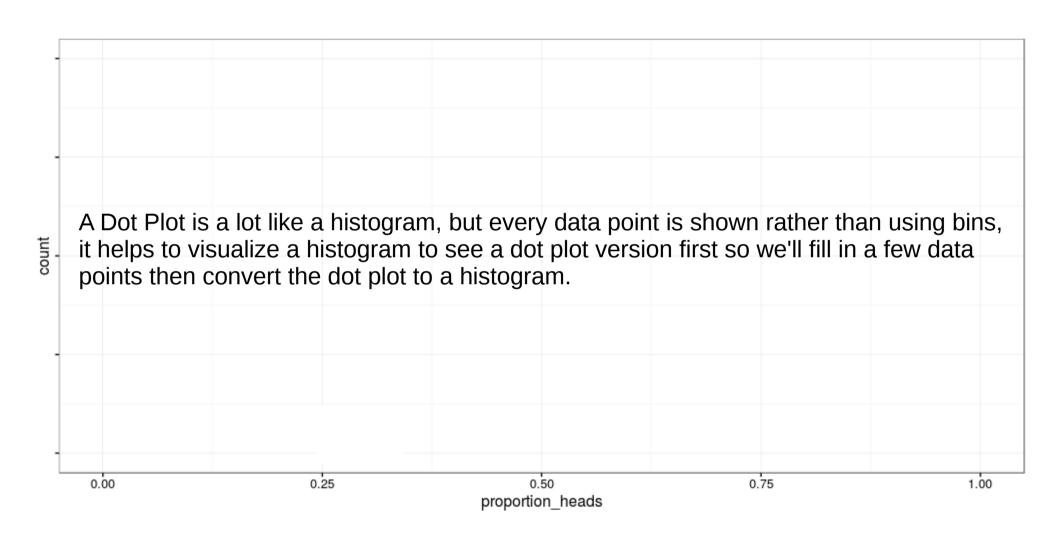
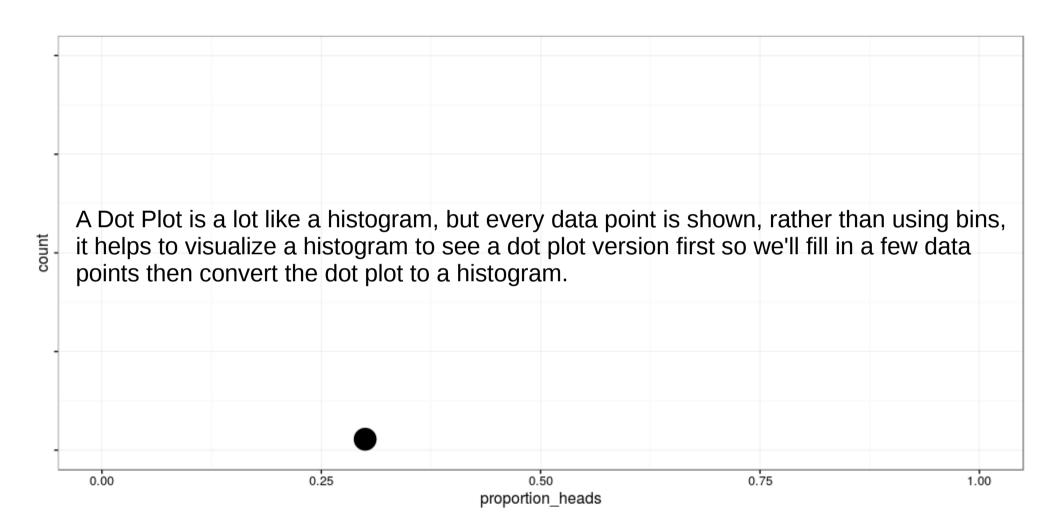
Histograms – show you the distribution of your data. They have a binned numerical variable on the X axis and count on the Y axis.

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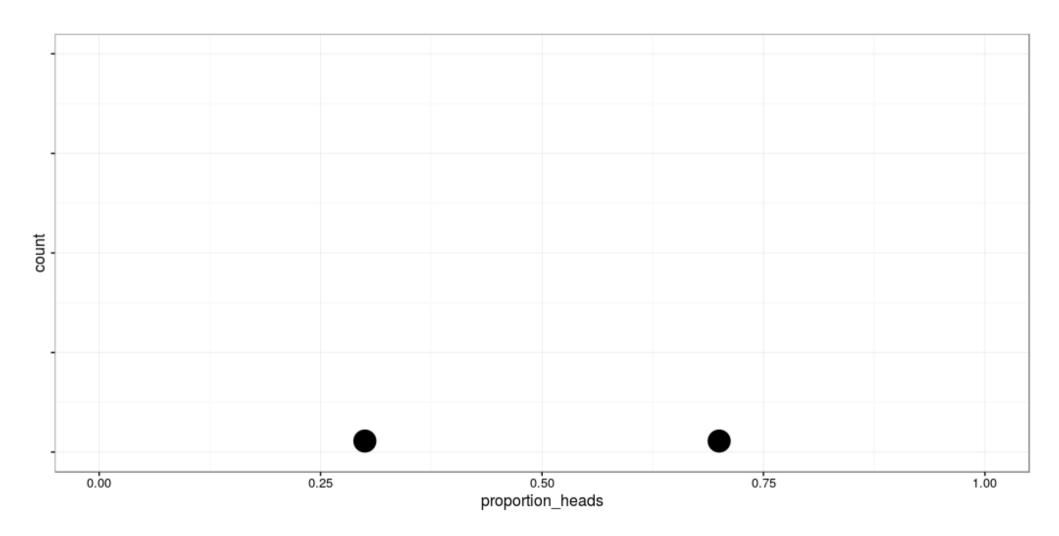


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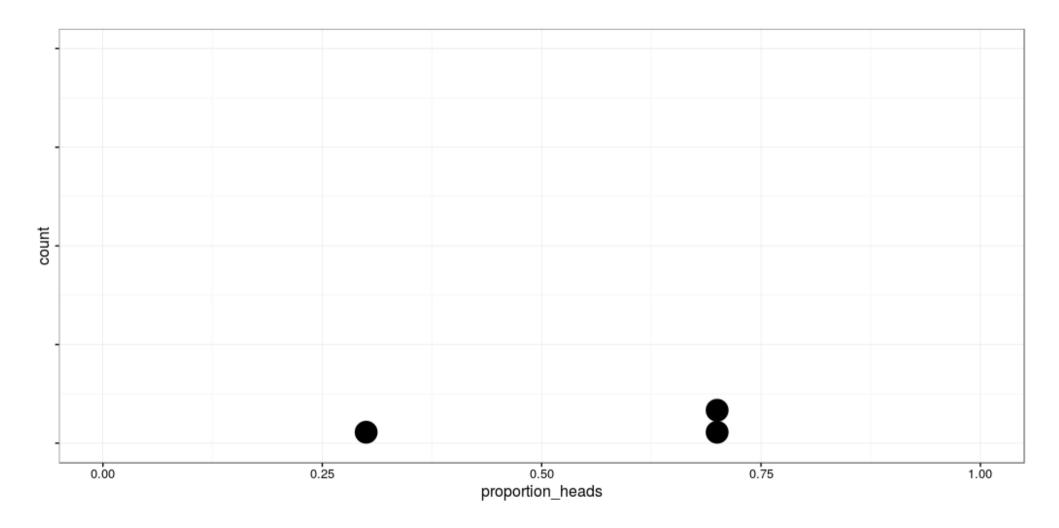
Plot our first data point.



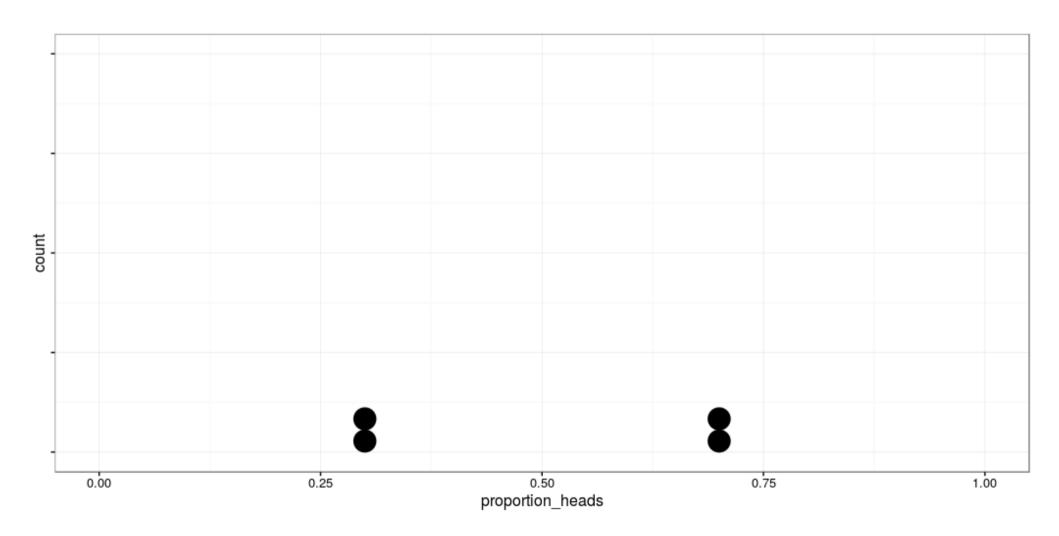
Second data point.



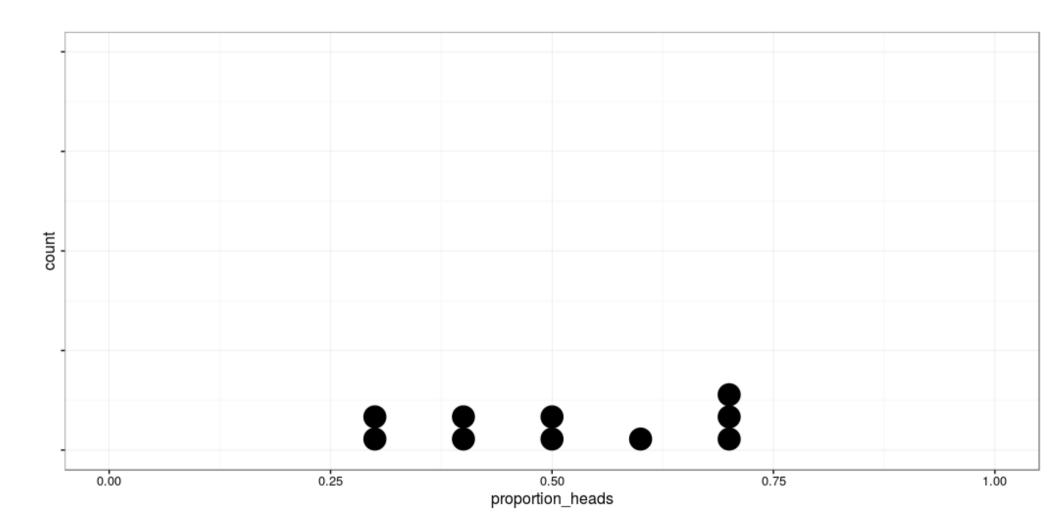
Third data point.



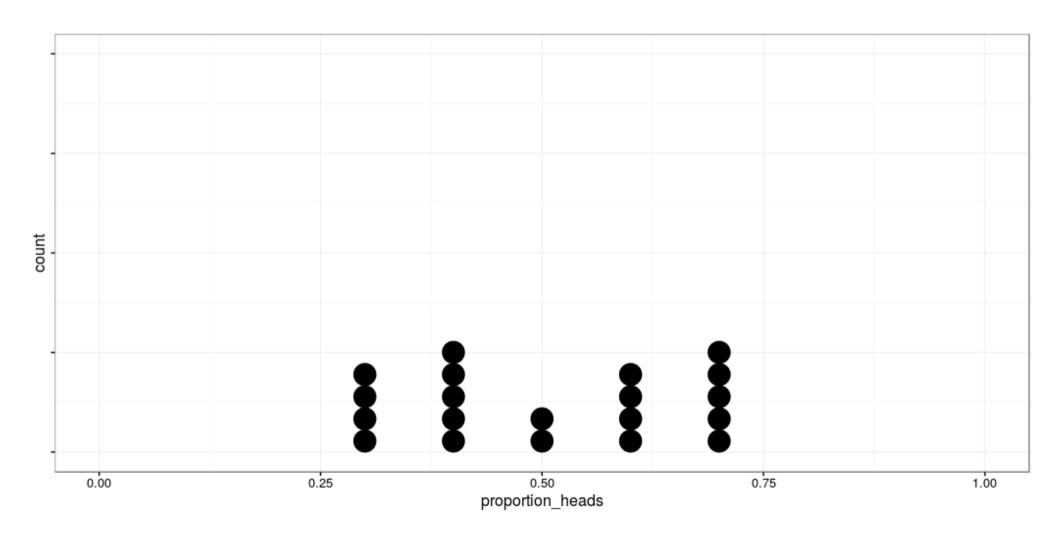
Fourth data point.



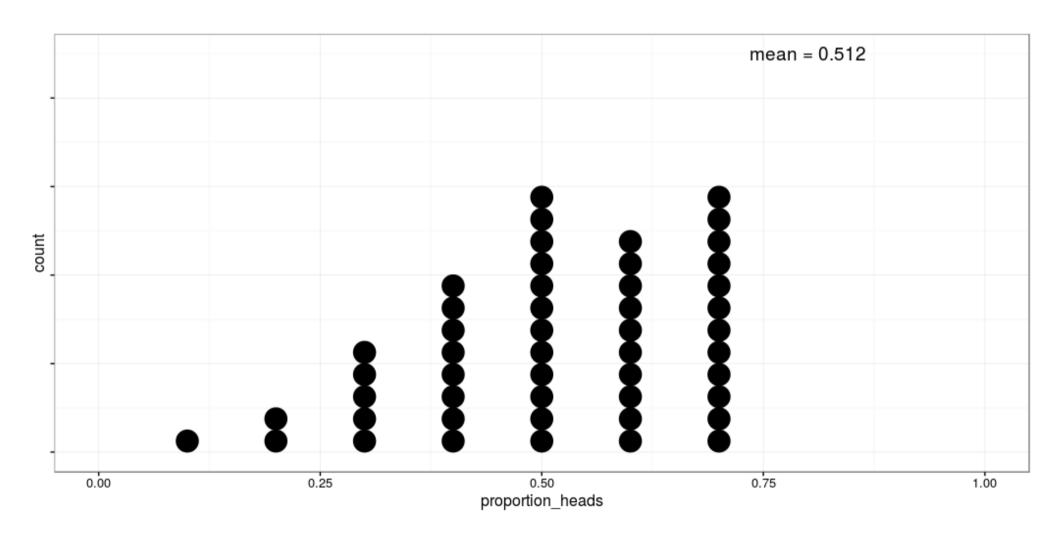
Ten data points.



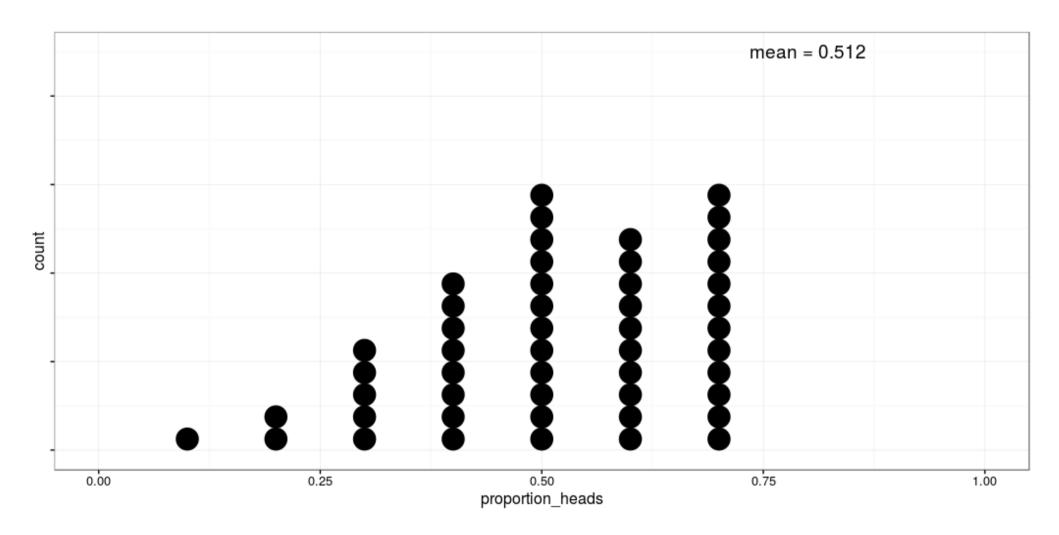
Twenty data points.



All fifty data points on a dot plot.

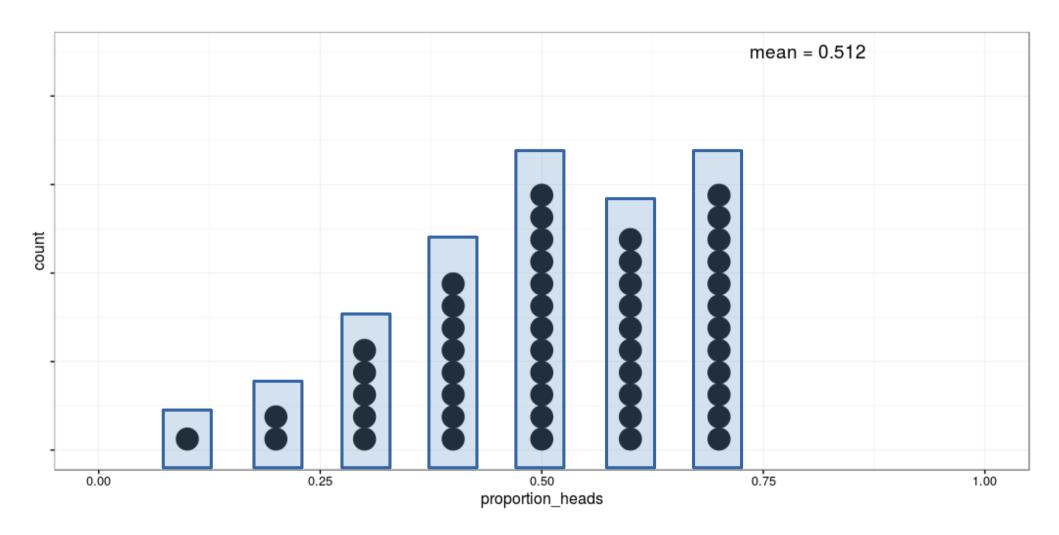


All fifty data points on a dot plot.



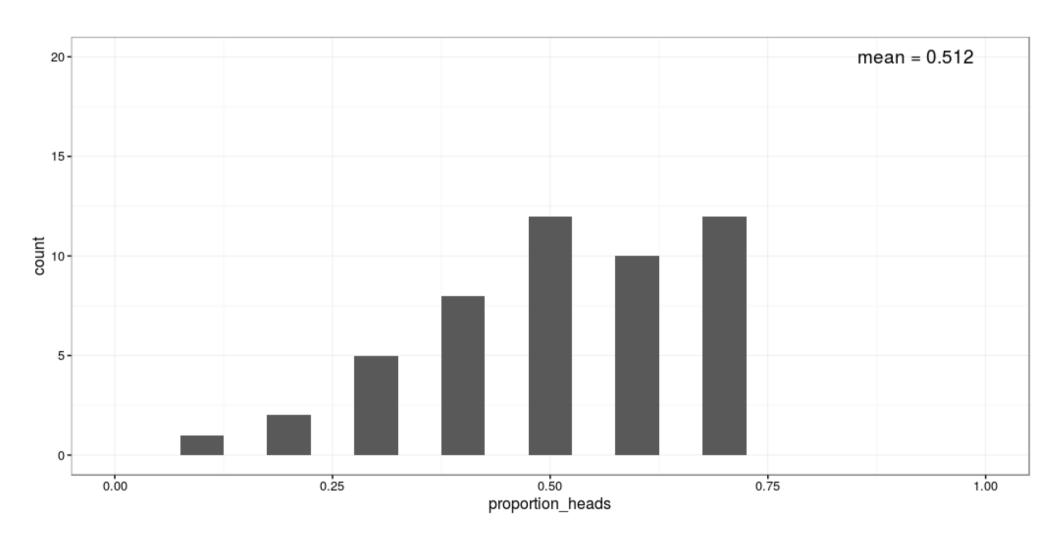
Now let's convert this to a histogram by binning the data. The nice thing is, that since this data was generated using 10 coin flips for each data point, there are a finite number of possibilities and the bins are pretty obvious.

All fifty data points on a dot plot.

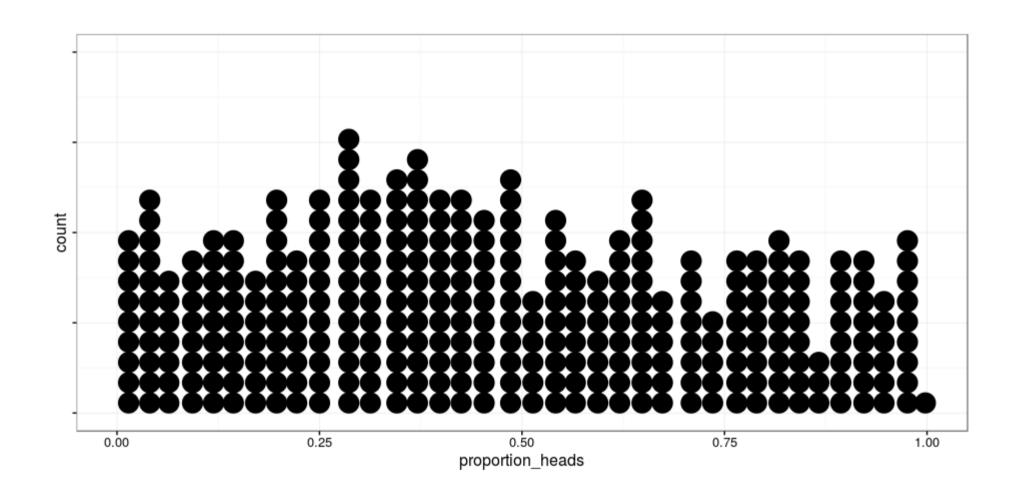


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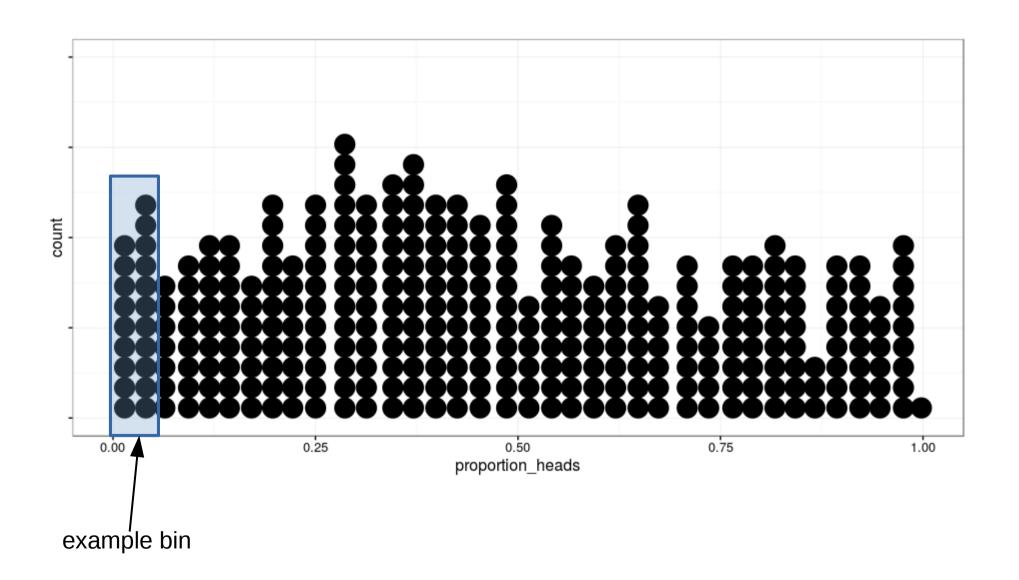
Final Histogram of the same data that is in the dot plot of 50 data points.



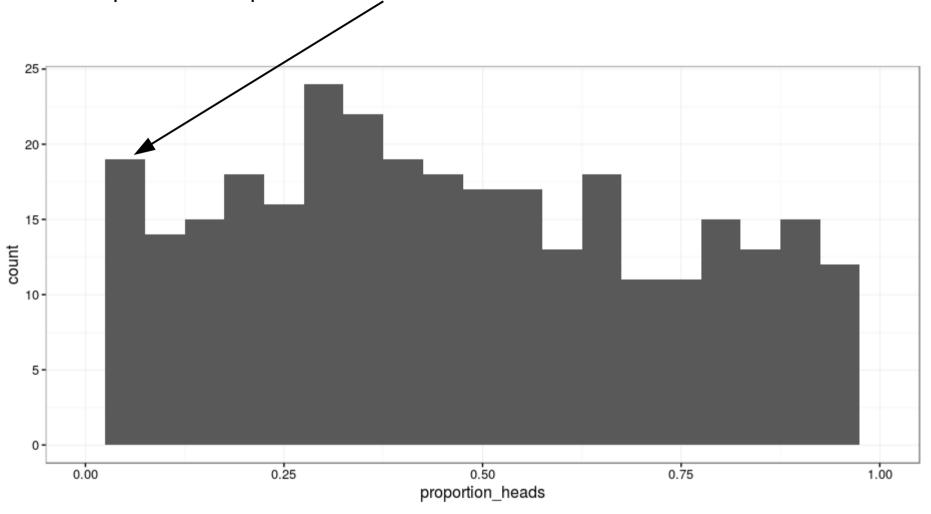
Dot plots start to get really busy, and binning to make a histogram is really better.



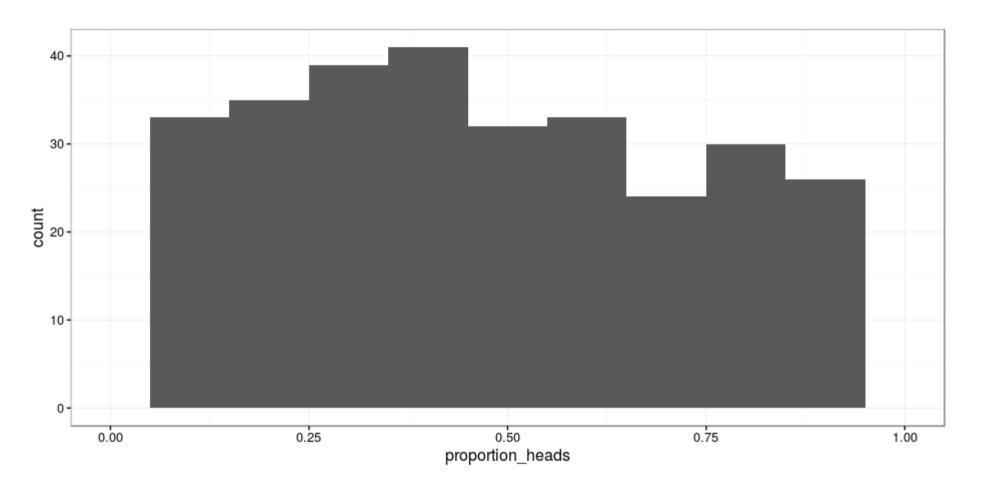
Dot plots start to get really busy, and binning to make a histogram is really better.



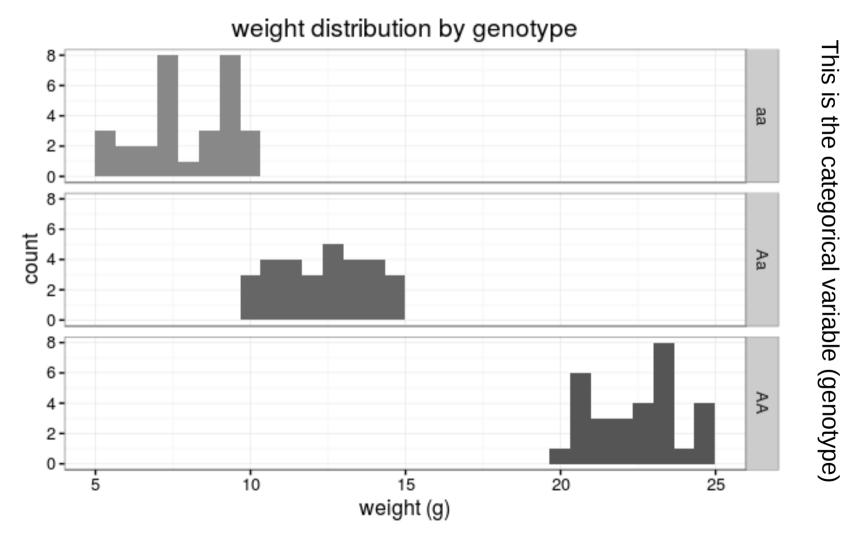
See? This histogram has a bin width of 0.05 and combines about 2 columns of dots from the previous dot plot into a bin.



This histogram has a bin width of 0.1 and combines more columns of dots from the previous dot plot into a bin. Choosing a bin width isn't always easy, but you want to accurately represent the distribution of your data. (these are a bit too big for me)



Stacked histogram – it's like 3 histograms for the price of one! And lets you compare the distributions of a numerical variable (weight) among the categories (genotype) All 3 of the categories share the same X axis, and the left hand Y axis (count) repeats on the same scale for each category



This is the numerical variable (weight)

Box Plot – if you want to provide a little more summary information about the numerical values for each category, a side-by-side box plot is nice. It also lets you compare the distributions and medians of a numerical variable (weight) among the categories (genotype). All 3 of the categories share the same Y axis (weight) with each category having its own box.

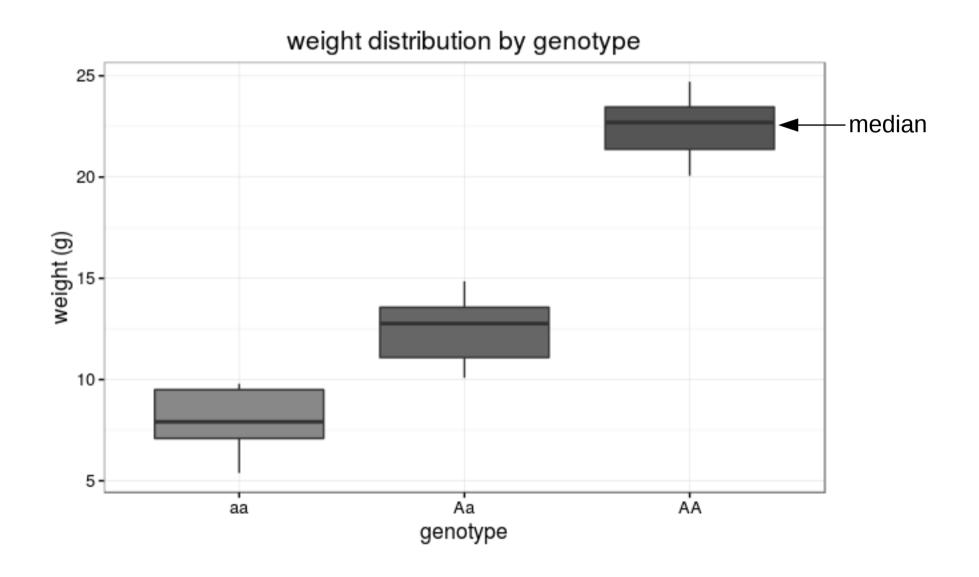


Fig 4.12 Description of a Boxplot

