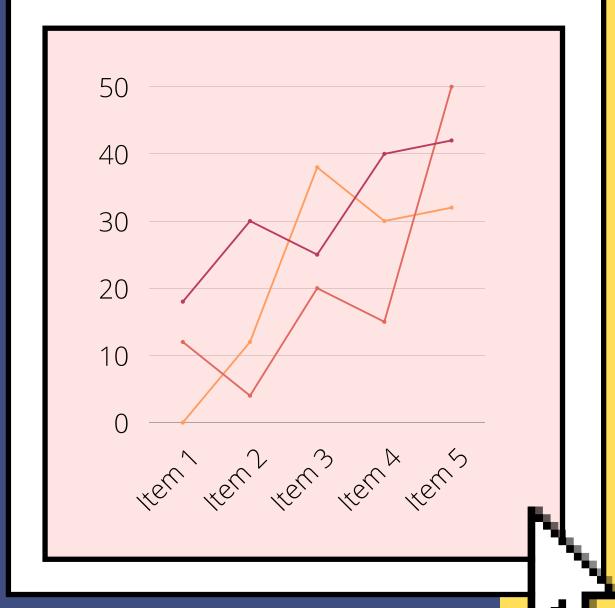
Data Visualisation



CORRELOGRAM

Based on Iris Data Set



Contributers

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About Correlogram



Things to note before you jump in

A correlogram or correlation matrix allows to analyse the relationship between each pair of numeric variables of a dataset. The relationship between each pair of variable is visualised through a scatterplot, or a symbol that represents the correlation (bubble, line, number). The diagonal often represents the distribution of each variable, using an histogram or a density plot.

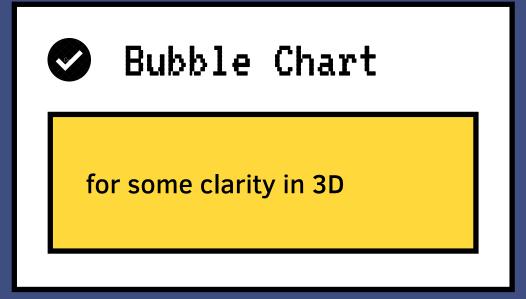
The correlogram is a commonly used tool for checking randomness in a data set.

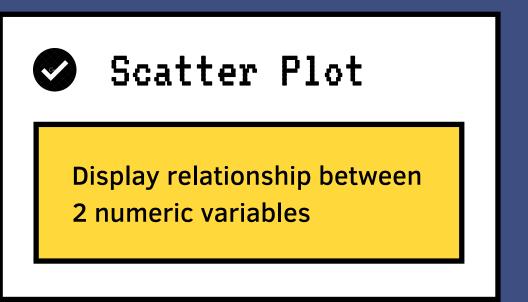
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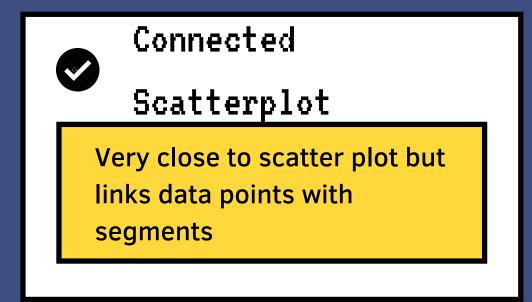


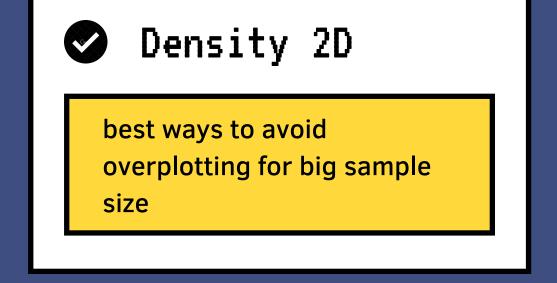


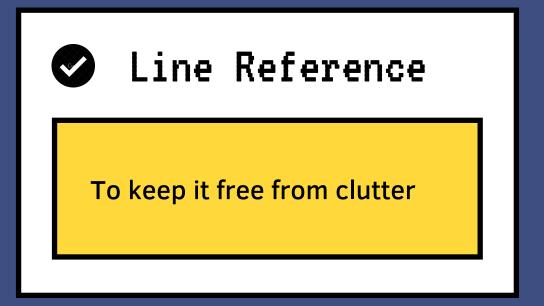
Related Charts:











Iris flower data set

The Iris flower data set is a multivariate data set



The Data:

The data set consists of 50 samples from each of three species of Iris (Iris setosa, Iris virginica and Iris versicolor). Four features were measured from each sample: the length and the width of the sepals and petals, in centimeters. Based on the combination of these four features, Fisher developed a linear discriminant model to distinguish the species from each other.

The Dataset:

The dataset contains a set of 150 records under five attributes - sepal length, sepal width, petal length, petal width and species.

Data Values:

https://docs.google.com/spreadsheets/d/1Kc3YI2F6TEF9D2yurz-7gkkTwVMwRkfo8h1ck1kQeAU/edit?usp=sharing

Code here





HTML

```
<head>
<!-- Load plotly.js into the DOM -->
<script src='https://cdn.plot.ly/plotly-
latest.min.js'></script>
</head>

<body>
<div id='myDiv'><!-- Plotly chart will be
drawn inside this DIV --></div>
</body>
```

```
JAVASCRIPT
Plotly.d3.csv('https://raw.githubusercontent.com/plotly/datasets/mas
ter/iris-data.csv', function(err, rows){
  function unpack(rows, key) {
    return rows.map(function(row) { return row[key.replace('.',' ')]; });
  colors = []
  for (i=0; i < unpack(rows, 'class').length; i++) {</pre>
  if (unpack(rows, 'class')[i] == "Iris-setosa") {
    colors.push(0)
   } else if (unpack(rows, 'class')[i] == "Iris-versicolor") {
    colors.push(0.5)
   } else if (unpack(rows, 'class')[i] == "Iris-virginica") {
    colors.push(1)
  var pl_colorscale=[
        [0.0, '#19d3f3'],
        [0.333, '#19d3f3'],
        [0.333, '#e763fa'],
        [0.666, '#e763fa'],
```

[0.666, '#636efa'],

[1, '#636efa']

Code here



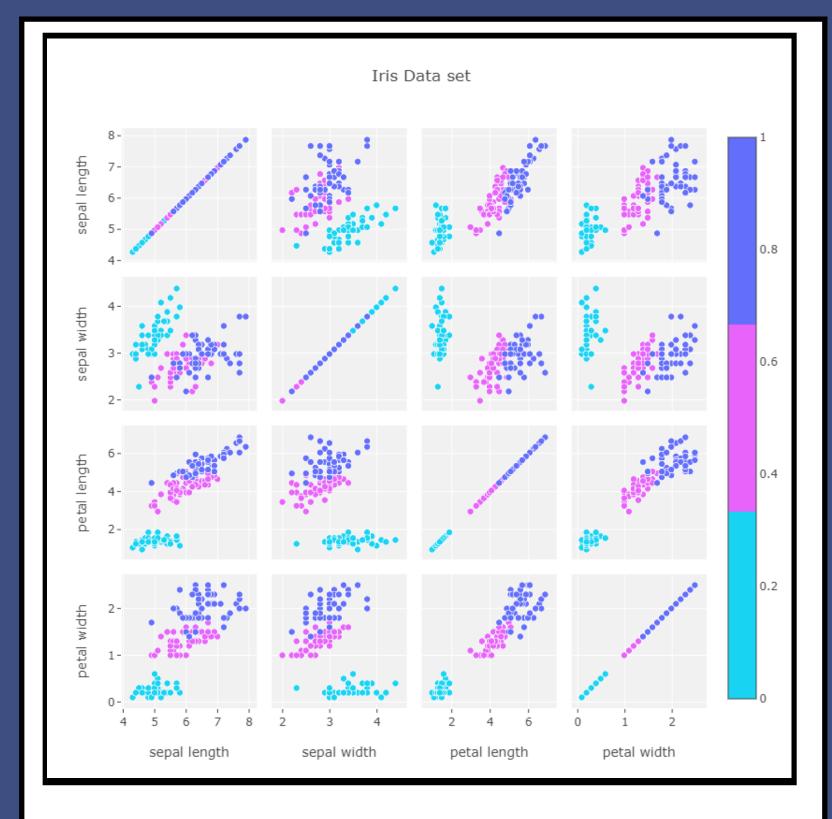


JAVASCRIPT (CONT.)

```
var axis = () => ({
   showline:false,
   zeroline:false,
   gridcolor: '#ffff',
   ticklen:4
  })
  var data = [{
   type: 'splom',
   dimensions: [
    {label:'sepal length',
values:unpack(rows,'sepal length')},
    {label:'sepal width',
values:unpack(rows,'sepal width')},
    {label:'petal length',
values:unpack(rows,'petal length')},
    {label:'petal width',
values:unpack(rows,'petal width')}
text: unpack(rows, 'class'),
   marker: {
    color: colors,
    showscale: true,
```

JAVASCRIPT

```
colorscale:pl_colorscale,
    size: 7,
   line: {
     color: 'white',
     width: 0.5
 var layout = {
   title:'Iris Data set'.
   height: 800,
   width: 800,
   autosize: false,
   hovermode: 'closest',
   dragmode: 'select',
   plot_bgcolor:'rgba(240,240,240, 0.95)',
   xaxis:axis(),
   yaxis:axis(),
   xaxis2:axis(),
   xaxis3:axis(),
   xaxis4:axis(),
   yaxis2:axis(),
   yaxis3:axis(),
   yaxis4:axis(),
   template: {
    data: {
     choroplethmapbox: [{
      marker: {
       colorbar: { ypad: 100},
       showscale: true,
```



THE OUTPUT

Correlogram are really handy for exploratory analysis. It allows to visualize the relationships of the whole dataset in a glimpse. For instance, the linear relationship between petal length and petal width is obvious here, as the one concerning sepal.



GRACIAS