STAT 4234/5234: Calculating the regression estimator for a population total

Consider the population $\{(x_i, y_i) : i = 1, ..., N\}$ and suppose we wish to estimate the population mean \bar{y}_U based on a simple random sample of size n. We further suppose the value of \bar{x}_U , the population mean for the auxiliary variable, is known. In regression estimation we estimate \bar{y}_U by

$$\hat{y}_{\text{reg}} = \bar{y} + \hat{B}_1(\bar{x}_U - \bar{x}) = \bar{y} + r \frac{s_y}{s_x}(\bar{x}_U - \bar{x})$$

where s_x and s_y are the sample standard deviations of x and y, respectively, and r denotes the sample correlation coefficient.

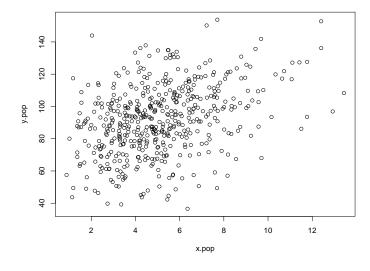
We have further seen that

SE
$$(\hat{y}_{reg}) = \sqrt{\frac{s_y^2(1-r^2)}{n} \left(1-\frac{n}{N}\right)}$$

gives an expression for the standard error of the regression estimator.

To illustrate the computing for regression estimation, we create a fictional population of (x_i, y_i) as follows.

```
> set.seed(5234)
> x.pop <- rgamma(500, shape=5, rate=1)
> y.pop <- rnorm(500, mean=75+3*x.pop, sd=20)
> plot(x.pop, y.pop)
> cor(x.pop, y.pop)
[1] 0.3611513
> xbar.U <- mean(x.pop); xbar.U;
[1] 5.095567</pre>
```



The population correlation is R = 0.36, and the auxiliary variable population mean is $\bar{x}_U = 5.10$.

Now we take a simple random sample of size n = 25.

```
> N <- 500; n <- 25;
> samp <- sample(N, n)</pre>
> x.samp <- x.pop[samp]; y.samp <- y.pop[samp];</pre>
Calculate the regression estimator \hat{\bar{y}}_{reg}.
> xbar <- mean(x.samp); ybar <- mean(y.samp);</pre>
> fit <- lsfit(x.samp, y.samp)</pre>
> names(fit)
[1] "coefficients" "residuals"
                                   "intercept"
                                                        "qr"
> fit$coefficients
Intercept
                    Χ
             2.12263
84.77118
> B1.hat <- as.numeric(fit$coefficients)[2]; B1.hat;</pre>
[1] 2.12263
> ybar.hat.reg <- ybar + B1.hat * (xbar.U - xbar)</pre>
> ybar.hat.reg
[1] 95.58719
And the standard error of our estimate.
> e <- fit$residuals</pre>
> V.hat <- var(e)/n * (1 - n/N)
> SE <- sqrt(V.hat); SE;</pre>
[1] 4.472836
Now a 95% confidence interval for \bar{y}_U is
> ybar.hat.reg + c(-1,1) * 1.96 * SE
[1] 86.82043 104.35395
And a 95% CI for the population total t_y is
> N * (ybar.hat.reg + c(-1,1) * 1.96 * SE)
[1] 43410.21 52176.97
```

Here is an R function that takes the sample data as inputs, along with N and \bar{x}_U , and returns the regression estimator of \bar{y}_U along with its standard error.

```
regression.estimator.mean <- function(x.samp, y.samp, N, xbar.U)
{
 n <- length(y.samp)</pre>
 xbar <- mean(x.samp); ybar <- mean(y.samp);</pre>
 fit <- lsfit(x.samp, y.samp)</pre>
 B1.hat <- as.numeric(fit$coefficients)[2]</pre>
 ybar.hat.reg <- ybar + B1.hat * (xbar.U - xbar)</pre>
 e <- fit$residuals
 V.hat <- var(e)/n * (1 - n/N)
 SE <- sqrt(V.hat)</pre>
 answer <- c(point.est=ybar.hat.reg, std.error=SE)</pre>
 return(answer)
}
You can use this function for your homework if you wish.
> result <- regression.estimator.mean(x.samp=x.samp,</pre>
    y.samp=y.samp, N=N, xbar.U=xbar.U)
> result
point.est std.error
95.587187 4.472836
A 95% confidence interval for \bar{y}_U is
> result[1] + c(-1,1) * 1.96 * result[2]
[1] 86.82043 104.35395
and a 95% CI for the population total t_y is
> N * (result[1] + c(-1,1) * 1.96 * result[2])
[1] 43410.21 52176.97
```