

## Assignment 2: Web-based Data Visualization using D3

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### 1. Table Visualization

The provided input data were loaded by using `d3.csv()` built-in parser. All scripts were written in 'visualization.js' file, and the web page is implemented by 'index.html' file. Since the table was too big, the area of table was limited in the width of 600px and the height of 500px. The output is as follows:

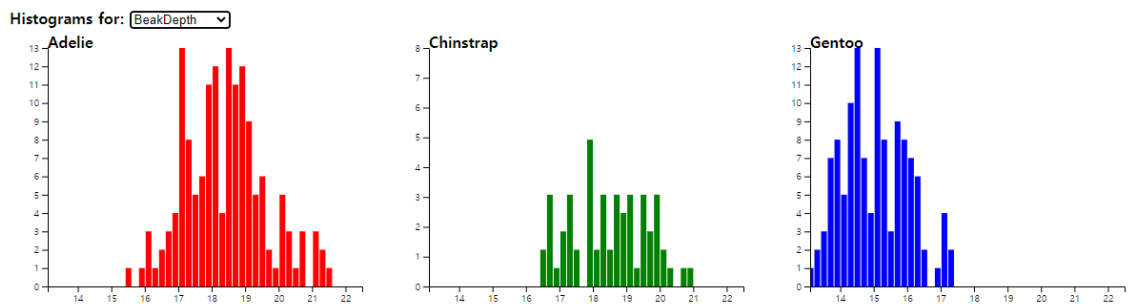
Table

Species	BeakLength	BeakDepth	FlipperLength	BodyMass
Adelie	39.1	18.7	181	3750
Adelie	39.5	17.4	186	3800
Adelie	40.3	18	195	3250
Adelie	36.7	19.3	193	3450
Adelie	39.3	20.6	190	3650
Adelie	38.9	17.8	181	3625
Adelie	39.2	19.6	195	4675
Adelie	34.1	18.1	193	3475
Adelie	42	20.2	190	4250
Adelie	37.8	17.1	186	3300
Adelie	37.8	17.3	180	3700
Adelie	41.1	17.6	182	3200
Adelie	38.6	21.2	191	3800
Adelie	34.6	21.1	190	4400

Source code reference: <https://keeper.tistory.com/13>

## 2. Histogram Plot

By selecting four attributes in a top-down menu, the number of each species were visualized in histogram. Adelie is represented as red, Chinstrap is green, and Gentoo is blue. The output is as follows:

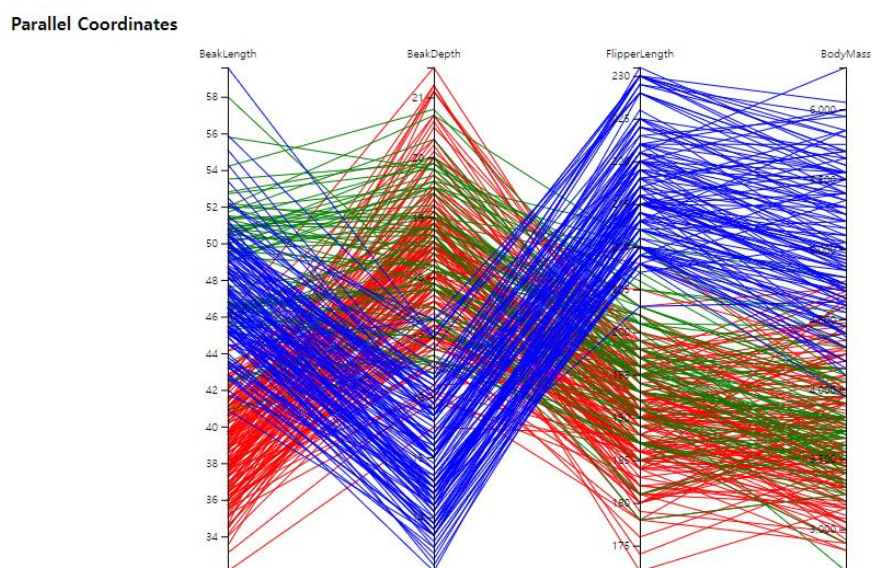


Source code reference: [https://d3-graph-gallery.com/graph/histogram\\_basic.html](https://d3-graph-gallery.com/graph/histogram_basic.html)

[https://d3-graph-gallery.com/graph/line\\_select.html](https://d3-graph-gallery.com/graph/line_select.html) (for top-down menu)

## 3. Parallel Coordinate Plot

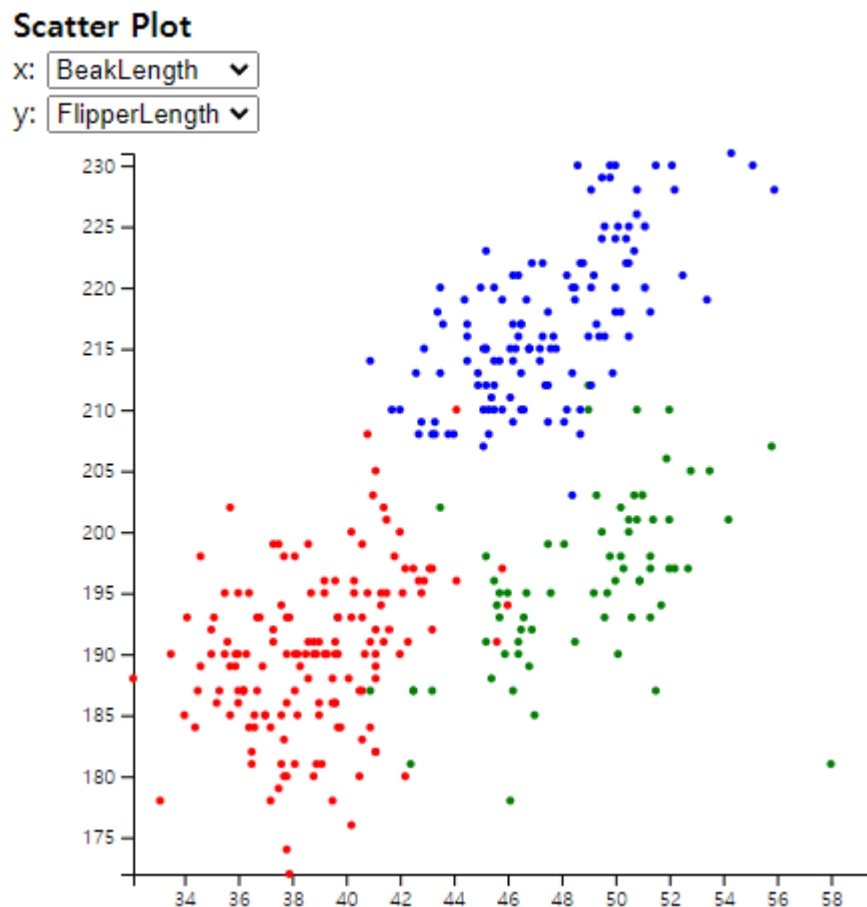
Parallel coordinate plot represents four different attributes, which are beak length, beak depth, flipper length, and body mass. Same as above, species were colored as red, green, and blue, which represent Adelie, Chinstrap, and Gentoo in same order. Each cluster of species is highlighted when the mouse is located on the lines. The output is as follows:



Source code reference: [https://d3-graph-gallery.com/graph/parallel\\_custom.html](https://d3-graph-gallery.com/graph/parallel_custom.html)

#### 4. Scatter Plot

By selecting each x and y axis attributes in top-down menus, each data is displayed as a point on a 2D plot with a corresponding color. Again, colors of species are same as above. The best combination of attributes that separates three penguin species was beak length and flipper length. The output is as follows:



Source code reference: [https://d3-graph-gallery.com/graph/scatter\\_grouped.html](https://d3-graph-gallery.com/graph/scatter_grouped.html)

#### 5. t-SNE Plot

To visualize 2D t-SNE embedding, 'tsne.min.js' script file was imported, which is an algorithm implemented in JavaScript. To calculate the output of t-SNE, raw data were parsed into nested array that only have numeric values of each penguin. After t-SNE is finished, the array contained t-SNE values was split into three arrays, which represent

each species of penguins. User-parameters such as perplexity, learning rate, and metric were available to change specific values. However, it was hard to find the best t-SNE that separates three penguin species. The output is as follows:

### t-SNE Plot

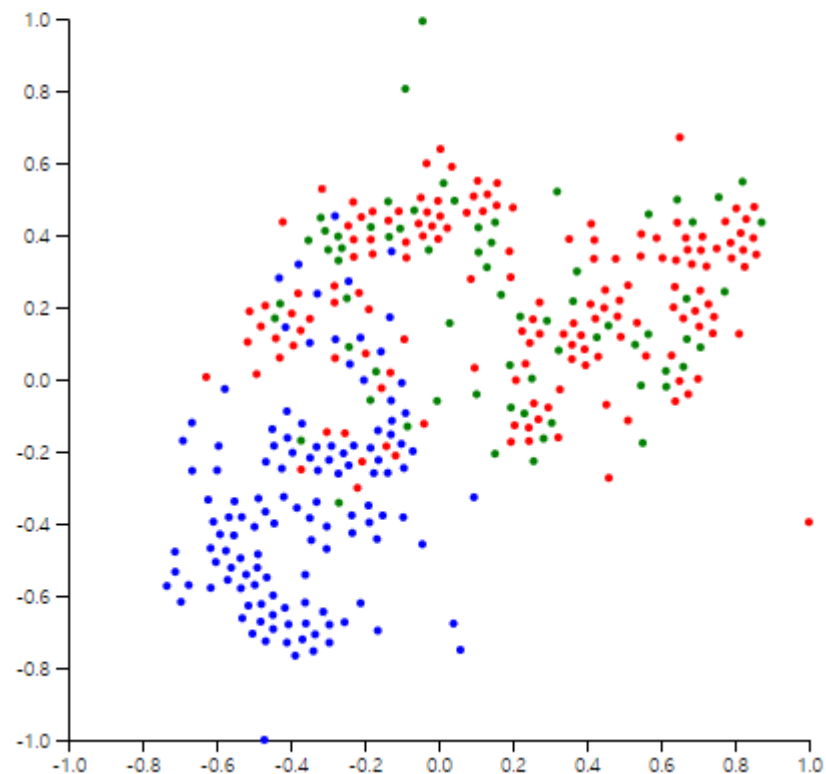
perplexity:

earlyExaggeration:

learningRate:

nIter:

metric:



Source code reference: <https://github.com/scienceai/tsne-js/tree/master>