Python: Data Visualization Notes

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1 Learning Objectives

What good is a data analysis if the answers or findings can't be conveyed properly? Or perhaps even more detrimental, what good is a dataset if the proper questions or hypotheses can't be formed in the first place? How can we properly tell the story the data is providing? Enter in data visualization! Both explaining the data and exploring the data can be significantly helped through the process of data visualization.

Data visualization is two-pronged:

- Exploratory Analysis: Helping to understand the data prior to analysis. Searching for relationships and insights.
- Explanatory Analysis: Helping to present analysis findings, helping to tell a story with the data. After insights were found.

And it's all a part of the entire data analysis process. We can also simplify the data analysis process into 5 steps:

- 1. Extract
- 2. Clean: Exploratory
- 3. Explore: Exploratory
- 4. Analyze: Exploratory OR Explanatory
- 5. Share: Explanatory

In this course, we'll be using the matplotlib, seaborn, and Pandas libraries to assist in the data analysis process.

Concepts

- Design of Visualization
- Exploration of Data

- Univariate Exploration of Data
- Bivariate Exploration of Data
- Multivariate Exploration of Data
- Explanatory Visualizations
- Visualization Case Study

2 Design of Visualization

To begin our discussion of visualization, we need to cover some basic vocabulary and distinctions.

Data can be broken into two main categories, each of which can be broken down further:

- Qualitative / Categorical:
 - Nominal: No order
 - Ordinal: Intrinsic Order
- Quantitative / Numerical:
 - Interval: Absolute differences are meaningful (addition and subtraction follows logic)
 - Ratio: relative differences are meaningful (multiplication and division follows logic)

It should be noted that the quantitative data type can be also be broken down into discrete and continuous variables.

What about those 3-dimensions charts or fun backgrounds that we used to add to our science experiment plots as kids? That used to add some fun to our projects, right? While fun for the youth, it turns out there is an empirical rule when figuring out how much the additional "junk" either adds or detracts from conveying the data.

• Data-Ink Ratio = data-ink / total ink used to print the graphic. The higher the Data-Ink Ratio, the better conveyed data is.

Can visualizations be purposefully misleading, even when using the data appropriately? Absolutely!

A great example of this is trying to over-inflate the difference or change between data points during different time periods. Say a presenter is trying to make a claim there was a very large change from one year to the next. We'll say in year

1 the y-value was 100, and in year 2 the y-value was 105. The presenter changes the window of the graph to display from a y-value of 99 to a y-value of 106, and the x-values are only the two years. Obviously, this is going to look like a massive change! In reality, had the reporter shown the data at a true scale, the visual shows in actuality that the change isn't so tremendous.

This concept also has an empirical rule:

- Lie Factor = size of effect shown in graphic / size of effect shown in data
 - = (change in visual / visual start) / (change in data / data start)

As was said in the example, this can be used to purposefully distort data. In fact, a Lie Factor 1 suggests a misleading visual, and even greater than that suggest an even greater disparity from the truth.

Away from the empirical side of visuals, and more into the logical, we come across the common mistake of using too many colors! Colors can be useful when separating categories, however, they it's very easy to cause redundancy with them. Here are some tips when using color:

- Get it right in black and white (and shades of grey)
- Use less intense colors such as natural or pastel, and higher grey colors. The eye can actually concentrate longer under these conditions.
- Color facilitates communication. Use color to separate the data into groups of interest, not just to color a visual.
- Design for Color Blindness. Stay away from red / green pallets, and use blue / orange pallets.

Don't want to overdo it on the color schemes? Don't forget about other visual queues such as shape and size.

Some tips on shape, size & other tools:

- Use different types of encodings, rather than using color (square / dot vs. colors to separate groups of interest).
- Color and shape are good for categorical variables.
- $\bullet\,$ Size of marker can assist in adding additional quantitative data.

3 Exploration of Data

We have a dataset on a topic or concept that has been deemed worthy for inspection! Surely, there are some insights to be gained from it. We load up the data, and then we hit a wall... Which columns are important? What questions can be answered? We can find the general statistics of the set, so what?

This section will help with the initial process of data exploration, helping to find what is actually useful and should be examined further in the data. We'll start with single variables from the data and move into visually pairing multiple data points at once.

We'll be using a few different Python libraries in this section:

```
import numpy as np
import pandas as pd
import matplotlib.pyplot as plt
import seaborn as sb
```

3.1 Univariate Exploration of Data

The first tools added into our data exploration toolkit will be for univariate data, or examining the data a single variable at a time. Even though we're going to be using several different libraries, we'll still try to keep track of new and essential commands and terminology:

Commands:

- df.head: Retrieves the first few rows of data from a Pandas DataFrame. Able to specify how many rows
- sb.countplot: Seaborn's command for generating a bar chart.
- df[column].value_counts().index: Will return an immutable sequence sorted number of categorical entries (highest to lowest).
- df.melt: Will unpivot a DataFrame from wide to long format (i.e. we can "melt" two entries together).
- plt.hist: Matplotlib's command for generating a histogram.
- sb.distplot: Seaborn's command for generating a histogram. Default command also includes a density curve estimation.

Terminology:

• bar chart: Useful to show counts across categories.

- relative frequency: Shows proportion of each category in population.
- pie chart: Use these to show how whole of data is broken down into parts, useful when plotting a small number of slices (usually no more than four).
- histogram: Useful to examine quantitative data.

We'll be using a Pokemon dataset for our examples. Let's get an idea of what our data looks like:

```
pokemon = pd.read_csv("pokemon.csv")
pokemon.shape
# pokemon.shape = (807, 14)
pokemon.head(10)
```

Pokemon.Head(10)

	id	id species		ge	generation_id		nt weight	base_experience		type_1
0	1	bu.	lbasaı	_	1		7 6.9	_	64	grass
1	2	ivy	saur			1 1.	0 13.0		142	grass
2	3	ver	nusau	r		1 2.	0 100.0		236	grass
3	4	cha	arman	der		1 0.	6 8.5		62	fire
4	5	cha	armele	eon		1 1.	1 19.0		142	fire
5	6	cha	arizar	d		1 1.	7 90.5		240	fire
6	7	squirtle				1 0.	5 9.0		63	water
7	8	wa	rtortle	e		1 1.	0 22.5		142	water
8	9	bla	stoise	;		1 1.	6 85.5		239	water
9	10	cat	erpie			1 0.	3 2.9		39	bug
	typ	e_2	hp	attack	defense	$_{\mathrm{speed}}$	special-at	-	al-defen	
0	pois	on	45	49	49	45		65		65
1	pois	on	60	62	63	60		80		80
2	poison		80	82	83	80		100		00
3	NaN		39	52	43	65	60			50
4	NaN		58	64	58	80		80		65
5	flying		78	84	78	100		109	85	
6	NaN		44	48	65	43		50	64	
7	NaN		59	63	80	58		65	80	
8	NaN	J	79	83	100	78		85	1	05
9	NaN	1	45	30	35	45		20		20

Now that we've had a peak at the data, let's get into some visualization.

A simple, yet effective visual trick is adding some order to your plots via sorting.

Let's combine the previous coloring and sorting to look at a new variable. Additionally, since we know the x-axis labels are somewhat long. Two favorable ways

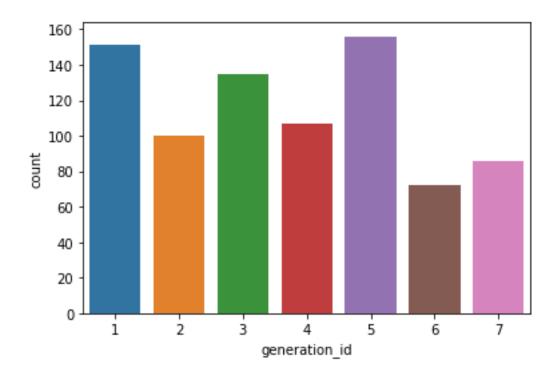


Figure 1: Generic bar chart.

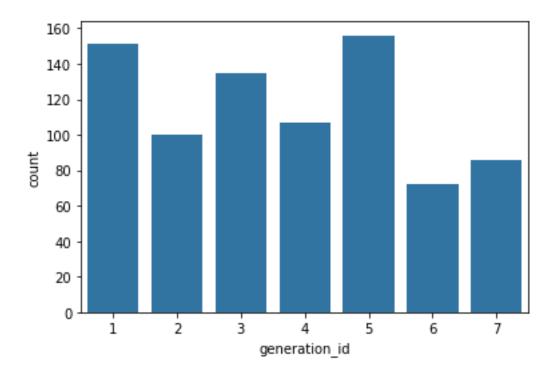


Figure 2: Bar chart with neutral coloring.

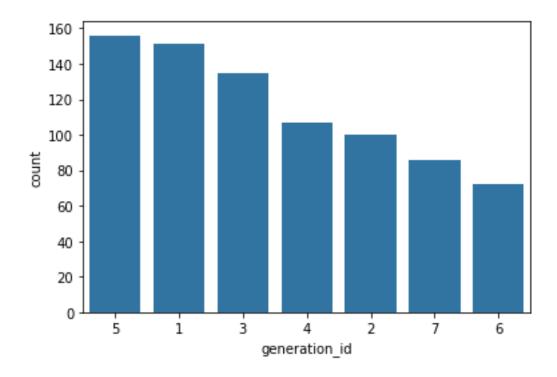


Figure 3: Bar chart with neutral coloring and ordering applied. $\,$

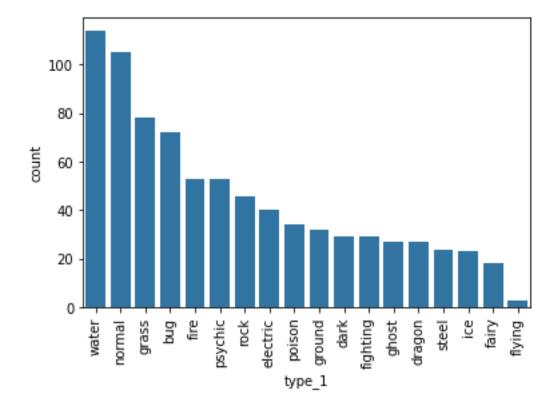


Figure 4: Standard bar chart format with x-axis labels rotated.

of dealing with this is to either rotate the x-axis labels, or create a horizontal bar chart.

The previous examples display actual counts in the bar charts, but it can also be useful to display data using relative frequency (proportion of each category in the population). This gives us an opportunity to introduce a few other concepts as well. Most importantly, we'll introduce the melt function, which we'll use to combine two categories (two distinct columns) into a single category.

```
pkmn_types = pokemon.melt(id_vars=['id', 'species'],
```

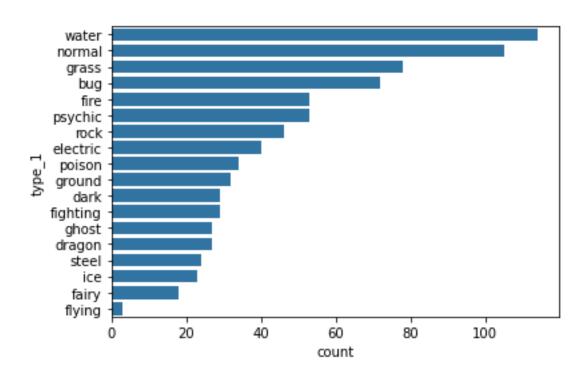


Figure 5: Standard bar chart format in vertical form.

```
value_vars=['type_1', 'type_2'],
       var_name='type_level',
       value_name='type')
Let's walk through how the melt function is being used here:
- id_vars: columns to utilize as identifier variables
- value_vars: columns to unpivot
- var_name: name of column to describe data being unpivoted
                                together
- value_name: name of column to match values from the unpivoted
                                labels
Note: the pokemon DataFrame has 807 rows (pokemon entries). This
                                use of melt puts two categories
                                together into a single category.
                                Thus, each pokemon now has two
                                entries each, making the
                                pkmn_types DataFrame have double
                                the rows as the original
                                DataFrame or 1614 rows.
....
base_color = sb.color_palette()[0]
type_counts = pkmn_types['type'].value_counts()
type_order = type_counts.index
sb.countplot(data = pkmn_types, y = 'type', color = base_color,
                                order = type_order)
# See Figure 6
# now we'll change the tick counts
n_pokemon = pokemon.shape[0]
max_type_count = type_counts[0]
max_prop = max_type_count / n_pokemon
\# note: max\_prop = 0.16
tick_props = np.arange(0, max_prop, 0.02)
tick_names = ['{:0.2f}'.format(v) for v in tick_props]
tick_props creates the numbers for an equally spaced axis, while
                                tick_names takes the numeric
                                version and returns a list of a
                                string values.
n n n
# The culmination of the above work into a chart:
sb.countplot(data = pkmn_types, y = 'type', color = base_color,
                                order = type_order)
plt.xticks(tick_props * n_pokemon, tick_names)
plt.xlabel('proportion')
for j in range(type_counts.shape[0]):
count = type_counts[j]
pct_string = '{:0.1f}%'.format(100*count/n_pokemon)
plt.text(count + j, j, pct_string, va = 'center')
# See Figure 7
```

Whereas bar charts are a great initial step in exploring categorical data, histogram are useful in the exploration of quantitative data. Let's take a look at

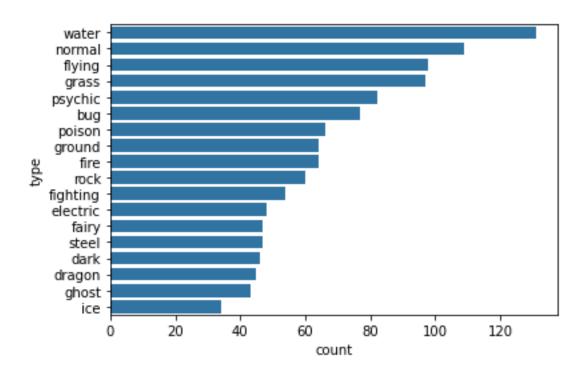


Figure 6: Bar chart of melted pokemon types.

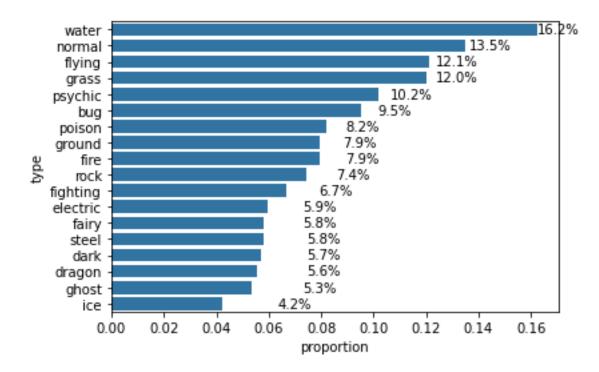


Figure 7: Bar chart of melted pokemon types shown in relative frequency.

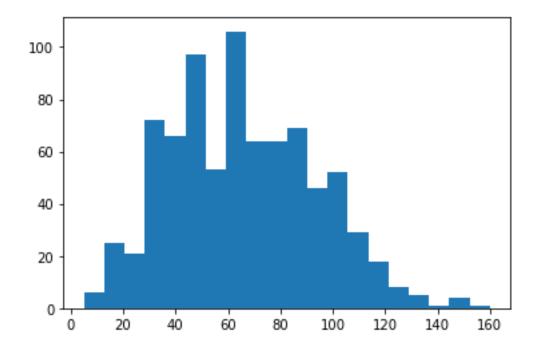


Figure 8: Basic histogram of Pokemon speeds.

our variables in the pokemon DataFrame which contain quantitative data.

Earlier in the course, we mentioned avoiding a high "lie factor". One case in which it may actually be beneficial to "zoom" in on data is when dealing with outliers, or when a large majority of the data is within certain axis limits.

For example, let's take a look at the height variable of Pokemon.

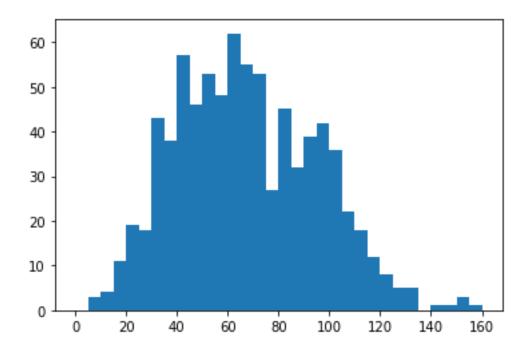


Figure 9: Histogram of Pokemon speeds with altered bins.

```
# even though this is a course on visuals, basic math and
                                 descriptive statistics can still
                                 be beneficial
pokemon['height'].describe()
         807.000000
count
           1.162454
mean
           1.081030
std
{\tt min}
           0.100000
           0.600000
25%
50%
           1.000000
           1.500000
75%
          14.500000
max
With the majority of our data between 0 and 1.5, it's acceptible
                                 to change our x-axis limits. The
                                 outlier of 14.5 would make our
                                 histogram and bin choice less
                                 effective from a visual sense.
bins = np.arange(0, pokemon['height'].max() + 0.2, 0.2)
plt.hist(data = pokemon, x = 'height', bins = bins)
plt.xlim((0, 6))
# See Figure 11
```

Another method when dealing with outliers or data points that have very large

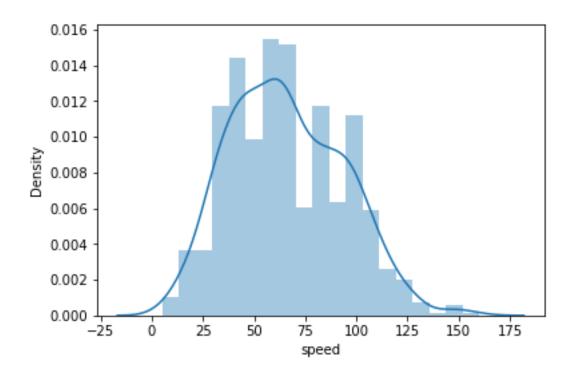


Figure 10: Seaborn's default histogram.

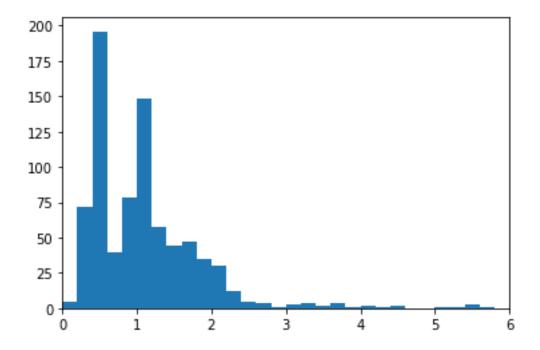


Figure 11: Histogram with altered x-axis limits.

differences between them is with scaling and transformations.

Let's take a look at the weight variable of Pokemon.

```
pokemon['weight'].describe()
count
         807.000000
mean
          61.771128
         111.519355
std
           0.100000
min
           9.000000
25%
50%
          27.000000
          63.000000
75%
         999.900000
max
Obviously, there exists some outlier(s) here. We should think
                                about using transformations and
                                tick-mark manipulation to
                                visualize the data this time
# Let's see what we're dealing with without any scaling or
                                transformations
bins = np.arange(0, pokemon['weight'].max() + 40, 40)
plt.hist(data = pokemon, x = 'weight', bins = bins)
```

```
# See Figure 12
Even with specifying the bins, the histrogram proves to not be
                                too insightful. Changing the bins
                                 alone likey won't solve this
# Let's try applying a transformation of the x-scale, itself
bins = np.arange(0, pokemon['weight'].max() + 40, 40)
plt.hist(data = pokemon, x = 'weight', bins = bins)
plt.xscale('log')
# See Figure 13
It could be argued this is a worse approach, but using a
                                logarithmic transformation could
                                steer us in the right direction.
np.log10(pokemon['weight'].describe())
        2.906874
count
mean
        1.790786
        2.047350
std
min
        -1.000000
25%
        0.954243
50%
        1.431364
75%
        1.799341
         2.999957
max
Applying a logarithmic transformation on the data, itself,
                                appears to shrinks the numbers
                                into a more manageable scale.
0.00
# Get the ticks for bins between [0 - maximum weight]
bins = 10 ** np.arange(-1, 3 + 0.1, 0.1)
# Generate the x-ticks we want to apply
ticks = [0.1, 0.3, 1, 3, 10, 30, 100, 300, 1000]
Important: note here how we are using differently spaced tick
                                marks in the original scale.
\# Convert ticks into string values, to be displayed along the x-
labels = ['{}'.format(v) for v in ticks]
# Plot the histogram
plt.hist(data=pokemon, x='weight', bins=bins);
# The argument in the xscale() represents the axis scale type to
                                apply.
# The possible values are: {"linear", "log", "symlog", "logit",
                                ...}
plt.xscale('log')
```

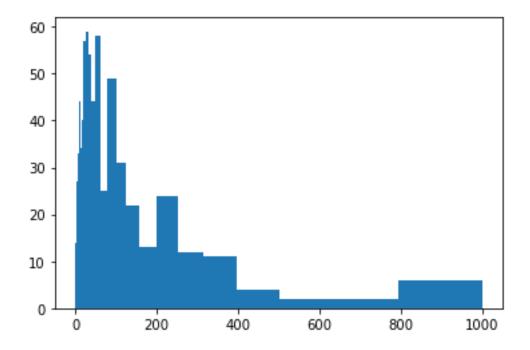


Figure 12: First try histogram of Pokemon's weight.

```
# Apply x-ticks
plt.xticks(ticks, labels)
# See Figure 14
```

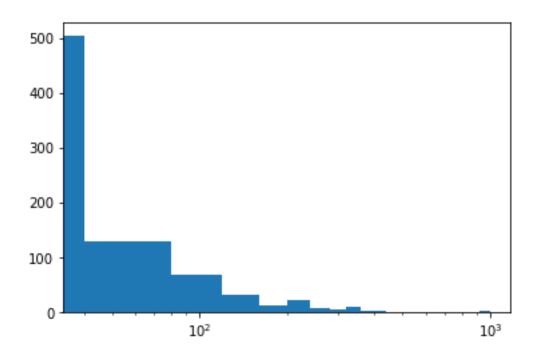


Figure 13: Histogram of Pokemon's weight with a log transformation applied to the x-scale.

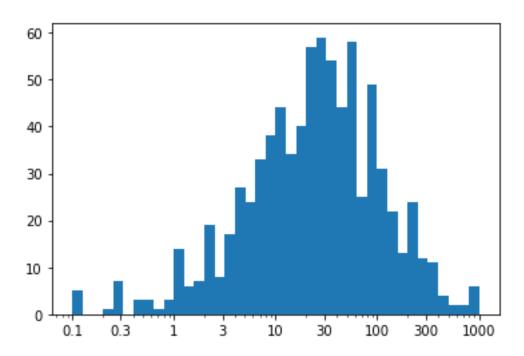


Figure 14: Histogram of Pokemon's weight with transformation and scaling applied.

- 3.2 Bivariate Exploration of Data
- 3.3 Multivariate Exploration of Data
- 4 Explanatory Visualizations
- 5 Visualization Case Study