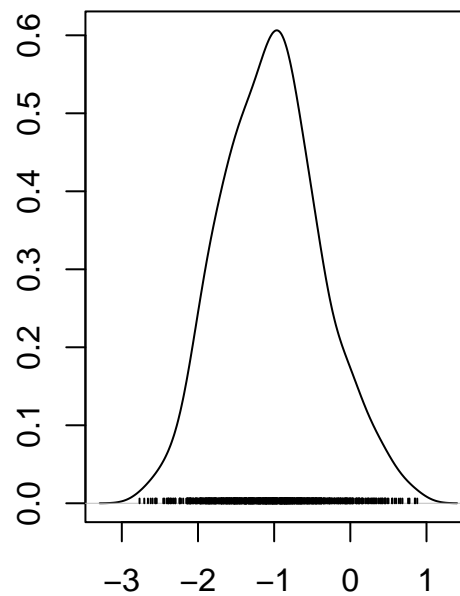
```
plot(flupost[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.1742

```
quantile(flupost[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%
## -2.2237639 -1.0404673  0.3412296
```

```
# wide
```

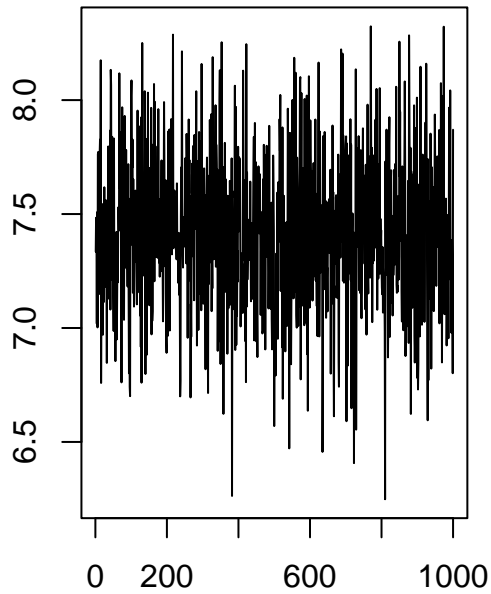
```
ttestBF(formula = DTFlu ~ Condition, data = dt.subset, rscale = "wide")
```

```
## Bayes factor analysis
## -----
## [1] Alt., r=1 : 0.5596128 ±0.03%
##
## Against denominator:
##   Null, mu1-mu2 = 0
## ---
## Bayes factor type: BFindepSample, JZS
```

```
flupost.w <- ttestBF(formula = DTFlu ~ Condition, data = dt.subset, rscale = "wide", posterior = TRUE, ...)
```

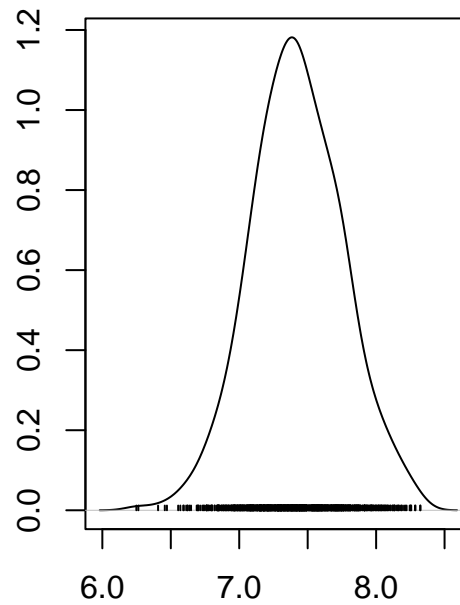
```
plot(flupost.w[, "mu"])
```

Trace of var1



Iterations

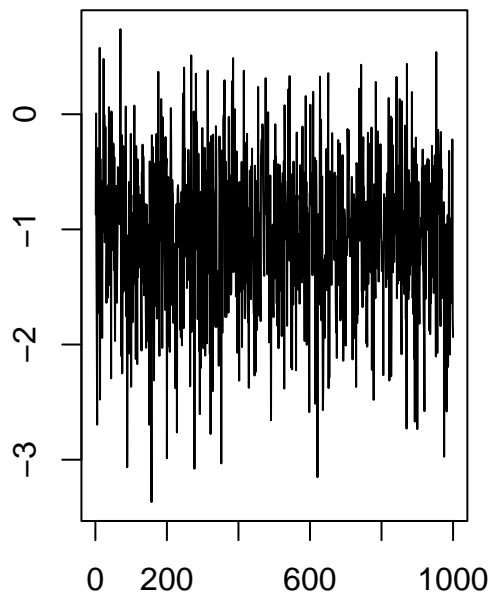
Density of var1



N = 1000 Bandwidth = 0.08925

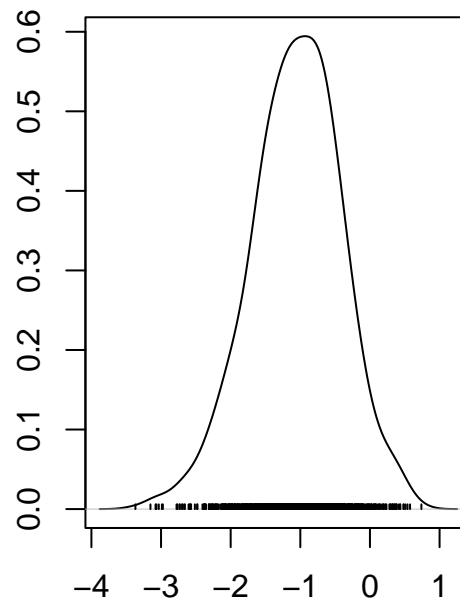
```
plot(flupost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.1709

```
quantile(flupost.w[, "beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## -2.3803232 -1.0393014  0.1930067
```

```
# ultrawide
```

```
ttestBF(formula = DTFlu ~ Condition, data = dt.subset, rscale = "ultrawide")
```

```
## Bayes factor analysis
```

```
## -----
```

```
## [1] Alt., r=1.414 : 0.4183677 ±0.05%
```

```
##
```

```
## Against denominator:
```

```
## Null, mu1-mu2 = 0
```

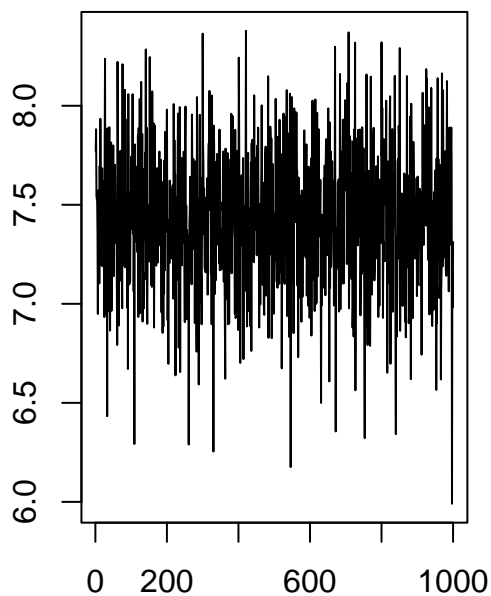
```
## ---
```

```
## Bayes factor type: BFindepSample, JZS
```

```
flupost.uw <- ttestBF(formula = DTFlu ~ Condition, data = dt.subset, rscale = "ultrawide", posterior = 'f')
```

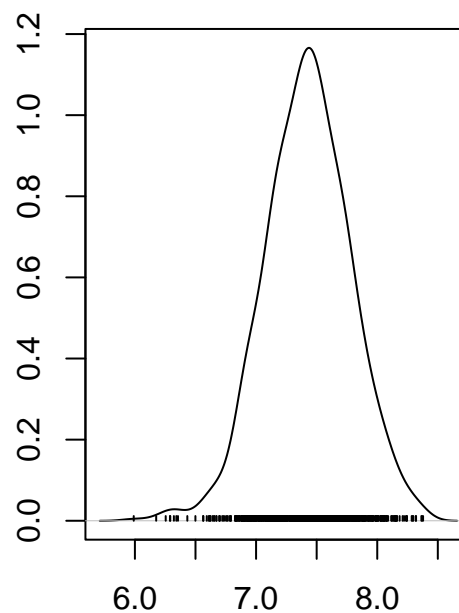
```
plot(flupost.uw[, "mu"])
```

Trace of var1



Iterations

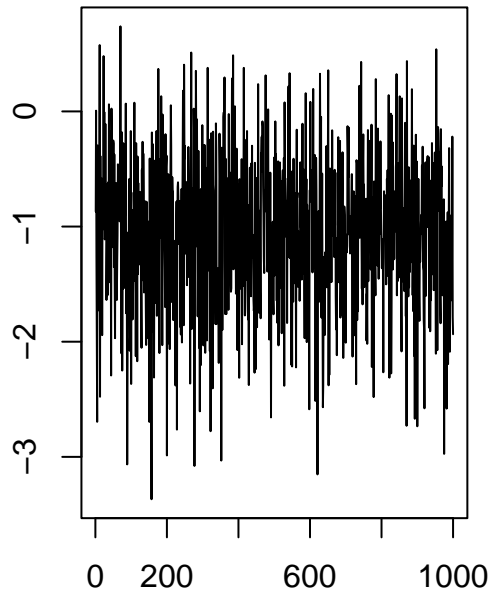
Density of var1



N = 1000 Bandwidth = 0.09356

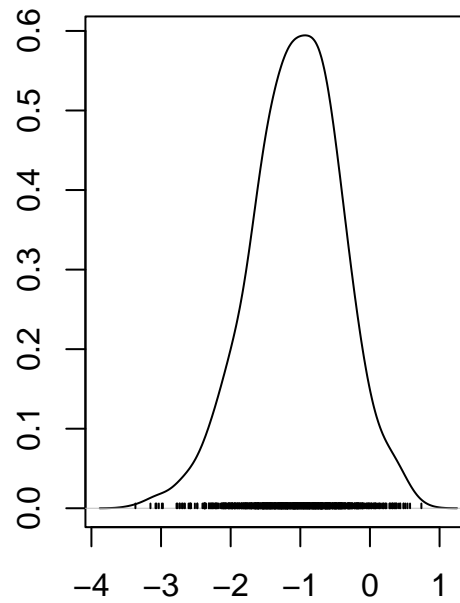
```
plot(flupost.w[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 1000 Bandwidth = 0.1709

```
quantile(flupost.uw["beta (Fixed - Growth)", probs = c(.025, .5, .975))
```

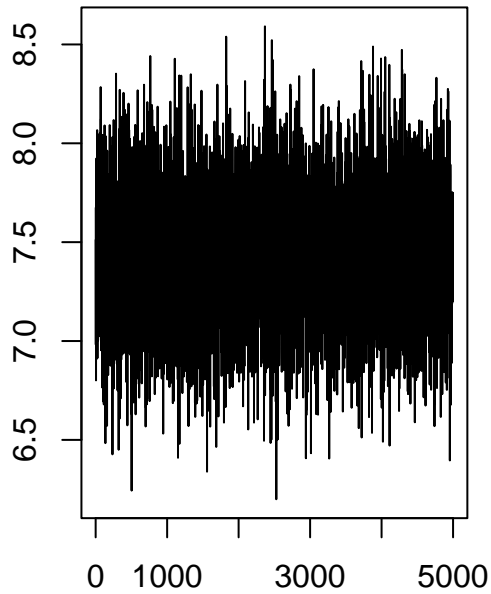
```
##      2.5%      50%      97.5%  
## -2.4107607 -1.1280036  0.0965674
```

```
## More iterations
```

```
flupost.check <- ttestBF(formula = DTFlu ~ Condition, data = dt.subset, posterior = TRUE, iterations = 10000)
```

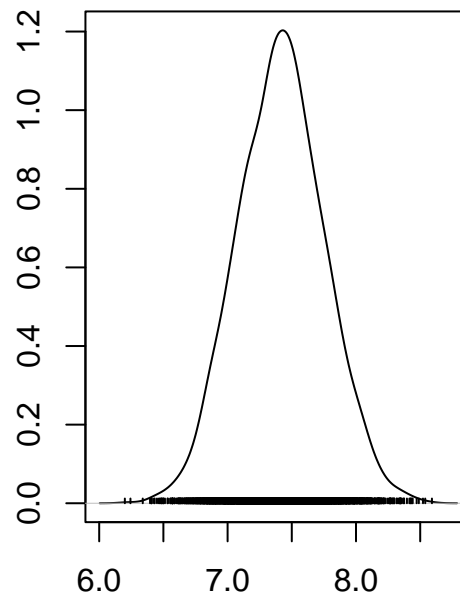
```
plot(flupost.check[, "mu"])
```

Trace of var1



Iterations

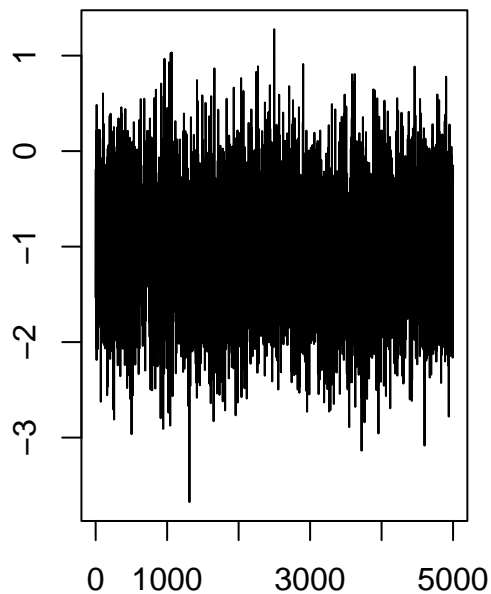
Density of var1



N = 5000 Bandwidth = 0.06521

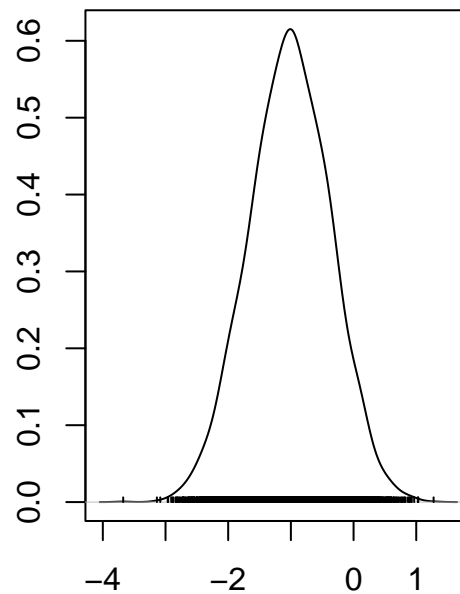
```
plot(flupost.check[, "beta (Fixed - Growth)"])
```

Trace of var1



Iterations

Density of var1



N = 5000 Bandwidth = 0.1247

```
quantile(flupost.check[, "beta (Fixed - Growth)"], probs = c(.025, .5, .975))
```

```
##      2.5%      50%      97.5%  
## -2.300783 -1.011913  0.212568
```

Cohen's D for all effects

```
# effect sizes
cohen.d(Fixed ~ Condition, data = d)

## Call: cohen.d(x = Fixed ~ Condition, data = d)
## Cohen d statistic of difference between two means
##      lower effect upper
## Fixed -0.92  -0.54 -0.16
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.54
## r equivalent of difference between two means
## Fixed
## -0.26

cohen.d(Growth ~ Condition, data = d)

## Call: cohen.d(x = Growth ~ Condition, data = d)
## Cohen d statistic of difference between two means
##      lower effect upper
## Growth  0.24   0.63  1.01
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.63
## r equivalent of difference between two means
## Growth
##    0.3

cohen.d(CSE ~ Condition, data = d)

## Call: cohen.d(x = CSE ~ Condition, data = d)
## Cohen d statistic of difference between two means
##      lower effect upper
## CSE -0.57  -0.19  0.18
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.19
## r equivalent of difference between two means
## CSE
## -0.1

cohen.d(JoanScore ~ Condition, data = d)

## Call: cohen.d(x = JoanScore ~ Condition, data = d)
## Cohen d statistic of difference between two means
##      lower effect upper
## JoanScore -0.59  -0.22  0.16
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.22
## r equivalent of difference between two means
## JoanScore
##    -0.11

cohen.d(DTScore ~ Condition, data = dt.subset)
```



```
## Call: cohen.d(x = DTScore ~ Condition, data = dt.subset)
## Cohen d statistic of difference between two means
##      lower effect upper
## DTScore -0.41  -0.03  0.34
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.033
## r equivalent of difference between two means
## DTScore
##   -0.02
```

```
cohen.d(DTFlu ~ Condition, data = dt.subset)
```

```
## Call: cohen.d(x = DTFlu ~ Condition, data = dt.subset)
## Cohen d statistic of difference between two means
##      lower effect upper
## DTFlu -0.05   0.33   0.7
##
## Multivariate (Mahalanobis) distance between groups
## [1] 0.33
## r equivalent of difference between two means
## DTFlu
##   0.16
```

Hypothesis 3 and 4

```
# correlations from BF including posterior
```

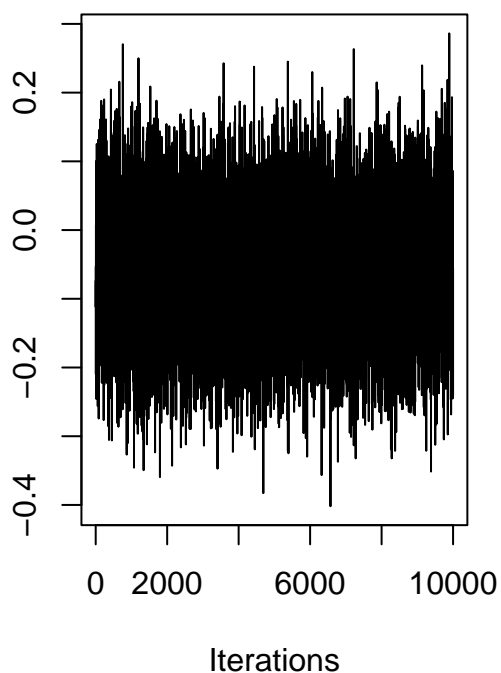
```
# hypothesis 3
```

```
samp3 <- correlationBF(y = d$JoanScore, x = d$Fixed, posterior = T, iterations = 10000)
```

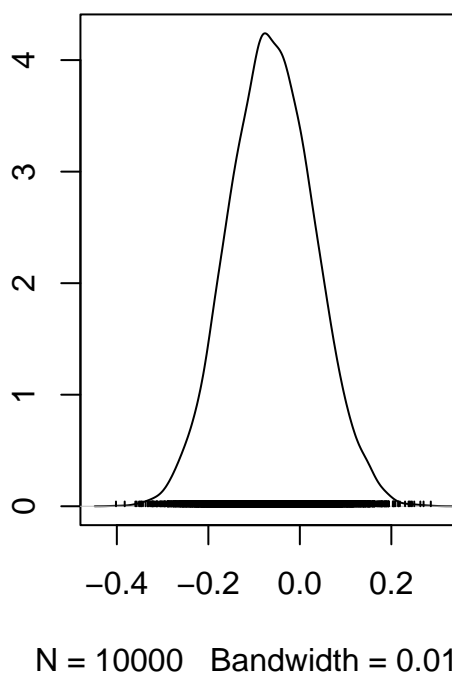
```
## Independent-candidate M-H acceptance rate: 98%
```

```
plot(samp3[, "rho"])
```

Trace of var1



Density of var1



```
summary(samp3)
```

```
##
## Iterations = 1:10000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 10000
##
## 1. Empirical mean and standard deviation for each variable,
##    plus standard error of the mean:
##
##      Mean      SD Naive SE Time-series SE
## rho  -0.06171 0.09215 0.0009215      0.0009824
## zeta  -0.06232 0.09328 0.0009328      0.0009943
##
## 2. Quantiles for each variable:
##
##      2.5%    25%    50%    75%  97.5%
## rho  -0.2396 -0.1253 -0.06311 0.0005138 0.1213
## zeta  -0.2444 -0.1259 -0.06320 0.0005138 0.1219
```

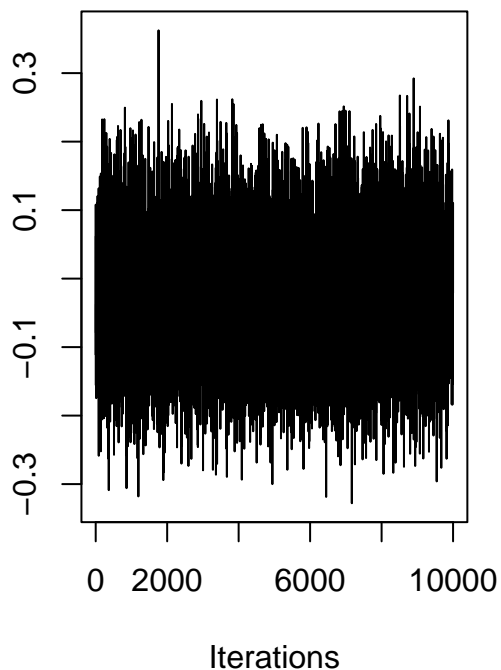
```
# hypothesis 4
```

```
samp4 <- correlationBF(y = d$DTFlu, x = d$CSE, posterior = T, iterations = 10000)
```

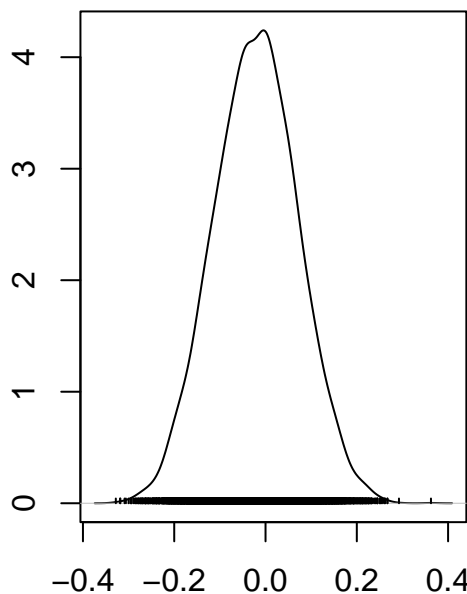
```
## Independent-candidate M-H acceptance rate: 98%
```

```
plot(samp4[, "rho"])
```

Trace of var1



Density of var1



N = 10000 Bandwidth = 0.01541

```
summary(samp4)
```

```
##
## Iterations = 1:10000
## Thinning interval = 1
## Number of chains = 1
## Sample size per chain = 10000
##
## 1. Empirical mean and standard deviation for each variable,
##    plus standard error of the mean:
##
##      Mean      SD Naive SE Time-series SE
## rho  -0.02098 0.09175 0.0009175      0.0009560
## zeta  -0.02116 0.09255 0.0009255      0.0009644
##
## 2. Quantiles for each variable:
##
##      2.5%      25%      50%      75%  97.5%
## rho  -0.1999 -0.08383 -0.02064 0.04139 0.1587
## zeta  -0.2026 -0.08403 -0.02064 0.04142 0.1601
```

For checking multiple choice answers

```
McheckF <- d[d$Condition == "Fixed",c("Condition","Q12","Q20")]
table(McheckF$Q12) / nrow(McheckF)
```

```
##
##      1      2      3
## 0.2727273 0.6181818 0.1090909
```

```
table(McheckF$Q20) / nrow(McheckF)
```

```
##  
##          1          2          3  
## 0.05454545 0.76363636 0.18181818
```

```
McheckG <- d[d$Condition == "Growth",c("Condition","Q21","Q11")]  
table(McheckG$Q21) / nrow(McheckG)
```

```
##  
##          1          2          3  
## 0.89285714 0.07142857 0.03571429
```

```
table(McheckG$Q11) / nrow(McheckG)
```

```
##  
##          1          2          3  
## 0.92857143 0.01785714 0.05357143
```