GOODNESS OF FIT

The goodness of fit of a model describes how well it fits the observations. There are several tools that can be used to evaluate it.

We start with the first "tooc": tests to assess whether the model is useful. In general, these tests evaluate the following system of hypotheses:

{ Ho: the model does not help to expension the vowability of Y

H1' the model helps to expecin the variobility of Y

· simple einear model: Yi= β1 + β2 xi + Ei (only one covariate x) The question becomes: does the inclusion of x help to explain the voriability of y?

Under to the inclusion of x is not useful: If Ho is true, the correct model is the null model Yi = P1 + E:

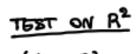
For this special case, we have already seen that we can answer to this question using a test to: $\beta_1 = 0$ vs $H_2: \beta_2 \neq 0$ (M) Test t)

To text the git of the model we can also use R2; we have seen that

 \cdot R^2 racks o : no linear relation between γ and the covariants x

· R2 × 1: strong linear relations between y and the covariate x

We can do a formal statistical Test:



$$\frac{1681^{\circ} \text{ ON N}}{\text{Ho}: R^{2}=0}$$
 \Rightarrow under the, including no under the 1 use the null model $Y_{i}=\beta_{2}+\epsilon_{i}$ \Rightarrow is not useful no under the 1 use the full model $Y_{i}=\beta_{2}+\beta_{2}$ is $+\epsilon_{i}$. Here $x_{i}^{2}\neq0$ (i.e. $x_{i}^{2}>0$)

1-R

Recall that
$$R^2 = \frac{SSR}{SST} = \frac{\sum_{i=1}^{M} (\hat{y}_i - \overline{y})^2}{\sum_{i=1}^{M} (\hat{y}_i - \overline{y})^2} = 1 - \frac{SSE}{SST}$$

$$\frac{R^{2}}{88T} = \frac{83R}{88T} \cdot \left(1 - \frac{83R}{58T}\right)^{-1} = \frac{83R}{58T} \cdot \frac{58T}{55T-55R} = \frac{55R}{55}$$

$$= \frac{\sum_{i=1}^{h} (y_i - \overline{y})^2}{\sum_{i=1}^{h} (y_i - \hat{y}_i)^2} - 1$$

what are the two quartities A and B?

(A) is the suk of the sawared residuals of the null hobel: model with oney the intercept Since the null model is the model assumed under to, (A) is the sun of southress Residuals under tho.

Recall that if Yi = \beta + &i => the estimate is \beta = \beta => the predicted values are gi = g for all i

Let's define the residuals yi-y = et sum of squared residuals is $\sum_{i=1}^{n}(y_i-\overline{y})^2 = \sum_{i=1}^{n}e_i^{n-1}$

(B) is the sum of squared restinals of the full kodel.

- the model is the unconstrained model (i.e., the model under H1).
- (B) is the sun of somered residuals under H1. Rodel Y:= P1+P2 xi+Ei
- Let's define the residuals yi- gi = ei
- sum of squared residuals is $\sum_{i=1}^{\infty} (y_i \hat{y}_i)^2 = \sum_{i=1}^{\infty} e_i^2$

Returning now to the TEST STATISFIC

$$\frac{R^{2}}{4-R^{2}} = \frac{SST}{SSE} - 1 = \frac{SSE_{H_{0}}}{SSE_{H_{1}}} - 1 = \frac{\sum_{i=1}^{n} e_{i}^{*4}}{\sum_{i=1}^{n} e_{i}^{*2}} - 1$$

and the residuals of the model that includes x. $\sum_{i=1}^{n} e_{i}^{2} = n e^{2}$ where $e^{2} = \frac{1}{n} \sum_{i=1}^{n} (y_{i} - \overline{y})^{2}$ is the entermate of the variance of the error

we are composing the residuals of the model we would estimate in the observe of information (ie, x)

under the model with only the intercept (Ho). The denomination is $\sum_{i=1}^{n} e^{it} = n\hat{\sigma}^2$ where $\hat{\sigma}^2 = \frac{1}{n} \sum_{i=1}^{n} (y_i - \hat{y}_i)^2$ is the eithrete of the variance of the error

under the gull model (H2).

Hence,
$$\frac{R^2}{4-R^2} = \frac{\sum_{i=1}^{n} e_i^{n+2}}{\sum_{i=1}^{n} e_i^{n+2}} - 4 = \frac{n^{\frac{n}{2}}}{n^{\frac{n}{2}}} - 4 = \frac{6^2}{6^2} - 4 = \frac{6^2}{6^2}$$

$$\rightarrow \text{ we are comparing the estimated Volumes of the error under the two models.}$$

Now, we need to study what values the test statistic can assume First of all, notice that the quantity is always positive

What values of the test statistic do we expect under the and the?

ine., how are the REJECT and ACCEPTANCE REGIONS defined?

- hence the models under the and the will have similar performances at predicting y. (the full model can not be worse in terms of prediction, at most is the same as the null model)

· IF HO IS TRUE, X is not useful in explaining y

- → If the predictions under the Two models are similar, also the residuals will be similar → the "total amount of error" of the two models will be similar
- \rightarrow the quantities $\sum_{i=1}^{n} e_i^{i+1}$ and $\sum_{i=1}^{n} e_i^{i+2}$ will be similar (hence also \hat{e}^{i+2} and \hat{e}^{i+2}).
- $\frac{\sum_{i=1}^{N} e^{ix}}{\sum_{i=1}^{N} e^{ix}} = \frac{\sum_{i=1}^{N} e^{ix}}{\sum_{i=1}^{N} e^{ix}}$ under the 1 expect this quantity to be done to see A the Acceptance region will be (0;k) What happens if the is not true?
- -s the predictions under Hz will be more accurate -> the total amount of error of the full model will be smaller

Preciminary result

It is possible to show that:

→ ∑ e*1 » ∑ ei² → & »&2

In this case, the full model (H2) is better than the null model (H0)

- $\frac{\sum_{i=1}^{N} e_i^{2}}{\sum_{i=1}^{N} e_i^{2}} = \frac{\sum_{i=1}^{N} e_i^{2}}{\sum_{i=1}^{N} e_i^{2}} > 0 \quad \text{under He I expect earge possibility values!}$ $\Rightarrow \text{ the RESECT REGION will be } (k_i + \infty)$

If $X \sim X_{v_1}^2$ and $W \sim X_{v_2}^2$ independent, $\frac{X/v_2}{W/v_2} \sim F_{v_2,v_2}$ Faistribution with (v_2,v_2) degrees of freedom.

 $F = \frac{SSR_{/4}}{SSF_{/(n-2)}} = \frac{\left(\frac{SSR}{6^2}\right)_{/1}}{\left(\frac{SSF}{6^2}\right)_{/n-2}} \stackrel{\text{Ho}}{\sim} F_{1, n-2}$

$$\frac{\text{SSR}}{\text{G2}} = \frac{\sum_{i=1}^{n} (\hat{\gamma}_i - \hat{Y}_i)^2}{\text{G2}} \stackrel{\text{Mr}}{\approx} \chi_1^2$$

$$\frac{\text{SSE}}{\text{G2}} = \frac{\sum_{i=1}^{n} (\gamma_i - \hat{Y}_i)^2}{\text{G2}} \stackrel{\text{Mr}}{\approx} \chi_{n-2}^2$$

$$\frac{\text{SSR}}{\text{SSR}} \perp \text{SSE}$$

Hence it holds

 $F = \frac{R^2}{4-R^2} \cdot (n-2) = \frac{SSR}{SSC} \cdot (n-2) =$

The text statistic is $\frac{R^2}{1-D^2} = \frac{ssR}{ssE}$

$$= \left(\frac{SST}{SSE} - 1\right) (N-2) =$$

 $= \left(\frac{\sum_{k=1}^{n} (Y_k - \overline{Y})^2}{\sum_{k=1}^{n} (Y_k - \widehat{Y}_k)^2} - 4 \right) (n-2) =$

To finish the Dest

 $=\frac{\frac{n^2}{2}-\frac{\Delta^2}{2}}{\frac{\Delta^2}{2}}$ (n-2) $\stackrel{\text{Ho}}{\sim}$ F_{4_1} h-2

1) FIXED SIGNIFICANCE LEVEL & a = 1P(reject Ho! Ho Drue)

FIXED SIGNIFICANCE LEVEL
$$\alpha$$
 $\alpha = |P(reject Ho | Ho Drue)$
the reject region is on the right Dail $\Rightarrow \alpha = |P_{Ho}(Fe(k_1+\infty))$
 $= |P_{Ho}(F>k)$
what is the value k that quarantees that the probability that F

will assume values larger whom k is exactly a?

values smaller than k is $1-\alpha$) K = f₁, n-2; 1-a quantile of level (1-x) of a F₄, n-2 distribution PHO (F> fin-2) 1-a) = a

(i.e., the value that guarantees that the probability that F assumes

acceptiona region A = (0, f1, n-2; 1-a) reject region R = (f1, n-2; 1-a, +00) if fobs < Fin-2; 1-4 = we do not reject to

if
$$pobS$$
; $F_{a_1n-2; a-n}$ we reject the A $F_{a_1n-2; a-n}$ R

2) P-VALUE $a_1 = a_1 + a_2 = a_2 = a_1 + a_2 = a_2 =$

- If I am testing Ho: R2=0 vs Ha: R2>0 I would reject for earge values of R2 Since F is a monotone increasing transformation, large values of R2 correspond to large value of F

Remark: To see what values lead to rejecting to, we could also do a reasoning about the values of

⇒ reject replon: (ki+∞)

(n-2). R2 directly.