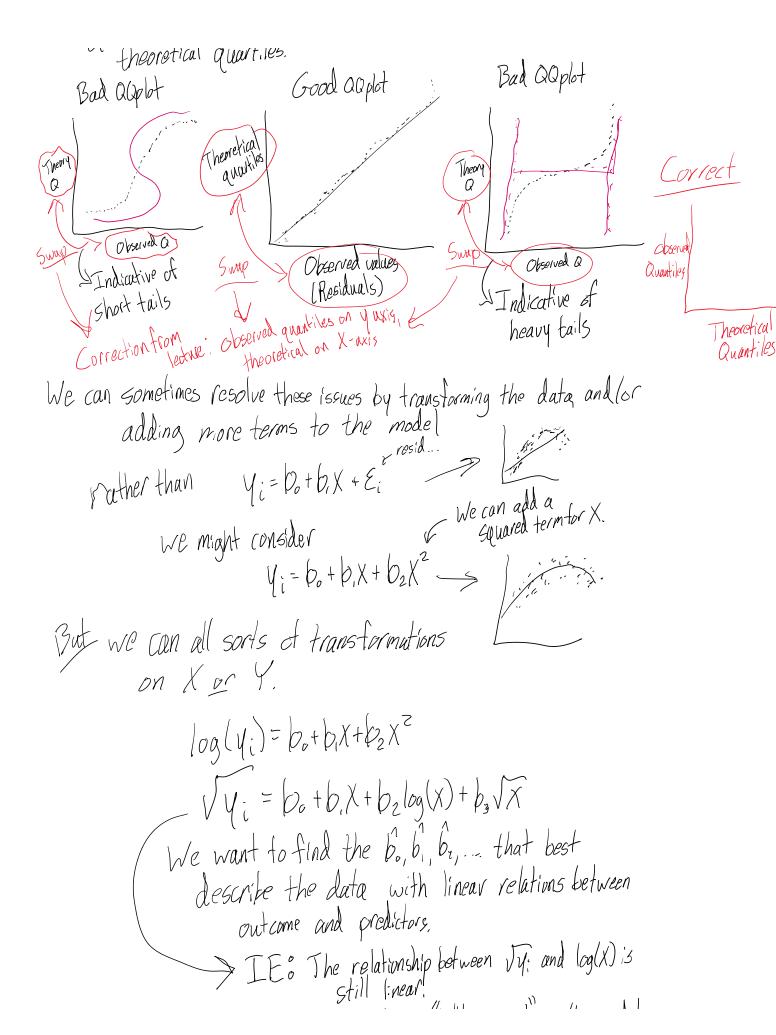
Diagnostics
When doing regression we assume follows a linear trend  When doing regression we assume follows a linear trend  and that residuals have no pattern and are normally  Alstributed. > (Yi - Yi) = resid;  residuals
and that residuals nave in pain
1/37/ Duted. (91 )- 1830, residual
One of the best ways to confirm  assumptions is to plot residuals against fitted value.  What if we have data that looks like non-homoscodastic heteroscodastic Sabasii
What if we have data that looks like non-homoscodastic
(non-linear) heteroscedastic
what if we have data that looks the non-homoscodastic neteroscodastic Sedasticity refers to randomness
Our residuals will look like this This pear shape is an indicator
Mon-constant
resided Marcator of resided Variance.
We want no pattern Japs are chay but maybe indicate
maybe indicate  different groups
Organising namality of residuals can be done with a simple histogram
Diagnosing namality of residuals can be done with a simple histogram or more methodically with a quartile-quartile plot  (qq plot)
A applot will find the quartiles of the data and plot them against theoretical quartiles.  But Opplot
Bad applot Good applot : Bad applot
Bad alphot (rood applet



Still [: near]
Still !: neal".
One transformation that is common but has little impact on the model
Tyle rounded them.
We can take X and Y and "normalize" or "standardize" them.
$Z_{y_i} = \underbrace{x_i - x}_{\text{mean 0, sd 1}}$ $Z_{x_i} = \underbrace{x_i - x}_{\text{mean 0, sd 1}}$
7 - Lit
Lyi Sy
mean 0, sd 1.
yneun U, su I
Benefit o Slope interpretation becomes: "For a 1 standard deviation increase
Genetit : Slope interpretation becomes: For a I standard deviation increase in Y
INCREASE IN Y
Benefit o 60 = 0
Standard Fe