

When the sample size is two, a 50% confidence interval is defined by the two values

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1 Sample two realisations from a normally distributed population

```
alpha <- 0.5           # 50% confidence

n <- 2                 # sample size
mu <- 0                # population mean
sigma <- 1             # population standard deviation

x <- rnorm(n, mu, sigma) # generate the sample

sort(x)
```

```
[1] -0.03521958  1.78251112
```

2 Calculate the confidence interval

```
error <- qt(1-alpha/2, df=n-1)*sd(x)/sqrt(n)
mean(x) + c(-1, +1) * error
```

```
[1] -0.03521958  1.78251112
```

3 Calculate again using one sample t-test function

```
t.test(x, conf.level = alpha)
```

One Sample t-test

```
data: x
t = 0.96125, df = 1, p-value = 0.5126
alternative hypothesis: true mean is not equal to 0
50 percent confidence interval:
 -0.03521958  1.78251112
sample estimates:
mean of x
0.8736458
```

4 References

<https://github.com/eamonn2014/programs/blob/master/50%20confidence%20interval.Rmd>

5 Computing Environment

R version 3.2.2 (2015-08-14)

Platform: x86_64-w64-mingw32/x64 (64-bit)

Running under: Windows 8 x64 (build 9200)

locale:

```
[1] LC_COLLATE=English_United Kingdom.1252
[2] LC_CTYPE=English_United Kingdom.1252
[3] LC_MONETARY=English_United Kingdom.1252
[4] LC_NUMERIC=C
[5] LC_TIME=English_United Kingdom.1252
```

attached base packages:

```
[1] stats      graphics  grDevices  utils      datasets  methods
[7] base
```

other attached packages:

```
[1] knitr_1.14
```

loaded via a namespace (and not attached):

```
[1] magrittr_1.5      formatR_1.4      tools_3.2.2      htmltools_0.3.5
[5] yaml_2.1.13       Rcpp_0.12.6      stringi_1.1.1    rmarkdown_1.0
[9] stringr_1.1.0     digest_0.6.10    evaluate_0.9
```

```
[1] "~/GIT\\50-percent-confidence-interval"
```

This took 0.21 seconds to execute.