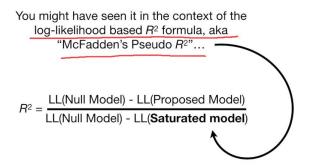
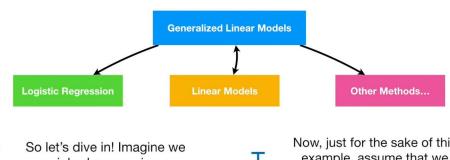
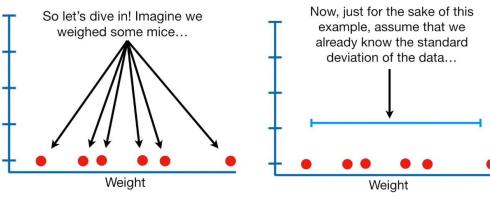
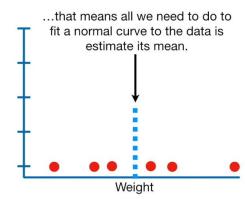
Saturated Models and Deviance Statistics

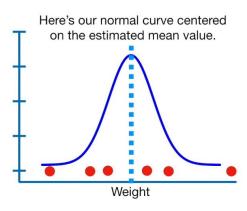


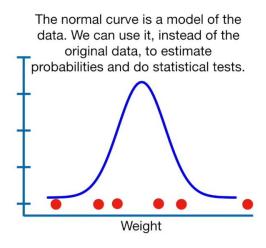
NOTE: The Saturated Model and Deviance Statistics are part of Generalized Linear Models and are intended to be used in a very wide variety of situations.

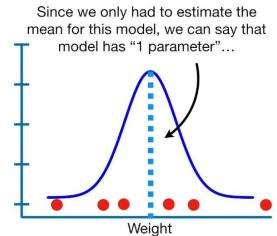


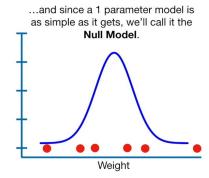


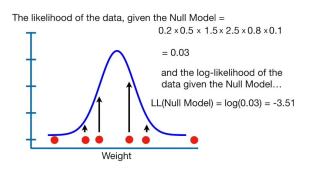


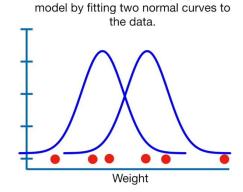




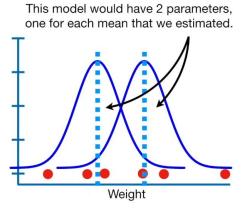


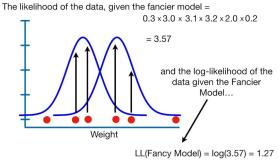


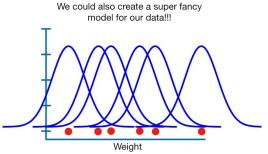


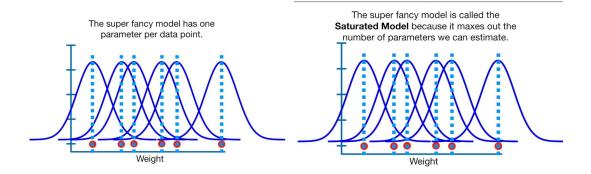


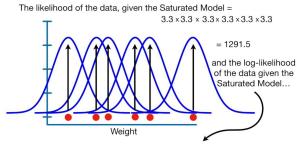
We could also create a slightly fancier





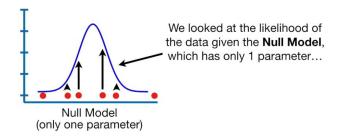




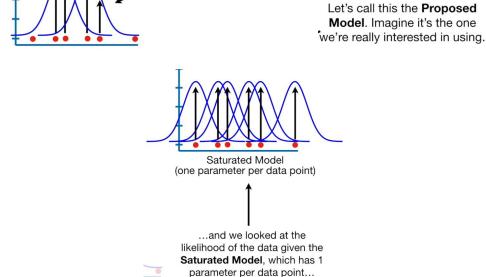


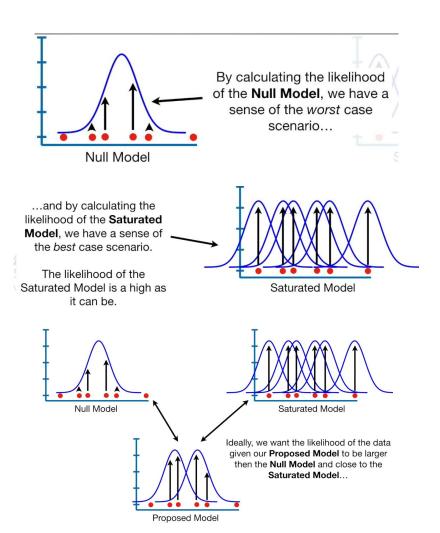
LL(Saturated Model) = log(1291.5) = 7.16

So far we've looked at the likelihood of the data using 3 different models.



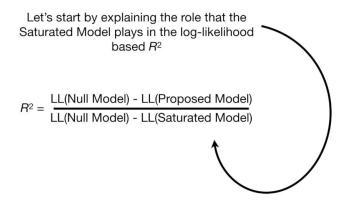
...we looked at the likelihood of the data given a model with 2 parameters.

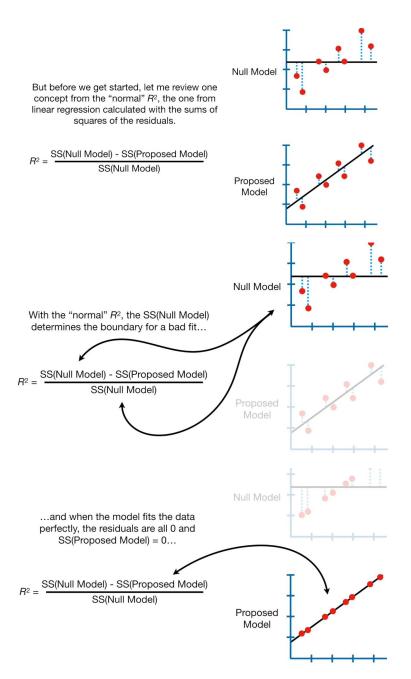




When using likelihoods, we use the **Null** and **Saturated models** to determine whether the **Proposed Model** does a good job with the data.

In other words, we use the **Null and Saturated Models** to calculate R^2 and its p-value for the **Proposed Model**.





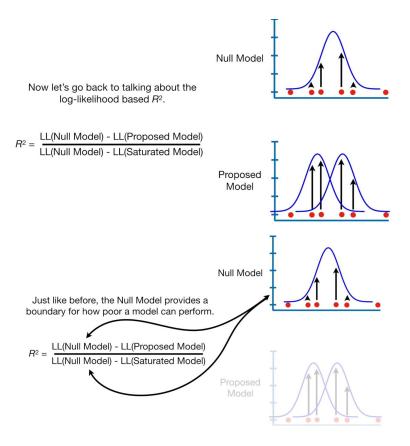
...in other words, there is a fixed value, 0, that represents a boundary for the best a model can do...

...we get
$$R^2 = 1$$

$$R^2 = \frac{\text{SS(Null Model)}}{\text{SS(Null Model)}} = \frac{1}{2}$$

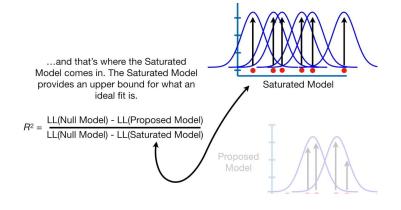
To summarize, the "normal" R^2 has a boundary for how poorly a model can perform, SS(Null Model), and a fixed value, 0, that represents the best possible model.

$$R^2 = \frac{SS(Null Model) - SS(Proposed Model)}{SS(Null Model)}$$



However, unlike the traditional R^2 , there is no fixed value for an "ideal fit" that works in every situation...

$$R^2 = \frac{\text{LL(Null Model)} - \text{LL(Proposed Model)}}{\text{LL(Null Model)} - \text{LL(Saturated Model)}}$$



Thus, if the Proposed Model is just as good as the Saturated Model, then LL(Proposed Model) = LL(Saturated Model)

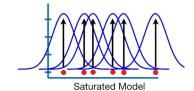
...and that makes the numerator the same as the denominator and $R^2 = 1$.

$$R^2 = \frac{\text{LL(Null Model)} - \text{LL(Proposed Model)}}{\text{LL(Null Model)} - \text{LL(Saturated Model)}} = \frac{1}{100}$$

In summary, the log-likelihood of the Saturated Model ensures that the R^2 value ranges between 0 and 1.

Now let's talk about **Deviance**, a concept related to the Saturated Model that will, ultimately, lead us to a *p*-value for the *R*².

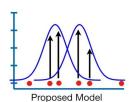
We're going to talk about two types of Deviance Residual Deviance and Null Deviance.

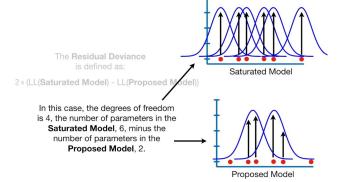


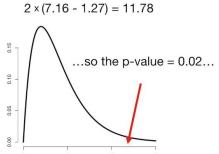
Residual Deviance is defined as:

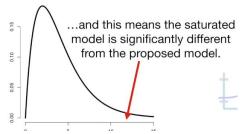
2 × (LL(Saturated Model) - LL(Proposed Model))

The "2 times" part makes the difference in these log-likelihoods have a Chi-squared distribution with degrees freedom equal to the difference in the number of parameters.



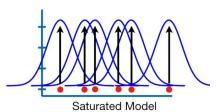






NOTE: For this Chi-square test to work correctly, the Proposed Model and Saturated Model have to be "nested".

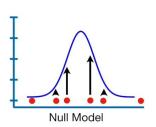
In other words, the Proposed Model has to be a simpler version of the Saturated Model (and not just any old model).



The **Null Deviance** is defined as:

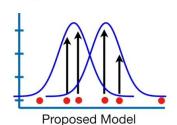
2 × (LL(Saturated Model) - LL(Null Model))

Again, the "2 times" part makes the difference in these loglikelihoods have a Chi-squared distribution with degrees freedom equal to the difference in the number of parameters.



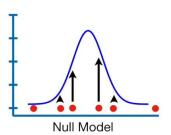
In this case, the degrees of freedom is 5, the number of parameters in the **Saturated Model**, 6, minus the number of parameters in the **Null Model**, 1.—

...you can calculate the *p*-value for the log-likelihood *R*² from the difference in the deviances...



Null Deviance - Residual Deviance

...this difference is a Chi-square value with degrees of freedom equal to the difference in the number of parameters for the Proposed Model and the Null Model.



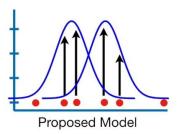
 $R^2 = \frac{\text{LL(Null Model)} - \text{LL(Proposed Model)}}{\text{LL(Null Model)} - \text{LL(Saturated Model)}}$



Thus, the *p*-value for the log-likelihood *R*² value we calculated before is 0.002.

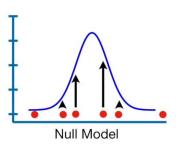
This means that not only does our model have a reasonable effect size (the R^2 value), we know that that value is not by chance.

Alternatively, we can calculate the significance of the R^2 directly by comparing the Proposed Model to the Null Model, since the Null Model is nested within the Proposed Model.



2 × (LL(Proposed Model) - LL(Null Model))

Again, the "2 times" part makes the difference in these log-likelihoods have a Chi-squared distribution with degrees freedom equal to the difference in the number of parameters.

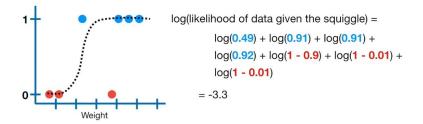


However, I mentioned the method that uses Deviances because the Null and Residual Deviances are very frequently reported. In contrast, the LL(Proposed Model) and the LL(Null Model) are not always reported.

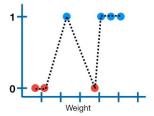
OK, we've talked about the Saturated Model, Residual Deviance and Null Deviance.

Now let's talk about why you can ignore the Saturated Model entirely when you do Logistic Regression.

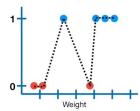
When you do logistic regression, you're fitting a squiggle to your data and calculating the log-likelihood of the data given that squiggle.



However, with the saturated model, the squiggle fits the data perfectly...



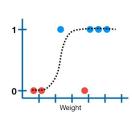
...and the log-likelihood of the data is 0, because log(1) = 0, so the log-likelihood is just a big sum of zeros.



log(likelihood of data given the saturated squiggle) =

$$log(1) + log(1) + log(1) + log(1) + log(1 - 0) + log(1 - 0) + log(1 - 0)$$

This means that the equation for the likelihood based R^2 is this...



 $R^2 = \frac{\text{LL(Null Model)} - \text{LL(Proposed Model)}}{\text{LL(Null Model)} - 0}$