Confidence Intervals

Using the Central Limit Theorem for large n we have

$$P(\mid rac{\sqrt{n}(ar{X}-\mu)}{\sigma}\mid \leq 1.96)pprox 0.95$$
 -

Assume CUDKUnknown $E(X) = \mu$ $Var(X) = \sigma^2 - Known$

which is the same as saying

$$P(\mu \in \left(-rac{1.96\sigma}{\sqrt{n}} + ar{X}, rac{1.96\sigma}{\sqrt{n}} + ar{X}
ight)) pprox 0.95$$

Sample X, Xy. Xn from

The interval $\left(-\frac{1.96\sigma}{\sqrt{n}} + \bar{X}, \frac{1.96\sigma}{\sqrt{n}} + \bar{X}\right)$ is called the 95% confidence interval for μ . μ is dependent on Sample. and is unlice if χ_{n} is called the 95% confidence interval for μ .

Confidence Intervals - Important - Previse meaning.

Meaning: for n large if we did m (large) repeated trials and computed the above interval for each trial then true mean would belong to approximately 95% of m intervals calculated.

The below is code for finding the confidence interval for a data x.

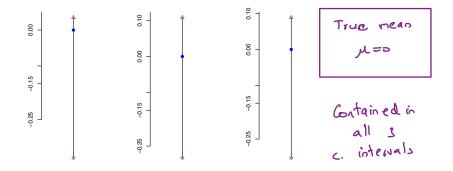
```
> cifn = function(x, alpha=0.95){
  + z = qnorm( (1-alpha)/2, lower.tail=FALSE)
  + sdx = sqrt(1/length(x))
  + c(mean(x) - z*sdx, mean(x) + z*sdx)
```

```
> x1 = rnorm(100, 0, 1); y = cifn(x1)
> y
[1] -0.35705304 0.03493976
> x^2 = rnorm(100, 0, 1); z = cifn(x^2)
> z
[1] -0.2832489 0.1087439
> x3 = rnorm(100,0,1); w = cifn(x3)
> w
[1] -0.30294682 0.08904598
```

Does 0 belong to all the three confidence intervals ?

rnorm(100,0,1) = data.

The below is a plot of the three confidence intervals computed in the previous slide.



We generate 10 trials of 100 samples from Normal(0,1) and compute the confidence intervals using the function defined earlier.

> normaldata = replicate(10, rnorm(100,0,1),

```
+ simplify=FALSE)
```

> cidata = sapply(normaldata, cifn)

It is easy to check how many of them contain 0.

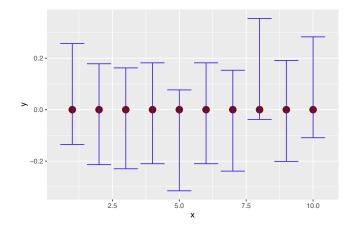
> TRUEIN = cidata[1,]*cidata[2,]<0</pre>

```
> table(TRUEIN)
```

(9,5) 70 jj a.b<0

```
TRUE
```

Confidence Intervals : 10 Trials



We generate 10 trials of 100 samples from Normal(0,1) and compute the confidence intervals using the function defined earlier.

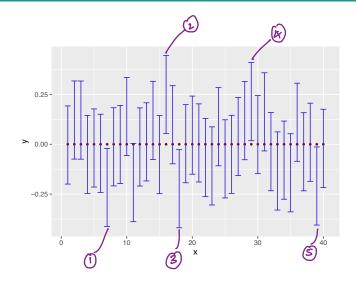
- > normaldata = replicate(40, rnorm(100,0,1),
- + simplify=FALSE)
- > cidata = sapply(normaldata, cifn)

It is easy to check how many of them contain 0.

- > TRUEIN = cidata[1,]*cidata[2,]<0</pre>
- > table(TRUEIN)

- FALSE TRUE
 - 5 35

Confidence Intervals: 40 trials Plot



We generate 100 trials of 100 samples from Normal(0,1) and compute the confidence intervals using the function defined earlier.

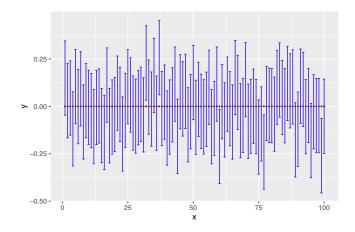
- > normaldata = replicate(100, rnorm(100,0,1),
- + simplify=FALSE)
- > cidata = sapply(normaldata, cifn)

It is easy to check how many of them contain 0.

- > TRUEIN = cidata[1,]*cidata[2,]<0</pre>
- > table(TRUEIN)

- FALSE TRUE
 - 5 95

Confidence Intervals : 100 Trials



We generate 1000 trials of 100 samples from Normal(0,1) and compute the confidence intervals using the function defined earlier.

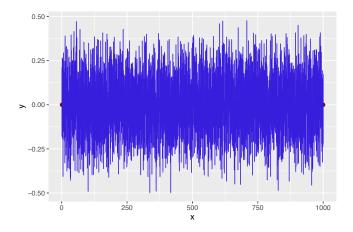
- > normaldata = replicate(1000, rnorm(100,0,1),
- + simplify=FALSE)
- > cidata = sapply(normaldata, cifn)

It is easy to check how many of them contain 0.

- > TRUEIN = cidata[1,]*cidata[2,]<0</pre>
- > table(TRUEIN)

- FALSE TRUE
 - 54 946

Confidence Intervals : 1000 Trials



95% confidence interval for μ is $\left(-\frac{1.96\sigma}{\sqrt{n}} + \bar{X}, \frac{1.96\sigma}{\sqrt{n}} + \bar{X}\right)$

Meaning: for n large if we did m (large) repeated trials and computed the above interval for each trial then true mean would belong to approximately 95% of m intervals calculated.

Thus numerically the above meaning seems to hold for a Normal population.