

dm25s1

Topic 04 : Exploratory Data Analysis

Part 02 : EDA Pass2

Dr Bernard Butler

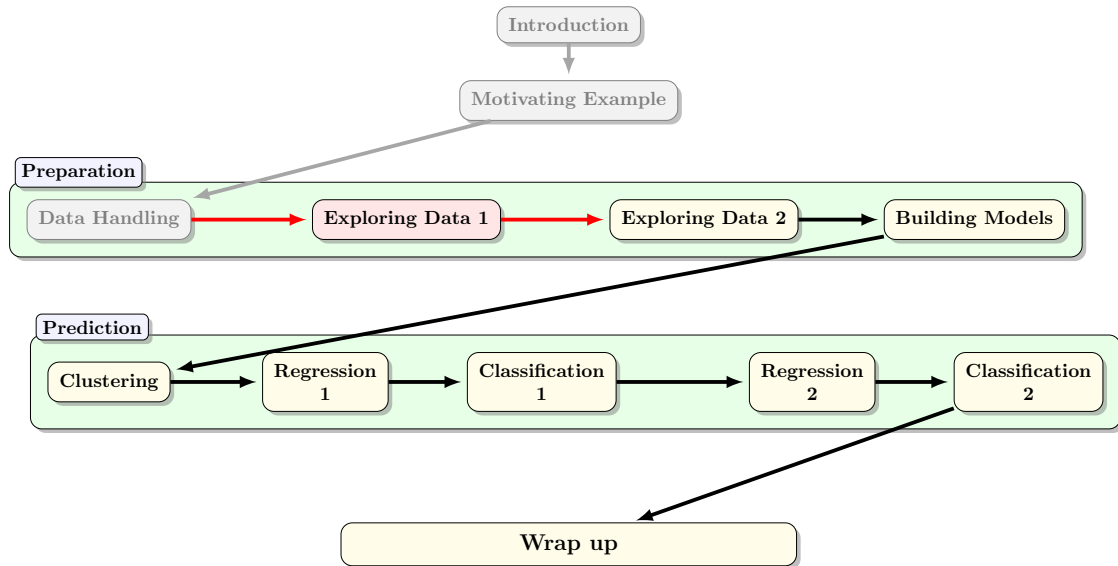
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Autumn Semester, 2025

Outline

- Feature and target distributions
- Plotting
- After EDA Pass 2

Data Mining (Week 4)



1. Introduction

2. A Selection of Statistical Visualisations and Metrics

2.1 Categorical Features

2.2 Numerical Features

3. Summary

Acknowledgment

A big thanks to Dr Kieran Murphy, who provided many of the slides for today's lecture.

Introducing EDA Pass2

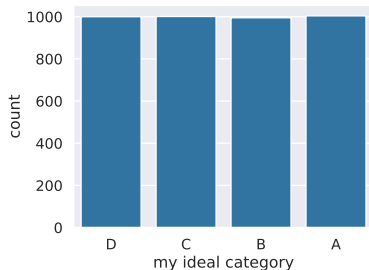
A Selection of Statistical Visualisations and Metrics

Relationship to Target		
	Feature	<div>Categorical</div> <div>Numerical</div>
Categorical	nunique, unique, describe, value_counts, ...	crosstab, ...
	countplot, ...	catplot, boxplot, ...
Numerical	describe, ...	groupby+describe, ...
	histplot, boxplot, displot, qqplot, ...	lmpplot, ...

Categorical Variables

The Ideal

- Each level equally likely.
- Not too many levels: 2–12(ish).



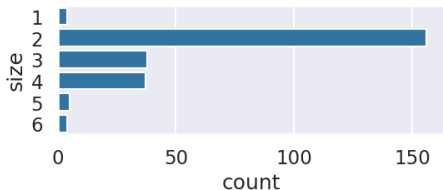
Tools

- `nunique`, `unique`, `value_counts`.
- `sns.countplot` — shows the counts of observations in each categorical level using bars.

Reality

```
sns.countplot(y="size", data=df);
```

Tips



- If size was the target, then most models will train towards the majority class (`size=2`).
- If size was a feature, then quality of predictor could vary greatly depending on the feature categorical level.
- Consider merge/drop rare category levels.

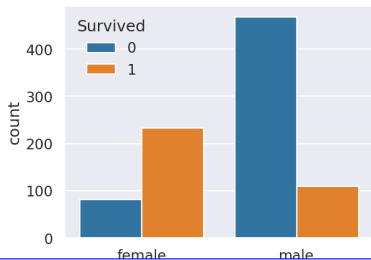
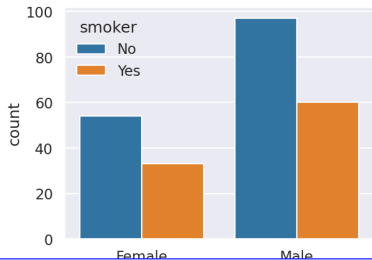
Categorical Variables — Relationship with (Categorical) Target

feature

target

Titanic

```
sns.countplot(x="Sex", hue="Survived", data=df);
```



```
pd.crosstab(columns=df.Sex, index=df.Survived, margins=True, normalize='columns') \
    .style.format("{:.2%}").background_gradient(cmap='summer_r')
```

sex	Female	Male	All
smoker			
No	62.07%	61.78%	61.89%
Yes	37.93%	38.22%	38.11%

No relationship between sex and smoker

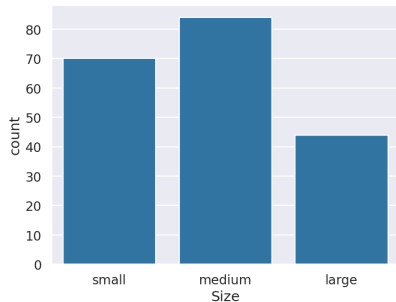
Sex	female	male	All
Survived			
0	25.80%	81.11%	61.62%
1	74.20%	18.89%	38.38%

Strong relationship between Sex and Survived

Categorical Variables — Relationship with (Numerical) Target

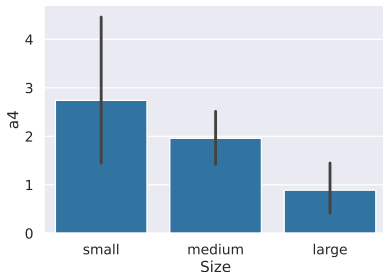
I

```
sns.countplot(x="Size", data=df);
```



- Shows the counts of observations in each categorical level using bar (height/width).

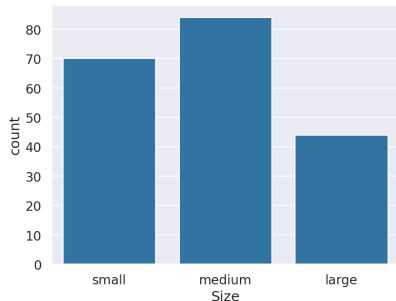
```
sns.catplot(x="Size", y="a4", data=df, kind='bar');
```



Categorical Variables — Relationship with (Numerical) Target

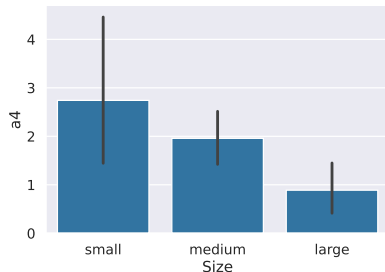
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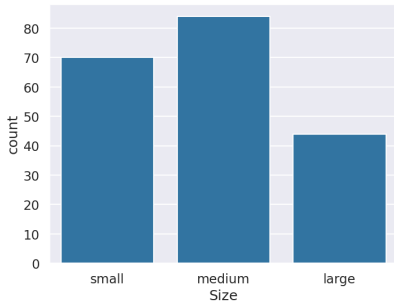


- Shows the average level (mean) and uncertainty (std) of the numerical target (a4) in each categorical level of the categorical variable.
- Vertical bar shows 95% confidence interval.

Categorical Variables — Relationship with (Numerical) Target

I

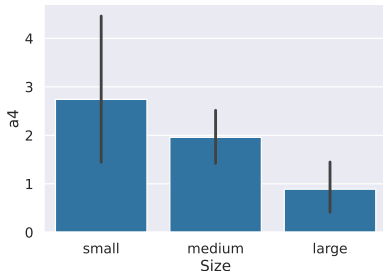
```
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```



- Shows the counts of observations in each categorical level using bar (height/width).

Is it usable?

```
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```



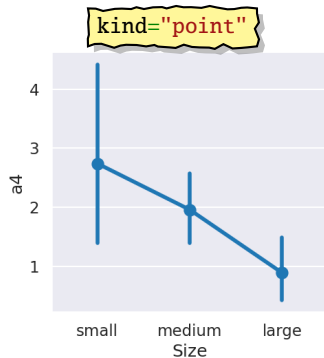
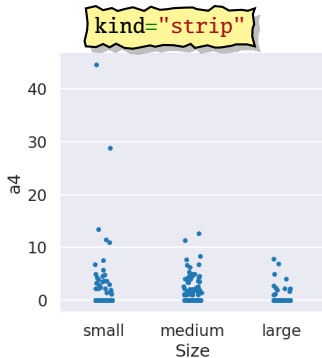
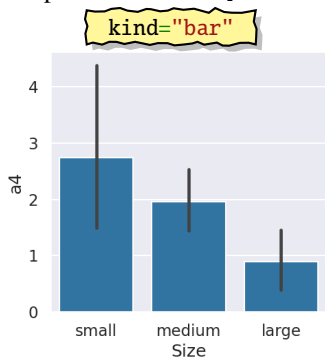
- Shows the average level (mean) and uncertainty (std) of the numerical target (a4) in each categorical level of the categorical variable.
- Vertical bar shows 95% confidence interval.

Is it useful?

Categorical Variables — Relationship with (Numerical) Target

II

The option `kind` in `catplot` can be:

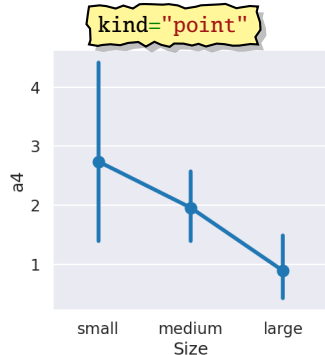
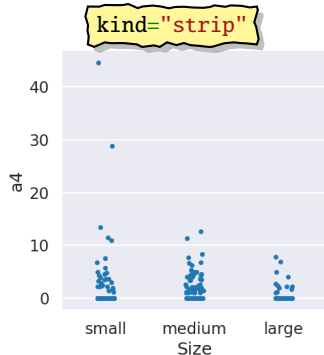
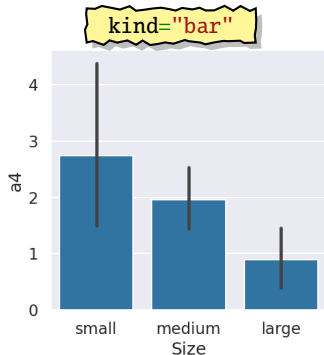


- bar and point show essentially the same information, but point is more compact when comparing multiple categorical features to a continuous target on the same plot.

Categorical Variables — Relationship with (Numerical) Target

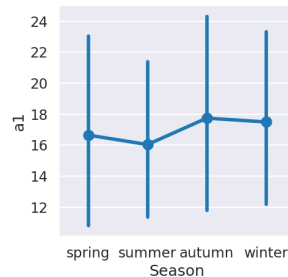
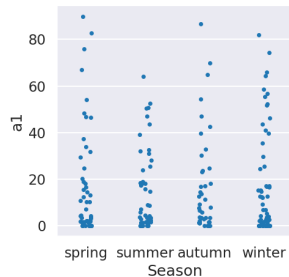
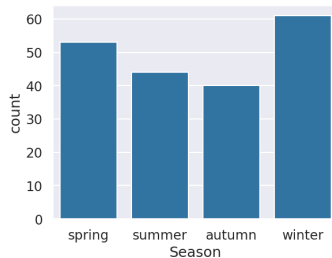
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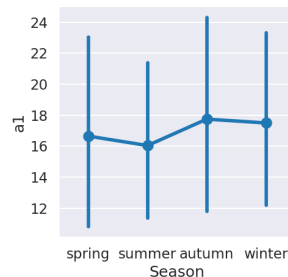
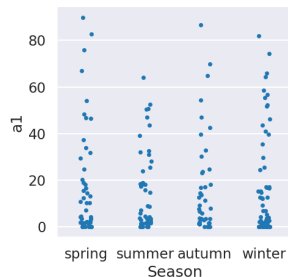
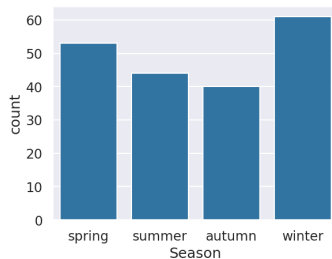


- bar and point show essentially the same information, but point is more compact when comparing multiple categorical features to a continuous target on the same plot.
- strip shows individual observations — useful (as in this case) to show that the larger uncertainty in `Size="small"` observations is mainly due to two outliers.

Example — Dataset: Algae Blooms, Feature: Season, Target: a1



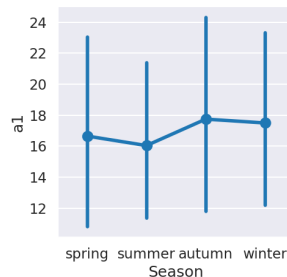
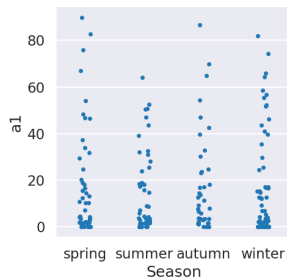
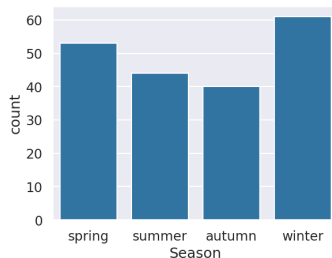
Example — Dataset: Algae Blooms, Feature: Season, Target: a1



```
df.groupby("Season")["a1"].agg(["mean", "count", "std"])
```

Season	mean count		std
	\bar{x}	n	σ
spring	16.649057	53	23.093786
summer	16.038636	44	17.920798
autumn	17.745000	40	21.611203
winter	17.498361	61	22.568256

Example — Dataset: Algae Blooms, Feature: Season, Target: a1

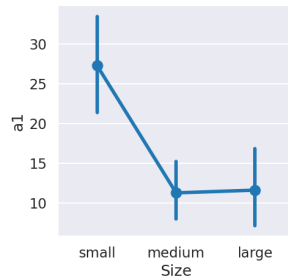
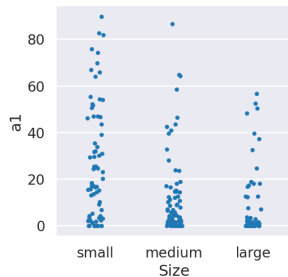
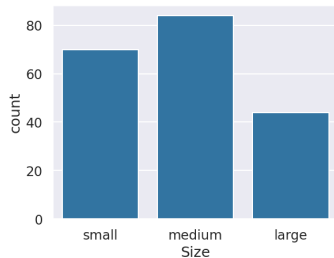


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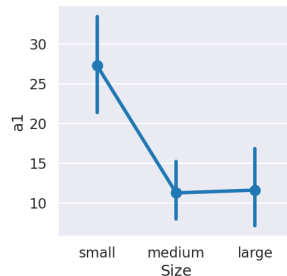
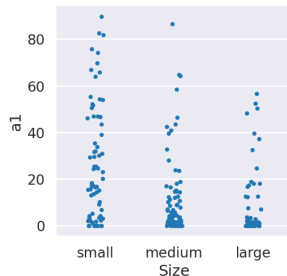
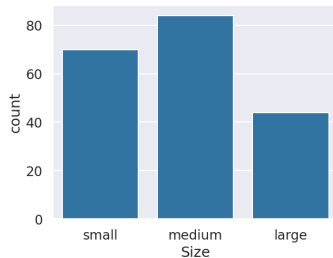
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- Countplot shows no issues with feature **Season** — all levels approximately equally represented.
 - Catplots show slightly less spread in **a1** for **Season="summer"** observations. (**strip** shows smaller range, **point** shows smaller standard deviation).
- ⇒ Mean levels of **a1** for different levels of **Season** are well within the 95% confidence intervals ($\bar{x} \pm \sigma 1.96/\sqrt{n}$), so no/weak relationship between categorical feature and numerical target.

Example — Dataset: Algae Blooms, Feature: Size, Target: a1



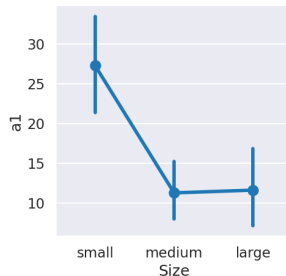
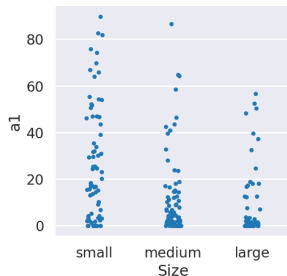
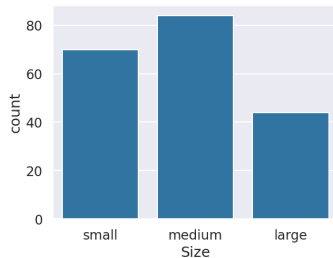
Example — Dataset: Algae Blooms, Feature: Size, Target: a1



```
df.groupby("Size")["a1"].agg(["min", "max", "mean", "count", "std"])
```

	min	max	mean	count	std
Size			\bar{x}	n	σ
small	0.0	89.8	27.255714	70	24.895426
medium	0.0	86.6	11.267857	84	17.163124
large	0.0	56.8	11.611364	44	16.556123

Example — Dataset: Algae Blooms, Feature: Size, Target: a1



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- Countplot shows no issues with feature Size.
 - Catplot (point) shows that levels of a1 are higher for Size="small" observations.
- ⇒ Confidence interval for Size="small" observations do not overlap with CI for other levels, so significant relationship between categorical feature and numerical target.

Numerical Variables

Things here are more complicated as a numerical variable could follow many different distributions. Here we look at data following the standard normal distribution. To start we generate 10,000 values and put in to new DataFrame, df2.

```
rv = stats.norm()  
data = rv.rvs(size=10_000)  
df2 = pd.DataFrame(data, columns=["x"])  
df2.head(5)
```

	x
0	-1.915795
1	0.641175
2	0.040114
3	0.725531
4	0.252675

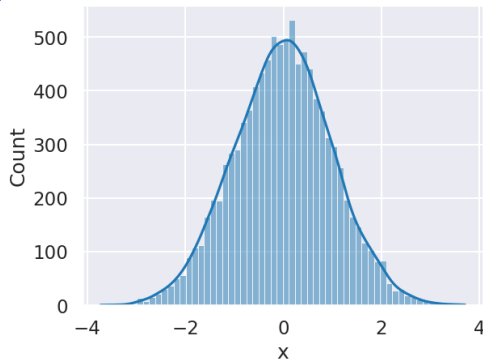
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```
sns.histplot(x="x", data=df2, kde=True);
```



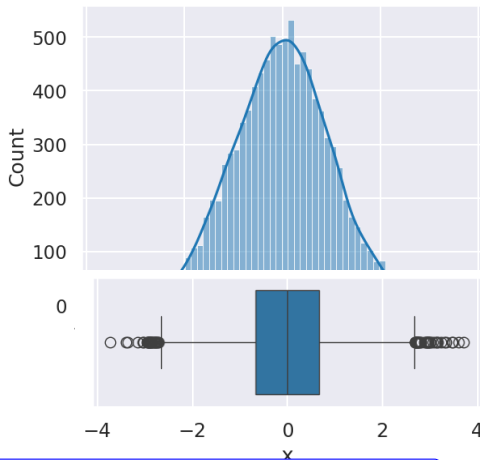
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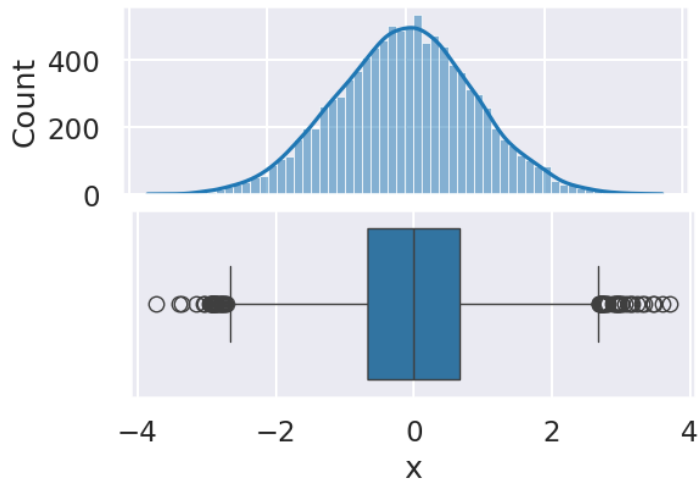
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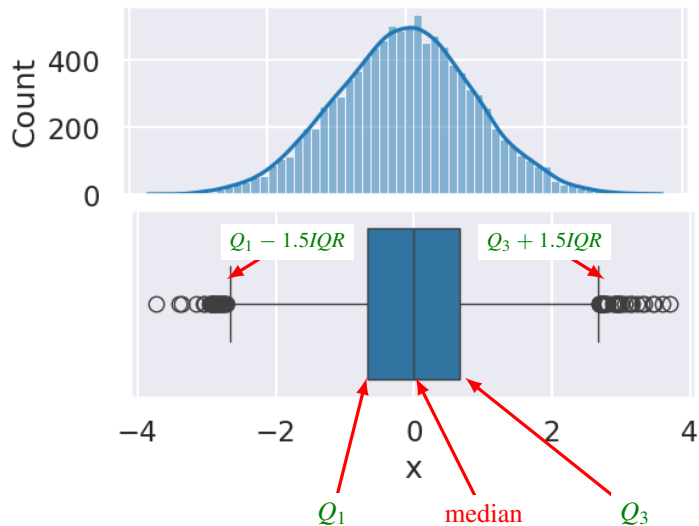


```
sns.boxplot(x="x", data=df2);
```

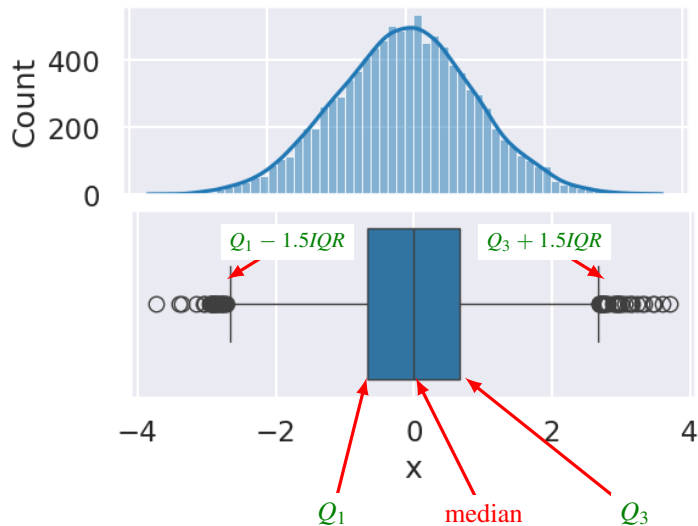
Histplot (Histogram) and Boxplot



Histplot (Histogram) and Boxplot

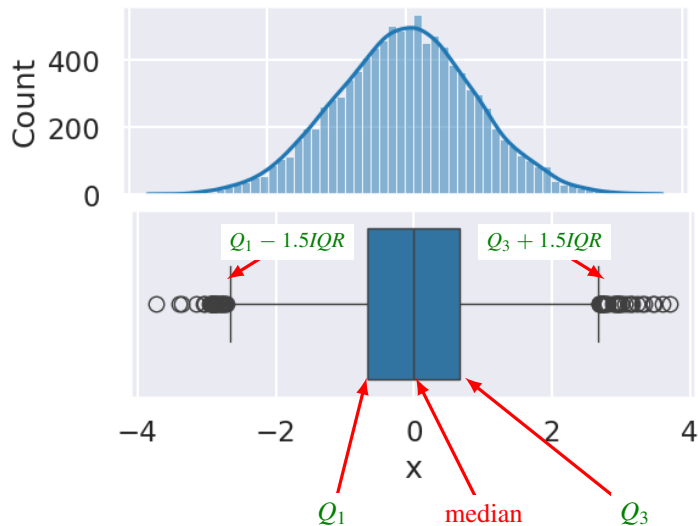


Histplot (Histogram) and Boxplot



- Histogram is useful in depicting location, spread and shape.
- Curve, is estimate of shape given infinite data and infinite number of bins.
- Boxplots also depicts location, spread and shape, but uses median for estimate of centre, and quartiles for spread.

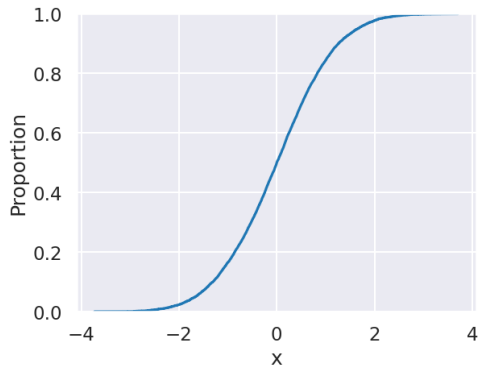
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- Curve, is estimate of shape given infinite data and infinite number of bins.
- Boxplots also depicts location, spread and shape, but uses median for estimate of centre, and quartiles for spread.
- Half the data is within the box, data points outside the whiskers (lines) are possible outliers, denoted by circles.

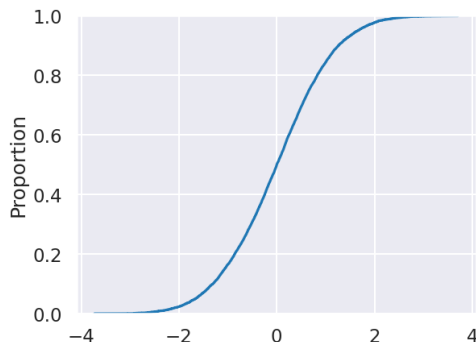
Cumulative Plot and QQ-Plot

```
sns.ecdfplot(data=df2, x="x");
```



Cumulative Plot and QQ-Plot

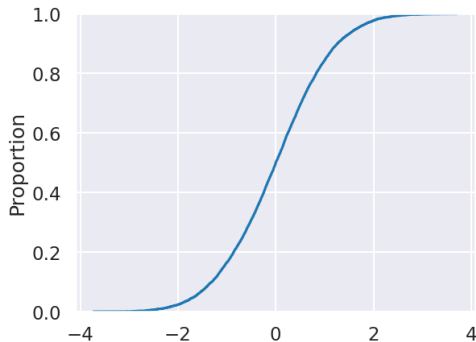
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- Represents the proportion of observations less than or equal to given value.

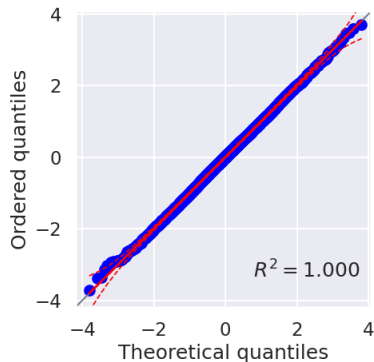
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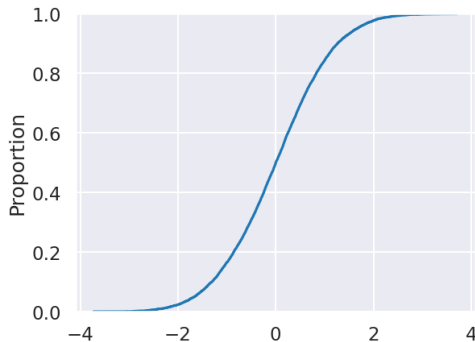
- Represents the proportion of observations less than or equal to given value.

```
import pingouin as pg  
pg.qqplot(df2.x);
```



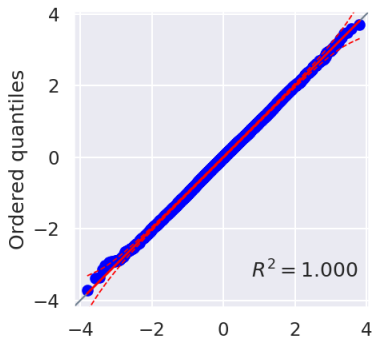
Cumulative Plot and QQ-Plot

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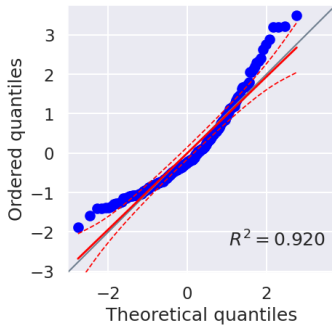
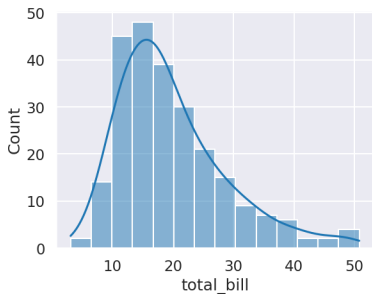
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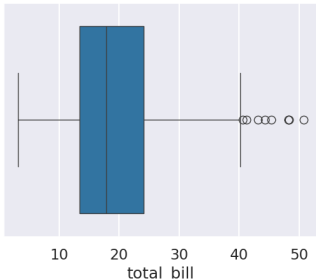
- Plot of observed quantiles against theoretical (assumed normal) quantiles. If both sets of quantiles came from the same distribution, the points lie on a line (approx.).

Example — Dataset: Tips, Feature: total_bill

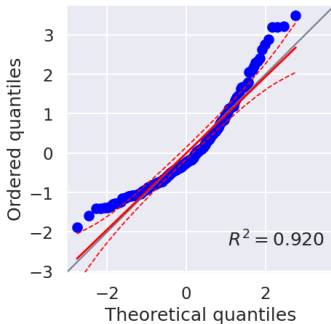
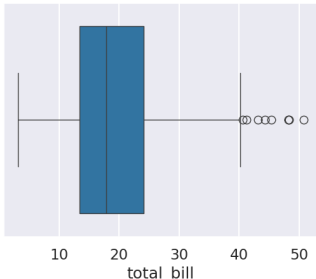
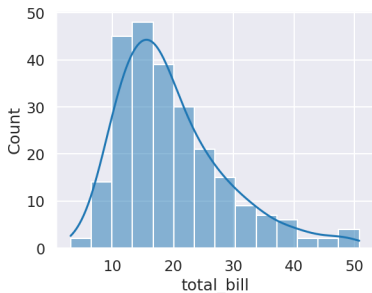


```
df.total_bill.describe()
```

```
count    244.000000
mean      19.785943
std       8.902412
min       3.070000
25%      13.347500
50%      17.795000
75%      24.127500
max      50.810000
Name: total_bill, dtype: float64
```



Example — Dataset: Tips, Feature: total_bill

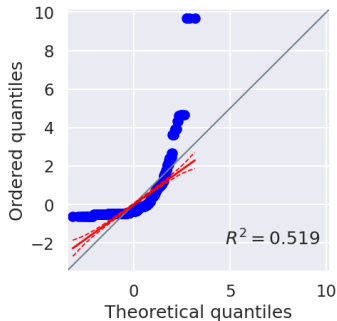
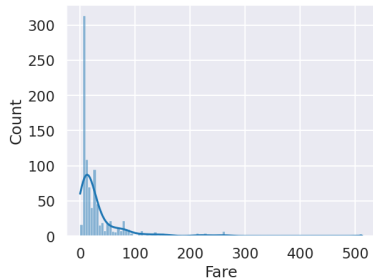


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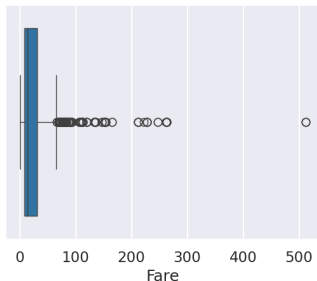
- Data is bell curve shaped, but right skewed (data is more spread out to the right).
- Outliers to the right.
- QQ-Plot indicate that data is not normal, but we could transform it to be more closer to normal.

Example — Dataset: Titanic, Feature: Fare

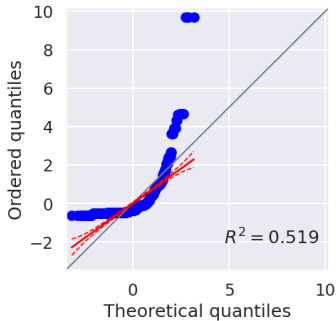
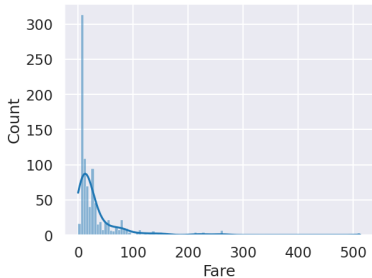


```
df.Fare.describe()
```

```
count    891.000000  
mean      32.204208  
std       49.693429  
min        0.000000  
25%       7.910400  
50%      14.454200  
75%      31.000000  
max      512.329200  
Name: Fare, dtype: float64
```

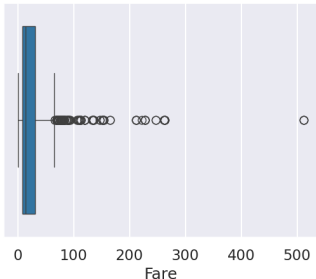


Example — Dataset: Titanic, Feature: Fare



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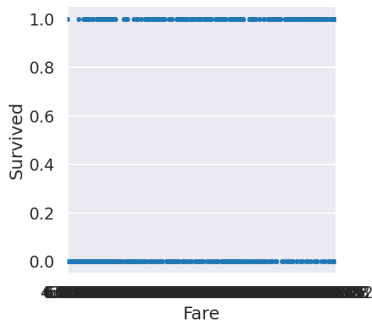
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- This variable is more skewed and dominated by its outliers which need to be resolved.

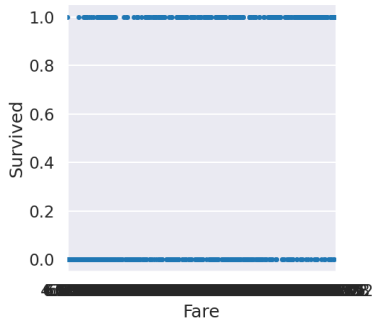
Warning — Plot Output Depends on Data Assumptions

```
df = pd.read_csv("data/train.csv")  
sns.catplot(data=df, x="Fare", y="Survived");
```

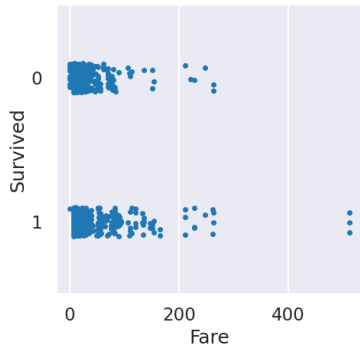


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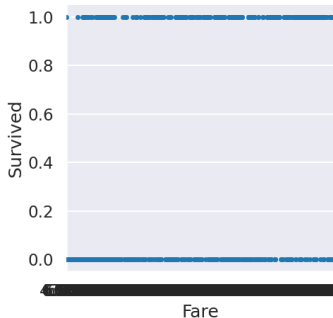


```
df = pd.read_csv("data/train.csv")  
df.Survived = df.Survived.astype(str)  
sns.catplot(data=df, x="Fare", y="Survived");
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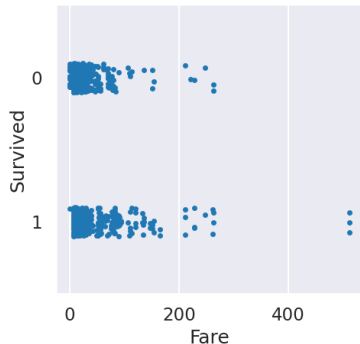
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- **seaborn** tries to infer the correct graph based on the data values/type, but it does not always get it correct.
- **Survived** stores 0 and 1 and has dtype **int**.
- Converting to a **Categorical** with numeric levels is not enough.
- **astype(str)** converts 0 and 1 to "0" and "1".

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 - ③ the visualisation objective: observe distributions or relationships within a column