

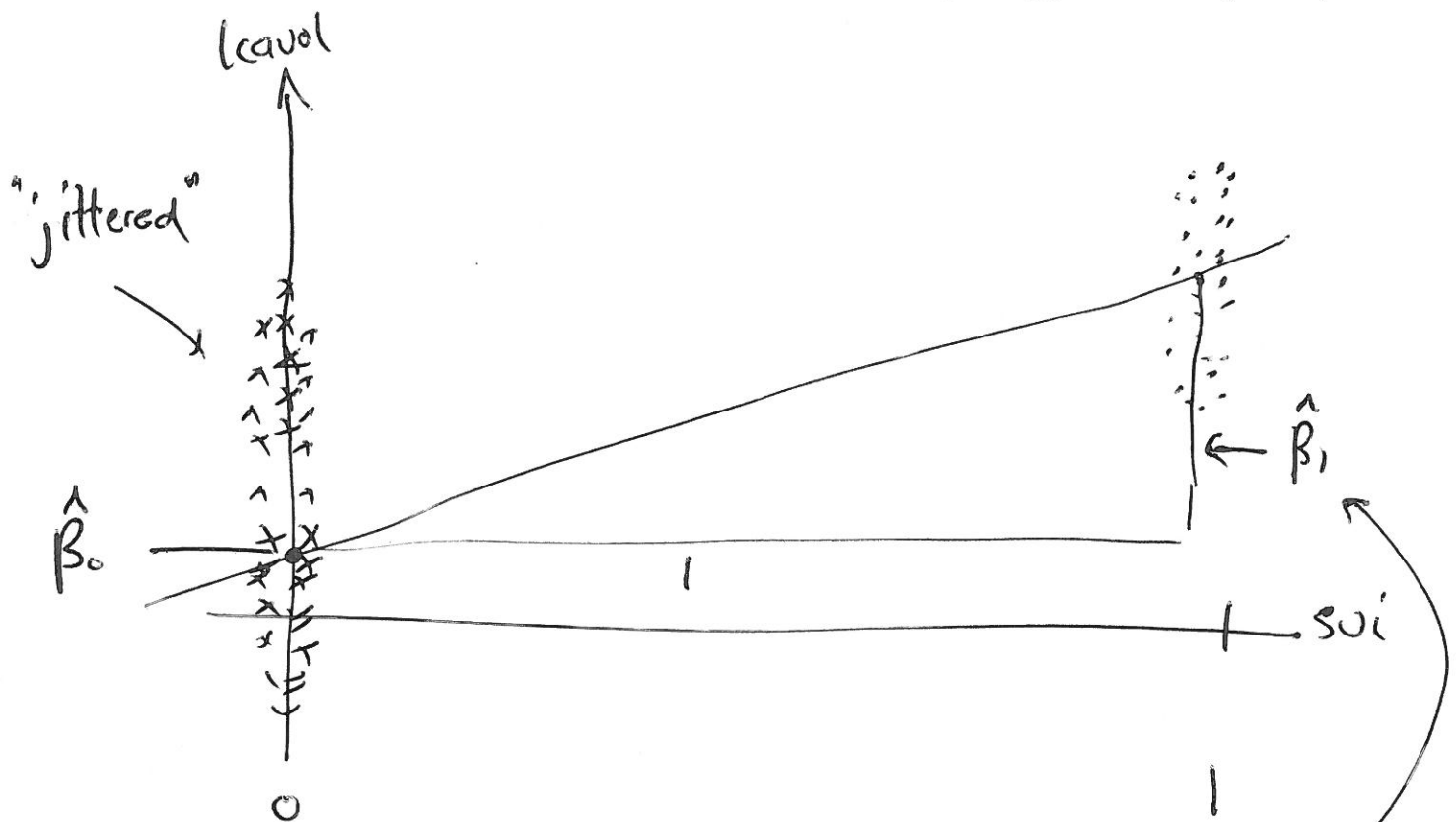
Indicator Variables

$$sui_i = \begin{cases} 1 & \text{if "seminal vesicle invasion" present for observation } i \\ 0 & \text{otherwise} \end{cases}$$

Model

$$lcauld_i = \beta_0 + \beta_1 sui_i + \varepsilon_i, \quad i=1, \dots, n$$

$$\varepsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$$



$$\hat{\beta}_1 = \text{difference in the means}$$

$$= \text{mean}(lcauld | sui=1) - \text{mean}(lcauld | sui=0)$$

Indicator variables (cont.)

New model: $Y = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \varepsilon; \varepsilon \stackrel{iid}{\sim} N(0, \sigma^2)$

where $Y = \text{laval}$

$\rightarrow X_1 = \text{svi}_i = \begin{cases} 1 & \text{if svi} = \text{Yes for obs } i \\ 0 & \text{otherwise} \end{cases}$
factor or treatment variable

$X_2 = \text{lpsa}$ (a continuous covariate)

\rightarrow an "analysis of covariance" model

when $\text{svi} = 0$, $X_1 = 0$

fitted model

$$\hat{Y} = \hat{\beta}_0 + 0 + \hat{\beta}_2 X_2 = \hat{\beta}_0 + \hat{\beta}_2 X_2$$

when $\text{svi} = 1$, $X_1 = 1$

$$\hat{Y} = \hat{\beta}_0 + \hat{\beta}_1 \cdot 1 + \hat{\beta}_2 X_2 = (\hat{\beta}_0 + \hat{\beta}_1) + \hat{\beta}_2 X_2$$

two parallel lines — one for $\text{svi} = 0$

& one for $\text{svi} = 1$